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EU464751.1 gg_id:300229 Hartmann's mountain zebra feces clone AFZEB_aaj70b10 EU464
CU927648.1 gg_id:564787 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU927
FJ167470.1 gg_id:324457 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1674
CU919583.1 gg_id:541370 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU919
GU455142.1 gg_id:530933 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU455

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EU801561.1 gg_id:332569 It's all ranking aquatic Chesapeake Bay MD clone 3C002885 EU801
EU800965.1 gg_id:328912 It's all ranking aquatic Delaware Bay NJ clone 2C229230 EU800
AY947947.1 gg_id:141740 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY9479
AY792257.1 gg_id:124653 Humic Lake clone CrystalBog021H12 AY7922
DQ501305.1 gg_id:195239 Seasonal dynamics free-living and particle- four lakes northeastern Germany freshw DQ501
EU801645.1 gg_id:341378 It's all ranking aquatic Chesapeake Bay MD clone 3C002991 EU801
AY948061.1 gg_id:141397 Synchrony and seasonality two temperate rivers USA: Massachusetts Parker River cl AY9480
EU800346.1 gg_id:310935 It's all ranking aquatic Delaware Bay NJ clone 2C228402 EU800
FJ612165.1 gg_id:512885 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP3.4.43 FJ6121
AM849435.1 gg_id:243487 Seasonality Piburger See clone PIB-24 AM849
EF667919.1 gg_id:250266 Long-term maintenance species-specific microbiota basal metazoan Hydra epitheliun EF6679
EF667927.1 gg_id:245084 Long-term maintenance species-specific microbiota basal metazoan Hydra epitheliun EF6679
FJ612313.1 gg_id:533183 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.1.5 FJ6123
AY863096.1 gg_id:112283 wastewater treatment pool clone SXAU059 AY863
AM501437.1 gg_id:240464 Pelomonas aquatica str. CCUG52631 AM501

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L33974.1 gg_id:7170 Leptothrix discophora str. SP-6	L33974
AM501440.1 gg_id:248405 Pelomonas puraquae str. CCUG52770 MG64	AM501
AF396932.1 gg_id:112368 Pseudomonas saccharophila str. PRO7	AF3969
DQ884343.1 gg_id:170667 Pseudomonas sp. str. JS-1	DQ884
AM039830.1 gg_id:133518 Polaromonas aquatica str. CCUG 39402	AM039
NC_008781.1 gg_id:194205 Polaromonas naphthalenivorans str. CJ2	NC_00
AAFQ02000005.1 gg_id:129807 Polaromonas sp. str. JS666	AAFQ0
DQ298150.1 gg_id:142773 -dependent and microbial assemblages silkworm midgut clone swm42	DQ298
AB464945.2 gg_id:323641 Altitudinal changes flora on Gulkana Glacier Alaska analyzed sequencing and -cycle c	AB4649
AF361203.1 gg_id:53922 mesoeutrophic reservoir clone 16	AF3612
AY166684.1 gg_id:76891 Polaromonas naphthalenivorans str. CJ2	AY1666
U14585.1 gg_id:7005 Polaromonas vacuolata str. 34-P	U14585
DQ341420.1 gg_id:150971 Acidovorax sp. str. Antarctic IS08	DQ341
AF407386.1 gg_id:33769 MCB-contaminated groundwater-treating reactor clone RB7C8	AF4073
AB252078.1 gg_id:152145 entrapped Puruogangri ice core clone PIC-D3	AB2520
EU801289.1 gg_id:323062 It's all ranking aquatic Chesapeake Bay MD clone 3C002543	EU8012
EU636027.1 gg_id:278933 Antarctic str. G020	EU6360
EU636024.1 gg_id:279595 Antarctic str. G076	EU6360
AY218637.1 gg_id:81748 penguin droppings sediments clone KD4-57	AY2186
EU636029.1 gg_id:278079 Antarctic str. G088	EU6360
FJ979854.1 gg_id:384326 Polaromonas sp. str. tsz24	FJ9798
FJ801199.1 gg_id:552577 Investigation Water -DGGE and Clone Library Zoige Alpine Wetland clone ZWB3-18	FJ8011
EU030285.1 gg_id:254221 Polaromonas jejuensis str. JS12-13	EU0302
EU801930.1 gg_id:355065 It's all ranking aquatic Chesapeake Bay MD clone 3C003311	EU8019
EF423322.1 gg_id:212990 Polaromonas sp. str. 1006	EF4233
EU703261.1 gg_id:448038 lakes on Eastern Tin Plateau oligosaline lake altitude 4718 masl clone XZNMC60	EU7032
DQ094183.1 gg_id:130292 Polaromonas sp. 'hydrogenovorans' str. hydrogenovorans	DQ094
EU636028.1 gg_id:279079 Antarctic str. CC10	EU6360
AY188325.1 gg_id:79327 penguin droppings sediment clone KD7-99	AY1883
FM161402.1 gg_id:291214 Polaromonas sp. str. 01WB02.3-26	FM161
AB167243.1 gg_id:107951 str. c115	AB1672
AY168739.1 gg_id:75143 arsenite-oxidizing biofilm clone HC-3	AY1687
GU213397.1 gg_id:540184 Polaromonas sp. str. 190	GU213
EF516819.1 gg_id:224825 grassland soil clone FCPN433	EF5168
GU208230.1 gg_id:588612 Study on Sediment Dongping Lake sediment clone 3S4-1	GU208
AY921863.1 gg_id:113128 farm soil clone AKYH1084	AY9218
EU881341.1 gg_id:340905 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM	EU8813
DQ984611.1 gg_id:237973 LY oil-contaminated soil clone LYC180	DQ984
GQ079635.1 gg_id:390755 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw1232d(	GQ079
AY439189.1 gg_id:104669 soil clone MeCl 24	AY4391
AJ532684.1 gg_id:76621 uranium mill tailings clone Gitt-KF-21	AJ5326

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DQ354711.1 gg_id:154792 subsurface water clone DR550SWSAEE7 DQ354
EF516224.1 gg_id:221803 grassland soil clone FCPU687 EF5162
AB487846.1 gg_id:591765 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4878
FJ478510.1 gg_id:578098 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4785
DQ451482.1 gg_id:156693 forest soil clone FAC43 DQ451
EU979062.1 gg_id:317424 Effects intercropping Rhizobium inoculation on microbial biomass and rhizosphere b EU979
FM178814.1 gg_id:343929 Monitoring abundance and expression 'dehalococcoides' species chloroethene-redu FM178
NC_007908.1 gg_id:153264 Rhodoferax ferrireducens str. DSM 15236 NC_00
D16212.1 gg_id:6979 Rhodoferax fermentans str. FR3 D16212
X95837.1 gg_id:6986 situ visualization genetic natural microbial ecies strain T3 X95837
FJ517716.1 gg_id:513243 Disturbing epithelial homeostasis metazoan Hydra leads drastic changes microbiota e FJ5177
AY214178.1 gg_id:82365 benzene-contaminated groundwater clone ZZ9C20 AY2141
DQ628924.1 gg_id:162964 sequences Arctic glacier subglacial water ice or sediment clone SIB1 2B DQ628
DQ856540.1 gg_id:254818 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C2M DQ856
AJ606333.1 gg_id:100951 Comamonadaceae SB1 AJ6063
DQ628925.1 gg_id:170859 sequences Arctic glacier subglacial water ice or sediment clone SIB2 1D DQ628
AB199580.1 gg_id:110990 river water Japan clone RVW-13 AB1995
EU907895.1 gg_id:315915 Identification hexahydro-135-trinitro-135-triazine (RDX)-degrading microorganisms v EU9078
AY218635.1 gg_id:81515 penguin droppings sediments clone KD4-45 AY2186
AF418942.1 gg_id:43083 USA: Colorado Fort collins Horsetooth Reservoir clone HTA10 AF4189
EF203797.1 gg_id:236168 Chlorochromatium aggregatum EF2037
AB154311.1 gg_id:104501 freshwater clone S9F-53 AB1543
EF451646.1 gg_id:216339 Rhodoferax sp. str. KAR16 EF4516
EF580989.1 gg_id:278249 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM7 EF5809
AJ421912.2 gg_id:27768 Germany:Elbe River clone Elb08 AJ4219
EF516373.1 gg_id:224371 grassland soil clone FCPT660 EF5163
AB237669.1 gg_id:140742 microorganisms deep subsurface groundwater sedimentary rock milieu clone HDBW AB2376
EU937983.1 gg_id:333026 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
AJ867929.1 gg_id:142024 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone D8 AJ8679
EU937978.1 gg_id:330894 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
EU937973.1 gg_id:303742 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
EU104042.1 gg_id:353830 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1040
GU233447.1 gg_id:516603 Rhodoferax antarcticus str. ANT.BR GU233
CU466889.1 gg_id:338390 Evry wastewater treatment plant anoxic basin clone CU466
AB475015.1 gg_id:566864 sequences located Engiadina Bassa Valley Switzerland iron containing freshwater so AB4750
AY214173.1 gg_id:81368 benzene-contaminated groundwater clone ZZ9C2 AY2141
AB478658.1 gg_id:534216 SSU sequences Alps Cadagno di Fuori submerged biofilm mats montane wetland clo AB4786
AB504936.1 gg_id:552998 structure microorganisms reddish-brown iron-rich snow clone oze05B33 AB5049
GQ340212.1 gg_id:552309 Prokaryotic Marathonas Reservoir Greece water column clone VE07-26-BAC GQ340
EF580979.1 gg_id:279495 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1 EF5809

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AF435948.2 gg_id:75345 Rhodoferax ferrireducens str. T118 AF435
EF520458.1 gg_id:233033 structure lakes: what controls ? acid-impacted lake clone ADK-HDe02-15 EF5204
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AB475020.1 gg_id:534700 sequences located Engiadina Bassa Valley Switzerland iron containing freshwater so AB475
CU466882.1 gg_id:347539 Evry wastewater treatment plant anoxic basin clone CU466
EU340210.1 gg_id:248312 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU340
AY013643.2 gg_id:23857 sample taken downstream landfill clone BVB72 landfill AY013
AY050600.1 gg_id:34176 MCB-contaminated groundwater-treating reactor clone GOUTB17 AY050
AF523029.1 gg_id:88367 natural mineral water clone G-5 AF523
EU104076.1 gg_id:340208 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
AM697182.1 gg_id:222656 sequencing libraries indoor dust clone BF0001C023 AM697
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FJ437765.1 gg_id:546951 Benthic okenone production Fayetteville New York Green Lake chemocline 20.5 m wa FJ4377
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GQ389172.1 gg_id:587804 quality deterioration drinking system during serious red water outbreak clone D42 GQ389
AJ421927.2 gg_id:20305 Germany:Elbe River clone Elb154 AJ4219
AF523038.1 gg_id:90178 natural mineral water clone H-5 AF523
EU104187.1 gg_id:331535 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
DQ228371.1 gg_id:144642 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
EU499697.1 gg_id:263522 Field-based laboratory stable isotope probing surveys identities both aerobic and an EU499
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EF520463.1 gg_id:231370 structure lakes: what controls ? acid-impacted lake clone ADK-MOh02-46 EF5204
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EU636030.1 gg_id:278005 Antarctic str. G079 EU636
DQ227795.1 gg_id:142898 Rhodoferax sp. PIC-C33 DQ227
AJ421938.2 gg_id:18730 Germany:Elbe River clone Elb263 AJ4219
DQ664242.1 gg_id:172764 Rhodoferax sp. str. IMCC1723 DQ664
DQ228394.1 gg_id:143719 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
DQ532326.1 gg_id:168650 Geographical Variation Microbial Burden and Spacecraft Assembly Kennedy Space C DQ532
AF407396.1 gg_id:50642 MCB-contaminated groundwater-treating reactor clone RB9C6 AF407
CU466916.1 gg_id:337770 Evry wastewater treatment plant anoxic basin clone CU466
AF407412.1 gg_id:44459 MCB-contaminated groundwater-treating reactor clone RB13C6 AF407
AM935036.1 gg_id:546472 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
AY214181.1 gg_id:82413 benzene-contaminated groundwater clone ZZ12C5 AY214
DQ521494.1 gg_id:198759 Antarctic lake ice cover clone ANTLV1_G06 DQ521

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CT574057.1 gg_id:151642 Evry municipal wastewater treatment plant clone 054F01_B_DI_P58 CT574
CR933282.1 gg_id:109578 Evry municipal wastewater treatment plant clone 061G11_B_SD_P93 CR933
AF422679.1 gg_id:37583 TCE-sediment clone d156 AF4226
AY218554.1 gg_id:82194 penguin droppings sediments clone KD1-134 AY218
FJ982926.1 gg_id:422910 Simplicispira limi str. ST3 FJ9829
AF078755.1 gg_id:7088 Aquaspirillum psychrophilum str. LMG 5408 AF0787
AY780904.1 gg_id:143192 Aquaspirillum metamorphum str. 416 AY780
Z93960.1 gg_id:7097 activated sludge clone T33 Z93960
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EU652480.1 gg_id:278324 Simplicispira sp. str. JPB-3.01 EU6524
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CU466844.1 gg_id:324604 Evry wastewater treatment plant anoxic basin clone CU466
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DQ640671.1 gg_id:233183 full-scale EBPR sludge clone Skagenf43 DQ640
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GQ024888.1 gg_id:484782 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ024

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DQ521517.1 gg_id:190163 Antarctic lake ice cover clone ANTLV2_E04 DQ521
FM873335.1 gg_id:367349 occupant source house floor dust clone FD01H06 FM873
EF660749.1 gg_id:245337 Pelomonas soli str. SYB4 EF6607
EF632937.1 gg_id:252287 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6329
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AB114618.1 gg_id:103415 rice plant clone ER-50 AB1146
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AJ422174.2 gg_id:23589 Germany:Spittelwasser River clone Spb283 AJ4221
DQ656489.1 gg_id:160616 Aquicola tertiaricarbonis str. L10 DQ656
FJ517015.1 gg_id:538250 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled water c FJ5170
EU881220.1 gg_id:355367 Karst region natural restoration land soil clone KNR200711-005 EU881
DQ413070.1 gg_id:152813 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
EF520457.1 gg_id:233054 structure lakes: what controls ? acid-impacted lake clone ADK-GRe02-95 EF5204
AB252908.1 gg_id:173323 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 187 AB252
DQ413123.1 gg_id:153307 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
EU528188.1 gg_id:282506 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU528
CU466856.1 gg_id:311170 Evry wastewater treatment plant anoxic basin clone CU466
NR_024710.1 gg_id:561541 Pelomonas saccharophila str. DSM 654 NR_02
DQ066963.1 gg_id:129997 Lake Washington sediment clone pLW-2 DQ066
CU466911.1 gg_id:312297 Evry wastewater treatment plant anoxic basin clone CU466
AY947999.1 gg_id:143722 Synchrony and seasonality two temperate rivers USA: Massachusetts Parker River cl AY947
AY921693.1 gg_id:113507 farm soil clone AKYG1800 AY921
FJ612162.1 gg_id:576928 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP3.4.31 FJ6121
EU638506.1 gg_id:340339 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU638
AB252904.1 gg_id:171114 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 233 AB252
AY212560.1 gg_id:102288 water 10 m downstream manure clone 109ds10 AY212
EF540425.1 gg_id:218435 oil-shale semi-coke solid waste clone MK4a EF5404
FJ484912.1 gg_id:561277 microbial autonomous rover world's deepest phreatic sinkhole water column El Zacat FJ4849
DQ327690.1 gg_id:167798 Biological filtration limits carbon availability affects downstream biofilm formation a DQ327
DQ337559.1 gg_id:145288 Pseudomonas sp. BBCT8 DQ337
AM990747.1 gg_id:278347 Leptothrix sp. MOLA 523 AM990
FJ902318.1 gg_id:541652 phreatic limestone sinkholes northeastern Mexico biomat rock outcrop cenote La Pal FJ9023
FJ902628.1 gg_id:541372 phreatic limestone sinkholes northeastern Mexico biofilm 1m deep hydrothermal ver FJ9026
AY212645.1 gg_id:107717 water 20 m upstream manure clone 192up AY212
EU244077.1 gg_id:365878 16SrRNA probes river Leine sediment clone sls1359 EU244
AY947934.1 gg_id:143876 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY947
AB240317.1 gg_id:144929 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
AF234681.1 gg_id:48162 sludge clone A2 AF234
AY622242.1 gg_id:106999 subsurface soil clone B-E7 AY622
AB015048.1 gg_id:7128 Leptothrix sp. str. MBIC3364 Y8 AB015
DQ066964.1 gg_id:130439 Lake Washington sediment clone pLW-3 DQ066
DQ017912.1 gg_id:136963 upland stream clone S-Btb7_08 DQ017
DQ241397.1 gg_id:143057 Leptothrix sp. str. S1.1 DQ241
AB240488.1 gg_id:142529 microbial structure rhizosphere biofilm Sapporo root-tip (0 40 mm) Phragmites Sosei AB240
EU340201.1 gg_id:250274 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU340
AF498703.1 gg_id:58127 soil isolate Ellin321 AF498
GQ093947.1 gg_id:404750 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093

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

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DQ854970.1 gg_id:164474 Hydrogenophaga sp. str. GPTSA14 DQ854
AF019037.1 gg_id:6953 Hydrogenophaga intermedia str. S1 AF0190
AF508103.1 gg_id:66240 Variovorax paradoxus TG27 AF5081
DQ256326.1 gg_id:141917 subsurface water clone EV818BHEB5102502DRLWq27f021 DQ256
AM777949.1 gg_id:237044 nonsaline . Influence pore-size filters determination structure alkaline groundwater AM777
DQ824747.1 gg_id:173960 human fecal clone RL185_aaj72f07 DQ824
AB195164.1 gg_id:108969 Comamonas sp. str. PD-10t AB1951
AJ853614.1 gg_id:111232 landfill leachate clone GZKB122 AJ8536
AB294326.1 gg_id:204877 Microbial structure deep coal seem groundwater stream clone SWB15 AB2943
DQ234232.2 gg_id:141836 determined library mangrove clone DS150 DQ234
AY218568.1 gg_id:81130 penguin droppings sediments clone KD2-104 AY2185
EU104128.1 gg_id:352511 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1041
EF469609.1 gg_id:224686 Pseudacidovorax intermedius str. CC-CC21 EF4696
EF580985.1 gg_id:279450 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM5(EF5809
AY823974.1 gg_id:148639 Investigation Acetate-Fed Denitrifying Microbial Stable Isotope Probing Full-Cycle an AY8239
EU037911.1 gg_id:235933 Metabolic Profiles Microbial Chlorinated Pesticides Contaminated Sites Three Geogr: EU0379
AJ585992.1 gg_id:97176 Hydrogenophaga atypica str. BSB 41.8 AJ5859
AB077037.1 gg_id:78049 Macromonas bipunctata str. IAM 14880 AB0770
FJ382319.1 gg_id:572939 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3823
AY863093.1 gg_id:112155 wastewater treatment pool clone SXAU055 AY8630
DQ337093.1 gg_id:146820 subsurface water clone EV818SWSAP75 DQ337
DQ824754.1 gg_id:184303 human fecal clone RL185_aaj72g06 DQ824
AY947966.1 gg_id:141254 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY9479
Z93993.1 gg_id:6940 activated sludge clone T60 Z93993
FJ598354.1 gg_id:552978 Study Prokaryotic Some Selected Niches Western Ghats India water clone BCWCWP1 FJ5983
AM110076.2 gg_id:250992 Hydrogenophaga sp. str. Rs71 AM110
AM042694.1 gg_id:161866 Tepidimonas thermarum str. AA2 AM042
AM777993.1 gg_id:251137 nonsaline . Influence pore-size filters determination structure alkaline groundwater AM777
AB271047.1 gg_id:171214 Hydrogenophaga sp. str. Gsoil 1545 AB2710
EF532793.1 gg_id:227420 Hydrogenophaga bisanensis str. K102 EF5327
FJ660572.1 gg_id:561367 Correlation functional stability and dynamics full-scale wastewater treatment plant a(FJ6605
GQ114408.1 gg_id:418439 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw59 GQ114
AF208386.1 gg_id:28592 Variovorax paradoxus str. WP1 AF2083
AY212671.1 gg_id:104415 water 20 m downstream manure clone 223ds20 AY2126
DQ824567.1 gg_id:191848 human fecal clone RL185_aaj69g05 DQ824
DQ349098.1 gg_id:148264 Variovorax limosa str. EMB320 DQ349
GQ008933.1 gg_id:500732 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ008
AB162104.1 gg_id:132598 trphc Dntrfng Hrbrd Sbsrfc Uplnd Sl Imprvd Cltrblt uplnd sl AB1621
AF509578.1 gg_id:91935 intestine clone AF5095

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EF157145.1 gg_id:253142 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1571
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AY947979.1 gg_id:144564 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY947
AY168733.2 gg_id:136160 Oxidizing Biofilms: Cultivation Approaches and Rates Arsenite Oxidation arsenite-oxi AY168
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GQ098862.1 gg_id:436788 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw558a GQ098
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EU801154.1 gg_id:314075 It's all ranking aquatic Delaware Bay NJ clone 2C229504 EU801
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AF502223.1 gg_id:59344 EBPR sludge lab scale clone HP1B35 AF502
EF516428.1 gg_id:220686 grassland soil clone FCPP550 EF5164
GQ009720.1 gg_id:499945 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw77 GQ009
DQ178978.1 gg_id:139068 Variovorax dokdonensis str. DS-43 DQ178
DQ432053.1 gg_id:154227 Variovorax koreensis str. GH 9-3 DQ432
AY160868.1 gg_id:95932 termite intestinal tract clone COB P5-8 AY1608
AJ420329.1 gg_id:35360 Variovorax paradoxus str. DSM 66 AJ4203
EU169184.1 gg_id:247107 Variovorax paradoxus str. B54 EU169
DQ257419.1 gg_id:141979 Variovorax paradoxus str. CAI-26 DQ257
DQ256486.1 gg_id:144358 Variovorax paradoxus Jm111 DQ256
AB269775.1 gg_id:209527 Variovorax paradoxus str. HYK0512-SK12 AB269
EF641108.1 gg_id:279410 Variovorax paradoxus str. TBEA6 EF6411
FJ214196.1 gg_id:342494 Co-speciation earthworms and their nephridial symbionts nephridia clone Ai3-13 FJ2141
FJ214173.1 gg_id:327535 Co-speciation earthworms and their nephridial symbionts nephridia clone Lf3-5 FJ2141
FJ214186.1 gg_id:315987 Verminephrobacter sp. str. At4 FJ2141
AJ543434.1 gg_id:83899 nephridia Octolasion lacteum clone Ol2-2 AJ5434

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DQ327663.1 gg_id:146840 Verminephrobacter foetidai str. EF01-2 DQ327
DQ327662.1 gg_id:149224 Verminephrobacter foetidai str. EF01-1 DQ327
NC_008786.1 gg_id:179440 Verminephrobacter eiseniae str. EF01-2 NC_00
GQ066930.1 gg_id:374294 Topographical and Temporal Human Skin Microbiome skin back clone nbw730f08c1 GQ066
FJ605423.1 gg_id:430937 Xenophilus aerolata str. PR-1 FJ6054
NR_025114.1 gg_id:571390 Xenophilus azovorans str. KF46F; DSM 13620 NR_02
EF079087.1 gg_id:247480 Ideonella sp. str. K5.1 EF079
DQ664241.1 gg_id:164847 Ideonella sp. str. IMCC1722 DQ664
AB476705.1 gg_id:547214 SSU sequences prokaryotes inside emergency Gotthard road tunnel Switzerland sulf AB476
AJ421931.2 gg_id:51718 Germany:Elbe River clone Elb197 AJ4219
EU104142.1 gg_id:336502 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
EF572731.1 gg_id:231452 Coco's Island site 23 marine clone S23_830 EF5727
CU466908.1 gg_id:356048 Evry wastewater treatment plant anoxic basin clone CU466
EU340221.1 gg_id:251319 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU340
NR_025916.1 gg_id:555018 Leptothrix discophora str. SS-1; ATCC 43182 NR_02
EU130970.1 gg_id:243909 Methylibium sp. BAC115 EU130
AM774413.1 gg_id:239835 Rivibacter subsaxonicus str. BF49 AM774
EU130974.1 gg_id:236153 Methylibium sp. BAC199 EU130
AJ422161.2 gg_id:25579 Germany:Spittelwasser River clone Spb111 AJ4221
EF516772.1 gg_id:225864 grassland soil clone FCPP576 EF5167
AM697195.1 gg_id:217417 sequencing libraries indoor dust clone BF0001C036 AM697
FM209333.1 gg_id:571602 Microbial Negev desert s sand soil clone 1250 FM209
AY382153.1 gg_id:136994 enriched nonylphenol ethoxylates activated sludge clone S1B10 AY382
AY218641.1 gg_id:81483 penguin droppings sediments clone KD4-64 AY218
EU536904.1 gg_id:287179 profile human microbiota antecubital fossa (inner elbow) skin clone nbt223f06 EU536
AM777983.1 gg_id:239229 nonsaline . Influence pore-size filters determination structure alkaline groundwater AM777
FM209332.1 gg_id:537782 Microbial Negev desert s sand soil clone 225 FM209
EU499698.1 gg_id:274988 Field-based laboratory stable isotope probing surveys identities both aerobic and an EU499
EF515904.1 gg_id:226906 grassland soil clone FCPP759 EF5159
AY771860.1 gg_id:108580 wastewater treatment pool clone SXAU009 AY7718
DQ837237.1 gg_id:248403 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_2B_11 DQ837
AB240506.1 gg_id:142476 microbial structure rhizosphere biofilm Sapporo root-base (80 160 mm) Phragmites AB240
FJ893771.1 gg_id:540262 Matriptase-deficient mice exhibit ichthyotic selective shift skin microbiota B1E; ear pu FJ8937
AY921694.1 gg_id:114102 farm soil clone AKYH1130 AY9216
GU208509.1 gg_id:580636 Study on Sediment Dongping Lake sediment clone Shf4-24 GU208
AF280853.1 gg_id:24241 bioreactor clone mle1-19 AF2808
AY218657.1 gg_id:103477 penguin droppings sediments clone KD9-144 AY218
EU907887.1 gg_id:309250 Identification hexahydro-135-trinitro-135-triazine (RDX)-degrading microorganisms v EU9078
EU362134.1 gg_id:305378 and Mesquite Flats Site Death Valley dune sand clone 5 EU362
EU491350.1 gg_id:259064 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X3 clone EU491

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s__Herbaspirillum rubrisubalbicans	14767
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s__Herbaspirillum seropedicae	14340

EF580947.1 gg_id:279576 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM7 EF5809
AM412128.1 gg_id:239980 Aquabacterium sp. str. P-130 AM412
AY212562.1 gg_id:102602 water 10 m downstream manure clone 110ds10 AY2125
AM697083.1 gg_id:226216 sequencing libraries indoor dust clone BF0002B033 AM697
EF999406.1 gg_id:246240 Vertical and Methanotrophs Estuarine Sediments Pearl River Estuary sediments 50cn EF9994
AY218638.1 gg_id:81258 penguin droppings sediments clone KD4-58 AY2186
GQ043432.1 gg_id:389491 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw96 GQ043
DQ676448.1 gg_id:221331 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
EU283391.1 gg_id:236412 consecutively aerated submerged treating domestic wastewater activated sludge m EU2833
AB252909.1 gg_id:164634 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 181 AB2529
FJ478501.1 gg_id:554081 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4785
EU639232.1 gg_id:317724 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU6392
DQ298309.1 gg_id:153016 studies on and nifH Lasiurus sindicus Sewan grass Thar Desert Rajasthan India rhizos DQ298
FM872725.1 gg_id:366825 occupant source house floor dust clone FB01B01 FM872
EF515947.1 gg_id:223606 grassland soil clone FCPT445 EF5159
EF522266.1 gg_id:217402 Endolithic Microbial Ecosystems Rocky Mountain endolithic sandstone clone FQSS04 EF5222
EU881306.1 gg_id:304030 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
FJ478963.1 gg_id:511130 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4789
DQ640690.1 gg_id:234716 full-scale EBPR sludge clone Skagenf14 DQ640
EF516703.1 gg_id:226711 grassland soil clone FCPT716 EF5167
FJ592525.1 gg_id:547327 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
EF020021.1 gg_id:214611 aspen rhizosphere clone Elev_16S_1414 EF0200
EF580981.1 gg_id:278694 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1 EF5809
DQ676337.1 gg_id:226112 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
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AY281140.1 gg_id:100480 Collimonas fungivorans str. CTO 166 AY2811
AY281150.1 gg_id:105543 Collimonas fungivorans str. NCCB 100029 AY2811
AJ496445.1 gg_id:57762 Collimonas fungivorans str. CTE227 = NCCB100027 AJ4964
GU441679.1 gg_id:514468 Collimonas sp. str. Cr9-30 GU441
GU213333.1 gg_id:560017 Herminiimonas sp. str. 4-96 GU213
AJ238357.1 gg_id:7300 Herbaspirillum frisingense str. 75B AJ2383
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AB094401.1 gg_id:69647 Herbaspirillum chlorophenolicum str. CPW301 AB0944
EU549858.1 gg_id:302270 Herbaspirillum frisingense str. AU14559 EU5498
EU399918.1 gg_id:275308 Herbaspirillum frisingense str. CAF265 EU3999
AB021424.1 gg_id:7295 Herbaspirillum rubrisubalbicans str. ATCC 19308T AB0214
AJ238356.1 gg_id:7299 Herbaspirillum rubrisubalbicans str. M4 AJ2383
AY191273.1 gg_id:80182 Herbaspirillum seropedicae str. Z152 AY1912

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AY191272.1 gg_id:80181 Herbaspirillum seropedicae str. ZA69 AY1912
AF361023.1 gg_id:74451 Herbaspirillum sp. str. NAH4 AF3610
AJ505863.1 gg_id:58574 Herbaspirillum sp. str. PIV-34-1 = DSM 14683 AJ5058
AY043371.1 gg_id:63615 Pseudomonas huttiensis str. AU2210A AY0433
GU433469.1 gg_id:528777 Herbaspirillum huttiense str. CCBAU 61429 GU433
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AJ583172.1 gg_id:94668 radioactive site ground water clone S15B-MN104 AJ5831
AY049039.1 gg_id:47876 Pseudomonas huttiensis str. D AY0490
AJ238359.1 gg_id:7302 Herbaspirillum frisingense str. Mb11 AJ2383
AB021366.1 gg_id:7292 Pseudomonas huttiensis str. ATCC 14670T AB0213
FJ266340.1 gg_id:534872 Herbaspirillum rubrisubalbicans str. RFNB30 FJ2663
FJ605419.1 gg_id:404943 Herbaspirillum rubrisubalbicans str. PPs-3 FJ6054
NR_028656.1 gg_id:584170 Herbaspirillum putei str. 7-2; IAM 15032 NR_028
AY191276.2 gg_id:80103 Herbaspirillum seropedicae str. M2 AY1912
AB512142.1 gg_id:564904 Herminiimonas fonticola str. da Costa S-94 (=CIP 108398 =LMG AB5121
AB512141.1 gg_id:546345 Herminiimonas saxobsidens str. NEU 1218 (=DSM 18748 =CCM 7436) AB5121
EU133873.1 gg_id:251868 tallgrass prarie soil clone FFCH17907 EU1338
GU244359.1 gg_id:593835 Janthinobacterium sp. str. Lc10-8 GU244
EU847234.1 gg_id:353532 Janthinobacterium sp. str. blxAG EU8472
FJ894020.1 gg_id:555629 Matriptase-deficient mice exhibit ichthyotic selective shift skin microbiota B1E; ear pu FJ8940
FJ193906.1 gg_id:313175 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1939
EU536428.1 gg_id:285908 profile human microbiota antecubital fossa (inner elbow) skin clone nbt213f12 EU5364
AF529096.1 gg_id:62419 trichloroethene-contaminated site clone ftl206 AF5290
EU982453.1 gg_id:341939 Enrichment and detection moderate thermophilic microorganisms vents Lau Basin h EU9824
AY792243.1 gg_id:119507 Humic Lake clone CrystalBog5B11 AY7922
FJ267554.1 gg_id:308624 Massilia sp. str. II_Gauze_W_12_10 FJ2675
DQ228653.1 gg_id:142568 Microbial colonizing mineral surfaces within sulfide-microbial incubator clone CH2b1 DQ228
FM209304.1 gg_id:571521 Microbial Negev desert s sand soil clone 2321 FM209
AM412135.1 gg_id:241097 Massilia sp. str. P-118 AM412
GQ354568.1 gg_id:556334 Naxibacter sp. str. HMD2167 GQ354
EU221406.1 gg_id:235543 Massilia timonae str. H2P8 EU2214
GQ015305.1 gg_id:494362 Topographical and Temporal Human Skin Microbiome skin palm clone nbw766g11c1 GQ015
FM209303.1 gg_id:536558 Microbial Negev desert s sand soil clone 2220 FM209
GQ098872.1 gg_id:406246 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw558b GQ098
AB487721.1 gg_id:539718 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4877
AY792245.1 gg_id:126195 Humic Lake clone CrystalBog2G3 AY7922
EU431716.1 gg_id:562896 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU4317
EU244075.1 gg_id:365805 16SrRNA probes river Leine sediment clone sl1354 EU2440

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FJ191812.1 gg_id:323439 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1918
AB488284.1 gg_id:569272 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4882
EU626407.1 gg_id:279407 Massilia sp. str. biosZ4 EU6264
EU373360.1 gg_id:270893 Massilia timonae str. HNL19 EU3733
FJ605429.1 gg_id:456558 Janthinobacterium sp. str. PR-2 FJ6054
FJ191801.1 gg_id:331807 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1918
EU365682.1 gg_id:269884 Massilia timonae str. M-10 EU3656
EU160504.1 gg_id:318413 diagnosis central-venous infections cultivation-dependent and techniques biofilm sc EU1605
EF682966.1 gg_id:246159 Eastern Mediterranean atmosphere clone F27_2E_FL EF6829
GQ021547.1 gg_id:488122 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbu27 GQ021
AB487246.1 gg_id:558053 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB4872
FJ375473.1 gg_id:510182 T-RFLP powerful tool rapid assessment cells microbial fuel cell sheet carbon paper an FJ3754
AB234525.1 gg_id:145411 termite gut homogenate clone MgMjD-103 AB2345
EU274637.2 gg_id:259460 Massilia timonae str. U2 EU2746
AY966001.1 gg_id:115346 Sinobacter luteus str. 101 AY9660
AB487486.1 gg_id:567594 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4874
EU431713.1 gg_id:540535 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU4317
EF516951.1 gg_id:223889 grassland soil clone FCPU439 EF5169
AJ871463.1 gg_id:110987 Massilia timonae CP183-9 AJ8714
GU377105.1 gg_id:518781 Janthinobacterium sp. str. sptzw14 GU377
AF443584.1 gg_id:88301 semiarid soil clone C-F-25 AF4435
AM412137.1 gg_id:236550 Massilia sp. str. P-127 AM412
AF408326.1 gg_id:34181 Massilia sp. clone KL-11-1-5 AF4083
AM412141.1 gg_id:235451 Oxalobacter sp. str. P-154 AM412
AY512824.1 gg_id:106310 Massilia timonae str. S11 AY5128
EU554441.1 gg_id:275614 Naxibacter haematophilus str. Ag15 EU5544
AY679161.1 gg_id:102382 Naxibacter alkalitolerans str. YIM 31775 AY6791
AB097172.1 gg_id:87579 Janthinobacterium sp. str. J3 AB0971
EU369116.1 gg_id:551576 Janthinobacterium sp. str. AR12 EU3691
AY599723.1 gg_id:102086 fungal ascocarp clone Cl-73-TB2-II AY5997
EU861944.1 gg_id:344291 effects chronic fertilization on alpine tundra microbial implications carbon and cyclin EU8619
DQ537532.1 gg_id:164523 manganese oxidation soil clone JH-GY25 DQ537
AJ582187.1 gg_id:93999 soil near uranium mill tailings clone Gitt-GS-70 AJ5821
AM942544.1 gg_id:517426 Duganella sp. str. dug2 CMI-17 AM942
EU431682.1 gg_id:557117 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU4316
EF575562.1 gg_id:234460 Duganella sp. str. BD-a14 EF5755
GQ339887.1 gg_id:580212 Massilia sp. str. MK02 GQ339
U49752.1 gg_id:70632 Oxalobacter formigenes str. OXWR wild rat cecum U49752
NR_029188.1 gg_id:571425 Oxalobacter formigenes str. OXB; ATCC 35274 NR_029

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U49751.1 gg_id:7367 Oxalobacter formigenes str. SOX-4 freshwater lake sediment U4975
U49749.2 gg_id:7365 Oxalobacter formigenes str. BA4 human feces U4974
U49758.1 gg_id:7372 Oxalobacter formigenes str. HC1 human feces U4975
U49756.1 gg_id:7371 Oxalobacter formigenes str. OXK human feces U4975
FJ264587.1 gg_id:331690 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedim FJ2645
AB488167.1 gg_id:559492 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR AB488
U49754.1 gg_id:7369 Oxalobacter formigenes str. OXCR laboratory rat cecum U4975
AB486719.1 gg_id:591499 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR AB486
FJ374254.1 gg_id:363874 Inter- and intraspecific humivorous scarab beetle larvae (Pachnoda spp.) larval hindg FJ3742
EF454731.2 gg_id:217779 termite hindgut clone 290cost002-P3L-2608 EF4547
EF434363.1 gg_id:216215 human infant gastrointestinal tract microbiota stool mother babies 13 and 14 7 mont EF4343
AB288876.1 gg_id:202850 termite gut homogenate clone BOf1-03 AB288
DQ796767.1 gg_id:174649 human fecal clone RL179_aao54b12 DQ796
FJ269064.1 gg_id:340557 dissimilatory ferric iron reducers arsenic contaminated Hunan south China paddy soil FJ2690
DQ676429.1 gg_id:224948 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
AB038368.1 gg_id:19906 Diaphorina citri symbiont AB038
FJ371291.1 gg_id:360508 feces adult twins and mothers; TS6 clone TS6_a03c02 FJ3712
AB234524.1 gg_id:145512 termite gut homogenate clone MgMjD-092 AB234
AF371862.1 gg_id:50320 swine intestine clone p-2518-18B5 AF371
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GQ069768.1 gg_id:401837 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ069
DQ248254.1 gg_id:141616 carbon tetrachloride contaminated soil clone UE7 DQ248
AY509446.2 gg_id:99154 freshwater clone LiUU-5-340 AY509
DQ351915.1 gg_id:172665 manganese oxidation soil clone JH-WH04 DQ351
GU213338.1 gg_id:593816 Janthinobacterium sp. str. 338 GU213
AM292626.1 gg_id:173162 Influence uranyl and sodium nitrate on natural soil uranium mining waste pile clone AM292
DQ537535.1 gg_id:162048 manganese oxidation soil clone JH-GY37 DQ537
AY167838.1 gg_id:101564 Janthinobacterium agaricidamnosum str. SAFR-022 AY167
AB240320.1 gg_id:144767 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
AB487490.1 gg_id:569658 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR AB487
EU801571.1 gg_id:317157 It's all ranking aquatic Chesapeake Bay MD clone 3C002897 EU801
AM888196.1 gg_id:247545 Herbaspirillum sp. str. THWCSN35 AM888
EF516890.1 gg_id:226310 grassland soil clone FCPU733 EF5168
DQ906892.1 gg_id:256121 Oman subsurface soil clone 12A-1 DQ906
AY080915.1 gg_id:61192 acidic forest soil clone UP9 AY0809
AM991238.1 gg_id:279160 Microbial groundwater and their potential use biomonitoring karst spring water clo AM991
AF468345.1 gg_id:79836 Arctic sea ice ARK10157 AF4683

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AM397629.1 gg_id:169981 Undibacterium sp. str. CCUG 49012 AM397
EU802136.1 gg_id:335152 It's all ranking aquatic Chesapeake Bay MD clone 3C003549 EU802
EF516112.1 gg_id:225949 grassland soil clone FCPT483 EF5161
FM874576.1 gg_id:357346 occupant source house mattress dust clone MD03G10 FM874
DQ351928.1 gg_id:171310 manganese oxidation soil clone JH-GY06 DQ351
EF516828.1 gg_id:225672 grassland soil clone FCPP661 EF5168
EF516978.1 gg_id:222289 grassland soil clone FCPU492 EF5169
EF580950.1 gg_id:278562 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM8 EF5809
AY082467.1 gg_id:67839 neutral pH mine biofilm clone 44a-B1-39 AY0824
FM209308.1 gg_id:566578 Microbial Negev desert s sand soil clone 3303 FM209
AM697265.1 gg_id:222537 sequencing libraries indoor dust clone BF0001C106 AM697
GU321360.1 gg_id:561332 Massilia sp. str. NMC24 GU321
GU295961.1 gg_id:585352 Oxalicibacterium sp. str. JCN-21 GU295
AB487226.1 gg_id:562136 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB487
DQ787695.1 gg_id:166885 holocene lake sediment clone SLB419 DQ787
DQ241394.1 gg_id:144017 simple biogeochemical process removing arsenic mine drainage water clone LOP-89 DQ241
FM872500.1 gg_id:362441 occupant source house floor dust clone FA01C01 FM872
AJ863414.1 gg_id:121741 poplar tree microcosm rhizosphere flooded clone 5RHF33 AJ8634
AB487777.1 gg_id:579702 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB487
GU377118.1 gg_id:532409 Herbaspirillum sp. str. sptzw31 GU377
DQ264588.1 gg_id:142425 universal microarray broader than typical library when sampling subsurface groundv DQ264
DQ490985.1 gg_id:157892 Pseudoburkholderia malthae str. CC-AFH3 DQ490
FM955855.1 gg_id:590236 Massilia sp. str. Asd M1A2 FM955
EU735672.1 gg_id:345327 archaeal structures oil contaminated and soils China pristine soil Jidong Oilfield clone EU735
EU636042.1 gg_id:279632 Antarctic str. N14 EU636
EU362137.1 gg_id:352209 and Mesquite Flats Site Death Valley dune sand clone 8 EU362
AY198108.1 gg_id:80296 melt pond on Arctic sea ice floe clone ARKMP92 AY198
FJ191604.1 gg_id:340957 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1916
AY367012.1 gg_id:93490 ultramicrobacterium str. Um1 AY367
EF682968.1 gg_id:255137 Eastern Mediterranean atmosphere clone F13_2C_FL EF6829
AM936250.1 gg_id:567104 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
GU124500.2 gg_id:570154 endosymbiotic (Stal) (rice brown planthopper) Endosymbiont Nilaparvata lugens clo GU124
EF552158.1 gg_id:233331 Spirillum sp. str. NOX EF5521
AJ556800.1 gg_id:85781 Oxalobacteraceae MWH73 AJ5568
AB486782.1 gg_id:592565 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR AB486
FM253595.1 gg_id:341548 Rock gold mine unusually microbial biofilm clone D06-1 FM253
AB488125.1 gg_id:548180 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB488
GQ098449.1 gg_id:401707 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw559g GQ098
GQ461666.1 gg_id:533088 Coupling -ITS Clone Libraries and Automated Intergenic Spacer Microbial Bioremedi GQ461
DQ787709.1 gg_id:163192 holocene lake sediment clone SLB530 DQ787

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DQ248255.1 gg_id:142344 carbon tetrachloride contaminated soil clone UA6 DQ248
AF358019.1 gg_id:26437 peat soil microcosm clone LO13.5 AF358
DQ469209.1 gg_id:157564 Microbial and ecology water Zhong Tiaoshan mine Shanxi province China AMD clone DQ469
AF529095.1 gg_id:62422 trichloroethene-contaminated site clone FTL205 AF529
EF620474.1 gg_id:248260 ultramicrobacterium str. ISSDS-831 EF6204
NR_028859.1 gg_id:566047 Herbaspirillum lusitanum str. P6-12; LMG 21710 NR_02
DQ017931.1 gg_id:133568 upland stream clone S-Jos_62 DQ017
DQ787707.1 gg_id:171969 holocene lake sediment clone SLB523 DQ787
EU676429.1 gg_id:287560 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals- EU676
EF592558.1 gg_id:232495 Duganella sp. str. B2 EF5925
AJ582194.1 gg_id:94238 soil near uranium mill tailings clone Gitt-GS-138 AJ5821
DQ177470.1 gg_id:140398 Janthinobacterium sp. str. Tibet-IIK43 DQ177
EU528198.1 gg_id:286535 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU528
DQ341421.1 gg_id:150786 Janthinobacterium sp. str. Antarctic IS09 DQ341
AB008503.1 gg_id:7288 ultramicrobacterium str. MY14 AB008
EF682967.1 gg_id:243222 Eastern Mediterranean atmosphere clone F14_4D_FL EF6829
EU669620.1 gg_id:329777 soil clone S22-8 EU669
DQ158104.1 gg_id:139019 Use Field-Based Stable Isotope Probing Identify Adapted and Track Carbon F throug DQ158
EU636048.1 gg_id:278716 Antarctic str. GAo51 EU636
EF019643.1 gg_id:215915 aspen rhizosphere clone Elev_16S_794 EF0196
EU489741.1 gg_id:277881 Herminiimonas sp. str. UMB49 EU489
EU636041.1 gg_id:278413 Antarctic str. GA091 EU636
AF407411.1 gg_id:21324 MCB-contaminated groundwater-treating reactor clone RB13C5 AF4074
AY214204.1 gg_id:81252 benzene-contaminated groundwater clone ZZ15C3 AY2142
EU057874.1 gg_id:253699 Duganella sp. 5B EU057
AB488131.1 gg_id:589716 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR AB488
EF515920.1 gg_id:218687 grassland soil clone FCPT636 EF5159
AY218774.1 gg_id:81640 penguin droppings sediments clone KD6-99 AY2187
DQ188985.1 gg_id:140045 Herbaspirillum rhizosphaerae str. UMS-40 DQ188
FJ391493.1 gg_id:364438 primers targeting Gallionella- neutrophilic oxidizers enrichments and soils iron rich irr FJ3914
GU205377.1 gg_id:583924 Massilia sp. str. DR-a8 GU205
AY214210.1 gg_id:81997 benzene-contaminated groundwater clone ZZ17C9 AY2142
AF468324.1 gg_id:80693 melt pond on Arctic sea ice floe clone ARKMP-20 AF4683
AJ867677.1 gg_id:109615 melted red snow water clone RS 8-Bact69 AJ8676
AJ295498.1 gg_id:27239 rape rhizosphere clone wr0040 AJ2954
EU194890.1 gg_id:359091 Duganella sp. str. B41 EU1948
AM085762.1 gg_id:138928 Herminiimonas aquatilis str. CCUG 36956 AM085
GU377114.1 gg_id:525330 Janthinobacterium sp. str. sptzw26 GU377
GQ451298.1 gg_id:585579 Fecal Yunnan snub-nosed monkey feces clone J308 GQ451
AB487141.1 gg_id:588705 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR AB487

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AJ867670.1 gg_id:111083 melted red snow water clone RS 8-Bact32 AJ8676
AB487627.1 gg_id:544493 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4876
AB487567.1 gg_id:588968 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4875
GU213421.1 gg_id:589183 Janthinobacterium sp. str. 286 GU213
EU150263.1 gg_id:278184 acidobacteria and soil Niwot Ridge LTER clone Ni_E6P2 EU150
AY198109.1 gg_id:79646 melt pond on Arctic sea ice floe clone ARKMP91 AY1981
EF018240.1 gg_id:207516 aspen rhizosphere clone Amb_16S_591 EF0182
EF019746.1 gg_id:216361 aspen rhizosphere clone Elev_16S_922 EF0197
AF468329.1 gg_id:80338 melt pond on Arctic sea ice floe clone ARKMP-56 AF4683
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EU636047.1 gg_id:278100 Antarctic str. GA045 EU636
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DQ264563.1 gg_id:141833 universal microarray broader than typical library when sampling subsurface groundv DQ264
EF574525.1 gg_id:232833 site S25 near Coco's Island marine clone S25_869 EF5745
AB487774.1 gg_id:583648 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4877
AY061962.2 gg_id:202769 Oxalicibacterium flavum str. TA17; NEU98; LMG21571 AY0619
EU881344.1 gg_id:329302 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
AB488296.1 gg_id:587787 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4882
DQ787694.1 gg_id:162906 holocene lake sediment clone SLB418 DQ787
EU160577.1 gg_id:310584 diagnosis central-venous infections cultivation-dependent and techniques biofilm sc EU1605
AB488003.1 gg_id:580561 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4880
AM697274.1 gg_id:222450 sequencing libraries indoor dust clone BF0001C115 AM697
AB074523.1 gg_id:66301 Aquaspirillum arcticum str. IAM 14963 AB0745
AJ867657.1 gg_id:110953 melted red snow water clone RS 8-Bact21 AJ8676
AJ532689.1 gg_id:76765 uranium mill tailings clone Gitt-KF-179 AJ5326
GQ080374.1 gg_id:373791 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ080
AB074524.1 gg_id:56608 Aquaspirillum autotrophicum str. IAM 14942 AB0745
AY792246.1 gg_id:122054 Humic Lake clone CrystalBog5F8 AY7922
FJ517670.1 gg_id:568152 Disturbing epithelial homeostasis metazoan Hydra leads drastic changes microbiota e FJ5176
GU213418.1 gg_id:572788 Janthinobacterium sp. str. 278 GU213
AB487887.1 gg_id:587586 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4878
GU113050.1 gg_id:583060 soil overlying mountain Manasi County Xinjiang soils influenced underground coal m GU113
AY515386.1 gg_id:111167 sediment Lake Tuusulanjarvi isolate str. F39 AY5153
AM849431.1 gg_id:236633 Seasonality Piburger See clone PIB-20 AM849
FJ535228.1 gg_id:572527 Contamination risk vegetables Arcobacter during post-harvest processing water carr FJ5352
AY227003.1 gg_id:81741 melt pool on Arctic sea ice floe clone ARKMP-94 proteobacterium AY2270
EF094990.1 gg_id:194580 Rubrivivax gelatinosus str. L31 EF0949
CU466884.1 gg_id:313285 Evry wastewater treatment plant anoxic basin clone CU466

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GQ396839.1 gg_id:587796 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
CU466928.1 gg_id:325979 Evry wastewater treatment plant anoxic basin clone CU466
GU454909.1 gg_id:528368 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
CU466864.1 gg_id:327001 Evry wastewater treatment plant anoxic basin clone CU466
CU466850.1 gg_id:338411 Evry wastewater treatment plant anoxic basin clone CU466
GQ249369.1 gg_id:553551 Research on biodiversity sludge bioflim Biological Aeration Filter clone B11 GQ249
GQ302537.1 gg_id:585468 prokaryotic sediment Shawan methods cold spring clone sw-xj209 GQ302
AB087565.1 gg_id:80525 Sphaerotilus sp. L8 AB087
AB087561.1 gg_id:80068 Sphaerotilus sp. L2 AB087
Z18534.1 gg_id:7139 Sphaerotilus natans str. 565 Z18534
AF309815.1 gg_id:73442 synthetic wastewater tmbr15-22 complete clone wastewater AF3098
AF177943.1 gg_id:42541 Tepidimonas ignava str. SPS-1037T AF177
EF648100.1 gg_id:255518 dynamics during produced water treatment aerobic activated sludge clone HB116 EF6481
AY845054.1 gg_id:110733 Tepidimonas sp. str. I1-1 AY845
AY594193.1 gg_id:124693 Tepidimonas arfidensis AY594
EU799903.1 gg_id:338160 It's all ranking aquatic Newport Harbour RI clone 1C227573 EU799
AY519354.1 gg_id:107473 Thiomonas sp. str. 51 AY519
FM992405.1 gg_id:536889 Thiomonas intermedia FM992
AY455808.1 gg_id:103970 Thiomonas perometabolis str. ATCC 23370 AY455
DQ146028.1 gg_id:138509 Thiomonas sp. ML1-60 DQ146
DQ146101.1 gg_id:138673 Thiomonas sp. ML2-48 DQ146
DQ146068.1 gg_id:139031 Thiomonas sp. ML2-1 DQ146
AB462557.1 gg_id:321818 Microbial production and energy source hyperthermophilic prokaryotes geothermal AB462
DQ146030.1 gg_id:138725 Thiomonas sp. ML1-62 DQ146
DQ146119.1 gg_id:138822 Thiomonas sp. ML2-67 DQ146
DQ146041.1 gg_id:138202 Thiomonas sp. ML1-74 DQ146
AB308270.1 gg_id:253106 Thiomonas sp. str. RAN4 AB308
DQ146097.1 gg_id:138841 Thiomonas sp. ML2-42 DQ146
DQ146015.1 gg_id:138792 Thiomonas sp. ML1-48 DQ146
DQ902578.1 gg_id:189483 Thiomonas sp. str. CS2 DQ902
DQ146103.1 gg_id:138853 Thiomonas sp. ML2-5 DQ146
AJ549220.1 gg_id:82391 Thiomonas sp. B3 AJ5492
DQ146013.1 gg_id:138541 Thiomonas sp. ML1-46 DQ146
DQ145996.1 gg_id:138828 Thiomonas sp. ML1-28 DQ145
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DQ092334.1 gg_id:181396 Thiomonas sp. str. S10 DQ092

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GU233443.1 gg_id:531748 Thiomonas thermosulfata str. 21071 32621 GU233
EU937876.1 gg_id:351331 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
AY050601.1 gg_id:53885 MCB-contaminated groundwater-treating reactor clone GOUTB18 AY050
GU208450.1 gg_id:536615 Study on Sediment Dongping Lake sediment clone Fr2-13 GU208
EU937989.1 gg_id:307219 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
EF520449.1 gg_id:229543 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-80 EF5204
GQ388859.1 gg_id:562874 quality deterioration drinking system during serious red water outbreak clone Z238 GQ388
GQ390167.1 gg_id:574746 oxic-anoxic interface meromictic crater lake Lac Pavin clone eub62A12 GQ390
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FJ391492.1 gg_id:367650 primers targeting Gallionella- neutrophilic oxidizers enrichments and soils iron rich irr FJ3914
AB475012.1 gg_id:540849 sequences located Engiadina Bassa Valley Switzerland iron containing freshwater so AB475
EU937953.1 gg_id:356589 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
FM204958.1 gg_id:576266 Prokaryote shifts and reductive dehalogenase redundancy response electron donor FM204
GU208476.1 gg_id:592888 Study on Sediment Dongping Lake sediment clone Shc3-16 GU208
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GQ264147.1 gg_id:589210 Structure Simulated Level Waste Site simulated level waste site clone WC3_88 GQ264
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GQ388776.1 gg_id:576649 quality deterioration drinking system during serious red water outbreak clone Z10 GQ388
EU937859.1 gg_id:327025 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
AY922005.1 gg_id:112903 farm soil clone AKYG404 AY922
GQ388790.1 gg_id:593879 quality deterioration drinking system during serious red water outbreak clone Z114 GQ388
AY766002.1 gg_id:108663 streamer acidic iron-rich spa water clone TrefC4 AY766
FJ037638.1 gg_id:310147 biofilms growing tropical warm soda spring Grenada West Indies clone 070218-Grena FJ0376
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FM174362.1 gg_id:541762 biotrickling filter treating loads H2S biology tools clone H2SRC133X FM174
GQ402778.1 gg_id:534649 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
EF562088.1 gg_id:241212 Subsurface microbial deep-granitic-fracture water colorado Henderson Molybdenum EF562
GQ339166.1 gg_id:520139 Comprehensive Investigation on Iron Cycling Freshwater Seep ing Microscopy Cultiv GQ339
EU283473.1 gg_id:278266 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A531 EU283
GU183583.1 gg_id:551555 Structures Reduced and Oxidized Aquifer lower Aqueous Arsenic Titas Bangladesh a GU183
DQ676395.1 gg_id:225924 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676

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

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

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

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

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DQ378255.1 gg_id:155163 oil-polluted soil clone M38_Pitesti DQ378
AY212692.1 gg_id:105727 water 10 m downstream manure clone 244ds10 AY2126
EF580978.1 gg_id:278346 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1: EF5809
EF019146.1 gg_id:206079 aspen rhizosphere clone Amb_16S_1787 EF0191
EU703371.1 gg_id:396964 lakes on Eastern Tin Plateau freshwater lake altitude 4298 masl clone XZTSH19 EU7033
AB464941.2 gg_id:343674 Altitudinal changes flora on Gulkana Glacier Alaska analyzed sequencing and -cycle c AB4649
AY486132.1 gg_id:103599 Methylovorus mays str. C AY4861
EU268114.1 gg_id:250802 summer Dalian and Liaodong bay determined sea water clone ZL1702 EU2681
EU799457.1 gg_id:330088 It's all ranking aquatic Newport Harbour RI clone 1C227059 EU7994
EU287055.1 gg_id:235281 Pacific arctic surface sediment clone B78-91 EU2870
EU591642.1 gg_id:264560 Microbial Ecology Focused-Pulsed Electric Field Sludge Pretreatment Methanogenic : EU5916
EU801464.1 gg_id:308817 It's all ranking aquatic Chesapeake Bay MD clone 3C002760 EU8014
AY238979.1 gg_id:105790 sludge dentrifying reactor fed with methanol clone DSBR-C043 AY2389
EU804023.1 gg_id:338998 It's all ranking aquatic Lake Gatun clone 5C231706 EU8040
EU236355.1 gg_id:289508 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg91B9 EU2363
DQ513082.1 gg_id:161213 Microbial life fluids ridge flank crustal fluid clone FS266-46B-03 DQ513
GQ350653.1 gg_id:543211 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
DQ066956.1 gg_id:135981 Lake Washington sediment clone pLW-13 DQ066
AJ400352.1 gg_id:29664 marine clone ZD0412 AJ4003
EF492937.1 gg_id:240233 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4929
AY678517.1 gg_id:103273 estuarine sediment clone C319a-R8C-D8 AY6785
EU801551.1 gg_id:329876 It's all ranking aquatic Chesapeake Bay MD clone 3C002872 EU8015
AY360530.1 gg_id:137280 Stable isotope probing dynamic methylotroph trophic interactions fungi and protozo AY3605
FJ612340.1 gg_id:548303 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.2.8 FJ6123
AY360525.1 gg_id:137243 Stable isotope probing dynamic methylotroph trophic interactions fungi and protozo AY3605
GQ340339.1 gg_id:568552 Prokaryotic Marathonas Reservoir Greece water column clone VE08-118-BAC GQ340
DQ810320.1 gg_id:170033 Oxygen Minimum Zone Surface- Lower Waters and Other Marine oxygen minimum : DQ810
AY546509.1 gg_id:101169 soda lake sediment clone 5H_52 AY5465
EU804036.1 gg_id:306270 It's all ranking aquatic Lake Gatun clone 5C231720 EU8040
AB294945.1 gg_id:219027 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma AB2949
EU800826.1 gg_id:328759 It's all ranking aquatic Delaware Bay NJ clone 2C229045 EU8008
EU037912.1 gg_id:247600 Metabolic Profiles Microbial Chlorinated Pesticides Contaminated Sites Three Geogr: EU0379
AY360533.1 gg_id:137226 Stable isotope probing dynamic methylotroph trophic interactions fungi and protozo AY3605
EU800277.1 gg_id:338875 It's all ranking aquatic Delaware Bay NJ clone 2C228317 EU8002
GQ350509.1 gg_id:580430 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
GQ472336.1 gg_id:560126 Microbial Dongping Lake lake sediment clone 2S4-17 GQ472
GU230465.1 gg_id:587327 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
AY792265.1 gg_id:125742 Humic Lake clone CrystalBog021G5 AY7922
EU287027.1 gg_id:255894 Pacific arctic surface sediment clone B78-63 EU2870
EF173353.1 gg_id:177340 Options situ remediation contaminated mixture chlorinated hydrocarbons hydrocarb EF1733
FJ612406.1 gg_id:584989 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.4.19 FJ6124

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g__Eikenella

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

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NZ_ACJW02000005.1 gg_id:470523 Kingella oralis str. ATCC 51147 NZ_AC
AY644511.1 gg_id:107215 Kingella sp. M03-232 AY6445
AY349388.1 gg_id:92601 human oral cavity clone JC012 AY3493
DQ003616.1 gg_id:125235 Kingella genomosp. P1 oral cone MB2_C20 DQ003
GU227231.1 gg_id:536311 Microbiological immunological Adolescent Moroccan Aggressive Periodontitis Patier GU227
EU118276.2 gg_id:248652 Laribacter hongkongensis str. LHHZ69f EU118
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DQ201330.1 gg_id:140124 Neisseria meningitidis str. CMCC(B)29034 DQ201
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AF382295.1 gg_id:65965 Neisseria meningitidis str. M7905 AF3822
AY132143.1 gg_id:59582 Neisseria meningitidis str. M8585 AY1321
AY132136.1 gg_id:59586 Neisseria meningitidis str. M8579 AY1321
AY132142.1 gg_id:59580 Neisseria meningitidis str. M8586 AY1321
AY146975.1 gg_id:76992 Neisseria meningitidis str. N.221/91 AY1469

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AF310540.1 gg_id:18187 Neisseria meningitidis str. M6199 AF3105
AJ239298.1 gg_id:6444 Neisseria polysaccharea str. P8 AJ2392
AJ239315.1 gg_id:6455 Neisseria polysaccharea str. P7 AJ2393
AJ239308.1 gg_id:6450 Neisseria meningitidis str. HF116 AJ2393
AJ239312.1 gg_id:6471 Neisseria meningitidis str. S3446 AJ2393
AJ239281.1 gg_id:6437 Neisseria pharyngis str. NCTC4590 AJ2392
AJ239282.1 gg_id:6401 Neisseria mucosa str. LNP405 AJ2392
AJ239294.1 gg_id:6443 Neisseria sicca str. Q29 AJ2392
AJ239301.1 gg_id:6447 Neisseria flava str. U40 AJ2393
AF479577.1 gg_id:53760 Neisseria subflava str. NJ9702 AF4795
EF511930.1 gg_id:217706 Loss During Antibiotic Treatment 1 Intubated Patients Colonized Pseudomonas aerug EF5119
AJ239291.1 gg_id:6429 Neisseria subflava str. U37 AJ2392
FJ823154.1 gg_id:569075 Neisseria mucosa str. TeTa FJ8231
AF479578.1 gg_id:16823 Neisseria subflava str. NJ9703 AF4795
L10738.1 gg_id:6396 Neisseria weaveri L10738
AF395911.1 gg_id:51956 Neisseria weaveri VA6362grgr_2001 AF3959
NZ_ABZO01000111.1 gg_id:440950 Neisseria gonorrhoeae str. PID332 NZ_AB
AF398310.1 gg_id:62766 Neisseria meningitidis str. M7724 AF3983
FJ191618.1 gg_id:354448 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1916
FM874743.1 gg_id:363843 occupant source house dust skin surface swab clone SA02H07 FM874
EF511172.1 gg_id:218785 Loss During Antibiotic Treatment 1 Intubated Patients Colonized Pseudomonas aerug EF5111
AF398299.1 gg_id:62778 Neisseria meningitidis str. M7744 AF3982
AF310565.1 gg_id:32023 Neisseria meningitidis str. M2786 AF3105
GU233442.1 gg_id:517381 Neisseria iguanae str. NVSL 85737 GU233
EF511120.1 gg_id:221884 Loss During Antibiotic Treatment 1 Intubated Patients Colonized Pseudomonas aerug EF5111
DQ201335.1 gg_id:139970 Neisseria meningitidis str. CMCC(B)29201 DQ201
AJ239299.1 gg_id:6445 Neisseria cinerea str. F1 AJ2392
FJ192856.1 gg_id:338605 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1928
FJ191574.1 gg_id:335421 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1915
AY648028.1 gg_id:104161 Neisseria meningitidis str. NZ90/21 AY648
AY648033.1 gg_id:99467 Neisseria meningitidis str. NZ00/189 AY648
FJ192099.1 gg_id:338921 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1920
AF409940.1 gg_id:60517 Neisseria meningitidis str. M4643 AF4099
NZ_ACQV01000002.1 gg_id:470738 Neisseria flavescens str. SK114 NZ_AC
AF398309.1 gg_id:62767 Neisseria meningitidis str. M7921 AF3983
EF511128.1 gg_id:221648 Loss During Antibiotic Treatment 1 Intubated Patients Colonized Pseudomonas aerug EF5111
NZ_ACEN01000027.1 gg_id:470720 Neisseria flavescens str. NRL30031/H210 NZ_AC
AY005029.1 gg_id:42684 subgingival dental plaque clone AK105 AY005
AF398308.1 gg_id:62770 Neisseria meningitidis str. M7509 AF3983

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DQ201336.1 gg_id:140609 Neisseria meningitidis str. CMCC(B)29202 DQ201
FJ191449.1 gg_id:316460 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1914
AF382307.1 gg_id:65953 Neisseria meningitidis str. M7925 AF3823
AF310344.1 gg_id:30575 Neisseria meningitidis str. M3291 AF3103
EF509732.1 gg_id:218613 Loss During Antibiotic Treatment 1 Intubated Patients Colonized Pseudomonas aerug EF5097
AF310551.1 gg_id:52834 Neisseria meningitidis str. M4797 AF3105
AF398275.1 gg_id:62802 Neisseria meningitidis str. M812 AF3982
AF409938.1 gg_id:60518 Neisseria meningitidis str. M4644 AF4099
FJ191958.1 gg_id:321506 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1919
DQ201323.1 gg_id:140409 Neisseria meningitidis str. CMCC(B)29019 DQ201
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AJ239284.1 gg_id:6465 Neisseria lactamica str. L20 AJ2392
AY132119.1 gg_id:59602 Neisseria meningitidis str. M8574 AY1321
FJ191979.1 gg_id:350810 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1919
AJ239293.1 gg_id:6442 Neisseria sicca str. Q28 AJ2392
AJ239280.1 gg_id:6428 Neisseria flavescens str. LNP444 AJ2392
FJ192352.1 gg_id:350507 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
FJ976263.1 gg_id:539814 Interindividual healthy human oral microbiota mouth clone 6BBB59 FJ9762
AY132085.1 gg_id:69908 Neisseria meningitidis str. M9097 AY1320
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FJ192317.1 gg_id:318935 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
AF398273.1 gg_id:62803 Neisseria meningitidis str. M1168 AF3982
AF398301.1 gg_id:62777 Neisseria meningitidis str. M7918 AF3983
FJ192355.1 gg_id:353666 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
DQ201334.1 gg_id:139856 Neisseria meningitidis str. CMCC(B)29046 DQ201
AY648034.1 gg_id:99627 Neisseria meningitidis str. NZ00/215 AY648
AF398325.1 gg_id:62755 Neisseria meningitidis str. N.33/00 AF3983
NZ_ACDX01000110.1 gg_id:470729 Neisseria mucosa str. ATCC 25996 NZ_AC
L06171.1 gg_id:6402 Neisseria elongata L06171
AF409941.1 gg_id:72478 Neisseria meningitidis str. M1417 AF4099
AF398252.1 gg_id:62824 Neisseria meningitidis str. M5283 AF3982
AB194302.1 gg_id:108299 Sivamonas terrae str. KM-45 AB1943
AF328146.1 gg_id:69998 Simonsiella muelleri str. ATCC29441 AF3281
FJ976380.1 gg_id:554114 Interindividual healthy human oral microbiota mouth clone 22B524 FJ9763
GQ102572.1 gg_id:447914 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ102
DQ003630.1 gg_id:126626 human mouth clone MB5_P15 DQ003
GQ106230.1 gg_id:468425 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ106
EF212919.1 gg_id:214313 Stenoxybacter acetivorans str. TAM-WH4 EF2129
DQ837616.1 gg_id:164644 structures honeybee intestines and their response insecticidal proteins gut clone HB DQ837
AB192065.1 gg_id:144630 termite gut homogenate clone M1NP1-30 AB1920

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EU370420.1 gg_id:264994 Neisseria sp. str. 93S1 EU370
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AY157705.1 gg_id:75888 Neisseria sp. str. CCUG 46910 AY157
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GQ068667.1 gg_id:466523 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
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AY721627.1 gg_id:104026 'New Zealand ' AY721
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FJ356021.1 gg_id:365006 Targeted FACS enrichment groups lab scale systems aerobic lab-scale EBPR activated FJ3560
FJ502346.1 gg_id:512007 Neisseria sp. str. GRW38 FJ5023
GQ068870.1 gg_id:426828 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
GQ094121.1 gg_id:457731 Topographical and Temporal Human Skin Microbiome skin alar crease clone nbw37(GQ094
FJ876273.1 gg_id:589124 Neisseria sp. str. CSL 7565 FJ8762
GQ081376.1 gg_id:414774 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ081
GQ068798.1 gg_id:391640 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
FJ356031.1 gg_id:368201 Targeted FACS enrichment groups lab scale systems anaerobic:aerobic lab-scale EBPF FJ3560
DQ836355.1 gg_id:168478 Prevotia chitinolytica str. JS11-7 DQ836
GQ081072.1 gg_id:384019 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ081
NR_025742.1 gg_id:558848 Chitinibacter tainanensis str. S1; BCRC 17254 NR_02
AY426973.1 gg_id:105593 Neisseria canis VA25810gw_03 AY426
GQ068787.1 gg_id:376358 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
AJ239288.1 gg_id:6399 Neisseria animalis str. NCTC10212 AJ2392
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GQ068942.1 gg_id:392219 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
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GQ081358.1 gg_id:438216 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ081
GQ068691.1 gg_id:466608 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
CU924421.1 gg_id:556357 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU924
Y17602.1 gg_id:6368 Formivibrio citricus str. DSM 6150 Y17602

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DQ314581.1 gg_id:148882 chitin-degrading Taiwan C14 DQ314
AY609199.2 gg_id:140478 Chromobacterium sp. str. 2002 AY609
EU536080.1 gg_id:280825 profile human microbiota antecubital fossa (inner elbow) skin clone nbt08d11 EU536
AB013258.1 gg_id:6372 deep-sea sediment clone NKB6 AB013
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AY904024.1 gg_id:146009 Aquaspirillum sp. str. 411 AY904

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AB076000.1 gg_id:64884 Aquaspirillum putridiconchylium str. IAM 14964 AB076
FJ375419.1 gg_id:589312 T-RFLP powerful tool rapid assessment cells microbial fuel cell sheet carbon cloth ano FJ3754
L06174.1 gg_id:6375 Vitreoscilla stercoraria L06174
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AF363289.1 gg_id:40156 Nitrosomonas sp. str. K1 AF363
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AF353164.1 gg_id:22792 Nitrosomonas sp. DYS323 AF353
AY123794.1 gg_id:69852 Nitrosomonas sp. str. Nm143 AY123
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EF016119.1 gg_id:182102 Nitrosomonas oligotropha str. AS1 EF0161
EF092216.1 gg_id:268524 Persistent Axinella corrugata Nitrosomonas clone B2505_H7 EF0922
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AY123795.1 gg_id:69328 Nitrosomonas eutropha str. Nm57 AY123
AY123807.1 gg_id:70906 Nitrosospira multiformis str. Nl13; C-71; ATCC 25196 AY123
AY856377.1 gg_id:110754 Nitrosospira sp. str. TCH711 AY856
AY856375.1 gg_id:109672 Nitrosospira sp. str. HBN8222A AY856
AF353156.1 gg_id:20624 Nitrosospira sp. TCH716 AF353
AY123804.1 gg_id:72266 Nitrosospira sp. str. Nsp17 AY123
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M96405.1 gg_id:6602 Nitrosovibrio tenuis str. Nv12 M9640
AY123800.1 gg_id:70700 Nitrosospira briensis str. Nsp10 AY123
AY631269.1 gg_id:102482 Nitrosovibrio sp. FJI82 AY631
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AF414581.1 gg_id:33825 uranium mine sediment clone P3OB-51 AF4145
AF080256.1 gg_id:6535 Nitrosospira sp. str. TYM9 AF080
AJ630749.1 gg_id:105991 marine surface water clone BET_1D11G67 AJ6307
EU734961.1 gg_id:311415 assemblages sediment station DBSE Northern Bering Sea clone 054D10 EU734
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L35505.1 gg_id:6517 Nitrosospira briensis str. C-128 L35505

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

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NC_007298.1 gg_id:203562 Dechloromonas aromatica str. RCB NC_00
FJ535057.1 gg_id:591896 Enhancement protein conversion volatile fatty acids accumulation during addition: ef FJ5350
DQ640664.1 gg_id:233934 full-scale EBPR sludge clone Skagenf42 DQ640
GQ340232.1 gg_id:551430 Prokaryotic Marathonas Reservoir Greece water column clone VE07-68-BAC GQ340
AM084133.1 gg_id:140097 Dechloromonas sp. str. R-28400 AM084
AB240507.1 gg_id:142587 microbial structure rhizosphere biofilm Sapporo root-base (80 160 mm) Phragmites S AB240
FJ719082.1 gg_id:531877 Protozoan grazers alter metabolically active aquifer aquifier sediments clone p04_E0 FJ7190
EF515635.1 gg_id:224951 Electricigen Enrichment MFC full-scale anaerobic bioreactor sludge treating brewery EF5156
DQ376553.1 gg_id:153255 Microbial ecology removing phosphate wastewater EBPR process aerobic sequencin DQ376
CU466914.1 gg_id:319502 Evry wastewater treatment plant anoxic basin clone CU466
CU920167.1 gg_id:572334 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU920
CU466920.1 gg_id:304463 Evry wastewater treatment plant anoxic basin clone CU466
AB240296.1 gg_id:144181 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clone AB240
CU466858.1 gg_id:344521 Evry wastewater treatment plant anoxic basin clone CU466
FJ444762.1 gg_id:542590 Cotton Rhizosphere cotton rhizosphere clone 4y-107 FJ4447
CU926684.1 gg_id:554053 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU926
Y17060.1 gg_id:6706 Ferribacterium limneticum str. cda-1 Y17060
DQ296471.1 gg_id:161279 Microbial dissection quinoline-degradation lab-scale anaerobic biofilm reactor quino DQ296
CU466895.1 gg_id:319055 Evry wastewater treatment plant anoxic basin clone CU466
DQ923115.1 gg_id:163080 Methyloversatilis universalis str. 500 DQ923
FJ167509.1 gg_id:333442 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1675
FJ902588.1 gg_id:568555 phreatic limestone sinkholes northeastern Mexico biofilm 1m deep hydrothermal ver FJ9025
FM207922.1 gg_id:305052 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
DQ088734.1 gg_id:137170 subsurface clone BE16FW031601GDW_hole1-11 DQ088
AB234251.1 gg_id:150027 Biotransformation Polychlorinated Dioxins Microbial Dynamics Sediment Microcosm AB234
DQ676301.1 gg_id:224877 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
FM207901.1 gg_id:328865 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
DQ327692.1 gg_id:166580 Biological filtration limits carbon availability affects downstream biofilm formation a DQ327
AY945886.1 gg_id:146463 Thauera and Azoarcus functionally important genera denitrifying quinoline-removal AY945
FJ484883.1 gg_id:511775 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4848
DQ088748.1 gg_id:129402 subsurface clone BE325FW032701CTS_hole1-85 DQ088
FM207903.1 gg_id:327175 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
FM207916.1 gg_id:351540 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
EF363051.1 gg_id:268998 Microbial life within saline fluid Continental Deep Drilling German continental deep g EF3630
AB252920.1 gg_id:171974 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 010 AB2529
AY879297.1 gg_id:111577 Denitratisoma oestradiolicum str. AcBE2-1 AY8792
FM207918.1 gg_id:344874 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
FJ485121.1 gg_id:533665 microbial autonomous rover world's deepest phreatic sinkhole water column El Zacat FJ4851
DQ676390.1 gg_id:222171 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
GQ340363.1 gg_id:538380 Prokaryotic Marathonas Reservoir Greece water column clone VE08-161-BAC GQ340
EF205275.1 gg_id:200507 Arctic gold mine fracture water clone Lupin-1130m-1-MDA-ubp1 EF2052

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

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AB089103.1 gg_id:78240 termite gut homogenate clone Rs-K54 AB0891
NR_024855.1 gg_id:545495 Propionivibrio pelophilus str. asp 66; DSM 12018 NR_024
AY064179.1 gg_id:65492 EBPR sludge clone UCTN173 AY0641
AF255631.1 gg_id:36331 EBPR sludge clone Ebpr16 AF2556
DQ087192.1 gg_id:134476 clone HE018 DQ087
AY691423.1 gg_id:105089 Rhodocyclus sp. str. HOD 5 AY6914
AF229868.1 gg_id:18623 Thauera aromatica str. 4FB2 AF2298
AF229867.1 gg_id:42293 Thauera aromatica str. 4FB1 AF2298
AF229882.1 gg_id:69798 Thauera aromatica str. 3CB3 AF2298
AF123264.1 gg_id:6759 Thauera chlorobenzoica str. 3CB-1 AF1232
AF229887.1 gg_id:36677 Thauera aromatica str. 3BB1 AF2298
AJ315678.1 gg_id:25264 Thauera phenylacetica str. B4P AJ3156
AJ315677.1 gg_id:40990 Thauera aminoaromatica str. S2 AJ3156
U95176.1 gg_id:6763 Thauera aromatica str. T1 U95176
AY548757.1 gg_id:111927 Thauera sp. str. Cin34 AY5487
EU434525.1 gg_id:273528 Thauera phenylacetica str. b208 EU4345
AY098637.2 gg_id:71611 industrial-phenol-degrading community clone MM1 sp. AY0986
DQ018816.1 gg_id:131190 anaerobic swine lagoon clone C-147 DQ018
AJ505850.1 gg_id:58572 Thauera sp. str. PIV-1 AJ5058
EU434574.1 gg_id:265716 Thauera aromatica str. b66 EU4345
FJ900965.1 gg_id:513787 Polymorphism microbial reservoirs water cut stage oil field clone G-3 FJ9009
AM982629.2 gg_id:279478 microbial and slurry pig saw dust spent bedding clone PISD-AJG10 AM982
AB210048.1 gg_id:122645 mini plant Run 5 Nippon Steel Research Engineering Center clone nsmpV06 AB2100
CU466853.1 gg_id:346949 Evry wastewater treatment plant anoxic basin clone CU4668
AJ853609.1 gg_id:111043 landfill leachate clone GZKB116 AJ8536
AY823976.1 gg_id:145956 Investigation Acetate-Fed Denitrifying Microbial Stable Isotope Probing Full-Cycle an AY8239
X68491.1 gg_id:6756 Thauera selenatis str. Ax X68491
EF205256.1 gg_id:197834 Thauera sp. 'CJAOPY3 (T-II)' str. CJAOPY3 (T-II) EF2052
FJ825447.1 gg_id:569266 Microbial during start-up mesophilic biogas plant clone 112_BE3_37 FJ8254
AY823967.1 gg_id:145778 Investigation Acetate-Fed Denitrifying Microbial Stable Isotope Probing Full-Cycle an AY8239
AY570693.1 gg_id:108565 Thauera sp. Al7 AY5706
Y17590.1 gg_id:6761 Thauera mechernichensis str. TL1 Y17590
EU434467.1 gg_id:271936 Thauera aminoaromatica str. a241 EU4344
AY945863.1 gg_id:149463 Thauera and Azoarcus functionally important genera denitrifying quinoline-removal AY9458
AY945915.1 gg_id:145574 Thauera and Azoarcus functionally important genera denitrifying quinoline-removal AY9459
Y17591.1 gg_id:6762 Thauera selenatis str. ATCC 55363T Y17591
AJ005816.1 gg_id:6747 Thauera linaloolentis str. 47Lol AJ0058
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EU104249.1 gg_id:335484 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1042
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GQ093262.1 gg_id:442735 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw42 GQ093
AB192185.1 gg_id:144844 termite gut homogenate clone RsTz-39 AB1921
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GQ100922.1 gg_id:463566 Topographical and Temporal Human Skin Microbiome skin retroauricular crease clor GQ100
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AY124797.1 gg_id:69212 Dechloromonas sp. str. LT-1 AY1247
DQ376566.1 gg_id:152042 Microbial ecology removing phosphate wastewater EBPR process aerobic sequencin DQ376
AB369190.1 gg_id:254556 Microbiological Assessment Circulation Mud Fluids During First Operation Drilling De AB3691
AF072921.1 gg_id:6799 nitrifying/denitrifying industrial activated sludge clone S23 S23 AF0729
AF072918.1 gg_id:6786 nitrifying/denitrifying industrial activated sludge clone S3 S3 AF0729
DQ640700.1 gg_id:231550 full-scale EBPR sludge clone Skagenf76 DQ640
FJ823914.1 gg_id:526353 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8239
GQ024542.1 gg_id:485128 Topographical and Temporal Human Skin Microbiome skin palm clone nbw123e11c1 GQ024
AY947989.1 gg_id:142103 Synchrony and seasonality two temperate rivers USA: Massachusetts Parker River cl AY9479
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GQ093869.1 gg_id:399333 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093
EU015084.1 gg_id:236679 Investigation estrogen-degrading bioreactors membrane bioreactor suspension and EU0150
EF648064.1 gg_id:241562 dynamics during produced water treatment aerobic activated sludge clone HB61 EF6480
EU133837.1 gg_id:240971 tallgrass prarie soil clone FFCH15186 EU1338
AY160865.1 gg_id:96448 termite intestinal tract clone COB P5-5 AY1608
AB024934.1 gg_id:21532 Sphingomonas sp. A1 AB0249
DQ836761.1 gg_id:167461 attached biomass denitrifying bioreactor clone DB-86 DQ836
FJ478739.1 gg_id:511268 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4787
DQ974114.1 gg_id:245301 Azospira restricta str. SUA2 DQ974
FJ374246.1 gg_id:362367 Inter- and intraspecific humivorous scarab beetle larvae (Pachnoda spp.) larval hindgu FJ3742
AF204246.1 gg_id:50481 EBPR sludge clone SBRH147 AF2042
GQ085455.1 gg_id:431512 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw275h11 GQ085
EU133861.1 gg_id:249511 tallgrass prarie soil clone FFCH8183 EU1338
AY007722.1 gg_id:29482 Quadricoccus australiensis str. Ben 117; NCIMB 13738; CIP 107055 AY0077
AY212577.1 gg_id:99384 water 10 m downstream manure clone 126ds10 AY2125
EU015096.1 gg_id:236222 Investigation estrogen-degrading bioreactors membrane bioreactor suspension and EU0150
AB288886.1 gg_id:201491 termite gut homogenate clone BOf2-09 AB2888
AB487233.1 gg_id:562019 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB4872
FJ167461.1 gg_id:353414 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1674
AB237674.1 gg_id:140535 microorganisms deep subsurface groundwater sedimentary rock milieu clone HDBW AB2376
AJ582195.1 gg_id:93931 soil near uranium mill tailings clone Gitt-GS-125 AJ5821

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AB166784.1 gg_id:107623 EBPR sludge clone UTFS-OF08-18 AB1667
AB456210.1 gg_id:553408 Microorganisms involved anaerobic phenol degradation treatment synthetic coke-o AB4562
AM982545.1 gg_id:278568 microbial and slurry pig straw spent bedding clone PIST-AGC10 AM982
AB198538.1 gg_id:135858 termite gut wall fraction clone RsW02-065 AB1985
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GU454933.1 gg_id:520944 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
AB192066.1 gg_id:143225 termite gut homogenate clone M1PL1-62 AB192
GQ065924.1 gg_id:397799 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw94e08 GQ065
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AY770497.1 gg_id:143561 and isolation that grow on 12-dichloroethane under nitrate reducing conditions sedi AY7704
AB486739.1 gg_id:585587 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR AB4867
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AF011343.1 gg_id:6783 Azoarcus communis str. SWub3 AF0113
FJ823940.1 gg_id:529251 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8239
AY662036.1 gg_id:136814 uranium-contaminated groundwater clone 300I-F02 AY6620
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GQ108087.1 gg_id:430940 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ108
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EF019494.1 gg_id:214657 aspen rhizosphere clone Elev_16S_763 EF0194
AB049763.1 gg_id:27927 Denitromonas aromaticus str. AS-7 AB0497

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GQ096373.1 gg_id:385515 Topographical and Temporal Human Skin Microbiome skin back clone nbw387d08c1 GQ096
GQ095282.1 gg_id:374709 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ095
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D14254.1 gg_id:6731 Zoogloea ramigera str. IAM12136 D14254

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

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

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FJ810580.1 gg_id:375861 Subsurface ecosystem resilience: Long-term attenuation subsurface contaminants su FJ8105
AY328747.1 gg_id:90051 drinking water system simulator clone DSSD49 AY3287
EF492933.1 gg_id:257753 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4929
AB294323.1 gg_id:210077 Microbial structure deep coal seem groundwater stream clone SWB12 AB2943
EF492932.1 gg_id:247931 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4929
DQ337089.1 gg_id:150172 subsurface water clone EV818SWSAP66 DQ337
DQ404884.1 gg_id:155647 contaminated sediment clone 654956 DQ404
AM935609.1 gg_id:561445 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
AB294324.1 gg_id:203879 Microbial structure deep coal seem groundwater stream clone SWB13 AB2943
AY922035.1 gg_id:113557 farm soil clone AKYG578 AY922(
DQ837224.1 gg_id:242598 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_1B_8 DQ837
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DQ066965.1 gg_id:136009 Lake Washington sediment clone pLW-9 DQ066
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EU133881.1 gg_id:256389 tallgrass prarie soil clone FFCH2157 EU1338
EF019243.1 gg_id:205571 aspen rhizosphere clone Elev_16S_420 EF0192
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EF019405.1 gg_id:216696 aspen rhizosphere clone Elev_16S_641 EF0194
AY221066.1 gg_id:86386 Mammoth cave clone CCM6a AY221(
FJ810590.1 gg_id:419579 Subsurface ecosystem resilience: Long-term attenuation subsurface contaminants su FJ8105
EF999347.1 gg_id:240278 Vertical and Methanotrophs Estuarine Sediments Pearl River Estuary sediments 6cm EF9993
EU133877.1 gg_id:237992 tallgrass prarie soil clone FFCH1801 EU1338
EF574185.1 gg_id:230954 site S25 near Coco's Island marine clone S25_529 EF5741
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EU477832.1 gg_id:267229 and validation prototype -based taxonomic microarray members genus Burkholderia EU4778
AY234747.1 gg_id:112317 soil isolate Ellin6095 AY2347
EU133863.1 gg_id:249993 tallgrass prarie soil clone FFCH8270 EU1338
EF516320.1 gg_id:223302 grassland soil clone FCPS762 EF5163
EF516424.1 gg_id:227269 grassland soil clone FCPU607 EF5164
EU881164.1 gg_id:352281 Karst region primary forest soil clone KPF200711-070 EU8811
EF516735.1 gg_id:219174 grassland soil clone FCPU655 EF5167
EF516835.1 gg_id:225073 grassland soil clone FCPS400 EF5168
EF019212.1 gg_id:209746 aspen rhizosphere clone Amb_16S_1908 EF0192
EF654711.1 gg_id:245246 suspended sludge anammox reactor clone KIST-JJY042 EF6547
FJ479088.1 gg_id:555896 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4790
DQ413068.1 gg_id:152782 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
EF020095.1 gg_id:208009 aspen rhizosphere clone Elev_16S_1505 EF0200
DQ297977.1 gg_id:142009 Microbial hydrocarbon contaminated soil clone UA5 DQ297

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EU669638.1 gg_id:307844 soil clone S8-40 EU6696
EF516006.1 gg_id:226748 grassland soil clone FCPT739 EF5160
AY921954.1 gg_id:113273 farm soil clone AKYG1130 AY9219
EF632895.1 gg_id:235519 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6328
EF516769.1 gg_id:224647 grassland soil clone FCPS759 EF5167
AB294328.1 gg_id:207655 Microbial structure deep coal seem groundwater stream clone SWB17 AB2943
EF516759.1 gg_id:222745 grassland soil clone FCPT632 EF5167
AJ421933.2 gg_id:32837 Germany:Elbe River clone Elb205 AJ4219
DQ413138.1 gg_id:155225 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
EU193045.1 gg_id:359069 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU1930
AB426185.1 gg_id:273001 -situ identification study benzene-degrading s under methanogenic condition anaerc AB4261
EF516394.1 gg_id:223219 grassland soil clone FCPP685 EF5163
EU528239.1 gg_id:281246 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU5282
EF516152.1 gg_id:217704 grassland soil clone FCPS619 EF5161
EU223955.1 gg_id:236215 -RFLP microbes Qinghai-Tin rail soil clone 124 EU2239
EF516815.1 gg_id:223079 grassland soil clone FCPU398 EF5168
AY221080.1 gg_id:84108 Mammoth cave clone CCU24 AY2210
EF019895.1 gg_id:210313 aspen rhizosphere clone Elev_16S_1261 EF0198
EU133850.1 gg_id:236450 tallgrass prarie soil clone FFCH16997 EU1338
EF516351.1 gg_id:226263 grassland soil clone FCPT411 EF5163
EF019531.1 gg_id:205301 aspen rhizosphere clone Elev_16S_1034 EF0195
EU881346.1 gg_id:336201 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
AY921981.1 gg_id:113247 farm soil clone AKYG1746 AY9219
EF520945.1 gg_id:221665 Stable isotope probing 15N2 noncultivated diazotrophs 30 year fal soil Caldwell field EF5209
DQ676312.1 gg_id:220761 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
AB294330.1 gg_id:207373 Microbial structure deep coal seem groundwater stream clone SWB19 AB2943
EF020158.1 gg_id:207982 aspen rhizosphere clone Elev_16S_1590 EF0201
GQ396975.1 gg_id:553957 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
FJ466126.1 gg_id:509733 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4661
FJ479428.1 gg_id:583255 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4794
EU528189.1 gg_id:285197 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU5281
EF580984.1 gg_id:277681 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM5 EF5809
EF516904.1 gg_id:218538 grassland soil clone FCPS542 EF5169
FJ444750.1 gg_id:551308 Cotton Rhizosphere cotton rhizosphere clone 4y-67 FJ4447
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AY921702.1 gg_id:114209 farm soil clone AKYG1579 AY9217

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EF516025.1 gg_id:218017 grassland soil clone FCPT567 EF516C
GU208364.1 gg_id:576056 Study on Sediment Dongping Lake sediment clone Td4-20 GU208
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EF540415.1 gg_id:220324 oil-shale semi-coke solid waste clone 1_20MK EF5404
EF540371.1 gg_id:226578 oil-shale semi-coke solid waste clone 17C_MJK EF5403
EF018962.1 gg_id:215229 aspen rhizosphere clone Amb_16S_1476 EF0189
AB488102.1 gg_id:551424 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB488
EU404008.1 gg_id:355651 Microbial septic tank effluent and soil biozone infiltrative surface clean water clone (EU404(
EF516891.1 gg_id:222729 grassland soil clone FCPP448 EF5168
AM935489.1 gg_id:548533 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
EF590004.2 gg_id:234053 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF590C
EF516321.1 gg_id:220011 grassland soil clone FCPN390 EF5163
AB286353.1 gg_id:204179 Reducing production and domination comamonadaceae reducing oxygen supply was AB286
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DQ310741.1 gg_id:200807 Recurring seasonal dynamics microbial stream habitats riverine sediment clone WC(DQ310
EF516268.1 gg_id:224900 grassland soil clone FCPP617 EF5162
GQ396994.1 gg_id:593511 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
EF580928.1 gg_id:278471 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1 EF5809
EU340157.1 gg_id:238343 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU340
EU340165.1 gg_id:237526 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU340
EU104188.1 gg_id:341168 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
CU924494.1 gg_id:561226 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU924
AF371865.1 gg_id:25562 swine intestine clone p-2552-9F5 AF3718
EU459462.1 gg_id:296722 capybara feces clone CAP_aah97f02 EU459
EU459519.1 gg_id:294489 capybara feces clone CAP_aah99a09 EU459
FJ535157.1 gg_id:574522 Contamination risk vegetables Arcobacter during post-harvest processing water carr FJ5351
AF418961.1 gg_id:38016 USA: Colorado Fort collins Horsetooth Reservoir clone HTG2 AF418
EU778115.1 gg_id:356250 feces clone RL388_aao93c10 EU778
EF620467.1 gg_id:257876 Ralstonia sp. str. ISSDS-784 EF6204
EU978820.1 gg_id:536916 and metabolic potential metagenome glacier ice clone glb332c EU978
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DQ631725.1 gg_id:245791 sp. OC32 DQ631
EU443934.1 gg_id:268574 sp. str. o6-5b EU443
EU652615.1 gg_id:277698 Seasonal variation microbial Yel Sea sediment clone B8S-8 EU652
EU443952.1 gg_id:263927 sp. str. s3-3 EU443
AF084862.1 gg_id:19894 Bdellovibrio sp. str. JS7 AF084
FJ203067.1 gg_id:324529 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
EU443943.1 gg_id:274526 sp. str. 2bg EU443
EU443963.1 gg_id:269969 sp. str. w5-3 EU443

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EF188429.1 gg_id:346586 Microbial Altamira Cave clone 236 EF1884
AY947967.1 gg_id:141636 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY9479
AJ288899.1 gg_id:49685 stolpii str. DSM 12778 AJ2888
EU805174.1 gg_id:311266 It's all ranking aquatic 250 miles Panama City clone 6C233166 EU8051
EU662557.1 gg_id:580005 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems floating microb EU6625
EU237338.1 gg_id:239851 Genes transcripts Eastern and Western Equatorial North Atlantic Ocean microbial as EU2373
DQ266898.1 gg_id:141116 Cloning and microorganism genes alkaline soil clone pGXAR13 DQ266
EF687517.1 gg_id:281539 Biogeochemistry iron- and sulfur-precipitating microbial mats (Nile Mediterranean) s EF6875
AB369189.1 gg_id:239066 Microbiological Assessment Circulation Mud Fluids During First Operation Drilling De AB3691
AY294221.1 gg_id:101065 sp. str. PNEc1 AY2942
EU101250.1 gg_id:254683 Sulfide/oxygen supply ratio predicts outcome competition among sulfur-oxidizing wa EU1012
GU119471.1 gg_id:564132 Threatened Corals Provide Microbial Habitats reef water clone Reef_F14 GU119
AB176175.1 gg_id:182905 Microbial Hydrothermal System Suiyo Seamount on Izu-Ogasawara Arc vent area wa AB1761
AJ278146.1 gg_id:22594 Bdellovibrio sp. str. SD1 sewage AJ2781
EF574402.1 gg_id:229914 site S25 near Coco's Island marine clone S25_746 EF5744
GQ351210.1 gg_id:590861 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
DQ676387.1 gg_id:218005 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
DQ133926.1 gg_id:168155 Dominant microbial biofilms frasassi cave system Italy limestone-corroding stream b DQ133
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GU369899.1 gg_id:518456 and mats vents Arc white filamentous microbial mat photic shal hydrothermal vent GU369
AJ292518.1 gg_id:38566 Bdellovibrio sp. W str. ATCC 27047 AJ2925
AF148939.1 gg_id:30112 Bdellovibrio str. BRP4 AF1489
EU050708.1 gg_id:247981 Microbial waters production water Daqing oil field clone DQ328-109 EU0507
AJ292759.1 gg_id:40800 Bdellovibrio HD100 str. HD 100 DSM 50701 AJ2927
EU050700.1 gg_id:244273 Microbial waters production water Daqing oil field clone DQ328-1 EU0507
DQ847445.1 gg_id:172283 human forearm skin clone GR2-101 DQ847
GQ113033.1 gg_id:410201 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ113
DQ413094.1 gg_id:152494 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
AB286607.1 gg_id:213370 Reducing production and domination comamonadaceae reducing oxygen supply was AB2866
FJ542862.1 gg_id:525171 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
AM233527.1 gg_id:320761 Multiple symbionts two species co-occurring gutless oligochaete worms Mediterrar AM233
CU466777.1 gg_id:338371 Evry wastewater treatment plant anoxic basin clone CU4667
EF687743.1 gg_id:243041 Bdellovibrio sp. str. JSS EF6877
GQ039893.1 gg_id:416757 Topographical and Temporal Human Skin Microbiome skin retroauricular crease clor GQ039
AY294215.1 gg_id:105561 Bdellovibrio sp. str. MPA AY2942
EU104139.1 gg_id:353499 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1041
DQ640659.1 gg_id:232532 full-scale EBPR sludge clone Skagenf81 DQ640

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AF148941.1 gg_id:74869 Bdellovibrio str. TRA2 AF1489
AY662018.1 gg_id:137132 uranium-contaminated groundwater clone 300A-H04 AY6620
EU134532.1 gg_id:238695 tallgrass prarie soil clone FFCH17748 EU1345
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EF516458.1 gg_id:227450 grassland soil clone FCPT629 EF5164
EU134406.1 gg_id:257948 tallgrass prarie soil clone FFCH7045 EU1344
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EU133256.1 gg_id:247777 tallgrass prarie soil clone FFCH18766 EU1332
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FJ542822.1 gg_id:511221 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
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FJ202533.1 gg_id:352758 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
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EF516682.1 gg_id:224255 grassland soil clone FCPS531 EF5166
DQ129144.2 gg_id:189151 forest soil clone CWT ST01_F02 DQ129
EF442982.1 gg_id:239332 Desulfobacterium catecholicum str. Kette EF4429
FJ516957.1 gg_id:511303 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled upper s FJ5169
U12253.1 gg_id:10829 Desulfobulbus rhabdoformis str. M16 U12253
GU304430.1 gg_id:581441 inhabiting epithelial tissue ruminal digesta -DGGE and sequencing rumen epithelium GU304
DQ831531.1 gg_id:162379 Activity and sulphate-reducing contrasting sediments River Colne estuary UK brackis DQ831
EU844167.1 gg_id:341591 metagenomics random sampling pyrosequencing rumen fluid fiber adherent microbi EU8441
AB189697.1 gg_id:114072 termite gut homogenate clone Tc-66 AB1896
DQ394978.1 gg_id:151814 harbor sediment clone VHS-B4-9 DQ394
AY548789.1 gg_id:105497 Desulfobulbus propionicus str. DSM 2032 AY5487
NR_029305.1 gg_id:590669 Desulfobulbus elongatus str. FP; DSM 2908 NR_029
FJ769451.1 gg_id:542644 Changes Methanogenic Microbial After Sludge Pre-treatment full-scale municipal ana FJ7694
AJ535238.1 gg_id:77808 marine sediment above hydrate ridge clone Hyd24-21 proteobacterium AJ5352
AY355297.2 gg_id:105894 epibiontic clone C11-G5 AY3552
Y13672.1 gg_id:10794 Desulfocapsa sulfexigens str. SB164P1 Y13672
EU662381.1 gg_id:571160 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems microbial mat (EU6623
AJ511275.1 gg_id:69671 Desulfocapsa sp. Cad626 AJ5112
FJ497574.1 gg_id:580889 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4975

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FJ901718.1 gg_id:567131 phreatic limestone sinkholes northeastern Mexico biomat sediment cenote La Palita (FJ9017
DQ925886.1 gg_id:235454 Microbes and Thermal Gradients Temperature Hydrothermal Chimneys hydrotherm DQ925
EU101288.1 gg_id:249303 Sulfide/oxygen supply ratio predicts outcome competition among sulfur-oxidizing wa EU101
X95181.1 gg_id:10792 Desulfocapsa thiozymogenes str. Bra2 X95181
EU662622.1 gg_id:583279 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems floating microb EU662
EU101286.1 gg_id:250699 Sulfide/oxygen supply ratio predicts outcome competition among sulfur-oxidizing wa EU101
DQ521816.1 gg_id:159745 Anaerobic Methane Oxidizing ANME-1b Archaea Hypersaline Sediments Gulf Mexico DQ521
DQ676351.1 gg_id:217639 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
FJ712571.1 gg_id:531935 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7125
EU265788.1 gg_id:264490 epibiotic assemblage recently described Yeti Kiwa hirsuta (Kiwidae) surface crab hyc EU265
GU118041.1 gg_id:549531 Threatened Corals Provide Microbial Habitats clone Apal_D11 GU118
GQ850564.1 gg_id:591035 bottom water northern Bering Sea clone d75 GQ850
AY177802.1 gg_id:78959 Antarctic sediment clone SB4_58 AY1778
GU118122.1 gg_id:557861 Threatened Corals Provide Microbial Habitats clone Apal_M04 GU118
AF118453.1 gg_id:10810 Desulforhopalus singaporensis str. S'pore T1 AF1184
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AY770713.1 gg_id:110727 sea urchin Paracentrotus lividus clone PlS93 AY770
NR_024949.1 gg_id:534504 Desulfotalea arctica str. LSv514; DSM 12342 NR_02
AY770715.1 gg_id:110001 sea urchin Paracentrotus lividus clone PlS95 AY770
GQ357011.1 gg_id:583377 Metal Dependent Marine Methane Oxidation methane seep sediment clone FeLive(GQ357
DQ831548.1 gg_id:170756 Activity and sulphate-reducing contrasting sediments River Colne estuary UK marine DQ831
FJ203002.1 gg_id:319044 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2030
AY177796.1 gg_id:78728 Antarctic sediment clone SB1_68 AY177
AM774321.2 gg_id:245906 Cultivation and situ quantification consistently demonstrate that two distinct group AM774
EU617845.1 gg_id:329633 Seasonal variation microbial Yel Sea sediment clone A13S-69 EU6178
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FN396662.1 gg_id:550643 constant flux thermophilic cold seabed Arctic marine surface sediment clone s5_0_V FN3966
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GQ246441.1 gg_id:583729 Temporal and spatial China North Yel Sea sediments clone Ld1-46 GQ246
AM745146.1 gg_id:290803 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM745
FJ712454.1 gg_id:517813 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7124
DQ831544.1 gg_id:169282 Activity and sulphate-reducing contrasting sediments River Colne estuary UK marine DQ831
EU981262.1 gg_id:342198 sulfate reducing living bench scale aquifer column water clone D1_SW EU981
AF354148.1 gg_id:71200 marine methane seep clone 1431 AF3541
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AB294977.1 gg_id:226332 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma AB2949

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GQ246285.1 gg_id:590606 Temporal and spatial China North Yel Sea sediments clone Sd1-4 GQ246
EU869410.1 gg_id:328130 Microbial hypersaline Sidi Ameur water Himalatt Salt Lake Algerian Sahara clone ARI EU869
EU287257.1 gg_id:257411 Pacific arctic surface sediment clone S11-74 EU287
AM745158.1 gg_id:296704 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM745
DQ831543.1 gg_id:168864 Activity and sulphate-reducing contrasting sediments River Colne estuary UK marine DQ831
EU734932.1 gg_id:311410 assemblages sediment station DBSE Northern Bering Sea clone 023A38 EU734
EU981265.1 gg_id:346330 sulfate reducing living bench scale aquifer column water clone D18_SW EU981
AY771968.1 gg_id:111554 marine surface sediment clone SD27 AY771
AY355305.2 gg_id:101832 epibiontic clone C6-B1 AY355
AM404380.1 gg_id:567796 Hydrocarbon-fueled sulfate reduction can sustain long-lived tubeworm aggregation AM404
AY586398.1 gg_id:108445 fatty acid-degrading consortium clone FA-PB11 AY586
AY216446.1 gg_id:82114 temperate estuarine mud clone KM15 AY216
AY193139.1 gg_id:80324 marine sediment clone Bol88 AY193
EU528183.1 gg_id:281431 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU528
DQ677861.1 gg_id:243494 Blood Falls Subglacial Outf Taylor Glacier Antarctica subglacial outf clone BF99_C161 DQ677
AB240495.1 gg_id:141610 microbial structure rhizosphere biofilm Sapporo root-tip (0 40 mm) Phragmites Sosei AB240
AF050536.1 gg_id:10793 contaminated aquifer clone WCHB1-67 AF050
AY998140.1 gg_id:126281 Continental deep subsurface geochemically structured within alternating shale-sand AY998
EU328056.1 gg_id:278259 dynamics during bioremediation crude oil contaminated moderate saline soil clone B EU328
EU652618.1 gg_id:279049 Seasonal variation microbial Yel Sea sediment clone C8S-16 EU652
GU302481.1 gg_id:569243 Spatial structure and activity sedimentary microbial underlying Beggiatoa spp. mat I GU302
DQ521817.1 gg_id:161410 Anaerobic Methane Oxidizing ANME-1b Archaea Hypersaline Sediments Gulf Mexico DQ521
EU662614.1 gg_id:550521 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems floating microb EU662
EU290696.1 gg_id:241222 microbial structure biogeochemical contrasting shelf sediments Namibian upwelling EU290
NR_026354.1 gg_id:571713 Desulfofustis glycolicus str. PerGlyS NR_02
EF632754.1 gg_id:249075 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6327
EU542499.1 gg_id:260510 Effects chemical structure concentration on pathways microbial during dechlorinatio EU542
FJ264778.1 gg_id:341813 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedime FJ2647
AB100012.1 gg_id:99697 inactive deep-sea hydrothermal vent chimneys clone IndB4-24 AB100
AJ866934.1 gg_id:130219 Desulfobulbus mediterraneus str. NA62 AJ8669
AB252437.1 gg_id:156393 Microbial extremely CO2 rich deep-sea sediment Marine sediments Yonaguni Knoll I AB252
L42613.1 gg_id:10809 Desulforhopalus vacuolatus str. ltk10 L42613
FJ712433.1 gg_id:516762 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) K FJ7124
AY771961.1 gg_id:112020 marine surface sediment clone SP22 AY771
DQ831556.1 gg_id:164944 Activity and sulphate-reducing contrasting sediments River Colne estuary UK marine DQ831
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AY050586.1 gg_id:19356 MCB-contaminated groundwater-treating reactor clone GOUTA17 AY050
AJ241011.1 gg_id:10837 Norway:(Svalbard)Hornsund clone Sva0631 AJ2410
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EU287217.1 gg_id:252970 Pacific arctic surface sediment clone S11-34 EU287
FJ712587.1 gg_id:519783 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7125
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AY327886.1 gg_id:90296 surface vent snail foot clone SF_C6-D1 AY3278
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DQ154829.1 gg_id:138936 Diel Migration Restructures Hypersaline Microbial Mat hypersaline microbial mat cl DQ154
AB294967.1 gg_id:223812 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma AB2949
EF687172.1 gg_id:282050 Biogeochemistry iron- and sulfur-precipitating microbial mats (Nile Mediterranean) ir EF6871
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AJ620509.1 gg_id:108099 Olavius crassitunicatus endosymbiont AJ6205
AF420335.1 gg_id:71031 hydrothermal sediment clone AF420335 AF4203
FJ264573.1 gg_id:335868 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedim FJ2645
EU981263.1 gg_id:335188 sulfate reducing living bench scale aquifer column water clone D5_SW EU981
AY771952.1 gg_id:112016 marine surface sediment clone SK1 AY7719
GQ249525.1 gg_id:581436 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 6-8 cm layer clor GQ249
EU869406.1 gg_id:307864 Microbial hypersaline Sidi Ameur water Himalatt Salt Lake Algerian Sahara clone ARI EU8694
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AB100004.1 gg_id:79377 inactive deep-sea hydrothermal vent chimneys clone IndB2-42 AB1000
FN396953.1 gg_id:561836 integrative isotopic approach study degradation mixed Mixed (US3-M) enriched grou FN3969
EU869423.1 gg_id:309672 Microbial hypersaline Sidi Ameur water Himalatt Salt Lake Algerian Sahara clone ARI EU8694
EU283455.1 gg_id:279354 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A817 EU2834
AB179691.1 gg_id:102062 groundwater 0.2 m-captured fraction clone KNA6-EB15 AB1796
GU061178.1 gg_id:592956 Evaluation culturing methods cultivate previously Yel Sea intertidal beach surface se GU061
EU244009.1 gg_id:367854 16SrRNA probes river Leine sediment clone sl1127 EU2440
AY771929.1 gg_id:111926 marine surface sediment clone LacK4 AY7719
GQ246288.1 gg_id:512929 Temporal and spatial China North Yel Sea sediments clone Sd1-7 GQ246
AM493255.1 gg_id:308423 Multiple symbionts two species co-occurring gutless oligochaete worms Mediterrar AM493
AJ007373.1 gg_id:10840 environmental clone directly AJ0073
DQ513028.1 gg_id:161392 Microbial life fluids ridge flank crustal fluid clone FS142-27B-02 DQ513
GQ337170.1 gg_id:582065 Hydrography shapes biogeography deep Arctic Ocean water depth 410m clone CB1 GQ337

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EU491186.1 gg_id:260159 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
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EU386020.1 gg_id:274277 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
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EU385688.1 gg_id:268129 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU385
FJ716318.1 gg_id:551732 Chlorobi dominate shal chemocline Bahamas Sawmill Sink column 10.3 m water dept FJ7163
FJ517130.1 gg_id:512958 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled water c FJ5171
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GQ487920.1 gg_id:554266 Microbial metal soil polluted heavy metals clone T1-75 GQ487
AB188771.1 gg_id:114153 cold seep sediment clone SB1-2 AB188
EU491434.1 gg_id:259879 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X4 clone EU491
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EU937853.1 gg_id:352286 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9378
FJ746300.1 gg_id:508914 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7463
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g__Desulfohalobium
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g__Desulfonatronovibrio
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s__Desulfovibrio africanus	57109
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AY592130.1 gg_id:100278 marine ANAMMOX sediment clone AY592130 AY592
FJ755771.1 gg_id:586023 vertical archaeal water and Lake Taihu sediment clone LT-SB-B77 FJ7557
X99235.1 gg_id:10401 Desulfohalobium retbaense str. HR100 (DSM 5692T) X99235
U48244.1 gg_id:10400 Desulfohalobium retbaense str. DSM 5692 U48244
EU296539.1 gg_id:241507 Desulfonatronospira aceticus str. AHT 6 EU2965
X99234.1 gg_id:10404 Desulfonatronovibrio hydrogenovorans str. Z-7935 (DSM 9292T) X99234
GU196828.1 gg_id:557238 Desulfonatronovibrio sp. str. AHT21 GU196
EF644847.1 gg_id:299192 genes and citrate lyase deep-sea microbial MAR Rainbow hydrothermal vent system EF6448
AF524933.1 gg_id:74638 Desulfonauticus submarinus str. 6N AF5249
FJ194951.1 gg_id:329685 Desulfonauticus sp. str. DSM 4206 FJ1949
EU245681.1 gg_id:241136 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M4-F1(EU2456
DQ103666.1 gg_id:136882 hypersaline endoevaporitic microbial mat clone E6bG07 DQ103
EU245471.1 gg_id:253107 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M2-A0: EU2454
EU245401.1 gg_id:237109 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M1-C0: EU2454
AB264617.1 gg_id:167257 Desulfothermus okinawensis str. TFISO9 (= JCM 13304 = DSM 17375) AB2646
NR_025378.1 gg_id:537936 Desulfomicrobium orale str. DSM 12838 NR_025
AY464939.1 gg_id:100488 Desulfocaldus terraneus str. P6.2 AY4649
DQ069204.1 gg_id:130717 gold mine clone MAB25 DQ069
AJ277886.1 gg_id:33526 Desulfomicrobium escambiense str. DSM 10707T AJ2778
DQ069227.1 gg_id:128063 gold mine clone SRB5 DQ069
Y14594.1 gg_id:10408 Desulfonatronum lacustre str. z-7951 Y14594
AY725424.1 gg_id:103231 Desulfonatronum cooperatoris str. Z-9799 AY7254
GQ863488.1 gg_id:550045 Desulfonatronum sp. str. HSRB-L1 GQ863
DQ088749.1 gg_id:129416 subsurface clone BE325FW032701CTS_hole1-63 DQ088
GQ493712.1 gg_id:546485 Effect Diet on Human Gut Microbiome: Metagenomic Humanized Gnotobiotic Mice GQ493
FJ363013.1 gg_id:361170 feces adult twins and mothers; TS148 clone TS148_a01h07 FJ3630
AY978908.1 gg_id:121944 human transverse colon mucosal biopsy clone NG04 AY9789
EF602909.1 gg_id:230078 Salmonella typhimurium Exploits Inflammation Compete Intestinal Microbiota mous EF6029
EU454891.1 gg_id:305445 Innate immunity and intestinal microbiota Type 1 diabetes mouse cecum clone K77S EU4548
EU505347.1 gg_id:258794 cecal contents Mus musculus strain C57BL/6J; myd4 clone myd4_aaa02h11 EU5053
EU452762.1 gg_id:309442 Innate immunity and intestinal microbiota Type 1 diabetes mouse cecum clone K80N EU4527
EF096212.1 gg_id:180715 mouse cecum clone obob1_aaa02f06 EF0962
AY993240.1 gg_id:129830 mouse cecum clone C21_k13 AY9932
EF097816.1 gg_id:184560 mouse cecum clone SWPT20_aaa04c06 EF0978
AY991087.1 gg_id:131193 mouse cecum clone C13_O03 AY9910
EU455583.1 gg_id:344389 Innate immunity and intestinal microbiota Type 1 diabetes mouse cecum clone H77S EU4555
EF097361.1 gg_id:199580 mouse cecum clone SWPT19_aaa01e10 EF0973
DQ071477.1 gg_id:234242 microbial cecum captive and free-living capercaillies (Tetrao urogallus) clone TuCc14 DQ071
DQ308598.1 gg_id:148623 sheep rumen enrichment clone 196.G06 DQ308
EU659693.1 gg_id:283392 Desulfovibrio africanus str. SR-1 EU6596
DQ365927.1 gg_id:154031 Desulfovibrio alaskensis str. IMP-7760 DQ365

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g__Desulfovibrio
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s__Desulfovibrio aminophilus	57198
s__Desulfovibrio bastinii	56460
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AF423185.1 gg_id:34507 Toluene-degrading methanogenic consortium clone Eub 5 consortium AF4231
EU251185.1 gg_id:248227 Desulfovibrio sp. str. ds2-2 EU2511
U53462.1 gg_id:10445 Desulfovibrio bastinii str. SEBR 4225 U53462
NR_025766.1 gg_id:533801 Desulfovibrio bastinii str. SRL4225; DSM 16055 NR_025
FJ024714.1 gg_id:330860 Copper affects reduction and genetic enriched sulfate reducing coastal sediments exp FJ0247
DQ109922.1 gg_id:142678 Sulfate-reducing mats: Changing paradigms new discoveries salt pond nonlithifying DQ109
AY281345.1 gg_id:87605 Desulfovibrio sp. str. ANP3 AY2813
U85475.1 gg_id:10430 Desulfovibrio sp. str. BG50 U85475
AF230530.1 gg_id:30839 Desulfovibrio dechloracetivorans AF2305
FJ823871.1 gg_id:516368 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8238
GQ503772.1 gg_id:577487 uranium mine tailing pond inlet water clone Jdgsrb019 GQ503
GQ503807.1 gg_id:585949 effulent treatment plant inlet clone Jdgsrb054 GQ503
AF098671.1 gg_id:10516 Desulfovibrio desulfuricans AF0986
AJ295680.1 gg_id:74474 Desulfovibrio intestinalis str. JG-G12 AJ2956
DQ526427.1 gg_id:160710 Desulfovibrio desulfuricans str. EFX-DES DQ526
GQ503792.1 gg_id:591259 uranium mine tailing pond inlet water clone Jdgsrb039 GQ503
GQ503771.1 gg_id:577327 uranium mine tailing pond inlet water clone Jdgsrb018 GQ503
GQ898871.1 gg_id:572458 strain Desulfovibrio enrichment medium DSMZ 163 inoculated water Jaduguda Urar GQ898
GQ503812.1 gg_id:539624 effulent treatment plant inlet clone Jdgsrb059 GQ503
GQ503798.1 gg_id:564326 uranium mine tailing pond inlet water clone Jdgsrb045 GQ503
FJ823874.1 gg_id:530538 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8238
FJ823870.1 gg_id:521886 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8238
AJ544687.2 gg_id:80009 Desulfovibrio brasiliensis str. LVform1 AJ5446
NR_025977.1 gg_id:570680 Desulfovibrio gabonensis str. SEBR 2840; DSM 10636 NR_025
AP010904.1 gg_id:384156 Desulfovibrio magneticus str. RS-1 AP0109
AJ582758.1 gg_id:96567 Desulfovibrio ferrireducens str. CY2 AJ5827
GU139267.1 gg_id:542453 lactate formate propionate hydrogen donors reductive dehalogenation trichloroeth GU139
GQ402775.1 gg_id:567650 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
EF562565.1 gg_id:235762 stable microbial consortia capable degrading complex organic matter paper pulp colu EF5625
NR_029118.1 gg_id:592620 Desulfovibrio putealis str. B7-43; DSM 16056 NR_029
CP001649.1 gg_id:516164 Desulfovibrio salexigens str. DSM 2638 CP0016
AF050100.1 gg_id:10448 Desulfovibrio senezii str. CVL; DSM 8436 AF0501
AY340820.1 gg_id:91518 sulfate-reducing fluidized-bed reactor clone SR_FBR_E5 AY3408
X86689.1 gg_id:10505 Desulfovibrio sp str. DMB X86689
GQ898881.1 gg_id:585282 strain Desulfovibrio enrichment medium DSMZ 163 inoculated water East Calcutta v GQ898
AB062810.1 gg_id:23507 termite gut homogenate clone BCf11-19 AB0628
GQ502605.1 gg_id:582695 Effects Geographic Location on Formosan Subterranean Termite (FST) gut clone Cf6- GQ502
X95230.2 gg_id:10429 Desulfovibrio aespoeensis str. Aspo2 X95230
Y18049.1 gg_id:10442 Desulfovibrio zosterae Y18049
EU885150.1 gg_id:327333 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone AB2 EU8851
FM178807.1 gg_id:351705 Monitoring abundance and expression 'dehalococcoides' species chloroethene-redu FM178

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GU174149.1 gg_id:558603 Microbial Bactrian camel estimated rumen clone XJC38 GU174
EU885102.1 gg_id:304312 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone A3C12 EU885
EU885154.1 gg_id:337642 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone AF9 EU885
FJ225426.1 gg_id:364203 Desulfovibrio sp. str. H1 FJ2254
EF577029.1 gg_id:264685 Desulfovibrio defluvii str. RB22 EF5770
FJ440030.1 gg_id:510683 Megamonas hypermegale colonization correlates Campylobacter jejuni exclusion pou FJ4400
AM418397.1 gg_id:179230 Desulfohimalaya abdulkalamii str. pwc AM418
AY746987.1 gg_id:100353 Desulfovibrio oxyvorans str. DvO5 AY7469
FJ172813.1 gg_id:354049 Qinghai Yak Ruminal and Fungi China rumen clone RB-4C1 FJ1728
EU850487.1 gg_id:319780 Genetic swamp buffalo (Bubalus bubalis) rumen fluid clone BF252 EU850
EU245278.1 gg_id:257276 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H5-B11 EU245
EU885153.1 gg_id:316313 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone AB12 EU885
U90726.1 gg_id:10427 Desulfovibrio profundus str. DSM 11384 500-1 U9072
AF053752.1 gg_id:10462 Desulfovibrio burkinensis str. HDv; DSM 6830 AF053
EU775070.1 gg_id:308451 Francois Langur feces clone FL_2aaa02h01 EU775
AB234535.1 gg_id:147874 termite gut homogenate clone MgMjR-025 AB2345
AB192287.1 gg_id:141741 termite gut homogenate clone RPK-26 AB192
AF056091.1 gg_id:10518 Desulfovibrio sp. str. UNSW3caefatS AF056
NR_026303.1 gg_id:562366 Desulfovibrio vietnamensis str. G3 100 ID95-371 NR_02
AB470954.1 gg_id:536904 Desulfovibrio sp. str. r49 AB470
EU885140.1 gg_id:350456 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone A2C5 EU885
EU885104.1 gg_id:348053 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone A3G4 EU885
AY649785.1 gg_id:102334 Desulfovibrio alkalitolerans str. RT2 AY649
CP000112.1 gg_id:143024 Desulfovibrio desulfuricans str. G20 CP000
FJ655908.1 gg_id:558435 Desulfovibrio sp. str. I9 FJ6559
FJ202471.1 gg_id:305033 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2024
AF053751.1 gg_id:10470 Desulfovibrio alcoholovorans str. DSM 5433 AF053
AF192152.1 gg_id:51535 Desulfovibrio piger str. ATCC29098 AF192
U53464.1 gg_id:10452 Desulfovibrio gracilis str. SEBR 6116 U53464
EU616635.1 gg_id:321186 digestive-tract microbiota Hirudo orientalis european medicinal leech Desulfovibrio (EU616
DQ447183.2 gg_id:156761 Desulfovibrio gigas str. DSM 1382; ATCC 19364 DQ447
EU885143.1 gg_id:335081 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone AB10 EU885
AY726757.3 gg_id:107875 Desulfovibrio sp. SB1 AY726
DQ422859.1 gg_id:161406 Desulfovibrio bizertensis str. MB3 DQ422
AF371871.1 gg_id:30569 swine intestine clone p-2918-6C5 AF371
EU885152.1 gg_id:341908 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone AE3 EU885
EU885101.1 gg_id:322341 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone A3F9 EU885
AJ132721.1 gg_id:111852 deep-sea hydrothermal chimney clone CHA3-114 AJ1327
NR_025038.1 gg_id:564993 Desulfovibrio inopinatus str. HHQ 20; DSM 10711 NR_02
AY274444.1 gg_id:104557 Desulfovibrio sp. str. HS2 AY2744

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DQ186200.2 gg_id:243561 Desulfovibrio carbinoliphilus str. D41 DQ186
Y11984.3 gg_id:28397 Desulfovibrio alaskensis str. NCIMB 13491 Y11984
DQ841177.2 gg_id:195717 Desulfovibrio sp. str. AS36 DQ841
EF453987.2 gg_id:225093 termite hindgut clone 290cost002-P3L-1410 EF4539
AM947130.1 gg_id:270241 Desulfovibrio marrakechensis str. DSM 19337 AM947
CP001197.1 gg_id:529124 Desulfovibrio vulgaris 'Miyazaki F' str. Miyazaki F CP001
EF123506.1 gg_id:257510 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1235
U85468.1 gg_id:10463 Desulfovibrio sp. str. BG6 U85468
AF228128.2 gg_id:43170 Desulfovibrio sp. str. Pr1.2 AF2281
GQ502595.1 gg_id:585650 Effects Geographic Location on Formosan Subterranean Termite (FST) gut clone Cf6- GQ502
NR_025765.1 gg_id:576099 Desulfovibrio longus str. SEBR 2582; DSM 6739 NR_025
AY274449.1 gg_id:105315 Desulfovibrio ferrophilus str. IS5 AY2744
X95231.1 gg_id:10444 Bacterial sp. X95231
NC_008011.1 gg_id:159002 Lawsonia intracellularis str. PHE/MN1-00 NC_008
AY976442.1 gg_id:118641 human ascending colon mucosal biopsy clone LZ39 AY9764
GQ867364.1 gg_id:588730 and prevalence blaTEM genes Arctic seals colon content clone WSp127 GQ867
EU381557.1 gg_id:265414 Rumen eubacterial implicates Bacteroidetes major particulate phylum rumen liquid p EU3815
FJ833256.1 gg_id:572886 Microbial Ecology Crop Folivorous Hoatzin crop contents clone A3Q302_22F03 FJ8332
EU468458.1 gg_id:287520 okapi feces clone OK1_a06 EU4684
EU461101.1 gg_id:303251 red kangaroo feces clone KO1_aai44b11 EU4611
AB234532.1 gg_id:146574 termite gut homogenate clone MgMjD-084 AB2345
AB089110.1 gg_id:78137 termite gut homogenate clone Rs-N35 proteobacterium AB0891
AB192288.1 gg_id:143786 termite gut homogenate clone RPK-66 AB1922
X99501.1 gg_id:10486 Desulfovibrio sp. str. STL1 X99501
FJ374255.1 gg_id:365000 Inter- and intraspecific humivorous scarab beetle larvae (Pachnoda spp.) larval hindgu FJ3742
AB234538.1 gg_id:149936 termite gut homogenate clone MgMjR-064 AB2345
AB089109.1 gg_id:78006 termite gut homogenate clone Rs-M89 proteobacterium AB0891
EF454896.2 gg_id:220622 termite hindgut clone 290cost002-P3L-618 EF4548
X99504.1 gg_id:10485 Desulfovibrio sp. str. STL6 X99504
FJ374258.1 gg_id:361169 Inter- and intraspecific humivorous scarab beetle larvae (Pachnoda spp.) larval hindgu FJ3742
AB089105.1 gg_id:78148 termite gut homogenate clone Rs-K75 AB0891
AY532164.1 gg_id:137367 Desulfovibrio sp. str. ZIRB-2 AY5321
AB234531.1 gg_id:148238 termite gut homogenate clone MgMjD-065 AB2345
Y16942.1 gg_id:10572 Desulfurella propionica str. U-8 (DSM 10410) Y16942
Y16941.1 gg_id:10573 Desulfurella kamchatkensis str. K-119 (DSM 10409) Y16941
NR_026109.1 gg_id:566496 Desulfurella acetivorans str. A63; DSM 5264 NR_026
Y18292.1 gg_id:51780 Hippea maritima str. MH2T Y18292
FJ264595.1 gg_id:343915 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedime FJ2645
AB294284.1 gg_id:207880 Microbial structure deep coal seem groundwater clone YWB15 AB2942
AY062880.1 gg_id:38853 environmental electrode surface clone d14 AY0628
EU052251.1 gg_id:240954 degradation kinetics and current efficiency cells supplied anode microbial fuel cell fe EU0522

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EU246142.1 gg_id:235638 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P4-B12 EU246
AY062875.1 gg_id:52449 environmental electrode surface clone d9 AY062
AJ519664.1 gg_id:76391 uranium mill tailings soil sample clone GuBH2-AG-114 proteobacterium AJ5196
AB294287.1 gg_id:210461 Microbial structure deep coal seem groundwater clone YWB18 AB294
X79415.1 gg_id:10692 Desulfuromusa succinoxidans str. Gylac (DSM 8270) X79415
GQ433952.1 gg_id:554572 Subseafloor ocean crust microbial observatories: (F-through Osmo Colonization Syst GQ433
EU542456.1 gg_id:263005 Effects chemical structure concentration on pathways microbial during dechlorinatio EU542
NR_026154.1 gg_id:552396 Pelobacter acidigallici str. MaGal12T; DSM 2377 NR_02
GU180167.1 gg_id:581956 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
NR_026479.1 gg_id:539100 Malonomonas rubra str. GraMal1; DSM 5091 NR_02
FJ638573.1 gg_id:538924 Temperature dependent succession sulfate reducing muddy southwestern Taiwan hc FJ6385
AY280424.1 gg_id:101417 exterior deep-sea hydrothermal vent sulfide chimney Juan de Fuca Ridge Pacific Oce AY280
EU386013.1 gg_id:262433 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
GQ246309.1 gg_id:511753 Temporal and spatial China North Yel Sea sediments clone Sd1-28 GQ246
EU052244.1 gg_id:237074 degradation kinetics and current efficiency cells supplied anode microbial fuel cell fe EU052
DQ677843.1 gg_id:245056 Blood Falls Subglacial Outf Taylor Glacier Antarctica subglacial outf clone BF99_C24 DQ677
DQ309326.1 gg_id:145062 Desulfuromonas alkaliphilus str. Z-0531 DQ309
AJ441215.1 gg_id:74843 hydrothermal vent polychaete mucous clone P. palm C/A 79 AJ4412
EU052243.1 gg_id:252121 degradation kinetics and current efficiency cells supplied anode microbial fuel cell fe EU052
AY187308.1 gg_id:102591 Pelobacter masseliensis AY1873
EU617856.1 gg_id:346218 Seasonal variation microbial Yel Sea sediment clone C13S-139 EU617
AB355057.1 gg_id:236199 Monitoring and archaeal two heavy polluted areas lake Egypt based on sediment Ma AB355
AY998187.1 gg_id:115871 Continental deep subsurface geochemically structured within alternating shale-sand AY998
AY515473.1 gg_id:103018 bulk water German Wadden Sea October 1999 clone GWS-K17 AY5154
EU531802.1 gg_id:264753 unculturable volcanos Xinjiang mud volcano (salty) clone 285 EU5318
AB189359.1 gg_id:113759 cold seep sediment clone JT75-105 AB189
AY187304.2 gg_id:108676 Geopsychrobacter electrodiphilus str. A2 AY1873
EU735643.1 gg_id:329004 archaeal structures and pristine soils China oil contaminated soil Jidong Oilfield clone EU735
FM179908.1 gg_id:353366 Biogeochemical processes microbial Gullfaks and Tommeliten methane seeps (Nortl FM179
AY155599.1 gg_id:81506 Geothermobacter ehrlichii str. SS015 AY1555
GQ249592.1 gg_id:554911 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 12-14 cm layer cl GQ249
EF687392.1 gg_id:286079 Biogeochemistry iron- and sulfur-precipitating microbial mats (Nile Mediterranean) s EF6873
AF458286.1 gg_id:77161 Mono Lake California clone AF4582
NR_026076.1 gg_id:561012 Geobacter bremensis str. Dfr1; ATCC BAA-607 NR_02
AAVG01000086.1 gg_id:179082 Geobacter lovleyi str. SZ AAVG0
NC_007517.1 gg_id:141183 Geobacter metallireducens str. GS-15 NC_00
AY653551.1 gg_id:129581 Geobacter psychrophilus str. P11 AY6535
AJ534676.1 gg_id:76599 ground water deep-well injection disposal site radioactive wastes Tomsk-7 clone S15A AJ5346
AB488058.1 gg_id:581180 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB488
AB486606.1 gg_id:576801 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR0 AB486

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NC_009483.1 gg_id:233864 Geobacter uraniumreducens str. Rf4 NC_00
NC_008609.1 gg_id:179753 Pelobacter propionicus str. DSM 2379 NC_00
AB307657.1 gg_id:262500 clay wall used Japanese traditional buildings clone R33 AB307
FJ793176.1 gg_id:532352 Archaea and Hot Spring Thailand Tao Dam hot spring clone TDB56 FJ7931
AJ617860.1 gg_id:109573 oxic-anoxic interphase flooded paddy soil clone D28213 AJ6178
AB487900.1 gg_id:590693 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB487
GQ487953.1 gg_id:563942 Microbial metal soil polluted heavy metals clone T7-90 GQ487
EF515449.1 gg_id:222702 Electricigen Enrichment MFC upf microbial fuel cell anode clone 24e09 EF5154
FM204965.1 gg_id:569815 Prokaryote shifts and reductive dehalogenase redundancy response electron donor FM204
FJ461924.1 gg_id:548027 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4619
FJ461960.1 gg_id:557399 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4619
AJ307942.1 gg_id:18952 derived microbial 'pearl'- community clone sipK44 AJ3079
AB488320.1 gg_id:576040 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB488
AB486133.1 gg_id:543299 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSBARC AB486
AY922116.1 gg_id:113721 farm soil clone AKYH1470 AY922
AB487641.1 gg_id:591167 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB487
FJ461907.1 gg_id:544277 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4619
AB486599.1 gg_id:589964 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR0 AB486
GQ355025.1 gg_id:555760 structure between two geochemically distinct springs Alum Rock Park spring clone M GQ355
DQ138957.1 gg_id:139194 chemical fertilizer paddy soil clone JG135 DQ138
FJ269083.1 gg_id:326474 dissimilatory ferric iron reducers arsenic contaminated Hunan south China paddy soil FJ2690
GQ263990.1 gg_id:582001 Structure Simulated Level Waste Site simulated level waste site clone WC2_80 GQ263
AB486446.1 gg_id:590701 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSBARC AB486
EF527427.1 gg_id:222124 Geobacter uraniumreducens str. Rf4 EF5274
AY947909.1 gg_id:142841 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY947
FJ461957.1 gg_id:585743 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4619
AM713386.1 gg_id:239296 Heterogeneity microorganisms sediments caused pH coal-mining lake sediment Fe(AM713
DQ086800.1 gg_id:143335 Geobacter sp. str. CLFeRB DQ086
AB240241.1 gg_id:144252 microbial structure rhizosphere Phragmites bulk soil reed bed reactor laboratory clo AB240
EU335196.1 gg_id:242051 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
NR_026077.1 gg_id:552049 Geobacter pelophilus str. Dfr2; ATCC BAA-603 NR_02
AB307656.1 gg_id:259279 clay wall used Japanese traditional buildings clone R40 AB307
AB234266.1 gg_id:149023 Biotransformation Polychlorinated Dioxins Microbial Dynamics Sediment Microcosm AB234
FJ393130.1 gg_id:358635 Fusing Biology Engineering Improve Power Output Microbial Fuel Cells microbial fuel FJ3931
Y19190.1 gg_id:10733 Geobacter sp. str. CdA-2 Y19190
EF014495.1 gg_id:236914 Geobacter sp. G02 EF0144
AB240367.1 gg_id:144809 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
DQ354751.1 gg_id:150927 subsurface water clone MS140BH1062003ms331 DQ354
CP001124.1 gg_id:312961 Geobacter bemidjiensis str. Bem CP001
AY013644.2 gg_id:23462 sample taken downstream landfill clone BVB71 landfill AY013

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DQ676355.1 gg_id:226377 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
AF335183.1 gg_id:29370 Geobacter grbiciae str. TACP-5 AF335
DQ145535.1 gg_id:142149 Geobacter pickeringii str. G13 DQ145
FM204949.1 gg_id:571587 Prokaryote shifts and reductive dehalogenase redundancy response electron donor FM204
EF515364.1 gg_id:225124 Electricigen Enrichment MFC upf microbial fuel cell anode clone 23d10 EF5153
EF527233.1 gg_id:220762 Geobacter sp. Ply1 str. PLY1 EF5272
AJ009496.1 gg_id:10720 anaerobic TCB-transforming consortium clone SJA-152 AJ0094
Y19191.1 gg_id:10708 Geobacter sp. str. CdA-3 Y19191
FJ479238.1 gg_id:593010 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4792
AM713384.1 gg_id:258079 Heterogeneity microorganisms sediments caused pH coal-mining lake sediment Fe(AM713
EF527234.1 gg_id:220531 Geobacter sp. Ply4 str. PLY4 EF5272
AY395392.1 gg_id:104157 pasture soil clone EB1073 AY3953
AB488092.1 gg_id:552927 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB488
EF515400.1 gg_id:224061 Electricigen Enrichment MFC upf microbial fuel cell anode clone 23h09 EF5154
AY737507.1 gg_id:105662 Geobacter hephaestius str. VeEG4 AY7375
NR_025895.1 gg_id:543004 Geobacter metallireducens str. GS-15; ATCC 53774 NR_02
AY214185.1 gg_id:82426 benzene-contaminated groundwater clone ZZ12C11 AY2141
EF515282.1 gg_id:218355 Electricigen Enrichment MFC upf microbial fuel cell anode clone 22d02 EF5152
FM204997.1 gg_id:547916 Prokaryote shifts and reductive dehalogenase redundancy response electron donor FM204
EF515564.1 gg_id:217229 Electricigen Enrichment MFC full-scale anaerobic bioreactor sludge treating brewery EF5155
EU335191.1 gg_id:251985 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
AB239707.1 gg_id:143933 microbial structure rhizosphere Phragmites bulk soil reed bed reactor laboratory clo AB239
AY297795.1 gg_id:89444 waterlogged archaeological wood clone pACH72 AY297
EF520540.1 gg_id:231136 structure lakes: what controls ? acid-impacted lake clone ADK-MOh02-53 EF5205
AY770971.1 gg_id:108346 water injection well Dagang oilfield clone W31 AY7709
AF223382.1 gg_id:32027 Trichlorobacter thiogenes AF2233
EF520542.1 gg_id:229376 structure lakes: what controls ? acid-impacted lake clone ADK-MOh02-26 EF5205
FM210335.1 gg_id:572312 Prokaryote shifts and reductive dehalogenase redundancy response electron donor FM210
EU335154.1 gg_id:244045 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
AB447710.1 gg_id:570781 Microbial Structure and Dynamics Granules Expanded Granular Sludge Bed (EGSB) R AB447
AB262721.1 gg_id:255281 microorganisms shal subsurface Sarobetsu mire peat soil clone HSM050P-B-10 AB262
EU244079.1 gg_id:356946 16SrRNA probes river Leine sediment clone sl1361 EU244
FJ478832.1 gg_id:511698 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4788
FJ461926.1 gg_id:541076 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4619
EU134368.1 gg_id:252584 tallgrass prarie soil clone FFCH5218 EU1343
EF515297.1 gg_id:223706 Electricigen Enrichment MFC upf microbial fuel cell anode clone 22e08 EF5152
EF192881.1 gg_id:202514 sediments temperate artificial lake Rapel reservoir sediment clone Rap1_22D EF1928
AY922036.1 gg_id:113772 farm soil clone AKYG597 AY922
AB186850.1 gg_id:107711 polychlorinated dioxin dechlorinating microcosm clone TSBW04 AB186
AB488109.1 gg_id:590962 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB488

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AB486722.1 gg_id:534035 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR AB486
AY947936.1 gg_id:141881 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY947
AB488329.1 gg_id:535240 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR AB488
CP000142.1 gg_id:140482 Pelobacter carbinolicus str. DSM 2380 CP000
EU528245.1 gg_id:287305 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU528
AF357914.2 gg_id:51450 Desulfuromonas michiganensis str. BRS1 AF357
U41562.1 gg_id:10667 Pelobacter venetianus U4156
DQ647155.1 gg_id:251658 Microbial structure -temperature oil-field produced water Troll formations North Se DQ647
AB243821.1 gg_id:240641 Methods Investigate Microbial Niigata oil well clone Niigata-10 AB243
AY570613.1 gg_id:108489 biodegraded Canadian oil reservoir clone PL-27B4 AY570
AB294283.1 gg_id:215360 Microbial structure deep coal seem groundwater clone YWB14 AB294
EU328027.1 gg_id:279210 dynamics during bioremediation crude oil contaminated moderate saline soil clone B EU328
GU180166.1 gg_id:537907 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
AM745207.1 gg_id:293270 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM745
U23140.1 gg_id:10684 "Desulfuromonas acetexigens" U2314
AB186848.1 gg_id:103899 polychlorinated dioxin dechlorinating microcosm clone TSAC13 AB186
AY185329.1 gg_id:107087 Desulfuromonas sp. clone AKS68 AY185
X70955.1 gg_id:10668 Pelobacter acetylenicus str. WoAcy1 DSM2348 X70955
GQ433929.1 gg_id:580755 Subseafloor ocean crust microbial observatories: (F-through Osmo Colonization Syst GQ433
EU491247.1 gg_id:276607 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
EU134335.1 gg_id:243283 tallgrass prarie soil clone FFCH4372 EU134
EU669605.1 gg_id:329697 soil clone S11-29 EU669
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EU335175.1 gg_id:243968 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
EU236382.1 gg_id:299505 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg91H5 EU236
AF130847.1 gg_id:24509 Candidatus Entotheonella palauensis AF130
AM697535.1 gg_id:223474 sequencing libraries indoor dust clone BF0002D046 AM697
FJ466208.1 gg_id:534885 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4662
FJ479408.1 gg_id:559177 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4794
FJ542849.1 gg_id:531899 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
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DQ450812.1 gg_id:159853 Microbial Alpine Tundra Wet Meadow Soil: Chloroflexi Cold Water-Saturated satura DQ450
AY792293.1 gg_id:119981 Humic Lake clone CrystalBog021E5 AY792
EU134440.1 gg_id:241214 tallgrass prarie soil clone FFCH11684 EU134
EF516730.1 gg_id:225970 grassland soil clone FCPP728 EF5167
EF516405.1 gg_id:226410 grassland soil clone FCPT673 EF5164
EU803523.1 gg_id:339137 It's all ranking aquatic Lake Gatun clone 5C231108 EU803
AJ233906.1 gg_id:10634 Cystobacter violaceus str. Cb vi34 AJ2339
DQ768113.1 gg_id:169959 Cystobacter minus str. DSM 14751 DQ768

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AJ233913.1 gg_id:10629 Archangium gephyra str. Ar g2 AJ2339
AJ233910.1 gg_id:10626 Angiococcus disciformis str. d4 AJ2339
AB218222.1 gg_id:148095 Archangium gephyra str. NBRC 100087 AB218
AJ233900.1 gg_id:10630 Cystobacter ferrugineus str. Cb fe13 AJ2339
DQ768109.1 gg_id:168891 Cystobacter fuscus str. DSM 2262 DQ768
DQ768115.1 gg_id:169963 Cystobacter velatus str. DSM 14718 DQ768
AJ233897.1 gg_id:10617 Cystobacter fuscus str. Cb f6 AJ2338
EU263004.1 gg_id:274606 Cystobacter fuscus str. Cb9737 EU263
AJ233902.1 gg_id:10621 Cystobacter ferrugineus str. Cb fe27 AJ2339
AM930267.1 gg_id:251539 Melittangium lichenicola str. DSM 14877 AM930
EU262995.1 gg_id:264353 Cystobacter fuscus str. Bj41 EU2629
DQ768107.1 gg_id:169848 Cystobacter armeniaca str. DSM 14710 DQ768
DQ768111.1 gg_id:163670 Cystobacter miniatus str. DSM 14712 DQ768
AJ970180.1 gg_id:133876 Stigmatella erecta str. DSM 16858 AJ9701
EU193069.1 gg_id:365027 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU193
AJ233911.1 gg_id:10627 Angiococcus disciformis str. d6 AJ2339
AJ233905.1 gg_id:10633 Cystobacter violaceus str. Cb vi29 AJ2339
AJ233944.1 gg_id:10652 Polyangium vitellinum str. Pl vt1 AJ2339
EF108312.1 gg_id:180118 Haliangium ochraceum str. GS1 EF1083
AB062751.1 gg_id:38718 Haliangium tepidum AB062
EU134537.1 gg_id:236984 tallgrass prarie soil clone FFCH18685 EU1345
FJ712545.1 gg_id:530479 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7125
AB286567.1 gg_id:204372 Reducing production and domination comamonadaceae reducing oxygen supply was AB2865
GQ264394.1 gg_id:574674 Structure Simulated Level Waste Site simulated level waste site clone WW2_84 GQ264
AM072418.1 gg_id:137564 uranium mining waste pile soil clone JG35-2-JT67 AM072
EF018702.1 gg_id:204556 aspen rhizosphere clone Amb_16S_999 EF0187
FJ479136.1 gg_id:578408 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4791
GQ396851.1 gg_id:566634 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
EU134437.1 gg_id:242508 tallgrass prarie soil clone FFCH11506 EU1344
EU134439.1 gg_id:236963 tallgrass prarie soil clone FFCH11594 EU1344
EU676441.1 gg_id:300085 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals· EU6764
AY921676.1 gg_id:113324 farm soil clone AKYH1423 AY9216
AJ518792.1 gg_id:76933 uranium mining waste pile clone JG37-AG-33 proteobacterium AJ5187
AJ519631.1 gg_id:74929 uranium mill tailings soil clone Sh765B-TzT-42 AJ5196
EU617834.1 gg_id:333636 Seasonal variation microbial Yel Sea sediment clone C13S-30 EU6178
EF516060.1 gg_id:217328 grassland soil clone FCPP560 EF5160
EU925860.1 gg_id:353947 sediments sediment station DBS1 northern Bering Sea clone 061F22 EU9258
AM490754.1 gg_id:542734 Linking and functional nutrient spiraling mats (USA) microbial mat sulfidic cave sprir AM490
DQ395973.1 gg_id:152024 deep-sea octacoral clone ctg_CGOGA72 DQ395
EU617837.1 gg_id:313196 Seasonal variation microbial Yel Sea sediment clone D13S-10 EU6178

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DQ640732.1 gg_id:228460 full-scale EBPR sludge clone Skagen129 DQ640
AY922085.1 gg_id:114341 farm soil clone AKYG989 AY922(
EU134548.1 gg_id:246356 tallgrass prarie soil clone FFCH2604 EU1345
EU335231.1 gg_id:256412 Changes microbial metabolic and along hydrogeochemically variable profile unsatura EU3352
AY862012.1 gg_id:112562 Queens Laundry berry community clone QLBB011 AY862(
FJ748771.1 gg_id:578632 Vertical and sulfate-reducing prokaryotes estuarine Southern China Pearl River Estuai FJ7487
EF515967.1 gg_id:224129 grassland soil clone FCPO510 EF5159
EF516802.1 gg_id:221219 grassland soil clone FCPS608 EF5168
EF516477.1 gg_id:218069 grassland soil clone FCPO701 EF5164
EU617835.1 gg_id:345106 Seasonal variation microbial Yel Sea sediment clone D13S-24 EU6178
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FJ466420.1 gg_id:566988 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4664
EU652608.1 gg_id:279253 Seasonal variation microbial Yel Sea sediment clone B8S-115 EU6526
GU119139.1 gg_id:570059 Threatened Corals Provide Microbial Habitats clone Past_J10 GU119
AB286332.1 gg_id:204177 Reducing production and domination comamonadaceae reducing oxygen supply was AB2863
AY622263.1 gg_id:99343 subsurface soil clone B-BK96 AY6222
FM253665.1 gg_id:305498 Rock gold mine unusually microbial biofilm clone F04-2 FM253
FJ478686.1 gg_id:570453 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
FJ478958.1 gg_id:543753 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4789
AJ241020.1 gg_id:10682 Norway:(Svalbard)Hornsund clone Sva0679 AJ2410
GU208498.1 gg_id:592792 Study on Sediment Dongping Lake sediment clone Shf1-5 GU208
FJ466408.1 gg_id:548168 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4664
AJ576419.1 gg_id:102303 midgut homogenate Pachnoda ephippiata larva clone PeM65 AJ5764
EF580975.1 gg_id:277714 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1: EF5809
EF999354.1 gg_id:244128 Vertical and Methanotrophs Estuarine Sediments Pearl River Estuary sediments 6cm EF9993
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EU445232.1 gg_id:563875 Isolation Weight Forest Topsoil Metagenomic tropical forest topsoil clone R84 EU445
GU118929.1 gg_id:546642 Threatened Corals Provide Microbial Habitats clone Past_O02 GU118
DQ906778.1 gg_id:254611 Oman subsurface soil clone 2A-7 DQ906
GQ264118.1 gg_id:551080 Structure Simulated Level Waste Site simulated level waste site clone WC3_53 GQ264
EU617833.1 gg_id:332464 Seasonal variation microbial Yel Sea sediment clone D13S-2 EU6178
AM997778.1 gg_id:572137 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EU134286.1 gg_id:256241 tallgrass prarie soil clone FFCH3218 EU1342
EU134430.1 gg_id:239367 tallgrass prarie soil clone FFCH10899 EU1344
AY921878.1 gg_id:113228 farm soil clone AKYH1170 AY9218
EF580944.1 gg_id:278622 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM6(EF5809
EF019351.1 gg_id:207878 aspen rhizosphere clone Elev_16S_566 EF0193
EU861852.1 gg_id:345643 effects chronic fertilization on alpine tundra microbial implications carbon and nitrog EU8618
NC_009675.1 gg_id:250283 Anaeromyxobacter sp. str. Fw109-5 NC_00

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GQ397088.1 gg_id:532843 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ397
EU134349.1 gg_id:236268 tallgrass prarie soil clone FFCH6412 EU1343
FJ190071.1 gg_id:335745 Strain-specific probing strain and heterogeneous U(VI)-reducing Anaeromyxobacter s FJ1900
FJ190063.1 gg_id:353729 Strain-specific probing strain and heterogeneous U(VI)-reducing Anaeromyxobacter s FJ1900
EU134508.1 gg_id:241430 tallgrass prarie soil clone FFCH17129 EU1345
AM162453.1 gg_id:151071 and situ identification acidic sphagnum peat bog clone B88 AM162
FJ190066.1 gg_id:324580 Strain-specific probing strain and heterogeneous U(VI)-reducing Anaeromyxobacter s FJ1900
EU331407.1 gg_id:247762 Anaeromyxobacter dehalogenans str. DCP15 EU3314
FJ190072.1 gg_id:317365 Strain-specific probing strain and heterogeneous U(VI)-reducing Anaeromyxobacter s FJ1900
GQ402739.1 gg_id:579915 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
GQ402712.1 gg_id:582110 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
EU307086.1 gg_id:249072 Changes microbial metabolic and along hydrogeochemically variable profile unsatura EU3070
AY921809.1 gg_id:113159 farm soil clone AKYG886 AY9218
EF520536.1 gg_id:234490 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-17 EF5205
AF382397.1 gg_id:24157 Anaeromyxobacter dehalogenans str. 2CP-5 AF3823
FJ190062.1 gg_id:312494 Anaeromyxobacter dehalogenans str. FRC-W FJ1900
AB486683.1 gg_id:571531 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR AB4866
AJ233915.1 gg_id:10592 Myxococcus coralloides str. Cc c806 AJ2339
GU139156.1 gg_id:587638 Corallococcus macrosporus str. HBUM91004 GU139
EU263003.1 gg_id:259044 Corallococcus coralloides str. 0185-2 EU2630
EU263002.1 gg_id:263917 Corallococcus coralloides str. 0198-2 EU2630
GQ246607.1 gg_id:550449 Corallococcus exiguus str. HBUM94202 GQ246
AJ233931.1 gg_id:10589 Corallococcus exiguus str. Cc e12b AJ2339
AM930268.1 gg_id:247897 Melittangium lichenicola str. ATCC 25944 AM930
AJ811614.1 gg_id:106360 Corallococcus coralloides str. DSM 51301 AJ8116
AJ811612.1 gg_id:106050 Corallococcus coralloides str. DSM 52500 AJ8116
AJ811595.1 gg_id:105288 Corallococcus coralloides str. DSM 51651 AJ8115
EU262999.1 gg_id:265680 Corallococcus coralloides str. 0007-3 EU2629
AJ811593.1 gg_id:104972 Corallococcus coralloides str. DSM 51548 AJ8115
AJ811601.1 gg_id:100844 Corallococcus coralloides str. DSM 51492 AJ8116
AJ811604.1 gg_id:101327 Corallococcus coralloides str. DSM 52502 AJ8116
AJ811615.1 gg_id:106528 Corallococcus coralloides str. DSM 51408 AJ8116
AY724800.1 gg_id:99811 Myxococcus xanthus str. UCDaV17 AY7248
AY724799.1 gg_id:105036 Myxococcus xanthus str. UCDaV16 AY7247
AY072737.1 gg_id:17656 Myxococcus macrosporus str. 125-10-3 AY0727
AY072740.1 gg_id:21317 Myxococcus macrosporus str. 125-10-1 AY0727
AY724798.1 gg_id:104894 Myxococcus xanthus str. UCDaV5 AY7247
EU271853.1 gg_id:251509 Myxococcus stipitatus str. KYC1101 EU2718
GQ246570.1 gg_id:590399 Myxococcus stipitatus str. HBUM92079 GQ246
EU262651.1 gg_id:243378 Myxococcus fulvus str. KYC1136 EU2626
EU262655.1 gg_id:237067 Myxococcus fulvus str. KYC1198 EU2626

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AJ233922.1 gg_id:10607 Myxococcus stipitatus str. Mx s33 AJ2339
AB218209.1 gg_id:150276 Myxococcus fulvus str. NBRC 100070 AB218
GU139153.1 gg_id:579552 Myxococcus stipitatus str. HBUM93501 GU139
AJ233929.1 gg_id:10614 Myxococcus xanthus str. DSM 435 (Mx x1) AJ2339
GQ246531.1 gg_id:576354 Myxococcus virescens str. HBUM93351 GQ246
AJ233927.1 gg_id:10612 Myxococcus virescens str. Mx v1 AJ2339
AJ233909.1 gg_id:10600 Angiococcus disciformis str. d1 AJ2339
EU262997.1 gg_id:276309 Myxococcus macrosporus str. Myxo9736 EU262
DQ768123.1 gg_id:163033 Pyxicoccus fallax str. DSM 14698 DQ768
GQ246547.1 gg_id:579922 Corallococcus macrosporus str. HBUM93406 GQ246
GQ246552.1 gg_id:534207 Corallococcus macrosporus str. HBUM93407 GQ246
GU139157.1 gg_id:589401 Myxococcus virescens str. HBUM91069 GU139
AJ233930.1 gg_id:10615 Myxococcus xanthus str. Mx x132 AJ2339
AJ811621.1 gg_id:102087 Corallococcus coralloides str. DSM 52496 AJ8116
GQ246541.1 gg_id:539091 Myxococcus virescens str. HBUM93307 GQ246
EU262992.1 gg_id:277585 Myxococcus fulvus str. 0189-5 EU262
GQ246525.1 gg_id:540091 Myxococcus fulvus str. HBUM93175 GQ246
AJ233917.1 gg_id:10602 Myxococcus fulvus str. Mx f2 AJ2339
AJ811622.1 gg_id:102241 Corallococcus coralloides str. DSM 52498 AJ8116
AJ233920.1 gg_id:10605 Myxococcus macrosporus str. Cc m7 AJ2339
AY724797.1 gg_id:104738 Myxococcus xanthus str. UCDaV1 AY724
AJ233925.1 gg_id:10610 Myxococcus virescens str. Mx v2 AJ2339
EU158297.1 gg_id:247490 Myxococcus sp. str. HBUM 91040 EU158
GQ246558.1 gg_id:532677 Myxococcus fulvus str. HBUM93132 GQ246
AF280855.1 gg_id:73433 bioreactor clone mle1-22 AF280
EU283398.1 gg_id:253563 consecutively aerated submerged treating domestic wastewater activated sludge m EU283
ABCS01000137.1 gg_id:235635 Plesiocystis pacifica str. SIR-1 ABCS0
EU134550.1 gg_id:256957 tallgrass prarie soil clone FFCH2652 EU134
AY921891.1 gg_id:135973 farm soil clone AKYH963 AY921
FJ152686.1 gg_id:304479 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX1A_ FJ1526
EU050860.1 gg_id:245344 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_05_87 EU050
CU466781.1 gg_id:327106 Evry wastewater treatment plant anoxic basin clone CU466
FJ203639.1 gg_id:313265 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
DQ298325.1 gg_id:154699 studies on and nifH Lasiurus sindicus Sewan grass Thar Desert Rajasthan India rhizos DQ298
AB252740.1 gg_id:152915 Antimicrobial Cyclic Depsipeptides SMH-27-4A coastal soil AB252
AM997796.1 gg_id:546538 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EU236309.1 gg_id:298699 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg1a2H2 EU236
AJ233941.1 gg_id:10642 Chondromyces robustus str. Cm a22 AJ2339
NR_025346.1 gg_id:565046 Chondromyces robustus str. Cm a13; DSM 14608 NR_02
DQ491070.1 gg_id:170538 Chondromyces pediculatus str. BICC 8609 DQ491

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AJ233938.1 gg_id:10639 Chondromyces apiculatus str. Cm a14 AJ2339
AY079453.1 gg_id:29731 Polyangium cellulosum str. So0089-1 AY0794
AF387627.1 gg_id:26373 Polyangium cellulosum str. 9741 AF3876
EU240490.1 gg_id:277159 Sorangium cellulosum str. So0184-1 EU2404
FJ479176.1 gg_id:580635 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4791
AY921933.1 gg_id:113748 farm soil clone AKYG909 AY9219
EU193074.1 gg_id:369028 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU1930
EU881272.1 gg_id:355920 (long-term grazing and burned annual) Karst region serious disturbance field soil clor EU8812
EU134398.1 gg_id:257391 tallgrass prarie soil clone FFCH6364 EU1343
FJ479473.1 gg_id:579806 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4794
DQ295908.1 gg_id:145144 Cohn's Crenothrix filamentous oxidizer unusual methane monooxygenase physical e DQ295
FJ660562.1 gg_id:551380 Correlation functional stability and dynamics full-scale wastewater treatment plant a FJ6605
EF516308.1 gg_id:218711 grassland soil clone FCPO632 EF5163
AJ617907.1 gg_id:109755 oxic-anoxic interphase flooded paddy soil clone D142119 AJ6179
EU134486.1 gg_id:245900 tallgrass prarie soil clone FFCH15612 EU1344
FJ478737.1 gg_id:535402 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4787
AJ833647.1 gg_id:108677 Byssophaga cruenta str. c2 = DSM 14553 AJ8336
AY493915.1 gg_id:136914 Fingerprinting aggregates soil clone 604 AY4939
AJ233948.1 gg_id:10657 Myxococcales str. NOSO-1 AJ2339
AM490752.1 gg_id:588131 Linking and functional nutrient spiraling mats (USA) microbial mat sulfidic cave sprir AM490
EU104167.1 gg_id:347524 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1041
EU979074.1 gg_id:326414 Effects intercropping Rhizobium inoculation on microbial biomass and rhizosphere b EU9790
AF280858.1 gg_id:71817 bioreactor clone mle1-29 AF2808
EF018417.1 gg_id:203878 aspen rhizosphere clone Amb_16S_1034 EF0184
AB238772.1 gg_id:146819 shal subsurface Sarobetsu Mire peat soil clone HSM-SS-009 AB2387
EF580966.1 gg_id:278697 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1 EF5809
EF112185.1 gg_id:187653 Stigmatella koreensis str. KYC-1019 EF1121
EU134339.1 gg_id:240506 tallgrass prarie soil clone FFCH3210 EU1343
EU134451.1 gg_id:250244 tallgrass prarie soil clone FFCH12289 EU1344
EU134540.1 gg_id:237206 tallgrass prarie soil clone FFCH18737 EU1345
FJ671175.1 gg_id:511742 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6711
AY725263.1 gg_id:102557 decayed velvetleaf seed clone S1-4-CL5 AY7252
DQ768124.1 gg_id:170014 Hyalangium minutum str. DSM 14724 DQ768
DQ129589.1 gg_id:137984 microarray level air two cities Texas urban aerosol clone AKIW811 DQ129
AM697153.1 gg_id:221410 sequencing libraries indoor dust clone BF0002B103 AM697
GQ007850.1 gg_id:501815 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ007
FJ152674.1 gg_id:336745 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX1A_ FJ1526
GQ264111.1 gg_id:568834 Structure Simulated Level Waste Site simulated level waste site clone WC3_40 GQ264
DQ768110.1 gg_id:172129 Cystobacter gracilis str. DSM 14753 DQ768

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AB278147.1 gg_id:178175 hydrothermal vent prokaryotes trapped 0.2- and 0.1-micro-m-filtrates obtained Mar AB2781
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EU734958.1 gg_id:328791 assemblages sediment station DBSE Northern Bering Sea clone 051C67 EU7349
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AF317744.1 gg_id:136925 Nullarbor cave clone wb1_A15 AF3177
EU287161.1 gg_id:235691 Pacific arctic surface sediment clone P13-68 EU2871
AB252955.1 gg_id:170623 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 127 AB2529
FM253643.1 gg_id:334324 Rock gold mine unusually microbial biofilm clone C01-2 FM253
EU048674.1 gg_id:250443 and archaeal surface marine sediment Slope South China Sea clone MD2896-B15 EU0486
FJ416121.1 gg_id:357893 sediments sediment station NEC5 Northern Bering Sea clone 107N50 FJ4161
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EU287148.1 gg_id:236118 Pacific arctic surface sediment clone P13-55 EU2871
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EU734974.1 gg_id:325961 assemblages sediment station DBSE Northern Bering Sea clone 068C78 EU7349
EU287128.1 gg_id:250580 Pacific arctic surface sediment clone P13-35 EU2871
EU290680.1 gg_id:253113 microbial structure biogeochemical contrasting shelf sediments Namibian upwelling EU2906
EF208656.1 gg_id:200746 Vertical and horizontal structure microbial sandy carbonate sediment clone CI35cm.2 EF2086
EU734976.1 gg_id:325676 assemblages sediment station DBSE Northern Bering Sea clone 070C81 EU7349

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EU050853.1 gg_id:246832 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_08_44 EU050
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EU050852.1 gg_id:236169 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_01_50 EU050
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AF154103.1 gg_id:10724 hydrocarbon seep clone GCA047 AF154
DQ331004.1 gg_id:159301 Guerrero Negro hypersaline microbial mat clone 10D2Z33 DQ331
AM490745.1 gg_id:592584 Linking and functional nutrient spiraling mats (USA) microbial mat sulfidic cave sprir AM490
GU362952.1 gg_id:555750 and archaeal biogeochemical cycling genes within Qiongdongnan basin Sea: pilot stu GU362
AF154094.1 gg_id:10677 hydrocarbon seep clone BPC065 AF154
EU246004.1 gg_id:253746 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P1-D07 EU246
EU360307.1 gg_id:277150 microbial fuel cell enriched sediment clone FWS 711 EU360
DQ109919.1 gg_id:143327 Sulfate-reducing mats: Changing paradigms new discoveries salt pond nonlithifying DQ109
DQ676442.1 gg_id:217352 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
CP001322.1 gg_id:514785 Desulfatibacillum alkenivorans str. AK-01 CP001
AY493562.1 gg_id:103090 Desulfatibacillum alkenivorans str. PF2803 AY493

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AY184360.2 gg_id:79514 Desulfatibacillum aliphaticivorans str. CV2803 AY1843
AF418175.1 gg_id:74171 Desulfobacter curvatus str. DSM 3379 AF4181
NR_025348.1 gg_id:572343 Desulfobacterium anilini str. Ani1; DSM 4660 NR_02
CP001087.1 gg_id:523103 Desulfobacterium autotrophicum str. HRM2 CP001
AY542197.1 gg_id:113166 Gulf Mexico seafloor sediments clone GoM HDB-06 AY542
AF026998.1 gg_id:10771 hot spring clone OPB16 AF0269
AY771951.1 gg_id:111847 marine surface sediment clone SI28 AY7719
GU362956.1 gg_id:533418 and archaeal biogeochemical cycling genes within Qiongdongnan basin Sea: pilot stu GU362
EU385666.1 gg_id:262547 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU3856
AF029045.1 gg_id:10904 Benzene mineralizing consortium complete clone SB-21 consortium AF0290
FJ716478.1 gg_id:566769 Uncultivated archaea sulfate-reducing biofilm Frasassi cave system anoxic lake water FJ7164
FJ485020.1 gg_id:566534 microbial autonomous rover world's deepest phreatic sinkhole biomat El Zacaton 114 FJ4850
GQ356929.1 gg_id:544180 Metal Dependent Marine Methane Oxidation methane seep sediment clone Fe_B_1 GQ356
AF029048.1 gg_id:10906 Benzene mineralizing consortium complete clone SB-30 consortium AF0290
DQ330997.1 gg_id:159166 Guerrero Negro hypersaline microbial mat clone 09D2Z19 DQ330
AY592183.1 gg_id:99181 marine ANAMMOX sediment clone AY592183 AY5921
FJ628304.1 gg_id:564870 transition and zone within near-shore basin: brackish water anoxic fjord Nitinat Lake FJ6283
FN396628.1 gg_id:565202 constant flux thermophilic cold seabed Arctic marine surface sediment clone s5_0_II FN3966
FJ437822.1 gg_id:535056 Benthic okenone production Fayetteville New York Green Lake chemocline 20.5 m wa FJ4378
DQ109878.1 gg_id:142790 Sulfate-reducing mats: Changing paradigms new discoveries salt pan lithifying micro DQ109
AB177158.1 gg_id:132816 seafloor methane hydrate clone ODP1230B17.18 AB1771
GU180164.1 gg_id:540242 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
AM086120.1 gg_id:145576 Abundance and archaea mesotrophic Lake Kinneret lake profundal sediment clone AM086
AM745206.1 gg_id:294510 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM745
EU385798.1 gg_id:274378 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU3857
EU385717.1 gg_id:276920 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU3857
EU925857.1 gg_id:328502 sediments sediment station DBS1 northern Bering Sea clone 052E88 EU9258
FM164951.2 gg_id:395222 two Bulgarian geothermal springs hot spring clone S1-53 FM164
DQ811822.1 gg_id:164969 mangrove soil clone MSB-5A4 DQ811
EU245381.1 gg_id:238545 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M1-A0 EU2453
AY542227.1 gg_id:113116 Gulf Mexico seafloor sediments clone GoM IDB-15 AY5422
AM229200.1 gg_id:575796 Microbial vicinity and within hydrates Hydrate Ridge (Cascadia Margin) marine gas h AM229
EU385917.1 gg_id:270552 Stratified microbes sediments Core MD05-2902 subseafloor sediment South China S EU3859
GU302479.1 gg_id:557390 Spatial structure and activity sedimentary microbial underlying Beggiatoa spp. mat h GU302
EU592476.1 gg_id:592826 Seasonal changes Salton Sea hypersaline sediment clone SSS17A EU5924
DQ521799.1 gg_id:159904 Anaerobic Methane Oxidizing ANME-1b Archaea Hypersaline Sediments Gulf Mexico DQ521
FN429786.1 gg_id:548583 Biogeochemistry -activity cold seep Larsen B area western Weddell Sea Antarctica m FN4297
FJ264650.1 gg_id:339335 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedim FJ2646
DQ676438.1 gg_id:221104 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
DQ925877.1 gg_id:252087 Microbes and Thermal Gradients Temperature Hydrothermal Chimneys hydrotherm DQ925
AB015244.1 gg_id:10939 deepest cold-seep area Japan Trench clone JTB133 proteobacterium AB0152

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AY197382.1 gg_id:86333 Guaymas Basin hydrothermal vent sediments clone B01R011 AY1973
AY197376.1 gg_id:86388 Guaymas Basin hydrothermal vent sediments clone B01R004 AY1973
AJ441316.1 gg_id:67542 Desulfobacula toluolica str. DSM 7467 AJ4413
CP000859.1 gg_id:255870 Desulfococcus oleovorans str. Hxd3 CP000
DQ103668.1 gg_id:132520 hypersaline endoevaporitic microbial mat clone E4aE11 DQ103
GQ249552.1 gg_id:546119 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 12-14 cm layer cl GQ249
EU246073.1 gg_id:238062 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P2-H01 EU246
DQ397423.1 gg_id:159502 Guerrero Negro hypersaline microbial mat clone 04D2Z68 DQ397
GQ249596.1 gg_id:544699 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 12-14 cm layer cl GQ249
FJ712553.1 gg_id:521361 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7125
AM176871.1 gg_id:160902 Recovery mangrove sediment clone SZB46 AM176
EU245312.1 gg_id:257695 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H5-H01 EU2453
EU245273.1 gg_id:256695 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H5-B05 EU2452
AF513951.1 gg_id:111956 hypersaline lake clone LA30-B27 AF5139
AJ620510.1 gg_id:108098 Olavius crassitunicatus endosymbiont AJ6205
EU245134.1 gg_id:241850 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H2-G09 EU2451
AF099063.1 gg_id:10900 Desulfofaba gelida str. PSv29 AF0990
AB015240.1 gg_id:10907 deepest cold-seep area Japan Trench clone JTB16 AB0152
NR_028732.1 gg_id:592871 Desulfofrigus fragile str. LSv21; DSM 12345 NR_02
AB189366.1 gg_id:113570 cold seep sediment clone JT75-121 AB1893
FM242377.1 gg_id:349097 sequences sediment clone 19 T9d+oil FM242
EF125385.1 gg_id:194587 soil naked tidal flat sediment near mangrove clone MSB-1ax8 EF1253
FM242229.1 gg_id:322751 sequences sediment clone 96 T0h-oil FM242
AB530178.1 gg_id:583955 Monitoring Suez and Tokyo bays based on analyses marine sediment clone Suez..Ba AB5301
DQ351771.1 gg_id:147847 Microbial Adherent Sediment Particles Heavy Metal Contaminated North Sea Surfac DQ351
EF442994.1 gg_id:248725 Desulfotignum sp. str. DSM 7120 EF4429
AF233370.1 gg_id:31826 Desulfotignum balticum str. Sax (DSM 7044) marine mud AF2333
EU047532.1 gg_id:251611 Anaerobic degradation benzene marine and cell hybridization dominant phylotype b EU0475
EU047529.1 gg_id:235729 Anaerobic degradation benzene marine and cell hybridization dominant phylotype b EU0475
AM882605.1 gg_id:330092 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
EU386050.1 gg_id:264473 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU3860
AB240476.1 gg_id:143077 microbial structure rhizosphere biofilm Sapporo root-tip (0 40 mm) Phragmites Sosei AB2404
EU048686.1 gg_id:245974 and archaeal surface marine sediment Slope South China Sea clone MD2896-B29 EU0486
EF454380.2 gg_id:222931 termite hindgut clone 290cost002-P3L-2033 EF4543
EU487868.1 gg_id:561085 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4878
EU885093.1 gg_id:305714 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone A2A5 EU8850
FJ712462.1 gg_id:526057 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7124
EF454653.2 gg_id:226987 termite hindgut clone 290cost002-P3L-2488 EF4546
EU622291.1 gg_id:289758 syntrophic partnerships deep-sea vents direct cell capture and metagenomics Eel Ri EU6222
AM490751.1 gg_id:537744 Linking and functional nutrient spiraling mats (USA) microbial mat sulfidic cave sprir AM490

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AY542569.1 gg_id:113399 Gulf Mexico seafloor sediment clone GoM GB425 12B-61 AY5425
FJ484450.1 gg_id:571089 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4844
AB234533.1 gg_id:145609 termite gut homogenate clone MgMjD-089 AB2345
AB240382.1 gg_id:150612 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB2403
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AM882591.1 gg_id:327370 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
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EU246034.1 gg_id:249930 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P1-H08 EU2460
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AM086124.1 gg_id:146208 Abundance and archaea mesotrophic Lake Kinneret lake profundal sediment clone AM086
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EU399670.1 gg_id:273059 and localization microbial consortium full-scale granular activated carbon anaerobic EU3996
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AB188779.1 gg_id:113838 cold seep sediment clone SB3-7 AB1887
FJ789839.1 gg_id:534671 Desulfobotulus sapovorans str. ATCC 33892; 1pa3 FJ7898
FM242270.1 gg_id:314257 sequences sediment clone 4 T0h-oil FM242
EU160523.1 gg_id:352459 diagnosis central-venous infections cultivation-dependent and techniques biofilm sc EU1605
FJ202141.1 gg_id:314793 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2021
EU385675.1 gg_id:272166 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU3856
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DQ397412.1 gg_id:159381 Guerrero Negro hypersaline microbial mat clone 02D2Z96 DQ397
DQ811836.1 gg_id:173385 mangrove soil clone MSB-5F12 DQ811
AJ937698.1 gg_id:143680 Microbiological investigation methane- and hydrocarbon-discharging volcanoes Carp AJ9376
AY222304.1 gg_id:114278 host hindgut caecum clone Del 1 AY2223
AM882599.1 gg_id:326082 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
AB355082.1 gg_id:237712 Monitoring and archaeal two heavy polluted areas lake Egypt based on sediment Ma AB3550
EU622295.1 gg_id:294941 syntrophic partnerships deep-sea vents direct cell capture and metagenomics Eel Riv EU6222
FJ716993.1 gg_id:544106 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7169
DQ630689.1 gg_id:159784 Unexpected many-celled magnetotactic prokaryote salt marsh clone mmp5 DQ630
EU570889.1 gg_id:556574 transition and zone within near-shore anoxic basin: Nitinat Lake depth 50 m clone Ni EU5708
U85470.1 gg_id:10934 Desulfobotulus sp. str. BG14 U85470

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DQ397424.1 gg_id:159185 Guerrero Negro hypersaline microbial mat clone 04D2Z24 DQ397
AM882642.1 gg_id:320845 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
EU592487.1 gg_id:563232 Seasonal changes Salton Sea hypersaline sediment clone SSS3A EU592
FJ712404.1 gg_id:520767 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) K FJ7124
NR_025406.1 gg_id:544630 Desulfococcus biacutus str. DSM 5651 NR_02
DQ831540.1 gg_id:165612 Activity and sulphate-reducing contrasting sediments River Colne estuary UK marine DQ831
AB294926.1 gg_id:225575 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma AB294
EU290724.1 gg_id:253758 microbial structure biogeochemical contrasting shelf sediments Namibian upwelling EU290
FM242446.1 gg_id:341474 sequences sediment clone 50 T12d+oil FM242
EU386062.1 gg_id:275331 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
FM165241.1 gg_id:557184 seep chitineous tube vestimentiferan tubeworm Lamellibrachia sp. cold seeps clone FM165
FJ973581.1 gg_id:378495 Periodic sulfide irruptions impact microbial structure and hypersaline lake water colu FJ9735
FJ712482.1 gg_id:528626 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) K FJ7124
FJ485367.1 gg_id:566855 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4853
AY082457.1 gg_id:67851 neutral pH mine biofilm clone 44a-B1-10 AY082
DQ630675.1 gg_id:160459 Unexpected many-celled magnetotactic prokaryote salt marsh clone mmp23 DQ630
EF125398.1 gg_id:189923 soil naked tidal flat sediment near mangrove clone MSB-1cx7 EF1253
AJ704678.1 gg_id:109279 marine sediment clone HMMVPog-2 AJ7046
DQ415842.1 gg_id:157733 Dominant microbial limestone-corroding biofilms frasassi system Italy Frasassi sulfid DQ415
FN396636.1 gg_id:541666 constant flux thermophilic cold seabed Arctic marine surface sediment clone s5_0_II FN396
GU230401.1 gg_id:554116 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal sediment clo GU230
AY216443.1 gg_id:81354 temperate estuarine mud clone KM88 AY216
DQ630707.1 gg_id:160788 Unexpected many-celled magnetotactic prokaryote salt marsh clone mmp2_2 DQ630
DQ521813.1 gg_id:160182 Anaerobic Methane Oxidizing ANME-1b Archaea Hypersaline Sediments Gulf Mexic DQ521
FM242231.1 gg_id:317203 sequences sediment clone 189 T0h-oil FM242
FJ716967.1 gg_id:545744 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7169
AJ704696.1 gg_id:109295 marine sediment clone HMMVBeg-44 AJ7046
GQ356986.1 gg_id:581992 Metal Dependent Marine Methane Oxidation methane seep sediment clone FeSO4_ GQ356
GQ249585.1 gg_id:587834 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 12-14 cm layer c GQ249
AM882600.1 gg_id:306625 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
FJ814756.1 gg_id:556875 Microbial assemblages Thioploca sheaths cold seep settings Monterey Bay Carmel Ca FJ8147
EU487979.1 gg_id:574128 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU487
AJ237607.1 gg_id:10870 Desulfobacterium indolicum str. DSM 3383 marine mud AJ2376
AY771939.1 gg_id:111869 marine surface sediment clone SI29 AY771
EU385785.1 gg_id:272732 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU385
GQ433957.1 gg_id:590151 Subseafloor ocean crust microbial observatories: (F-through Osmo Colonization Syst GQ433
AY592200.1 gg_id:99952 marine ANAMMOX sediment clone AY592200 AY592
FJ517109.1 gg_id:549171 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled water c FJ5171
EU385687.1 gg_id:274595 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU385
AM882587.1 gg_id:340294 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882

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EF125423.1 gg_id:184819 mangrove soil clone MSB-2A5 EF1254
AY053490.1 gg_id:39986 gas hydrate clone AT425_EubD9 AY0534
AJ620501.1 gg_id:353759 Olavius 3 endosymbiont O. ilvae Individual 5 AJ6205
FJ746111.1 gg_id:513493 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7461
DQ811817.1 gg_id:172929 mangrove soil clone MSB-4G8 DQ811
FM242225.1 gg_id:337177 sequences sediment clone 342 T0h-oil FM242
AB237684.1 gg_id:139771 microorganisms deep subsurface groundwater sedimentary rock milieu clone HDBW AB237
AY082461.1 gg_id:67845 neutral pH mine biofilm clone 44a-B1-17 AY0824
GU302442.1 gg_id:552223 Spatial structure and activity sedimentary microbial underlying Beggiatoa spp. mat GU302
AM774314.1 gg_id:243934 Cultivation and situ quantification consistently demonstrate that two distinct group AM774
EF999398.1 gg_id:237632 Vertical and Methanotrophs Estuarine Sediments Pearl River Estuary sediments 50cn EF9993
DQ630706.1 gg_id:161002 Unexpected many-celled magnetotactic prokaryote salt marsh clone mmp2_28 DQ630
AJ241022.1 gg_id:10909 Norway:(Svalbard)Hornsund clone Sva0405 AJ2410
DQ831553.1 gg_id:163459 Activity and sulphate-reducing contrasting sediments River Colne estuary UK marine DQ831
AJ831748.1 gg_id:102310 chemocline clone 166 AJ8317
DQ146482.1 gg_id:164141 Desulfobacterium sp. MB-2005 DQ146
DQ630702.1 gg_id:160395 Unexpected many-celled magnetotactic prokaryote salt marsh clone mmp2_20 DQ630
FJ716406.1 gg_id:571102 Uncultivated archaea sulfate-reducing biofilm Frasassi cave system anoxic lake water FJ7164
AY851291.1 gg_id:148326 Algidimarina propionica str. AK-P AY8512
EF688206.1 gg_id:347101 Detection microbial sludge anaerobic wastewater treatment system clone 5A4_cons EF6882
EU717681.1 gg_id:425208 Ultrastructure tactic behaviour and potential sulfate reduction multicellular magnet EU717
U81720.2 gg_id:10893 anaerobic digestor clone vadinHA60 U8172
GQ356954.1 gg_id:576546 Metal Dependent Marine Methane Oxidation methane seep sediment clone Fe_B_1 GQ356
DQ811793.1 gg_id:173340 mangrove soil clone MSB-3ax11 DQ811
AB177130.1 gg_id:130033 seafloor methane hydrate clone ODP1230B1.06 AB177
AM774320.1 gg_id:248513 Cultivation and situ quantification consistently demonstrate that two distinct group AM774
EU160501.1 gg_id:325508 diagnosis central-venous infections cultivation-dependent and techniques biofilm sc EU160
U45991.1 gg_id:10897 Desulfonema ishimotonii str. DSM 9679 anaerobic intertidal mud flats Jadebusen (North U4599
EU617853.1 gg_id:345458 Seasonal variation microbial Yel Sea sediment clone D13S-20 EU6178
EU246163.1 gg_id:248972 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P4-E10 EU246
AM404371.1 gg_id:558344 Hydrocarbon-fueled sulfate reduction can sustain long-lived tubeworm aggregation AM404
AJ620497.1 gg_id:354146 Olavius 4 endosymbiont O. algarvensis Individual 2 AJ6204
DQ811815.1 gg_id:167112 mangrove soil clone MSB-4G4 DQ811
DQ811807.1 gg_id:166757 mangrove soil clone MSB-4C12 DQ811
DQ397413.1 gg_id:159387 Guerrero Negro hypersaline microbial mat clone 02D2Z34 DQ397
AJ831753.1 gg_id:106553 chemocline clone 293 AJ8317
DQ004679.1 gg_id:158084 Microbial Reduce Methane Efflux Mud Volcano Sediments Gulf Cadiz marine hydroc DQ004
U85477.1 gg_id:10775 Desulfoarculus sp. str. BG74 U8547
EF061952.1 gg_id:182353 Xiamen China mangrove sediment clone XME29 EF0619
AM882597.1 gg_id:342977 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882

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DQ394960.1 gg_id:155401 harbor sediment clone VHS-B3-73 DQ394
EU542523.1 gg_id:267799 Effects chemical structure concentration on pathways microbial during dechlorinatio EU542
EF125387.1 gg_id:193460 soil naked tidal flat sediment near mangrove clone MSB-1B10 EF1253
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AJ535249.1 gg_id:77873 marine sediment above hydrate ridge clone Hyd89-61 proteobacterium AJ5352
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AF022936.1 gg_id:10867 Desulfocella halophila str. GSL-But2 AF022
CU927196.1 gg_id:554689 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU927
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EU488036.1 gg_id:560020 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
AM745204.1 gg_id:287932 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM745
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DQ811797.1 gg_id:167399 mangrove soil clone MSB-3C3 DQ811
GU080089.1 gg_id:578329 Combined and proteomic approaches identify involved anaerobic 2-methylnaphthal GU080
DQ397436.1 gg_id:159112 Guerrero Negro hypersaline microbial mat clone 062DZ56 DQ397
AF418174.1 gg_id:71864 Desulfovibrio baarsii str. DSM 2075 AF418
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AM490746.1 gg_id:591624 Linking and functional nutrient spiraling mats (USA) microbial mat sulfidic cave sprir AM490
AM176857.1 gg_id:160904 Recovery mangrove sediment clone SZB57 AM176
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AM040123.1 gg_id:147720 sandy sediments clone Sylt 27 AM040
EU617854.1 gg_id:316203 Seasonal variation microbial Yel Sea sediment clone C13S-20 EU617
EU048693.1 gg_id:250681 and archaeal surface marine sediment Slope South China Sea clone MD2896-B42 EU048
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FJ484778.1 gg_id:542328 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4847
DQ811819.1 gg_id:169367 mangrove soil clone MSB-4H12 DQ811
DQ811816.1 gg_id:166688 mangrove soil clone MSB-4G6 DQ811
DQ404782.1 gg_id:155938 contaminated sediment clone 661234 DQ404
AF002671.1 gg_id:10776 Desulfobacca acetoxidans AF002
FJ902442.1 gg_id:548667 phreatic limestone sinkholes northeastern Mexico water column surface cenote Cara FJ9024
FJ810575.1 gg_id:373867 Subsurface ecosystem resilience: Long-term attenuation subsurface contaminants su FJ8105

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AF523963.1 gg_id:101995 forested wetland clone FW106 AF523
FJ902422.1 gg_id:586511 phreatic limestone sinkholes northeastern Mexico water column surface cenote Cara FJ9024
FJ484032.1 gg_id:537190 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4840
FJ484530.1 gg_id:554210 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4845
DQ404621.1 gg_id:152758 contaminated sediment clone 656066 DQ404
AB426195.1 gg_id:268751 -situ identification study benzene-degrading s under methanogenic condition anaerc AB426
AY340836.1 gg_id:91627 sulfate-reducing fluidized-bed reactor clone SR_FBR_L13 AY340
AB262720.1 gg_id:239069 microorganisms shal subsurface Sarobetsu mire peat soil clone HSM050P-B-9 AB262
FJ516878.1 gg_id:558853 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled upper s FJ5168
DQ676381.1 gg_id:221935 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
AM490653.1 gg_id:580869 Linking and functional nutrient spiraling mats (USA) microbial mat sulfidic cave sprir AM490
FJ437843.1 gg_id:511708 Benthic okenone production Fayetteville New York Green Lake 25 m water depth clor FJ4378
AF282177.1 gg_id:30060 Desulfomonile limimaris str. DCB-F AF282
AF282178.1 gg_id:31004 marine sediment clone Btol AF282
AJ316021.1 gg_id:42269 Desulfomonile sp. clone 624 AJ316
AM086646.1 gg_id:140135 Desulfomonile tiedjei str. DSM 6799 AM086
DQ133909.1 gg_id:164530 Dominant microbial biofilms frasassi cave system Italy limestone-corroding stream b DQ133
CT574193.1 gg_id:151947 Evry municipal wastewater treatment plant clone 060B12_B_SD_P93 CT5741
CU917968.1 gg_id:562603 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU917
CT573954.1 gg_id:153501 Evry municipal wastewater treatment plant clone 009F10_B_SD_P15 CT5739
CT573960.1 gg_id:155759 Evry municipal wastewater treatment plant clone 009F03_B_SD_P15 CT5739
FJ462108.1 gg_id:581591 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4621
EU407209.1 gg_id:275861 household biogas digester clone GW-26 EU407
CT574120.1 gg_id:155635 Evry municipal wastewater treatment plant clone 061D05_B_SD_P93 CT5741
AJ133795.1 gg_id:10858 Syntrophus sp AJ1337
EU104840.1 gg_id:265855 sulfate-reducing and methanogen sulfate-fed reactors libraries anaerobic biofilm rea EU1048
CT574027.1 gg_id:151989 Evry municipal wastewater treatment plant clone 008C04_B_SD_P15 CT5740
CT574029.1 gg_id:156244 Evry municipal wastewater treatment plant clone 008C02_B_SD_P15 CT5740
AB252682.1 gg_id:168756 and activities propionate-oxidizing that affected propionate concentration anaerobic AB2526
CR933148.1 gg_id:110803 Evry municipal wastewater treatment plant clone 012B10_B_SD_P15 CR9331
CU919519.1 gg_id:547047 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU919
DQ676391.1 gg_id:227149 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
CT574164.1 gg_id:152984 Evry municipal wastewater treatment plant clone 060F04_B_SD_P93 CT5741
DQ501304.1 gg_id:187961 Seasonal dynamics free-living and particle- four lakes northeastern Germany freshw DQ501
GU202945.1 gg_id:592840 Effect pH on anode microbes MFC microbial fuel cell inoculated activated sludge wa: GU202
CT573944.1 gg_id:154331 Evry municipal wastewater treatment plant clone 009H01_B_SD_P15 CT5739
EU407210.1 gg_id:276971 household biogas digester clone GW-27 EU407
EU522631.1 gg_id:262404 hydrocarbon-degrading methanogenic microbial consortia settling basin Northern Al EU5226
AY661413.1 gg_id:107448 anaerobic hybrid reactor clone TTA_H37 AY6614
AB234260.1 gg_id:145632 Biotransformation Polychlorinated Dioxins Microbial Dynamics Sediment Microcosm AB2342

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EU399668.1 gg_id:271079 and localization microbial consortium full-scale granular activated carbon anaerobic EU399
AB186854.1 gg_id:99369 polychlorinated dioxin dechlorinating microcosm clone TSBZ03 AB186
AF229793.1 gg_id:37165 terephthalate-degrading consortium clone TA20 AF229
AJ306771.1 gg_id:66928 DCP-dechlorinating consortium clone SHA-42 AJ3067
AB288891.1 gg_id:203169 termite gut homogenate clone BOf2-22 AB288
FJ462058.1 gg_id:593849 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4620
EU522634.1 gg_id:269542 hydrocarbon-degrading methanogenic microbial consortia settling basin Northern Al EU522
DQ521796.1 gg_id:159708 Anaerobic Methane Oxidizing ANME-1b Archaea Hypersaline Sediments Gulf Mexic DQ521
AB179694.1 gg_id:102521 groundwater 0.2 m-captured fraction clone KNA6-EB24 AB179
DQ521792.1 gg_id:160618 Anaerobic Methane Oxidizing ANME-1b Archaea Hypersaline Sediments Gulf Mexic DQ521
FJ902117.1 gg_id:579169 phreatic limestone sinkholes northeastern Mexico orange biomat 8m deep cenote Ca FJ9021
AJ535254.1 gg_id:77896 marine sediment above hydrate ridge clone Hyd89-52 proteobacterium AJ5352
FJ462062.1 gg_id:587689 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4620
AM746089.1 gg_id:299409 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM746
DQ811814.1 gg_id:173034 mangrove soil clone MSB-4G3 DQ811
CU922340.1 gg_id:534010 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU922
FN429788.1 gg_id:546409 Biogeochemistry -activity cold seep Larsen B area western Weddell Sea Antarctica m FN429
AB252432.1 gg_id:153221 Microbial extremely CO2 rich deep-sea sediment Marine sediments Yonaguni Knoll I AB252
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DQ067032.1 gg_id:133542 Lake Washington sediment clone pLW-41 DQ067
DQ404844.1 gg_id:150930 contaminated sediment clone 655974 DQ404
FJ902148.1 gg_id:592355 phreatic limestone sinkholes northeastern Mexico orange biomat 8m deep cenote Ca FJ9021
DQ415841.1 gg_id:158424 Dominant microbial limestone-corroding biofilms frasassi system Italy Frasassi sulfid DQ415
AF420337.1 gg_id:67800 hydrothermal sediment clone AF420337 AF4203
GU180161.1 gg_id:538685 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
EU101287.1 gg_id:237813 Sulfide/oxygen supply ratio predicts outcome competition among sulfur-oxidizing wa EU101
EU542439.1 gg_id:274550 Effects chemical structure concentration on pathways microbial during dechlorinatio EU542
FN429791.1 gg_id:586738 Biogeochemistry -activity cold seep Larsen B area western Weddell Sea Antarctica m FN429
AY053494.1 gg_id:34985 gas hydrate clone AT425_EubF5 AY0534
FJ437824.1 gg_id:509369 Benthic okenone production Fayetteville New York Green Lake chemocline 20.5 m wa FJ4378
DQ988259.1 gg_id:185953 Microbial lab-scale A1-A2-O fixed biofilm system coking wastewater treatment with DQ988
FJ462006.1 gg_id:569259 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4620
AF354149.1 gg_id:71121 marine methane seep clone 1513 AF3541
AM712339.1 gg_id:252529 detection and ly microbial Brothers Seamount Kermadec Arc New Zealand biofilm h AM712
EU522632.1 gg_id:269237 hydrocarbon-degrading methanogenic microbial consortia settling basin Northern Al EU522
AB177179.1 gg_id:128221 seafloor methane hydrate clone ODP1230B23.08 AB177
CR933196.1 gg_id:110185 Evry municipal wastewater treatment plant clone 008B10_B_SD_P15 CR9331
AY542210.1 gg_id:114116 Gulf Mexico seafloor sediments clone GoM HDB-32 AY5422
AB426198.1 gg_id:259940 -situ identification study benzene-degrading s under methanogenic condition anaerc AB4261

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EU592411.1 gg_id:535455 Seasonal changes Salton Sea hypersaline sediment clone SSS15N EU592
EU644111.1 gg_id:280179 and Samoylov Island Lena active layer -centred polygon near permafrost table; Rim_ EU6441
AY211657.1 gg_id:95801 Gulf Mexico seafloor sediment clone GoM GC185 503E AY2116
AY211680.1 gg_id:96542 Gulf Mexico seafloor sediment clone GoM GC234 621E AY2116
AF482437.1 gg_id:77432 granular sludge clone R2b29 AF4824
FJ902092.1 gg_id:566804 phreatic limestone sinkholes northeastern Mexico orange biomat 8m deep cenote Ca FJ9020
DQ521786.1 gg_id:160206 Anaerobic Methane Oxidizing ANME-1b Archaea Hypersaline Sediments Gulf Mexic DQ521
FJ462010.1 gg_id:593726 Microbial ecology industrial digestor mesophilic anaerobic reactor fed effluent chemi FJ4620
AJ583188.1 gg_id:94853 radioactive site ground water clone S15B-MN12 AJ5831
EU644229.1 gg_id:282165 and Samoylov Island Lena active layer -centred polygon near surface horizon; Rim_T EU6442
AB525446.1 gg_id:570354 Segregated ANME-1 and -2 cold-seep marine sediments deep-sea methane seep sed AB5254
DQ787721.1 gg_id:162776 holocene lake sediment clone SLB628 DQ787
FJ902107.1 gg_id:580738 phreatic limestone sinkholes northeastern Mexico orange biomat 8m deep cenote Ca FJ9021
AM745217.1 gg_id:294031 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM745
AF385080.1 gg_id:28552 Desulfacinum subterraneum str. 101 AF3850
AF170418.1 gg_id:73225 Desulfacinum sp. str. M40/2 CIV-2.3 AF1704
DQ006288.1 gg_id:123413 Desulfatimicrobium mahresensis str. SA1 DQ006
FN356276.1 gg_id:558010 Prokaryotic structure sulfate reducer activity -temperature reservoirs and without ni FN3562
FN356315.1 gg_id:570270 Prokaryotic structure sulfate reducer activity -temperature reservoirs and without ni FN3563
NR_027598.1 gg_id:544627 Syntrophobacter fumaroxidans MPOB str. MPOB; DSM 10017 NR_02
AAJF01000012.1 gg_id:134597 Syntrophobacter fumaroxidans str. MPOB AAJF01
GQ261306.1 gg_id:543826 Study on Sediment sediment Dongping Lake clone 1S2-18 GQ261
AB232801.1 gg_id:138526 Microbial mesophilic chemostat cultivation propionate-degrading methanogenic con AB2328
FJ535519.1 gg_id:538615 Microbial digestion swine wastewater anaerobic UASB reactor clone SWADLP2-18 FJ5355
GU208398.1 gg_id:581775 Study on Sediment Dongping Lake sediment clone Fh1-17 GU208
GQ402720.1 gg_id:583853 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
CU920959.1 gg_id:535086 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU920
CU924062.1 gg_id:573309 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU924
CU921674.1 gg_id:543856 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU921
FJ484661.1 gg_id:571785 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4846
X70905.1 gg_id:10750 Syntrophobacter wolinii str. defined co Desulfovibrio sp. DSM 2805 X70905
EF515568.1 gg_id:222770 Electricigen Enrichment MFC full-scale anaerobic bioreactor sludge treating brewery EF5155
AB232820.1 gg_id:139158 Microbial mesophilic chemostat cultivation propionate-degrading methanogenic con AB2328
DQ415751.1 gg_id:157212 Dominant microbial limestone-corroding biofilms frasassi system Italy Frasassi sulfid DQ415
CU921434.1 gg_id:549201 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU921
FN646446.1 gg_id:589878 candidate division OP3 cells limonene Limonene degrading methanogenic s clone LiN FN6464
X94911.1 gg_id:10755 Syntrophobacter sp X94911
GQ402673.1 gg_id:585966 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
EF688226.1 gg_id:329449 Detection microbial sludge anaerobic wastewater treatment system clone 5E3_cons EF6882
DQ404834.1 gg_id:155072 contaminated sediment clone 655963 DQ404
U25627.1 gg_id:10762 Thermodesulforhabdus norvegica str. A8444 U2562

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EU134394.1 gg_id:237260 tallgrass prarie soil clone FFCH612 EU1343
GQ389113.1 gg_id:535437 quality deterioration drinking system during serious red water outbreak clone S67 GQ389
FJ205268.1 gg_id:558961 Prokaryotic deep-sea region East Lau Spreading Centre active hydrothermal field sedi FJ2052
EU373905.1 gg_id:267944 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment clc EU3739
GU208360.1 gg_id:560302 Study on Sediment Dongping Lake sediment clone Td4-7 GU208
EF019335.1 gg_id:206015 aspen rhizosphere clone Elev_16S_545 EF0193
FJ478988.1 gg_id:560191 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4789
EU335317.1 gg_id:248449 Changes microbial metabolic and along hydrogeochemically variable profile unsatura EU3353
AM936205.1 gg_id:559827 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
GQ472446.1 gg_id:568881 Microbial Dongping Lake lake sediment clone 5T1-3 GQ472
AF523967.1 gg_id:63257 forested wetland clone FW136 AF5239
AY963427.1 gg_id:152097 structures and correlated ecological factors soils two evergreen broad-leaved forests AY9634
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FJ479469.1 gg_id:572414 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4794
AB426233.1 gg_id:275505 -situ identification study benzene-degrading s under methanogenic condition anaerc AB4262
GU208380.1 gg_id:565064 Study on Sediment Dongping Lake sediment clone Td2-11 GU208
AB128874.1 gg_id:103274 deep marine sediments Nankai Trough Japan; ODP Leg 190 clone 1174-1091-4 AB1288
EU134559.1 gg_id:243792 tallgrass prarie soil clone FFCH3288 EU1345
AY921794.1 gg_id:113824 farm soil clone AKYG681 AY9217
FJ478660.1 gg_id:573531 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
GQ259289.1 gg_id:564500 Microbial activities coastal marine sediments and water column Svalbard deep sedir GQ259
AF523886.1 gg_id:55388 forested wetland clone RCP2-54 AF5238
FJ479115.1 gg_id:540681 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4791
FJ479389.1 gg_id:559874 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4793
AM072417.1 gg_id:138410 Untersuchung Verschiebung der bakteriellen Populationen mit U(VI) inokulierten Bc AM072
FJ479315.1 gg_id:575779 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4793
GQ302546.1 gg_id:581532 prokaryotic sediment Shawan methods cold spring clone sw-xj100 GQ302
GU056068.1 gg_id:552030 Biologic Techniques Applied Microbial Prospecting China: Ban 876 Gas Oil Field Bein GU056
FJ479231.1 gg_id:593037 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4792
AJ296569.1 gg_id:38614 uranium mining mill tailing clone GR-WP33-30 AJ2965
CU921195.1 gg_id:543214 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU921
GU118124.1 gg_id:547351 Threatened Corals Provide Microbial Habitats clone Apal_A09 GU118
EF600558.1 gg_id:302892 soil clone E5-2 EF6005
EU202780.1 gg_id:326405 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2027
FJ479565.1 gg_id:510105 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4795
AJ581602.1 gg_id:93483 soil near uranium mill tailings clone KCM-B-53 AJ5816
EF032748.1 gg_id:191331 cyanobacterial mat in lava cave clone HAVOmat120 EF0327
AM997325.1 gg_id:555867 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
FJ478489.1 gg_id:572174 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4784
GQ500719.1 gg_id:581516 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Charon's Cascade sa GQ500

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DQ811806.1 gg_id:165767 mangrove soil clone MSB-4C10 DQ811
GQ262847.1 gg_id:549915 Structure Simulated Level Waste Site simulated level waste site clone F1_28X2 GQ262
EF018612.1 gg_id:215602 aspen rhizosphere clone Amb_16S_891 EF0186
EF032749.1 gg_id:192369 cyanobacterial mat in lava cave clone HAVOmat133 EF0327
EF590024.2 gg_id:228859 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF5900
EU617849.1 gg_id:331603 Seasonal variation microbial Yel Sea sediment clone D13S-11 EU6178
GU061491.1 gg_id:560172 and function microbial mesoscale eddy perturbations South China Sea clone CE1-DC GU061
FJ478639.1 gg_id:593827 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
GU230392.1 gg_id:586588 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal sediment clo GU230
EF032751.1 gg_id:195011 cyanobacterial mat in lava cave clone HAVOmat82 EF0327
AB240314.1 gg_id:144627 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB2403
AJ519632.1 gg_id:77057 uranium mill tailings soil sample clone Sh765B-AG-71 proteobacterium AJ5196
AY289452.1 gg_id:89599 soil clone ABS-13 AY2894
FJ152850.1 gg_id:315633 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1528
AM997396.1 gg_id:572529 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
AM936195.1 gg_id:548041 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
EU050854.1 gg_id:254522 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_02_28 EU0508
AJ518789.1 gg_id:75247 uranium mining waste pile near Johanngeorgenstadt soil clone JG37-AG-2 AJ5187
EF018554.1 gg_id:205913 aspen rhizosphere clone Amb_16S_816 EF0185
EU386119.1 gg_id:276230 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU3861
EU134520.1 gg_id:240393 tallgrass prarie soil clone FFCH8074 EU1345
FJ755756.1 gg_id:578972 vertical archaeal water and Lake Taihu sediment clone LT-SB-B12 FJ7557
EU134328.1 gg_id:249749 tallgrass prarie soil clone FFCH11550 EU1343
EU202763.1 gg_id:324982 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2027
AB234261.1 gg_id:147174 Biotransformation Polychlorinated Dioxins Microbial Dynamics Sediment Microcosm AB2342
AB252957.1 gg_id:168547 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 023 AB2529
EU881148.1 gg_id:323943 Karst region primary forest soil clone KPF200711-152 EU8811
EF471464.1 gg_id:262091 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay EF4714
EF019530.1 gg_id:214844 aspen rhizosphere clone Elev_16S_1033 EF0195
GQ246353.1 gg_id:509231 Temporal and spatial China North Yel Sea sediments clone Cm1-16 GQ246
FJ479302.1 gg_id:535350 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4793
AB252435.1 gg_id:152635 Microbial extremely CO2 rich deep-sea sediment Marine sediments Yonaguni Knoll I' AB2524
EF076107.1 gg_id:244235 Sponge- microorganisms: ecology and biotechnological potential Bahamas: Little San EF0761
FJ478944.1 gg_id:512156 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4789
AY963383.1 gg_id:152964 structures and correlated ecological factors soils two evergreen broad-leaved forests AY9633
AF234748.1 gg_id:48147 sludge clone H16 AF2347
EF516526.1 gg_id:226074 grassland soil clone FCPP428 EF5165
AM935117.1 gg_id:565668 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
AM889194.1 gg_id:593414 structure superficial sediments Bizerte (Tunisia) southern Mediterranean coastal an AM889
EF125461.1 gg_id:182959 mangrove soil clone MSB-2H1 EF1254

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AF026987.1 gg_id:10763 hot spring clone OPB33 AF0269
EU700154.1 gg_id:294756 Study on Shantou Bay Marine Sediments Microorganisms sediment clone STU12 EU7001
EF442978.1 gg_id:257733 Desulforhabdus sp. str. DDT EF4429
EU542438.1 gg_id:262663 Effects chemical structure concentration on pathways microbial during dechlorinatio EU5424
DQ984662.1 gg_id:242560 Anaerobic degradation unsaturated or saturated long chain fatty acids (LCFA) prese DQ984
GQ487978.1 gg_id:542670 Microbial metal soil polluted heavy metals clone G19-236 GQ487
X83274.1 gg_id:10759 Desulforhabdus amnigena str. ASRB1 X83274
AJ237605.1 gg_id:10754 Syntrophobacter sp str. TsuA1 AJ2376
AM176868.1 gg_id:160457 Recovery mangrove sediment clone SZB17 AM176
GU180163.1 gg_id:578505 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
DQ303457.1 gg_id:151758 Desulfoglaeba alkanexedens str. ALDC DQ303
DQ676446.1 gg_id:219894 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
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AJ306776.1 gg_id:66924 DCP-dechlorinating consortium clone SHA-117 AJ3067
AM176858.1 gg_id:161119 Recovery mangrove sediment clone SZB9 AM176
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CU919420.1 gg_id:538867 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU9194
AY297973.1 gg_id:89577 thermophilic anaerobic terephthalate-degrading sludge clone TTA_B25 AY2979
CT573926.1 gg_id:153737 Evry municipal wastewater treatment plant clone 012B11_B_SD_P15 CT5739
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AF229787.1 gg_id:29387 terephthalate-degrading consortium clone TA14 AF2297
CT573826.1 gg_id:152222 Evry municipal wastewater treatment plant clone 013H09_B_SD_P15 CT5738
EU542505.1 gg_id:275595 Effects chemical structure concentration on pathways microbial during dechlorinatio EU5425
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EU399678.1 gg_id:275018 and localization microbial consortium full-scale granular activated carbon anaerobic EU3996
AJ009465.1 gg_id:10718 TCB-transforming consortium clone SJA-51 AJ0094
EF198029.1 gg_id:180225 Microbial structures and thermophilic methanogenic consortia anaerobic mesophilic EF1980
AF523969.1 gg_id:63254 forested wetland clone FW121 AF5239
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DQ513073.1 gg_id:160937 Microbial life fluids ridge flank crustal fluid clone FS266-12B-03 DQ513
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GQ348270.1 gg_id:546068 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
GQ348264.1 gg_id:590521 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
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GQ351086.1 gg_id:590469 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ351
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GQ348641.1 gg_id:571912 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
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AY048892.1 gg_id:23990 rhizosphere clone Q3-6C17 AY048
DQ501327.1 gg_id:198258 Seasonal dynamics free-living and particle- four lakes northeastern Germany freshw DQ501
EU050847.1 gg_id:237789 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_03_45 EU050
EU487900.1 gg_id:574097 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU487
EU491765.1 gg_id:261919 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU491
EF573310.1 gg_id:233830 Coco's Island site 23 marine clone S23_1409 EF5733
FJ155019.1 gg_id:330963 seasonal fluctuation coastal Bohai Bay sequencing and RFLP ocean water clone BX FJ1550
GQ348578.1 gg_id:576386 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
AF371949.1 gg_id:23625 swine intestine clone p-4193-6Wa5 AF371
FJ484808.1 gg_id:509157 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4848
EU220836.1 gg_id:274794 Spirobacillus cienkowskii EU220
GQ337121.1 gg_id:577297 Hydrography shapes biogeography deep Arctic Ocean water depth 395m clone CB0 GQ337
CU466767.1 gg_id:336493 Evry wastewater treatment plant anoxic basin clone CU466
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U70713.1 gg_id:4709 Cape Hatteras picoplankton clone OM27 U7071
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EU101270.1 gg_id:257793 Sulfide/oxygen supply ratio predicts outcome competition among sulfur-oxidizing wa EU101
DQ330714.1 gg_id:159120 Guerrero Negro hypersaline microbial mat clone 01D1q147 DQ330
EU491257.1 gg_id:269593 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
DQ395799.1 gg_id:155720 deep-sea octacoral clone ctg_CGOF235 DQ395
EU925894.1 gg_id:346201 sediments sediment station DBS1 northern Bering Sea clone 115G73 EU925
GQ263929.1 gg_id:513958 Structure Simulated Level Waste Site simulated level waste site clone WC2_181 GQ263

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FJ484648.1 gg_id:572347 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4846
GQ472419.1 gg_id:547121 Microbial Dongping Lake lake sediment clone 4F2-23 GQ472
AB238778.1 gg_id:147775 shal subsurface Sarobetsu Mire peat soil clone HSM-SS-015 AB238
AM086107.1 gg_id:149503 Abundance and archaea mesotrophic Lake Kinneret lake profundal sediment clone (AM086
AJ519663.1 gg_id:76094 uranium mill tailings soil sample clone GuBH2-AD/TzT-67 proteobacterium AJ5196
FJ716435.1 gg_id:509967 Uncultivated archaea sulfate-reducing biofilm Frasassi cave system anoxic lake water FJ7164
EF590018.2 gg_id:231749 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF590
FJ484742.1 gg_id:534747 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4847
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DQ067029.1 gg_id:129752 Lake Washington sediment clone pLW-45 DQ067
FJ538146.1 gg_id:566857 metagenomics study rice treated milk vetch and no-tillage paddy field soil clone MBN FJ5381
EU335163.1 gg_id:239473 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
CU922370.1 gg_id:552630 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU922
GU180168.1 gg_id:590108 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
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CU921396.1 gg_id:561911 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU921
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CU921615.1 gg_id:549289 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU921
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EF575257.1 gg_id:228296 site S25 near Coco's Island marine clone S25_1601 EF5752
FJ155026.1 gg_id:330159 seasonal fluctuation coastal Bohai Bay sequencing and RFLP ocean water clone W FJ1550
EF573912.1 gg_id:228667 site S25 near Coco's Island marine clone S25_256 EF5739
EU592442.1 gg_id:537106 Seasonal changes Salton Sea hypersaline sediment clone SSS66N EU5924
EU592440.1 gg_id:589629 Seasonal changes Salton Sea hypersaline sediment clone SSS64N EU5924
EU488121.1 gg_id:560324 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
EU488297.1 gg_id:556385 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
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DQ394901.1 gg_id:156343 harbor sediment clone VHS-B1-26 DQ394
GQ261790.1 gg_id:544072 shifts taxa and response localized organic loading deep sea sediment whale falls clo GQ261

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DQ811792.1 gg_id:167519 mangrove soil clone MSB-3A9 DQ811
EF520534.1 gg_id:232980 structure lakes: what controls ? acid-impacted lake clone ADK-BTe02-06 EF5205
EU861868.1 gg_id:343349 effects chronic fertilization on alpine tundra microbial implications carbon and nitrog EU8618
EU803844.1 gg_id:337066 It's all ranking aquatic Lake Gatun clone 5C231481 EU8038
DQ451526.1 gg_id:156878 forest soil clone FAC87 DQ451
DQ811827.1 gg_id:170103 mangrove soil clone MSB-5C4 DQ811
EF208655.1 gg_id:202296 Vertical and horizontal structure microbial sandy carbonate sediment clone CI35cm.2 EF2086
EF125419.1 gg_id:199002 soil naked tidal flat sediment near mangrove clone MSB-1F8 EF1254
EU491252.1 gg_id:276800 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU4912
DQ925889.1 gg_id:255773 Microbes and Thermal Gradients Temperature Hydrothermal Chimneys hydrotherm DQ925
EU050850.1 gg_id:238559 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_09_96 EU0508
FJ535320.1 gg_id:536335 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc sedi FJ5353
AM712338.1 gg_id:255844 detection and ly microbial Brothers Seamount Kermadec Arc New Zealand biofilm h AM712
AF420341.2 gg_id:67796 hydrothermal sediment clone AF420341 AF4203
AJ441235.1 gg_id:57453 hydrothermal vent polychaete mucous clone P. palm A 205 AJ4412
AF420340.1 gg_id:67797 hydrothermal sediment clone AF420340 AF4203
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EU134544.1 gg_id:253026 tallgrass prarie soil clone FFCH2390 EU1345
FJ592579.1 gg_id:509119 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
EF573992.1 gg_id:230171 site S25 near Coco's Island marine clone S25_336 EF5739
EF574948.1 gg_id:230813 site S25 near Coco's Island marine clone S25_1292 EF5749
GQ347789.1 gg_id:541766 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
EU048624.1 gg_id:236563 Microbial surface marine sediment Xisha Trough South China Sea clone MD2902-B18 EU0486
EU386045.1 gg_id:274122 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU3860
EU373913.1 gg_id:275503 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment cl EU3739
FJ745210.1 gg_id:570846 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7452
GQ346753.1 gg_id:568507 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
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AB486345.1 gg_id:559850 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSBARC AB4863
DQ267495.1 gg_id:149878 Metagenomic group plankton 500m depth mesopelagic Antarctic waters clone DQ267
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FJ485412.1 gg_id:558030 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4854
AJ241001.1 gg_id:10681 Norway:(Svalbard)Hornsund clone Sva0485 AJ2410
AM882628.1 gg_id:350620 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
EF632708.1 gg_id:256201 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) wate EF6327
DQ103652.1 gg_id:131834 hypersaline endoevaporitic microbial mat clone E6aH12 DQ103
AM040101.1 gg_id:149432 sandy sediments clone Sylt 5 AM040

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AF382143.1 gg_id:79548 bacterioplankton clone ZA3735c AF382
GQ337096.1 gg_id:548348 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone BB(GQ337
FM242399.1 gg_id:328281 sequences sediment clone 86 T12d-oil FM242
EU735646.1 gg_id:322881 archaeal structures and pristine soils China oil contaminated soil Jidong Oilfield clone EU735
EU734937.1 gg_id:327296 assemblages sediment station DBSE Northern Bering Sea clone 028C19 EU734
AY375082.1 gg_id:98535 deep sea sediment clone B109 AY375
AJ240985.1 gg_id:10661 Norway:(Svalbard)Hornsund clone Sva0853 AJ2409
AM882631.1 gg_id:306845 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
DQ464344.1 gg_id:157072 Arcobacter skirrowii str. 449/80 DQ464
AY314755.1 gg_id:105549 Arcobacter cryaerophilus CCUG 17802 AY314
EU106662.1 gg_id:260782 Arcobacter nitrofigilis str. F2176 EU106
FJ203148.1 gg_id:334326 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
AB262383.1 gg_id:161324 Microbial Ishigaki-jima Island Japan sea water Ishigaki Port clone IG-1-21 AB2623
AB262379.1 gg_id:160905 Microbial Ishigaki-jima Island Japan sea water Ishigaki Port clone IG-1-16 AB2623
AY922188.1 gg_id:113246 whalefall clone 131599 AY922
GQ348908.1 gg_id:558608 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
FJ497477.1 gg_id:589078 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4974
AF513455.1 gg_id:60384 Arcobacter sp. str. LA31B AF5134
DQ071284.1 gg_id:139697 dominate filamentous sulfur mat 13 degree north field East Pacific Rise deep-sea hy(DQ071
AF314538.1 gg_id:36386 Arcobacter butzleri oN_107_00 AF3145
AB013832.1 gg_id:11163 NB1-k clone NB1-k AB0138
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AY225610.1 gg_id:96366 pumice fragments exposed to Mid-Atlantic Ridge vent clone AT-pp13 AY2256
GU061286.1 gg_id:546186 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061
AJ607391.1 gg_id:102614 Arcobacter cibarius str. LMG 21996 AJ6073
EF515495.1 gg_id:222170 Electricigen Enrichment MFC full-scale anaerobic bioreactor sludge treating brewery EF5154
AY922205.1 gg_id:113754 whalefall clone 131644 AY922
FJ202783.1 gg_id:349369 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2027
FJ403064.1 gg_id:357886 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4030
EF123613.1 gg_id:239833 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1236
AJ271654.1 gg_id:51975 Arcobacter sp. D1a1 AJ2716
AM084124.1 gg_id:140216 Arcobacter sp. str. R-28214 AM084
AY570641.1 gg_id:108516 biodegraded Canadian oil reservoir clone PL-8D4 AY5706
FN429802.1 gg_id:534177 Biogeochemistry -activity cold seep Larsen B area western Weddell Sea Antarctica m FN4298
FJ202326.1 gg_id:315205 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2023
DQ917897.1 gg_id:227146 Microbial profiles bleached and wild-type Muricea elongata Arcobacter clone ME15 DQ917
AY035822.1 gg_id:70950 enriched coastal seawater overlying anoxic marine sediment sulfidicus AY0358
EU050947.1 gg_id:255076 libraries sediment Kings Bay Svalbard Arctic clone SS1_B_02_37 EU0509

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DQ357825.1 gg_id:150082 Marine Hatchery: Balance Between Pathogenic and Potentially Probiotic Strains adu DQ357
DQ676433.1 gg_id:224791 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
AB030594.1 gg_id:50542 groundwater clone 1021 AB0305
EU265967.1 gg_id:567689 transition and zone within near-shore anoxic basin: Nitinat Lake depth 50 m clone Ni EU2659
FJ660508.1 gg_id:561979 Correlation functional stability and dynamics full-scale wastewater treatment plant a FJ6605
EF688595.1 gg_id:284348 Biogeochemistry iron- and sulfur-precipitating microbial mats (Nile Mediterranean) s EF6885
GU061298.1 gg_id:560130 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061
FJ717168.1 gg_id:571505 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7171
AY621116.1 gg_id:106004 Arcobacter butzleri str. ATCC 49616 AY6211
EF031092.1 gg_id:186810 Prokaryotic Tirez La Macha Spain endorheic hypersaline atalasohaline lake clone SalB EF0310
AY697901.1 gg_id:105639 Antarctica clone F3C43 AY6979
AB234539.1 gg_id:149689 termite gut homogenate clone MgMjD-024 AB2345
FJ203095.1 gg_id:346863 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
AB262386.1 gg_id:159735 Microbial Ishigaki-jima Island Japan sea water Ishigaki Port clone IG-1-27 AB2623
EU236320.1 gg_id:292623 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg1bF7 EU2363
AB015259.1 gg_id:11183 deepest cold-seep area Japan Trench clone JTB360 proteobacterium AB0152
AJ810558.1 gg_id:105006 superficial sediments clone AJ8105
DQ464340.1 gg_id:158050 Arcobacter butzleri str. 520H-2004 DQ464
GQ340204.1 gg_id:583784 Prokaryotic Marathonas Reservoir Greece water column clone VE07-08-BAC GQ340
AB013833.1 gg_id:11181 NB1-l clone NB1-l AB0138
GQ349268.1 gg_id:581309 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
EU781617.1 gg_id:322902 Bacteroides ureolyticus 4 EU7816
GQ167663.1 gg_id:581365 Campylobacter concisus str. ATCC 51561 GQ167
EU869312.1 gg_id:334581 Campylobacter cuniculorum str. 119/07 EU8693
NC_009715.1 gg_id:242986 Campylobacter curvus str. 525.92 NC_009
AY621301.1 gg_id:103129 Campylobacter fetus str. 23D AY6213
AF219233.1 gg_id:69901 Campylobacter fetus MGH 97-2126 AF2192
DQ174168.1 gg_id:151146 Campylobacter gracilis str. ATCC 33236 DQ174
DQ174161.1 gg_id:155173 Campylobacter helveticus str. NCTC 12470 DQ174
DQ174163.1 gg_id:154260 Campylobacter helveticus str. NCTC 12472 DQ174
U03022.1 gg_id:11247 Campylobacter helveticus U03022
CP000776.1 gg_id:257761 Campylobacter hominis str. ATCC BAA-381 CP0007
AF043422.1 gg_id:11239 Campylobacter cf. lanienae str. UB 992 AF0434
AF550661.1 gg_id:57347 Campylobacter mucosalis str. ATCC 49353 AF5506
FJ983015.1 gg_id:430191 Aspects Oral Microbiome root canal clone FJ9830
NZ_ACFU01000050.1 gg_id:469893 Campylobacter rectus str. RM3267 NZ_AC
NZ_ACVQ01000030.1 gg_id:470642 Campylobacter showae str. RM3277 NZ_AC
EU781616.1 gg_id:326311 Campylobacter showae 3 EU7816
X67775.1 gg_id:11227 Campylobacter sputorum str. biovar sputorum;LMG 7795 X67775
AB181364.1 gg_id:100818 Campylobacter lari str. A3 AB1813
AB181358.1 gg_id:105325 Campylobacter lari str. NCTC12893 AB1813

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AB181357.1 gg_id:105165 Campylobacter lari str. NCTC12892 AB1813
AB181359.1 gg_id:105482 Campylobacter lari str. NCTC12894 AB1813
AF550640.1 gg_id:57368 Campylobacter upsaliensis str. LMG 8851 AF5506
AF550632.1 gg_id:102123 Campylobacter lari str. LMG 11251 AF5506
GQ167658.1 gg_id:586732 Campylobacter upsaliensis str. RM3195 GQ167
DQ174156.1 gg_id:154704 Campylobacter showae str. CCUG 11641 DQ174
EU289113.1 gg_id:238608 microbiome cloacal openings urogenital and anal tracts tammar wallaby Macropus e EU2891
FJ832837.1 gg_id:542539 Microbial Ecology Crop Folivorous Hoatzin crop contents clone J3Q301_10B02 FJ8328
AF219235.1 gg_id:70687 Campylobacter hyointestinalis MGH 97-2652 AF2192
AB301962.1 gg_id:219863 Campylobacter subsp. hyointestinalis str. 2831-1 AB3019
AF550641.1 gg_id:57366 Campylobacter upsaliensis str. LMG 8853 AF5506
EU289065.1 gg_id:253584 microbiome cloacal openings urogenital and anal tracts tammar wallaby Macropus e EU2890
AY554142.1 gg_id:113002 Campylobacter sp. str. BTP1 AY5541
AY554144.1 gg_id:113756 Campylobacter sp. str. WB1 AY5541
GQ167676.1 gg_id:542181 Campylobacter coli str. ATCC 33559 GQ167
AF372091.1 gg_id:66989 Campylobacter jejuni str. NCTC 11351 AF3720
EU559332.1 gg_id:549039 Campylobacter sp. str. MIT 97-5311 EU5593
AJ276874.1 gg_id:69947 Campylobacter faecalis str. 1433 stool sample isolate AJ2768
AB301961.1 gg_id:219965 Campylobacter subsp. hyointestinalis str. 2728-1 AB3019
AF550628.1 gg_id:55836 Campylobacter jejuni str. H99/240 AF5506
AF550647.1 gg_id:57361 Campylobacter helveticus str. CCUG 30563 AF5506
DQ174159.1 gg_id:154512 Campylobacter upsaliensis str. NZ5564-94 DQ174
AM933372.1 gg_id:541426 Campylobacter subantarcticus str. LMG 24374 AM933
GQ451185.1 gg_id:533613 Fecal Yunnan snub-nosed monkey feces clone J16 GQ451
AF097688.1 gg_id:11237 Campylobacter hyointestinalis subsp. lawsonii str. CHY8 subsp. AF0976
AB066098.1 gg_id:54599 Campylobacter lari str. CF89-12 Japanese strain AB0660
AJ620504.1 gg_id:108770 Campylobacter insulaenigrae str. NCTC 12927 AJ6205
DQ174164.1 gg_id:152100 Campylobacter helveticus str. E504-90 DQ174
EU623474.1 gg_id:278725 Campylobacter sp. str. 24/06 EU6234
AF550620.1 gg_id:57384 Campylobacter coli str. LMG 9220 AF5506
DQ174146.1 gg_id:151930 Campylobacter lari str. NZ145-95 DQ174
FJ609419.1 gg_id:573432 Campylobacter jejuni str. PRCC-3 FJ6094
EF621895.1 gg_id:243035 Campylobacter canadensis str. L268 EF6218
AF550655.1 gg_id:57353 Campylobacter showae str. LMG 12636 AF5506
EU470464.1 gg_id:302158 Grevy's zebra feces clone GZ_aaa03d07 EU4704
AY288304.1 gg_id:86812 Campylobacter lanienae str. L52 AY2883
EF621896.1 gg_id:240175 Campylobacter canadensis str. L267 EF6218
NR_028692.1 gg_id:561090 Sulfurospirillum barnesii str. SES-3; ATCC 700032 NR_028
EF462744.1 gg_id:295702 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4627
AB355036.1 gg_id:244687 Monitoring and archaeal two heavy polluted areas lake Egypt based on surface wate AB3550

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FJ792436.1 gg_id:546339 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7924
FJ791877.1 gg_id:573109 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7918
FJ792027.1 gg_id:534646 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7920
U85964.1 gg_id:11189 Sulfurospirillum arsenophilum str. MIT-13 U8596
AB286550.1 gg_id:206504 Reducing production and domination comamonadaceae reducing oxygen supply was AB286
AY548994.2 gg_id:103740 marine bone clone boneC3F4 AY548
FJ792107.1 gg_id:534811 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7921
FJ792301.1 gg_id:562673 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7923
AY548973.1 gg_id:102465 marine bone clone boneC15D1 AY548
DQ295747.1 gg_id:160847 Epsilonproteobacteria terrestrial caves and springs microbial mat sulfidic Warm spri DQ295
DQ925871.1 gg_id:249367 Microbes and Thermal Gradients Temperature Hydrothermal Chimneys hydrotherm DQ925
DQ228579.1 gg_id:141569 and physiological significance attached microorganisms rock-hosted deep-sea carbo DQ228
AF357199.1 gg_id:22148 Sulfurospirillum sp. str. 18.1 AF357
FJ792478.1 gg_id:534881 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7924
AY672509.1 gg_id:99157 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH1_28_BAC_16SrR AY672
AY327877.2 gg_id:88092 surface vent snail foot clone SF_C11-F3 AY327
FJ640810.1 gg_id:512603 GeoChip-based metabolic microbial hydrothermal vent chimney 4143-1 Mothra field FJ6408
AY280408.1 gg_id:100828 exterior deep-sea hydrothermal vent sulfide chimney Juan de Fuca Ridge Pacific Oce AY280
AB234540.1 gg_id:146284 termite gut homogenate clone MgMjR-060 AB234
AY280434.1 gg_id:106449 exterior deep-sea hydrothermal vent sulfide chimney Juan de Fuca Ridge Pacific Oce AY280
AB192073.1 gg_id:143275 termite gut homogenate clone M2PT2-04 AB192
FN356324.1 gg_id:580201 Prokaryotic structure sulfate reducer activity -temperature reservoirs and without ni FN356
FJ485111.1 gg_id:541385 microbial autonomous rover world's deepest phreatic sinkhole water column El Zacat FJ4851
AY266270.1 gg_id:103575 oil production water clone CT1 AY266
DQ925875.1 gg_id:244236 Microbes and Thermal Gradients Temperature Hydrothermal Chimneys hydrotherm DQ925
FJ156092.1 gg_id:461508 Arcobacter mytili str. T234 FJ1560
AJ969453.1 gg_id:159487 Microbial black smoker s Mid Atlantic Ridge ecological implications rock fragments ch AJ9694
AB231076.1 gg_id:147107 termite gut wall clone NkW01-035 AB231
AJ441214.1 gg_id:106099 hydrothermal vent polychaete mucous clone P. palm C/A 39 AJ4412
AJ576349.1 gg_id:102626 hindgut homogenate Pachnoda ephippiata larva clone PeH32 AJ5763
AY192528.1 gg_id:111196 Flexispira rappini str. 205-2 AY192
AY192526.1 gg_id:109700 Flexispira rappini str. 204-1 AY192
AY126479.1 gg_id:131724 Flexispira rappini str. 205-1 AY1264
AF396078.1 gg_id:20816 Helicobacter cinaedi 30327 AF396
AF396081.1 gg_id:49270 Helicobacter cinaedi 31111 AF396
AB275325.1 gg_id:196588 Helicobacter cinaedi str. PAGU0621 (= 56) AB275
AF302108.1 gg_id:69782 Helicobacter sp. str. HC15880 AF302
AF118017.1 gg_id:11022 Flexispira rappini str. H1353 AF118
AF497810.1 gg_id:56169 Helicobacter cinaedi str. 94105 AF497
AF497807.1 gg_id:60772 Flexispira rappini str. FR106 AF497
AF072471.1 gg_id:10989 Helicobacter mesocricetorum str. MU97-1514 AF0724

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DQ438113.1 gg_id:154548 Helicobacter canadensis str. L172
AY631946.1 gg_id:106403 Helicobacter canis str. MIT 51402
L13464.1 gg_id:11062 Helicobacter canis
DQ307736.1 gg_id:154375 Helicobacter equorum str. EqF2
AF506785.1 gg_id:72292 Helicobacter heilmannii str. WD1
AY686607.1 gg_id:111336 Helicobacter felis str. Lee DS2
M57398.2 gg_id:10994 Helicobacter felis str. CS1; ATCC 49179
AF103882.1 gg_id:11030 Helicobacter felis str. 1136-7
AY561831.1 gg_id:106123 Helicobacter ganmani str. ES-5
AY631951.1 gg_id:101810 Helicobacter ganmani str. MIT 99-5102
U07573.2 gg_id:11042 Helicobacter hepaticus str. Hh-1
AY554124.1 gg_id:114186 Helicobacter kirbridei str. BTP1F
AF333339.1 gg_id:27055 Helicobacter sp. str. MIT 99-5504
AF010140.1 gg_id:11040 Helicobacter muridarum str. UNSW 1.7st
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DQ062214.1 gg_id:132728 Helicobacter sp. AGN-1
X67854.1 gg_id:11080 Helicobacter pylori str. ATCC 49396T
Z25741.1 gg_id:11013 Helicobacter pylori str. NCTC 11637
DQ062215.1 gg_id:132837 Helicobacter sp. GA-3
AY593989.1 gg_id:101118 Helicobacter pylori str. LPB 424-01
AF057163.1 gg_id:10974 Helicobacter sp
AF535198.1 gg_id:57708 Helicobacter pylori str. Lit81
M88137.2 gg_id:10945 Helicobacter trogontum str. ATCC 43966
AY034820.1 gg_id:47556 Helicobacter rappini str. H 95 S
AF225548.1 gg_id:28994 Helicobacter trogontum str. ATCC 49310
U96296.1 gg_id:10983 Helicobacter rodentium str. MIT 95-1707
AF506791.1 gg_id:64310 Helicobacter heilmannii str. P5
AF506790.1 gg_id:71558 Helicobacter heilmannii str. P4
AF506787.1 gg_id:73122 Helicobacter heilmannii str. P1
AY554121.1 gg_id:114199 Helicobacter vulpecula str. BTP7C
FJ667779.1 gg_id:548699 Helicobacter sp. 'pancake tortoise' str. FL-2007
AF292378.1 gg_id:72792 Helicobacter cetorum str. MIT 99-5656
AY554137.1 gg_id:113171 Helicobacter sp. str. BTP3S
M88152.2 gg_id:11054 Helicobacter sp. '91-266-11 Fox' str. 91-266-11 Fox
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L36141.1 gg_id:10982 Helicobacter pullorum str. NCTC 12824
AF058768.1 gg_id:11034 Helicobacter heilmannii 1
AF506794.1 gg_id:64304 Helicobacter heilmannii str. TD1

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AF237612.1 gg_id:26677 Helicobacter sp. str. Wee Tee AF2376
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Y18028.1 gg_id:10969 Helicobacter heilmannii str. AF53 Y18028
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AF333340.1 gg_id:49735 Helicobacter sp. str. MIT 99-5507 AF3333
AF506780.1 gg_id:72724 Helicobacter heilmannii str. CM1 AF5067
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AF506782.1 gg_id:64314 Helicobacter heilmannii str. MM1 AF5067
AY554140.1 gg_id:113412 Helicobacter sp. str. WB1F AY5541
U46129.1 gg_id:11050 Helicobacter cholecystus str. Hkb-1 U46129
DQ438119.1 gg_id:152253 Helicobacter canadensis str. L222 DQ438
AF297868.1 gg_id:48289 Helicobacter aurati str. MIT 97-5075c AF2978
AF302107.1 gg_id:47191 Helicobacter bizzozeronii str. ATCC700031 AF3021
AY742308.1 gg_id:102634 Helicobacter sp. str. MIT 02-6899 AY7423
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AY191468.1 gg_id:93668 microbial mat sulfidic cave spring clone LKC1.21_5 AY1914
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AB240521.1 gg_id:143003 microbial structure rhizosphere biofilm Sapporo root-base (80 160 mm) Phragmites S AB240
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DQ787698.1 gg_id:162117 holocene lake sediment clone SLB422 DQ787
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AF367495.1 gg_id:43869 deep-sea hydrothermal vent clone VC1.2-cl50 AF3674
AF367492.1 gg_id:70730 deep-sea hydrothermal vent clone VC1.2-cl32 AF3674
GQ433935.1 gg_id:541504 Subseafloor ocean crust microbial observatories: (F-through Osmo Colonization Syst GQ433

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U15101.1 gg_id:11096 hydrothermal vent clone PVB_OTU_2 U1510
NC_007575.1 gg_id:144857 Thiomicrospira denitrificans str. ATCC 33889 NC_00
FJ437888.1 gg_id:547212 Benthic okenone production Fayetteville New York Green Lake 25 m water depth clor FJ4378
AF121884.1 gg_id:11109 phenol-degrading consortium clone Phenol-1 AF1218
DQ295573.1 gg_id:160062 Epsilonproteobacteria terrestrial caves and springs microbial mat sulfidic cave spring DQ295
FJ615170.1 gg_id:593488 Microbial dynamics seasonally fjord: British Columbia Saanich Inlet depth 10m anoxic FJ6151
EU491812.1 gg_id:268142 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918
DQ234141.2 gg_id:142794 determined library mangrove clone DS057 DQ234
FJ497618.1 gg_id:563303 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4976
FJ717090.1 gg_id:551059 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7170
DQ676325.1 gg_id:224044 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
AY211683.1 gg_id:96203 Gulf Mexico seafloor sediment clone GoM GC234 633E AY2116
AY225615.1 gg_id:97077 iron-rich substrate exposed to Mid-Atlantic Ridge vent clone AT-cs10 AY2256
FN397808.1 gg_id:424252 geomicrobiological investigation sediments Gulf Cadiz GeoB 9072-1 anoxic sub-surfa FN3978
AJ441199.1 gg_id:57482 hydrothermal vent polychaete mucous clone P. palm C/A 78 AJ4411
AF355050.1 gg_id:48010 marine clone Arctic96B-13 AF3550
GQ356982.1 gg_id:592000 Metal Dependent Marine Methane Oxidation methane seep sediment clone FeSO4_ GQ356
AY225612.1 gg_id:97721 organic-rich substrate exposed to Mid-Atlantic Ridge vent clone AT-co23 AY2256
DQ676368.1 gg_id:222158 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
FJ497273.1 gg_id:590030 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4972
EF218993.1 gg_id:215778 and Eukaryotic Hydrothermal Site East Rise Ridge Flank Abyssal Hill hydrothermal site EF2189
DQ228650.1 gg_id:141130 Microbial colonizing mineral surfaces within sulfide-microbial incubator clone CH2b1 DQ228
EF219001.1 gg_id:209624 and Eukaryotic Hydrothermal Site East Rise Ridge Flank Abyssal Hill hydrothermal site EF2190
U15106.1 gg_id:11103 hydrothermal vent clone PVB_OTU_6 U1510
AY280406.1 gg_id:100516 exterior deep-sea hydrothermal vent sulfide chimney Juan de Fuca Ridge Pacific Oce AY2804
GQ848465.1 gg_id:578863 Microbial sediments vents along East Lau Spreading Centre deep sea hydrothermal GQ848
FJ535290.1 gg_id:546926 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
FJ717130.1 gg_id:551352 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7171
FJ717171.1 gg_id:533593 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7171
EF379628.1 gg_id:205139 sea water Victoria Harbor (Hong Kong) area clone VH-PA7-38 EF3796
EU652649.1 gg_id:279233 Seasonal variation microbial Yel Sea sediment clone B8S-4 EU6526
DQ836759.1 gg_id:169835 attached biomass denitrifying bioreactor clone DB-83 DQ836
AY218555.1 gg_id:82461 penguin droppings sediments clone KD1-140 AY2185
EU652651.1 gg_id:278362 Seasonal variation microbial Yel Sea sediment clone A8S-94 EU6526
FM242221.1 gg_id:343001 sequences sediment clone 259 T0h-oil FM242
EU652650.1 gg_id:277911 Seasonal variation microbial Yel Sea sediment clone C8S-4 EU6526
FJ717108.1 gg_id:569301 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7171
FJ535347.1 gg_id:510699 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc sedi FJ5353
AB530237.1 gg_id:584301 Monitoring Suez and Tokyo bays based on analyses marine sediment clone Tokyo..B AB5302
AB188787.1 gg_id:113750 cold seep sediment clone SB5-24 AB1887
AB193899.1 gg_id:108823 deep-sea vent clone Mb-NB05 AB1938

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GQ261804.1 gg_id:560545 shifts taxa and response localized organic loading deep sea sediment whale falls clo GQ261
AY922199.1 gg_id:112883 whalefall clone 131631 AY9221
DQ228651.1 gg_id:142365 Microbial colonizing mineral surfaces within sulfide-microbial incubator clone CH2b1 DQ228
FJ717087.1 gg_id:562754 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7170
AY216438.1 gg_id:82215 temperate estuarine mud clone KM51 AY2164
EU491805.1 gg_id:273299 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918
AY225613.1 gg_id:98692 pumice fragments exposed to Mid-Atlantic Ridge vent clone AT-pp6 AY2256
AB189375.1 gg_id:113264 cold seep sediment clone JT75-307 AB1893
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AJ441200.1 gg_id:57484 hydrothermal vent polychaete mucous clone P. palm C 92 AJ4412
EF999352.1 gg_id:237892 Vertical and Methanotrophs Estuarine Sediments Pearl River Estuary sediments 6cm EF9993
AM982624.1 gg_id:278960 microbial and slurry pig saw dust spent bedding clone PISD-AJC10 AM982
FJ535284.1 gg_id:512607 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
EF218991.1 gg_id:206907 and Eukaryotic Hydrothermal Site East Rise Ridge Flank Abyssal Hill hydrothermal site EF2189
AY532570.1 gg_id:105327 uranium-contaminated aquifer clone 1013-28-CG21 AY5325
FJ497261.1 gg_id:566005 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4972
AB235234.1 gg_id:141016 symbiont AB2352
FJ437839.1 gg_id:513719 Benthic okenone production Fayetteville New York Green Lake 25 m water depth clo FJ4378
AB176111.1 gg_id:175999 Microbial Hydrothermal System Suiyo Seamount on Izu-Ogasawara Arc vent area wa AB1761
FM203379.1 gg_id:370488 Biogeography filamentous and epsilonproteobacterial ectosymbionts vent shrimp e FM203
AJ515714.1 gg_id:94683 deep-sea shrimp gut clone 4 AJ5157
FM203398.1 gg_id:371282 Biogeography filamentous and epsilonproteobacterial ectosymbionts vent shrimp e FM203
AP009179.1 gg_id:249880 Sulfurovum sp. str. NBC37-1 AP0091
EF462799.1 gg_id:293907 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrothern EF4627
AJ515716.1 gg_id:94327 deep-sea shrimp gut clone 15 AJ5157
EU555122.1 gg_id:259007 Microbial Sulfide Hydrothermal Vent Field Juan de Fuca Ridge Dudley hydrothermal EU5551
AY672497.1 gg_id:102782 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH1_1_BAC_16SrR AY6724
FJ792125.1 gg_id:510296 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7921
FJ535278.1 gg_id:554288 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
AY531596.1 gg_id:100070 Indian Ocean hydrothermal vent clone SF_C23-C4_shell AY5315
DQ228647.1 gg_id:144474 Microbial colonizing mineral surfaces within sulfide-microbial incubator clone CH2b1 DQ228
AY542560.1 gg_id:113844 Gulf Mexico seafloor sediment clone GoM GB425 02B-28 AY5425
GQ249521.1 gg_id:580475 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 6-8 cm layer clo GQ249
GQ357025.1 gg_id:533206 Metal Dependent Marine Methane Oxidation methane seep sediment clone FeOrig_ GQ357
AF449231.1 gg_id:61878 Riftia pachyptila's tube clone R103-B15 AF4492
AB476184.1 gg_id:536849 Compositional and functional transition epibiotic microbial hydrothermal vent galat AB4761
AY531576.1 gg_id:107839 Indian Ocean hydrothermal vent clone SF_C23-G3 AY5315
FJ535277.1 gg_id:540587 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352

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AB440165.1 gg_id:552622 Farming and harvesting filamentous vent- galatheid crab (Decapoda; Anomura) vent AB4401
EU487998.1 gg_id:570269 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4879
FJ497296.1 gg_id:541606 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4972
GU369887.1 gg_id:519900 and mats vents Arc white filamentous microbial mat photic shal hydrothermal vent GU369
FJ535276.1 gg_id:511548 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
AY549001.2 gg_id:101316 marine bone clone sacC13A3 AY5490
FJ535285.1 gg_id:554836 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
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AY531587.1 gg_id:104132 Indian Ocean hydrothermal vent clone SF_C23-D11_shell AY5315
AY672538.1 gg_id:105257 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH9_1_BAC_16SrR AY6725
AB440171.1 gg_id:569838 Farming and harvesting filamentous vent- galatheid crab (Decapoda; Anomura) vent AB4401
AY531601.1 gg_id:104446 Indian Ocean hydrothermal vent clone SF_C23-C7_shell AY5316
FJ535304.1 gg_id:554479 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5353
EU652643.1 gg_id:278085 Seasonal variation microbial Yel Sea sediment clone B8S-12 EU6526
EU555129.1 gg_id:268986 Microbial Sulfide Hydrothermal Vent Field Juan de Fuca Ridge Dudley hydrothermal EU5551
DQ071277.1 gg_id:139667 dominate filamentous sulfur mat 13 degree north field East Pacific Rise deep-sea hy DQ071
EF462802.1 gg_id:289264 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4628
AJ441206.1 gg_id:57471 hydrothermal vent polychaete mucous clone P. palm C/A 113 AJ4412
EF462624.1 gg_id:296130 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4626
AB530172.1 gg_id:559058 Monitoring Suez and Tokyo bays based on analyses marine sediment clone Suez..Ba AB5301
FM165251.1 gg_id:558481 seep chitineous tube vestimentiferan tubeworm Lamellibrachia sp. cold seeps clone FM165
AF420359.1 gg_id:67782 hydrothermal sediment clone AF420359 AF4203
AB205405.1 gg_id:115002 Alviniconcha aff. hessleri gill endosymbiont AB2054
EF462766.1 gg_id:300444 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4627
AY531572.1 gg_id:107247 Indian Ocean hydrothermal vent clone SF_C23-G9 AY5315
FJ712419.1 gg_id:526671 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) K FJ7124
AF029044.1 gg_id:11139 Benzene mineralizing consortium complete clone SB-17 AF0290
EF462668.1 gg_id:299697 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4626
AY531593.1 gg_id:99597 Indian Ocean hydrothermal vent clone SF_C23-E5_shell AY5315
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AY672511.1 gg_id:103076 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH1_3f_BAC_16SrF AY6725
DQ270643.1 gg_id:142040 hydrothermal carbonate chimney clone LC1133B-58 DQ270
EU107475.1 gg_id:349595 Identification Ectosymbiont on Hair Inhabiting Hydrothermal Vents hair Shinkaia cro EU1074
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FM165253.1 gg_id:558667 seep chitineous tube vestimentiferan tubeworm Lamellibrachia sp. cold seeps clone FM165
AB099998.1 gg_id:79591 inactive deep-sea hydrothermal vent chimneys clone IndB1-30 AB0999
DQ394973.1 gg_id:153118 harbor sediment clone VHS-B3-94 DQ394
EF462777.1 gg_id:288783 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4627
AY531595.1 gg_id:99906 Indian Ocean hydrothermal vent clone SF_C23-E10_shell AY5315
EU265786.1 gg_id:258528 epibiotic assemblage recently described Yeti Kiwa hirsuta (Kiwidae) surface crab hyc EU2657

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AY280405.1 gg_id:100364 exterior deep-sea hydrothermal vent sulfide chimney Juan de Fuca Ridge Pacific Oce AY2804
AJ575963.1 gg_id:95437 deep sea hydrothermal vent clone T3-Hot3-706 AJ5759
AY531597.1 gg_id:100222 Indian Ocean hydrothermal vent clone SF_C23-B7_shell AY5315
AM176867.1 gg_id:161081 Recovery mangrove sediment clone SZB13 AM176
EU265795.1 gg_id:262084 epibiotic assemblage recently described Yeti Kiwa hirsuta (Kiwidae) surface crab hyc EU2657
EU662602.1 gg_id:577333 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems floating microb EU6626
AB476188.1 gg_id:536599 Compositional and functional transition epibiotic microbial hydrothermal vent galath AB4761
AY197378.1 gg_id:83351 Guaymas Basin hydrothermal vent sediments clone B01R006 AY1973
FJ535271.1 gg_id:544300 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
EF462600.1 gg_id:303123 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4626
EU265785.1 gg_id:277410 epibiotic assemblage recently described Yeti Kiwa hirsuta (Kiwidae) surface crab hyc EU2657
DQ394894.1 gg_id:151332 harbor sediment clone VHS-B1-11 DQ394
AY531575.1 gg_id:107689 Indian Ocean hydrothermal vent clone SF_C23-G4 AY5315
AY531585.1 gg_id:103824 Indian Ocean hydrothermal vent clone SF_C23-C11_shell AY5315
AY225614.1 gg_id:96015 iron-rich substrate exposed to Mid-Atlantic Ridge vent clone AT-cs3 AY2256
AB232530.1 gg_id:151369 Prokaryotic gills deep-sea hydrothermal vent shrimp gill Alvinocaris longirostris clone AB2325
DQ270645.1 gg_id:143843 hydrothermal carbonate chimney clone LC1133B-68 DQ270
AY531600.1 gg_id:104287 Indian Ocean hydrothermal vent clone SF_C23-C8_shell AY5316
AY531591.1 gg_id:99285 Indian Ocean hydrothermal vent clone SF_C23-B1_shell AY5315
AY327882.1 gg_id:90455 surface vent snail foot clone SF_C4-D5 AY3278
AY225624.1 gg_id:97289 pumice fragments exposed to Mid-Atlantic Ridge vent clone AT-pp27 AY2256
AF420351.1 gg_id:67787 hydrothermal sediment clone AF420351 AF4203
EU617859.1 gg_id:345356 Seasonal variation microbial Yel Sea sediment clone D13S-42 EU6178
FJ799124.1 gg_id:510696 Microbial Responsible Intermediates Degradation Anaerobic Digestion System Deterr FJ7991
AY225622.1 gg_id:97434 iron-rich substrate exposed to Mid-Atlantic Ridge vent clone AT-cs8 AY2256
EF462801.1 gg_id:299935 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4628
DQ270655.1 gg_id:141448 hydrothermal carbonate chimney clone LC1404B-6 DQ270
FJ535259.1 gg_id:542712 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
GQ167319.1 gg_id:562624 Microbial Life -13C Brine Ice-Sealed Antarctic ice-entrained brine 16.5 m Lake Vida c GQ167
FJ535292.1 gg_id:551198 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
AF449253.1 gg_id:61855 Riftia pachyptila's tube clone R76-B151 AF4492
AY672522.1 gg_id:99252 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH3_9_BAC_16SrRN AY6725
DQ071276.1 gg_id:140018 dominate filamentous sulfur mat 13 degree north field East Pacific Rise deep-sea hy DQ071
EF462733.1 gg_id:291526 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4627
AY704398.1 gg_id:112176 oceanic crust clone FS275-77B-03 AY7043
AB440161.1 gg_id:549495 Farming and harvesting filamentous vent- galatheid crab (Decapoda; Anomura) vent AB4401
EF462696.1 gg_id:289073 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4626
AY531570.1 gg_id:106942 Indian Ocean hydrothermal vent clone SF_C23-H11 AY5315
AB239758.1 gg_id:149235 filamentous epibiont Vulcanolepas osheai AB2397
FJ535270.1 gg_id:533833 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352

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EU265789.1 gg_id:258893 epibiotic assemblage recently described Yeti Kiwa hirsuta (Kiwidae) surface crab hyc EU265
AM268719.1 gg_id:232342 Fluid chemistry and microbial southern hydrothermal field (Lilliput) on Mid-Atlantic AM268
AF154091.1 gg_id:11137 hydrocarbon seep clone BPC056 AF154
DQ071288.1 gg_id:139782 dominate filamentous sulfur mat 13 degree north field East Pacific Rise deep-sea hy DQ071
AB440162.1 gg_id:557961 Farming and harvesting filamentous vent- galatheid crab (Decapoda; Anomura) vent AB440
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EF462627.1 gg_id:297366 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4626
EU487914.1 gg_id:538726 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU487
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AB193955.1 gg_id:108919 deep-sea vent clone Sd-EB04 AB193
EF462827.1 gg_id:292616 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4628
AY531571.1 gg_id:107097 Indian Ocean hydrothermal vent clone SF_C23-H10 AY5315
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DQ394936.1 gg_id:153573 harbor sediment clone VHS-B3-29 DQ394
AB188786.1 gg_id:113953 cold seep sediment clone SB5-11 AB188
FJ497361.1 gg_id:542661 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4973
AB278150.1 gg_id:177809 hydrothermal vent prokaryotes trapped 0.2- and 0.1-micro-m-filtrates obtained Mar AB278
AF121887.1 gg_id:11128 2-bromophenol-dehalogenating consortium clone 2BP-7 AF1218
FM203396.1 gg_id:423805 Biogeography filamentous and epsilonproteobacterial ectosymbionts vent shrimp e FM203
EF061978.1 gg_id:190937 Xiamen China mangrove sediment clone XME34 EF0619
AY327881.1 gg_id:88572 surface vent snail foot clone SF_C4-C4 AY3278
FJ497343.1 gg_id:549911 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4973
FJ497347.1 gg_id:582544 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4973
GU180158.1 gg_id:537179 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
EU107476.1 gg_id:343167 Identification Ectosymbiont on Hair Inhabiting Hydrothermal Vents hair Shinkaia cro EU107
EU662505.1 gg_id:571991 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems microbial mat EU662
EF462800.1 gg_id:293062 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4628
FJ497375.1 gg_id:581955 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4973
GQ249549.1 gg_id:575344 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 12-14 cm layer c GQ249
FJ264679.1 gg_id:328542 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedim FJ2646
EF687491.1 gg_id:284524 Biogeochemistry iron- and sulfur-precipitating microbial mats (Nile Mediterranean) s EF6874
FJ905647.1 gg_id:545620 s Tonga-Kermadec iron oxide chimney-like structure on Volcano 19 Tonga Arc clone V FJ9056
AY531592.1 gg_id:99437 Indian Ocean hydrothermal vent clone SF_C23-B12_shell AY5315
FJ497326.1 gg_id:571424 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4973
FJ717162.1 gg_id:554142 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7171
FJ497317.1 gg_id:574124 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4973
FJ535287.1 gg_id:544443 within iron-containing flocculent mats submarine volcanoes along Kermadec Arc iron FJ5352
FJ497276.1 gg_id:586842 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4972
EF061975.1 gg_id:175519 Xiamen China mangrove sediment clone XME6 EF0619
AY672498.1 gg_id:102927 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH1_10_BAC_16Sr AY6724

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AF449251.1 gg_id:61858 Riftia pachyptila's tube clone R76-B78 AF449
L35520.1 gg_id:11121 Alvinella pompejana L3552
GU369898.1 gg_id:520798 and mats vents Arc white filamentous microbial mat photic shal hydrothermal vent GU369
AY327880.1 gg_id:88951 surface vent snail foot clone SF_C5-F8 AY327
FJ712566.1 gg_id:508922 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) FJ7125
DQ925872.1 gg_id:241522 Microbes and Thermal Gradients Temperature Hydrothermal Chimneys hydrotherm DQ925
EF462629.1 gg_id:295292 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF462
GQ261802.1 gg_id:543184 shifts taxa and response localized organic loading deep sea sediment whale falls clo GQ261
AF420344.1 gg_id:67793 hydrothermal sediment clone AF420344 AF420
DQ295602.2 gg_id:159750 Epsilonproteobacteria terrestrial caves and springs microbial mat sulfidic spring Drin DQ295
FJ497396.1 gg_id:542332 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4973
EF462824.1 gg_id:302928 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF462
AB187014.1 gg_id:101863 Suiyo seamount hydrothermal vent water 0.2micro-m filterable fraction clone SSM-E AB187
AY542550.1 gg_id:114277 Gulf Mexico seafloor sediment clone GoM GB425 02B-9 AY542
FJ902253.1 gg_id:571547 phreatic limestone sinkholes northeastern Mexico water column 32m deep cenote C FJ9022
FJ264760.1 gg_id:339147 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedim FJ2647
AY355304.2 gg_id:101665 epibiontic clone C11-C9 AY355
AB091292.1 gg_id:84230 Sulfurovum lithotrophicum str. 42BKT AB091
AY672512.1 gg_id:103225 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH3_1_BAC_16SrR AY672
AY531604.1 gg_id:104899 Indian Ocean hydrothermal vent clone SF_C23-F6 AY531
AJ132737.1 gg_id:20059 hydrothermal vent clone Top-147 AJ1327
AY672506.1 gg_id:107659 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH1_23_BAC_16Sr AY672
EU245507.1 gg_id:252330 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M2-D1 EU245
FJ165504.1 gg_id:539775 Microbially-enhanced Dissolution Karst Carbonates microbial biofilm Sonterra #8 well FJ1655
DQ295697.1 gg_id:160880 Epsilonproteobacteria terrestrial caves and springs floating microbial mat sulfidic gr DQ295
AF449236.1 gg_id:61874 Riftia pachyptila's tube clone R103-B37 AF449
AY672537.1 gg_id:105096 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH9_6_BAC_16SrR AY672
AJ969446.1 gg_id:158924 Microbial black smoker s Mid Atlantic Ridge ecological implications rock fragments c AJ9694
GU390848.1 gg_id:530358 Dark hot springs: structure biofilms caves Acquasanta Terme (Italy) sulfidic cave stre GU390
AM745162.1 gg_id:292405 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM745
AF420360.1 gg_id:67780 hydrothermal sediment clone AF420360.1 AF420
AF420342.1 gg_id:67795 hydrothermal sediment clone AF420342 AF420
AB105048.1 gg_id:93761 Hydrogenimonas thermophila str. EP1-55-1% AB105
AY261811.1 gg_id:83596 UASB reactor granular sludge clone PD-UASB-2 proteobacterium AY261
DQ295692.1 gg_id:161262 Epsilonproteobacteria terrestrial caves and springs floating microbial mat sulfidic gr DQ295
AY531574.1 gg_id:107543 Indian Ocean hydrothermal vent clone SF_C23-G7 AY531
AB278149.1 gg_id:184732 hydrothermal vent prokaryotes trapped 0.2- and 0.1-micro-m-filtrates obtained Mar AB278
AB175498.1 gg_id:106632 Thioreductor micantisoli str. BKB25Ts-Y AB175
EF467531.1 gg_id:213162 situ microelectrode voltammetry sulfur-oxidizing biofilms Frasassi sulfidic cave strear EF467
FJ497432.1 gg_id:556468 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4974
EF467559.1 gg_id:211322 situ microelectrode voltammetry sulfur-oxidizing biofilms Frasassi sulfidic cave strear EF467

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EF467581.1 gg_id:216198 situ microelectrode voltammetry sulfur-oxidizing biofilms Frasassi sulfidic cave strear EF4675
AM268717.1 gg_id:233206 Fluid chemistry and microbial southern hydrothermal field (Lilliput) on Mid-Atlantic AM268
AY280433.1 gg_id:106294 exterior deep-sea hydrothermal vent sulfide chimney Juan de Fuca Ridge Pacific Oce AY2804
DQ295651.1 gg_id:161606 Epsilonproteobacteria terrestrial caves and springs microbial mat sulfidic cave strea DQ295
EF462765.1 gg_id:299712 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4627
GU196200.1 gg_id:579980 Bioaugmentation improved recovery digesters after toxicant exposure lab-scale ana GU196
AB278151.1 gg_id:180340 hydrothermal vent prokaryotes trapped 0.2- and 0.1-micro-m-filtrates obtained Mar AB278
AY531573.1 gg_id:107393 Indian Ocean hydrothermal vent clone SF_C23-G8 AY5315
FJ799123.1 gg_id:584284 Microbial Responsible Intermediates Degradation Anaerobic Digestion System Deterr FJ7991
AJ441202.1 gg_id:57475 hydrothermal vent polychaete mucous clone P. palm C 140 AJ4412
DQ295708.1 gg_id:160328 Epsilonproteobacteria terrestrial caves and springs microbial mat sulfidic stream Mil DQ295
EF467511.1 gg_id:215667 situ microelectrode voltammetry sulfur-oxidizing biofilms Frasassi sulfidic cave strear EF4675
FJ485355.1 gg_id:545841 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4853
DQ228655.1 gg_id:141137 Microbial colonizing mineral surfaces within sulfide-microbial incubator clone CH2b1 DQ228
EF462722.1 gg_id:296261 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4627
EU234099.1 gg_id:246280 receiving river and un techniques anaerobic process WWTP treating penicillin G prod EU2340
AJ576002.1 gg_id:98839 deep sea hydrothermal vent clone T6-Ph07-983 AJ5760
GU390742.1 gg_id:517462 Dark hot springs: structure biofilms caves Acquasanta Terme (Italy) sulfidic cave stre GU390
AY082468.1 gg_id:67838 neutral pH mine biofilm clone 44a-B1-40 AY0824
AY280427.1 gg_id:101920 exterior deep-sea hydrothermal vent sulfide chimney Juan de Fuca Ridge Pacific Oce AY2804
EF467460.1 gg_id:205100 situ microelectrode voltammetry sulfur-oxidizing biofilms Frasassi sulfidic cave strear EF4674
AB198587.1 gg_id:135519 termite gut fluid and protists fraction clone RsC01-014 AB1985
DQ295656.1 gg_id:160561 Epsilonproteobacteria terrestrial caves and springs microbial mat sulfidic cave strea DQ295
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AY571498.1 gg_id:114816 termite gut clone RsaHf427 AY5714
GU369897.1 gg_id:528299 and mats vents Arc white filamentous microbial mat photic shal hydrothermal vent GU369
AY355299.2 gg_id:106198 epibiontic clone C11-E8 AY3552
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AY578096.1 gg_id:105995 Helicobacter bilis str. KI900 AY5780
AY251062.1 gg_id:103492 deep-sea hydrothermal biofilm Edmond vent field on Central Indian Ridge clone FT17 AY2510
EF462645.1 gg_id:297781 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4626
AF449237.1 gg_id:61872 Riftia pachyptila's tube clone R103-B43 AF4492
EF467591.1 gg_id:214381 situ microelectrode voltammetry sulfur-oxidizing biofilms Frasassi sulfidic cave strear EF4675
AY672507.1 gg_id:107804 hydrothermal vent 9 degrees North East Rise Pacific Ocean clone CH1_25_BAC_16Sr AY6725
EF462782.1 gg_id:290493 meta-genome level extreme microbial symbiosis dorsal surface deep sea hydrotherm EF4627
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U51874.2 gg_id:10979 Helicobacter sp. Dog-4 str. Dog-4; Eaton 1302 U51874
DQ071280.1 gg_id:140620 dominate filamentous sulfur mat 13 degree north field East Pacific Rise deep-sea hy DQ071
AY531583.1 gg_id:103545 Indian Ocean hydrothermal vent clone SF_C23-F3 AY5315
GU390798.1 gg_id:516081 Dark hot springs: structure biofilms caves Acquasanta Terme (Italy) sulfidic cave stre GU390

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AF420347.1 gg_id:67790 hydrothermal sediment clone AF420347.1 AF4203
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AF420343.1 gg_id:67794 hydrothermal sediment clone AF420343 AF4203
AF449232.1 gg_id:61877 Riftia pachyptila's tube clone R103-B18 AF4492
GQ355026.1 gg_id:565854 structure between two geochemically distinct springs Alum Rock Park spring clone M GQ355
L35522.1 gg_id:11150 Alvinella pompejana epibiont L35522
AJ969465.1 gg_id:157898 Microbial black smoker s Mid Atlantic Ridge ecological implications rock fragments ch AJ9694
EU864466.1 gg_id:334109 structures under selection pressure levels antibiotics Xiao River receiving treated ox\ EU8644
EF644836.1 gg_id:300292 genes and citrate lyase deep-sea microbial MAR Rainbow hydrothermal vent system EF6448
EF467464.1 gg_id:213273 situ microelectrode voltammetry sulfur-oxidizing biofilms Frasassi sulfidic cave strear EF4674
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GU390782.1 gg_id:516622 Dark hot springs: structure biofilms caves Acquasanta Terme (Italy) sulfidic cave stre GU390
AY766315.1 gg_id:108123 VC2.2 hydrothermal chimney Snake Pit site on mid Atlantic ridge clone 3 AY7663
GU390711.1 gg_id:515047 Dark hot springs: structure biofilms caves Acquasanta Terme (Italy) sulfidic cave stre GU390
AY280435.1 gg_id:106612 exterior deep-sea hydrothermal vent sulfide chimney Juan de Fuca Ridge Pacific Oce: AY2804
NC_005090.1 gg_id:93903 Wolinella succinogenes str. DSM 1740 NC_005
AF273252.1 gg_id:29152 Wolinella succinogenes str. SeR-1 AF2732
AF068812.2 gg_id:11156 hydrothermal vent clone VC2.1 Bac12 AF0688
AY691429.1 gg_id:106012 Caminibacter sp. str. TB-1 AY6914
NR_028943.1 gg_id:571654 Caminibacter hydrogeniphilus str. AM1116; DSM 14510 NR_028
AF299128.1 gg_id:40818 S17sBac34 complete clone AF2991
AF068804.2 gg_id:11155 hydrothermal vent clone VC2.1 Bac30 AF0688
ABCJ01000004.1 gg_id:243079 Caminibacter mediatlanticus str. TB-2 ABCJ01
NR_028999.1 gg_id:579615 Caminibacter profundus str. CR; DSM 15016 NR_028
CP001279.1 gg_id:528797 Nautilia profundicola str. AmH CP0012
NR_025497.1 gg_id:566284 Nautilia lithotrophica str. 525; DSM 13520 NR_025
AM937002.1 gg_id:260087 Nautilia abyssi str. PH1209 AM937
AP009178.1 gg_id:247747 Nitratiruptor sp. str. SB155-2 AP0091
AB175499.1 gg_id:106788 Nitratiruptor tergarcus str. MI55-1 AB1754
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DQ431197.1 gg_id:156297 Acidithiobacillus caldus str. YN06 DQ431
FJ154522.1 gg_id:556258 Acidithiobacillus caldus str. Da FJ1545
AF329205.1 gg_id:33408 Acidithiobacillus ferrooxidans str. ATCC23270 AF3292
DQ646482.1 gg_id:257953 Acidithiobacillus ferrooxidans str. DX-AF-1 DQ646
DQ646485.1 gg_id:238863 Acidithiobacillus ferrooxidans str. DX-AF-4 DQ646
EU079375.1 gg_id:243219 Acidithiobacillus ferrooxidans str. NF-01 EU0793
AB039820.1 gg_id:19750 Acidithiobacillus ferrooxidans str. NASF-1 AB0398

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GU167981.1 gg_id:583008 Acidithiobacillus ferrooxidans GU167
AF359940.1 gg_id:21015 Acidithiobacillus thiooxidans str. KCTC 8929P AF3599
AJ879995.1 gg_id:127417 Acidithiobacillus ferrooxidans str. B4 AJ8799
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EU250259.1 gg_id:241484 acid mine drainage clone GXDC-45 EU250
EF446188.1 gg_id:210453 Prokaryotic and Ecology Floating Macroscopic Filaments Extreme Acidic River Rio Tint EF4461
FJ946877.1 gg_id:567607 Acidithiobacillus ferrooxidans str. NK-16 FJ9468
AY552091.1 gg_id:147725 Acidithiobacillus sp. str. CASQ22 AY552
FJ890908.1 gg_id:574423 Microbial heap: implications model and control process -grade copper bioleaching he FJ8909
EF612419.1 gg_id:248238 historic semiarid lead-zinc site mine tailings clone K6-C19 EF6124
EF059761.1 gg_id:188237 Acidithiobacillus ferrooxidans str. CHMS EF0597
X75266.1 gg_id:7484 Acidithiobacillus ferrooxidans X75266
FJ154519.1 gg_id:539257 Acidithiobacillus ferrooxidans str. F6 FJ1545
AM502930.1 gg_id:241149 Acidithiobacillus ferrooxidans str. LMT1 AM502
EU084698.1 gg_id:254158 Acidithiobacillus ferrooxidans str. DGS EU084
DQ159176.1 gg_id:139484 Wetland System Treating Acid Coal Mine Drainage wetland clone AMD 636 DQ159
EU250217.1 gg_id:253906 acid mine drainage clone GXDC-3 EU250
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AJ458408.1 gg_id:67017 Aeromonas encheleia str. LMG 16328 AJ4584
AJ458416.1 gg_id:67011 Aeromonas encheleia str. E195 = S 191 AJ4584
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FJ940843.1 gg_id:465898 Aeromonas enteropelogenes str. CRX105 FJ9408
FJ940844.1 gg_id:457890 Aeromonas enteropelogenes str. CRX203 FJ9408
FJ940845.1 gg_id:394895 Aeromonas enteropelogenes str. CRX207 FJ9408
FJ966347.1 gg_id:411204 Aeromonas veronii str. 5CB1 FJ9663
FJ940820.1 gg_id:455969 Aeromonas veronii str. LNC205 FJ9408
FJ940816.1 gg_id:401894 Aeromonas jandaei str. LC208 FJ9408
EU770282.1 gg_id:320191 Aeromonas veronii str. IH118 EU770
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AF157690.1 gg_id:9454 Aeromonas media str. 86352 AF157
AY900170.1 gg_id:143274 Aeromonas hydrophila str. PCA0081 AY900
FJ940831.1 gg_id:449999 Aeromonas media str. pW28 FJ9408
EU488683.1 gg_id:545213 Aeromonas media str. 974a EU488
FJ168773.1 gg_id:559246 Aeromonas media str. 480a FJ1687
AF418218.1 gg_id:59117 Aeromonas media 5 AF418
GQ470996.1 gg_id:546104 Aeromonas hydrophila str. HQ070610B-1 GQ470
AM262152.1 gg_id:158666 Aeromonas media str. CDC 0862-83 AM262

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FJ494900.1 gg_id:424847 Aeromonas hydrophila str. B27
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AM262150.1 gg_id:157051 Aeromonas media str. V69
EU669667.1 gg_id:306838 Aeromonas hydrophila
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AY532688.1 gg_id:99610 Aeromonas molluscorum str. 93M
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FJ233860.1 gg_id:313618 Aeromonas jandaei str. AE25
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AF170914.4 gg_id:9385 Aeromonas culicicola str. MTCC 3249
EU918692.1 gg_id:347086 Pasteurella multocida str. SXXG0802
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EU770278.1 gg_id:327136 Aeromonas veronii str. IH101
EU770304.1 gg_id:316400 Aeromonas veronii str. MW17
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AY987741.1 gg_id:198899 Aeromonas hydrophila str. -32
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EU770292.1 gg_id:328368 Aeromonas veronii str. IQ309
EU770270.1 gg_id:308207 Aeromonas veronii str. HH104
EU770296.1 gg_id:316179 Aeromonas punctata str. ME15
AY928478.1 gg_id:138908 Aeromonas veronii bv. Sobria str. A155
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AY928475.1 gg_id:138989 Aeromonas veronii bv. Sobria str. CDC0437-84
FJ233858.1 gg_id:323934 Aeromonas allosaccharophila str. AE35
EU770271.1 gg_id:317346 Aeromonas veronii str. HQ103
FJ940797.1 gg_id:414411 Aeromonas punctata str. 4LNC310

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FJ940830.1 gg_id:371339 Aeromonas jandaei str. pW23
EF077527.1 gg_id:193001 Aeromonas hydrophila

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AM184224.1 gg_id:161336 Aeromonas veronii bv. Veronii str. WAB1882
X74674.1 gg_id:9452 Aeromonas punctata str. ATCC 15468T
AY928477.1 gg_id:151083 Aeromonas veronii bv. Sobria str. A155
FJ940796.1 gg_id:433024 Aeromonas punctata str. 4LNC309
AY987745.1 gg_id:188420 Aeromonas jandaei str. -51
FM999978.1 gg_id:531043 Aeromonas popoffii str. LMG 17545
FJ233863.1 gg_id:333045 Aeromonas bv. veronii str. AE34
GU296672.1 gg_id:591386 Aeromonas hydrophila str. ANSE2
AF427150.1 gg_id:25552 Aeromonas sp. PAR2A
FJ544354.1 gg_id:433080 Aeromonas punctata str. ji14
AY987739.1 gg_id:191655 Aeromonas hydrophila str. -25
AY987749.1 gg_id:188082 Aeromonas bestiarum str. LMG 13448
EU770273.1 gg_id:326038 Aeromonas veronii str. HQ119
EU770286.1 gg_id:340453 Aeromonas veronii str. IH325
FJ940804.1 gg_id:372890 Aeromonas jandaei str. 4pM28
EU700096.1 gg_id:296747 Aeromonas hydrophila str. SWCH-6
AJ458411.1 gg_id:67014 Aeromonas eucrenophila str. LMG 17059
EU770293.1 gg_id:324097 Aeromonas veronii str. IQ318
EU488685.1 gg_id:549142 Aeromonas punctata str. 121c
EU770302.1 gg_id:315651 Aeromonas veronii str. MS15
EU488693.1 gg_id:575182 Aeromonas veronii str. 633c
EU770303.1 gg_id:334972 Aeromonas veronii str. MW15
FJ407187.1 gg_id:361976 Aeromonas punctata str. W20060720
AJ224309.1 gg_id:9396 Aeromonas encheleia str. 1881
AF418214.1 gg_id:56067 Aeromonas media 1
FJ544362.1 gg_id:411774 Aeromonas punctata str. ji31
FJ940810.1 gg_id:390498 Aeromonas veronii str. 4pW23
AB211227.1 gg_id:115297 Aeromonas salmonicida str. NKCM8904
FJ940832.1 gg_id:423276 Aeromonas jandaei str. CCC204
AY987724.1 gg_id:180428 Aeromonas veronii bv. Sobria str. AE-21
GQ262778.1 gg_id:592654 Aeromonas hydrophila str. S1
AB034761.1 gg_id:74870 Aeromonas hydrophila str. 28SA
FJ940833.1 gg_id:433791 Aeromonas hydrophila str. CHS101
DQ539497.1 gg_id:169858 Aeromonas hydrophila SBRD9
DQ816568.1 gg_id:161901 mouse cecum clone aab19g05

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FJ026735.1 gg_id:338819 Aeromonas punctata str. ZZF071029
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EU770290.1 gg_id:326583 Aeromonas veronii str. IQ304
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EU770285.1 gg_id:332437 Aeromonas veronii str. IH323
EU770288.1 gg_id:353344 Aeromonas veronii str. IQ117
EU730930.1 gg_id:311549 Aeromonas eucrenophila str. 189
EU770274.1 gg_id:327072 Aeromonas hydrophila str. IB101
DQ837028.1 gg_id:162354 Aeromonas jandaei sf3-14
FJ230076.1 gg_id:587653 Aeromonas sanarellii str. A2-67
AY987723.1 gg_id:198790 Aeromonas punctata str. AE-11
FJ940822.1 gg_id:389565 Aeromonas veronii str. QC102
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FJ976606.1 gg_id:545344 Aeromonas hydrophila
EF446897.1 gg_id:215466 Aeromonas veronii str. Philippines-An11
FJ544406.1 gg_id:447507 Aeromonas punctata str. w4
AY987733.1 gg_id:192245 Aeromonas hydrophila str. AE-55
EU770289.1 gg_id:343054 Aeromonas veronii str. IQ120
FJ947060.1 gg_id:508769 Aeromonas punctata str. nA3
AJ223181.1 gg_id:9390 Aeromonas popoffii str. LMG 17543
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FJ940815.1 gg_id:413385 Aeromonas jandaei str. LC206
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FJ230077.1 gg_id:589204 Aeromonas taiwanensis str. A2-50
AY987726.1 gg_id:190833 Aeromonas enteropelogenes str. AE-31
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CP001616.1 gg_id:526816 Tolumonas auensis str. DSM 9187 CP0016
EU569287.1 gg_id:274900 Zobellella denitrificans str. MW1 EU5692
EU109731.1 gg_id:237124 Shewanella sp. str. NH21 EU1097
FJ230836.1 gg_id:346053 Serratia sp. str. AN4 FJ2308
DQ195676.1 gg_id:140125 Zobellella taiwanensis str. ZT1T DQ195
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EU776462.1 gg_id:320643 Red Kangaroo feces clone KO2_aai17h02 EU7764
EU794084.1 gg_id:309826 bovine microbiota via oligonucleotide fingerprinting genes fecal clone EMP_E16 EU7940
GU303104.1 gg_id:577587 inhabiting epithelial tissue ruminal digesta -DGGE and sequencing rumen epithelium GU303
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FJ673136.1 gg_id:513301 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6731
FJ681014.1 gg_id:514935 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6810
FJ678936.1 gg_id:529684 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6789
FJ680138.1 gg_id:514703 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6801
FJ673291.1 gg_id:522804 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6732
FJ681705.1 gg_id:517924 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6817
FJ680474.1 gg_id:517852 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6804
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FJ680476.1 gg_id:521800 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6804
FJ679937.1 gg_id:525643 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6799
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FJ681292.1 gg_id:530658 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6812

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EF445277.1 gg_id:205408 dairy cow rumen clone NED4F7 EF4452

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FJ680116.1 gg_id:526488 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6801
FJ679573.1 gg_id:530904 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6795
EU794184.1 gg_id:344867 bovine microbiota via oligonucleotide fingerprinting genes fecal clone EMP_B23 EU7941
FJ679565.1 gg_id:527828 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6795
FJ679188.1 gg_id:520121 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6791
FJ673005.1 gg_id:530535 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6730
Y17599.1 gg_id:9351 Succinimonas amylolytica str. DSM 2873 Y17599
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FJ440064.1 gg_id:511871 Megamonas hypermegale colonization correlates Campylobacter jejuni exclusion pou FJ4400
AJ420985.1 gg_id:29865 Anaerobiospirillum thomasii str. DSM 11806T AJ4209
FJ832419.1 gg_id:577393 Microbial Ecology Crop Folivorous Hoatzin crop contents clone P1Q401_75C10 FJ8324
GU303829.1 gg_id:539467 inhabiting epithelial tissue ruminal digesta -DGGE and sequencing rumen contents c GU303
AF018457.1 gg_id:9354 Aeromonas subgroup subdivision clone 12-11 AF0184
FJ832403.1 gg_id:548716 Microbial Ecology Crop Folivorous Hoatzin crop contents clone P1Q401_45E06 FJ8324
FJ832430.1 gg_id:559309 Microbial Ecology Crop Folivorous Hoatzin crop contents clone P1Q401_91C12 FJ8324
DQ456106.1 gg_id:158543 pre-adolescent turkey cecum clone CFT114E2 DQ456
EU381934.1 gg_id:271179 Rumen eubacterial implicates Bacteroidetes major phylum rumen particulate phase EU381
FJ952818.1 gg_id:443894 Alteromonas sp. str. 2tb4 FJ9528
GQ391974.1 gg_id:519442 Aestuariibacter sp. str. PaD1.02 GQ391
AB076557.1 gg_id:60317 Agarivorans albus str. MKT 82 AB0765
GQ200591.1 gg_id:544653 Agarivorans sp. str. WH0801 GQ200
DQ416550.1 gg_id:153331 s1cb18 DQ416
AF290480.1 gg_id:27315 cf. Alishewanella fetalis clone Llangefni 75 mite AF2904
AF144407.1 gg_id:33192 Alishewanella fetalis str. CCUG 30811 AF1444
AY093457.1 gg_id:55811 deep marine sediment clone MB-A2-102 AY0934
DQ256339.1 gg_id:142481 subsurface water clone EV818BHEB5102702SAS10 DQ256
AY820719.1 gg_id:109054 drilling fluid clone CCSD_DF2030_B15 AY8207
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EU189225.1 gg_id:239877 Alteromonas marina str. MB 3 EU189
AY881234.1 gg_id:145384 Alteromonas marina SCH0402 AY8812
AF288360.1 gg_id:71626 Alteromonas alvinellae str. HY757 AF2883
AF386742.1 gg_id:44702 Alteromonas sp. str. 4BT AF3867
AJ319537.1 gg_id:71568 Alteromonas macleodii subsp. abyssi str. HYD657 subsp. AJ3195
AM931040.1 gg_id:239418 Alteromonas sp. str. BUS 07-8 AM931
DQ836766.1 gg_id:163484 Alteromonas simiduii str. BCRC 17572 DQ836
AF114499.1 gg_id:8878 attached marine recovered surface clone 17 proteobacterium AF1144
EF061429.1 gg_id:190836 Alteromonas sp. NJSX37 EF0614
EU982324.1 gg_id:322486 Alteromonas sp. str. UST010723-013 EU9823
FJ210814.1 gg_id:309174 Alteromonas sp. str. QDHT-19 FJ2108
DQ486510.1 gg_id:227411 Alteromonas sp. str. DG1302 DQ486
AF114505.1 gg_id:8883 attached marine recovered deep chlorophyll maximum depth clone 2 proteobacterium AF1145

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AF114498.1 gg_id:8877 attached marine recovered surface clone 14 proteobacterium AF1144
AF114504.1 gg_id:8882 attached marine recovered depth 400m clone 25 proteobacterium AF1145
AF114508.1 gg_id:8886 attached marine recovered surface clone 8 AF1145
AY428573.1 gg_id:105550 Alteromonas litorea str. TF-22 AY4285
Y18232.1 gg_id:19130 Alteromonas macleodii str. CH-518 Y18232
AY344429.1 gg_id:91492 Lake Kauhako 30 m clone K2-30-18 AY3444
AM885869.1 gg_id:240896 Alteromonas macleodii R-28696 AM885
AF114501.1 gg_id:8880 attached marine recovered surface clone 18 proteobacterium AF1145
AF114496.1 gg_id:8875 attached marine recovered deep chlorophyll maximum depth clone proteobacterium AF1144
EF574195.1 gg_id:228587 site S25 near Coco's Island marine clone S25_539 EF5741
DQ071162.1 gg_id:136631 marine surface water clone Surf1.42 DQ071
AM882576.1 gg_id:311934 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
AY165578.1 gg_id:81399 Antarctic pack ice Lasarev Sea Southern Ocean clone ANTXI/4_14-11 AY1655
AF468279.1 gg_id:80086 Arctic pack ice; northern Fram Strait; 80 31.1 N; 01 deg 59.7 min E clone ARKIA-118 AF4682
CP000388.1 gg_id:160590 Pseudoalteromonas atlantica str. T6c CP0003
AY697880.1 gg_id:104267 Antarctica clone F2C28 AY6978
U85854.1 gg_id:8873 Glaciecola pallidula str. ACAM 615T IC079 U85854
AY167284.2 gg_id:81210 Glaciecola sp. str. ANT9142 AY1672
AB238790.1 gg_id:140832 Glaciecola sp. #105 AB2387
AM941178.1 gg_id:284078 Microbes enriched seawater after addition coral mucus and reef waters mixture1-8 AM941
AM913910.1 gg_id:439090 antibiotic-active brown alga Laminaria saccharina Baltic Sea Saccharina latissima AM913
AJ240915.1 gg_id:73940 5' clone CHAB-XII-8 AJ2409
AY771709.1 gg_id:108683 Glaciecola mesophila str. S2-38 AY7717
AJ488501.1 gg_id:65357 Glaciecola mesophila str. KMM 241 AJ4885
Y18230.1 gg_id:202397 macleodii- and cellular rDNAs Mediterranean seawater mesocosm experiment Alterom Y18230
AJ548479.1 gg_id:89023 Glaciecola mesophila str. LMG 21855 AJ5484
AF173965.1 gg_id:71399 Alteromonas macleodii str. KT1113 AF1739
AY349463.1 gg_id:91221 Glaciecola sp. WHOI JT-6b AY3494
U73724.1 gg_id:8898 gas vacuolate str. 214.6 U73724
AJ293820.1 gg_id:39858 Glaciecola polaris str. LMG 21857 AJ2938
DQ784575.1 gg_id:162089 Glaciecola agarolytica str. NO2 DQ784
AY553295.1 gg_id:106840 Salinimonas chungwhensis str. BH030046 AY5532
EU583725.1 gg_id:272031 Alteromonas sp. str. JSM 073008 EU5837
AJ240916.1 gg_id:74015 5' clone CHAB-V-29 AJ2409
AB040465.1 gg_id:38060 Alteromonas sp. str. K AB0404
FJ024319.1 gg_id:368249 Prokaryotic and their role biogeochemical cycling marine basalts basalt Juan de Fuca I FJ0243
AB473549.1 gg_id:419632 Aestuariibacter sp. NT-2008 str. Sd 2-38 (= KMM 3894) AB4735
AY207502.1 gg_id:84007 Aestuariibacter salexigens str. JC2042 AY2075
AY207503.1 gg_id:83231 Aestuariibacter halophilus str. JC2043 AY2075
AB040466.1 gg_id:45496 Alteromonas sp. str. D AB0404
FJ202664.1 gg_id:350427 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2026

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EU188146.1 gg_id:347006 Uncultivated oceanic unicellular N2-fixing cyanobacteria are picoplankton unusual 7(EU188
FJ425616.1 gg_id:366875 communnities healthy and diseased colonies Caribbean coral Montastrea faveolata A FJ4256
GU289642.1 gg_id:552344 Alteromonas sp. str. MEBiC06033 GU289
EU907922.1 gg_id:304111 Alteromonas sp. str. SKUK MB1007 EU907
DQ343294.1 gg_id:149488 Bowmanella denitrificans str. BD1 DQ343
EF092177.1 gg_id:275572 Persistent Axinella corrugata Alteromonas clone AC3_H2 EF0921
AJ295715.1 gg_id:73259 Alteromonas stellipolaris str. LMG 21861 AJ2957
AM882575.1 gg_id:316071 Structure along hydrocarbon contamination coastal sediment oil polluted water pet AM882
EU287412.1 gg_id:256595 Pacific arctic surface sediment clone S26-112 EU287
AM990642.1 gg_id:277954 Alteromonas sp. MOLA 365 AM990
AY926461.1 gg_id:113825 Alteromonas hispanica str. B-35 AY9264
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AB526336.1 gg_id:573848 Alteromonas sp. str. JAM-GA13 AB5263
Y18233.1 gg_id:8895 Alteromonas macleodii str. CH-460 Y18233
EU188189.1 gg_id:317008 Uncultivated oceanic unicellular N2-fixing cyanobacteria are picoplankton unusual 7(EU188
GQ495024.1 gg_id:581173 Pseudoalteromonas sp. str. W-14-2 GQ495
AM183347.1 gg_id:155988 Alishewanella baltica str. BA131 AM183
EU841485.1 gg_id:310357 Alishewanella sp. str. 205 EU8414
AB295494.1 gg_id:216998 Alteromonas sp. str. A14 AB2954
NR_027234.1 gg_id:540876 Algicola sagamiensis str. B-10-31; DSM 14643 NR_02
GU108538.1 gg_id:591045 active chemolithoautotrophic prokaryotes sulfidic zone Black pelagic redoxcline dete GU108
EU714266.1 gg_id:332103 Alteromonas sp. str. BG20 EU714
DQ913886.1 gg_id:276341 Celerinatantimonas diazotrophica J-G1-2 DQ913
AJ002006.1 gg_id:8823 Rheinheimera baltica str. OS140 Baltic # 166 AJ0020
DQ985066.1 gg_id:177157 Rheinheimera sp. str. JL1145 DQ985
GQ009534.1 gg_id:500133 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw77 GQ009
GQ280117.1 gg_id:570209 Rheinheimera aquimaris str. JS-47 GQ280
FJ170027.1 gg_id:325282 Rheinheimera sp. str. CF12-10 FJ1700
FJ821606.1 gg_id:569335 Rheinheimera sp. str. TPS16 FJ8216
DQ298025.2 gg_id:143812 Rheinheimera chironomi str. K19414 DQ298
FJ902308.1 gg_id:509338 phreatic limestone sinkholes northeastern Mexico biomat rock outcrop cenote La Pal FJ9023
AM117933.1 gg_id:143326 marine boreal sponge Geodia baretti Norway:Korsfjord AM117
GQ240227.1 gg_id:576918 Rheinheimera sp. str. BZ19 GQ240
EU662623.1 gg_id:594183 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems floating microb EU6626
GQ130023.1 gg_id:440958 Genetic inventory 'JPL-SAF' during MSL mission spacecraft assembly clean room floc GQ130
AY887010.1 gg_id:114447 fresh water clone 1.34.5 AY8870
FJ902639.1 gg_id:593719 phreatic limestone sinkholes northeastern Mexico biofilm 1m deep hydrothermal ver FJ9026
AY701891.1 gg_id:140048 Rheinheimera texana str. A62-14B AY7018
GU108532.1 gg_id:548750 active chemolithoautotrophic prokaryotes sulfidic zone Black pelagic redoxcline dete GU108
FN397813.1 gg_id:405654 geomicrobiological investigation sediments Gulf Cadiz GeoB 9072-1 anoxic sub-surfa FN3978
AY167340.1 gg_id:81146 Psychrobacter sp. str. ARK10133 AY1673

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FN397812.1 gg_id:392867 geomicrobiological investigation sediments Gulf Cadiz GeoB 9072-1 anoxic sub-surfa FN397
GU061308.1 gg_id:560509 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061
U85841.1 gg_id:8840 Colwellia psychrerythraea str. ACAM 604 U8584
AF001375.1 gg_id:8836 Colwellia psychrerythraea str. ATCC27364; ACAM 550T AF001
AF530145.1 gg_id:61116 Colwellia psychrerythraea str. MGP-41 AF530
AY167332.1 gg_id:82309 Colwellia rossensis str. ANT9271 AY167
AY167269.1 gg_id:82300 Colwellia rossensis str. ANT8283 AY167
FJ905674.1 gg_id:560989 s Tonga-Kermadec iron oxide chimney-like structure on Volcano 19 Tonga Arc clone V FJ9056
DQ027051.1 gg_id:130223 Colwellia sp. str. MT41 DQ027
EF551378.1 gg_id:225725 Colwellia sp. str. BSw20516 EF5513
EU050773.1 gg_id:257619 libraries sediment Kings Bay Svalbard Arctic clone SS1_B_01_78 EU050
DQ837294.1 gg_id:235556 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_1B_55 DQ837
DQ490038.1 gg_id:158539 Microbial life ridge flank crustal fluids clone ODP-70B-02 DQ490
AY167330.1 gg_id:81643 Colwellia sp. str. ANT9407 AY167
DQ351740.1 gg_id:146764 Microbial Adherent Sediment Particles Heavy Metal Contaminated North Sea Surfac DQ351
GQ350647.1 gg_id:559927 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
GQ848466.1 gg_id:581903 Microbial sediments vents along East Lau Spreading Centre deep sea hydrothermal GQ848
L10950.1 gg_id:8854 marine snow clone agg53 L10950
AY829231.2 gg_id:179606 Colwellia sp. IE7-5 AY829
EU394582.1 gg_id:309555 Coccolithophore bloom dynamics shape (NE-Atl) northern Bay Biscay pelagic particle EU394
DQ027052.1 gg_id:131520 Colwellia sp. str. KT27 DQ027
GU391219.1 gg_id:531807 Thalassomonas sp. str. JYr2 GU391
AY165571.1 gg_id:81437 Antarctic pack ice Lasarev Sea Southern Ocean clone ANTXI/4_14-151 sea AY165
AB094412.1 gg_id:97603 Colwellia piezophila str. Y223G AB094
AF419350.1 gg_id:74479 Colwellia sp. str. Ant5-9 AF419
EU491932.1 gg_id:271119 Abundance and microbial life ocean crust deep seawater East Pacific Rise clone EPR4 EU491
EU599214.3 gg_id:319654 Colwellia sp. str. KMD002 EU599
EU491916.1 gg_id:258657 Abundance and microbial life ocean crust deep seawater East Pacific Rise clone EPR4 EU491
DQ492707.1 gg_id:158794 Colwellia sp. str. BSi20061 DQ492
NR_024635.1 gg_id:548903 Colwellia maris str. ABE-1 NR_02
EU491896.1 gg_id:259552 Abundance and microbial life ocean crust deep seawater East Pacific Rise clone EPR4 EU491
AY165566.1 gg_id:81127 Antarctic pack ice Lasarev Sea Southern Ocean clone ANTXI/13-90 sea AY165
AF530146.1 gg_id:103969 Colwellia sp. str. MGP-59 AF530
GU108544.1 gg_id:589927 active chemolithoautotrophic prokaryotes sulfidic zone Black pelagic redoxcline dete GU108
AY167264.1 gg_id:81818 Colwellia sp. str. ANT8203 AY167
DQ923440.1 gg_id:179604 Alteromonas sp. W-6 DQ923
DQ228659.1 gg_id:141317 Microbial colonizing mineral surfaces within sulfide-microbial incubator clone CH2b7 DQ228
DQ212914.1 gg_id:140089 Thalassomonas agarivorans str. TMA1 DQ212

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FJ717227.1 gg_id:553622 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7172
FJ202297.1 gg_id:322290 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2022
GU230351.1 gg_id:594414 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal sediment clo GU230
FJ203045.1 gg_id:322845 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
AY643537.2 gg_id:131602 White plague like disease Eilat coral reef AY6435
AY194066.1 gg_id:81968 Thalassomonas ganghwensis str. JC2041 AY1940
FJ203217.1 gg_id:306276 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
EU491328.1 gg_id:269454 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU4913
U15115.1 gg_id:8864 hydrothermal vent clone PVB_OTU_12 U15115
FJ791872.1 gg_id:513615 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7918
FJ203091.1 gg_id:324976 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
AJ294747.1 gg_id:26868 Thalassomonas viridans str. XOM5 CECT 5082 DSM 13753 AJ2947
AY770724.1 gg_id:110997 sea urchin Paracentrotus lividus clone PIS53 AY7707
FJ952826.1 gg_id:382239 Colwellia sp. str. 1tb5 FJ9528
AB193748.1 gg_id:111458 Ferrimonas marina str. A2A-30 AB1937
AB252733.1 gg_id:243052 Paraferrimonas sedimenticola str. Mok-142 AB2527
GU391222.1 gg_id:528038 Ferrimonas sp. str. JYr13 GU391
FJ203654.1 gg_id:341133 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
EU620413.1 gg_id:278677 Ferrimonas balearica str. S8-05 EU6204
AB193752.1 gg_id:112098 Ferrimonas balearica str. A2A-18 AB1937
AB193755.1 gg_id:111912 Ferrimonas sp. str. A3B-58 AB1937
NR_027560.1 gg_id:549106 Idiomarina baltica OS145 str. OS145; DSM 15154 NR_027
NC_006512.1 gg_id:128686 Idiomarina loihiensis str. L2TR NC_006
AE017340.1 gg_id:128915 Idiomarina loihiensis str. L2TR AE0173
AY092077.1 gg_id:67499 Idiomarina loihiensis str. 53 AY0920
AY635468.1 gg_id:107632 Idiomarina seosinensis str. CL-SP19 AY6354
EU603445.1 gg_id:275762 Idiomarina seosinensis str. PTG4-5 EU6034
AY553081.1 gg_id:107227 Idiomarina loihiensis str. GSP41 AY5530
NR_029115.1 gg_id:557590 Idiomarina fontislapidosi str. F23; LMG 22169 NR_029
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DQ235566.1 gg_id:142194 biofilm water pipeline biofilms steel pipelines Gulf Mexico clone 20 DQ235
AY526862.1 gg_id:102489 Idiomarina ramblicola str. R22 AY5268
AY505529.1 gg_id:101950 Idiomarina loihiensis str. GSP13 AY5055
FM877003.1 gg_id:574998 Idiomarina sp. IT-2008 str. CVS-6 FM877
DQ234155.2 gg_id:144284 determined library mangrove clone DS071 DQ234
EF486355.1 gg_id:222466 Idiomarina salinae str. ISL-52 EF4863
GQ180187.1 gg_id:591011 Pseudidiomarina sp. str. SW5 GQ180
GQ246209.1 gg_id:577128 Pseudidiomarina sp. str. YCSA4 GQ246
DQ416777.1 gg_id:268666 Idiomarina sp. G2DM-8 DQ416
DQ118948.1 gg_id:131585 Pseudoidiomarina taiwanensis str. PIT1 DQ118

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FJ581032.1 gg_id:541977 Idiomarina sp. str. IITRM14 FJ5810
EF212001.1 gg_id:214514 Pseudidiomarina sp. str. c121 EF2120
EU600202.2 gg_id:260949 Pseudidiomarina sp. str. 8C EU600
EU722654.1 gg_id:327394 Microbial athalassohaline Tirez lagoon (Spain) sediment clone SB43 EU722
EF554907.1 gg_id:222862 Idiomarina sp. str. G-He8 EF5549
GU458902.1 gg_id:562858 microbiota human male female forehead forearm and influence make-up skin clone GU458
X92129.1 gg_id:44150 Crater Lake isolate 11C1 X92129
GU471247.1 gg_id:517013 Idiomarina sp. str. 10-D-4 GU471
EF554896.1 gg_id:226059 Idiomarina sp. str. JK38 EF5548
FJ152984.1 gg_id:347471 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1529
EF554906.1 gg_id:223511 Idiomarina sp. str. G-He7 EF5549
X92128.1 gg_id:32525 Lake Nakuru isolate 20N1 X92128
D21224.1 gg_id:8975 Moritella japonica str. DSK1 D21224
AF530133.1 gg_id:70982 Moritella sp. str. MGP-6 AF5301
AJ132226.1 gg_id:8977 Moritella viscosa str. NVI 88/478T AJ1322
AB189343.1 gg_id:112983 cold seep sediment clone JT58-18 AB1893
AJ297540.1 gg_id:48310 Moritella marina str. NCIMB 2263 AJ2975
AY700605.1 gg_id:109434 Australia: Great Barrier Reef clone PDA-OTU7 AY7006
AB183504.1 gg_id:105913 Moritella sp. str. 47A1 AB1835
AB008797.1 gg_id:8989 Moritella yayanosii str. DB21MT-5 AB0087
AY382585.1 gg_id:96373 Moritella sp. str. 2-2-2-1 AY3825
AB038033.1 gg_id:33284 Moritella marina str. ATCC15381T AB0380
NR_025380.1 gg_id:573974 Moritella abyssi str. 2693; JCM 11436 NR_025
AJ252023.2 gg_id:72172 Moritella profunda str. 2674 AJ2520
AB183499.1 gg_id:101616 Moritella sp. str. 38C1 AB1834
AB183496.1 gg_id:101122 Moritella sp. str. 36C1 AB1834
AB013841.1 gg_id:8993 Moritella sp. 9-11.F AB0138
AB013842.1 gg_id:8994 Shewanella sp. 7-1.E AB0138
AB040467.1 gg_id:20333 Pseudoalteromonas sp. str. R AB0404
DQ349123.1 gg_id:147282 Pseudoalteromonas aurantia DQ349
CR954246.1 gg_id:140486 Pseudoalteromonas haloplanktis str. TAC125 CR9542
CR954246.1 gg_id:140406 Pseudoalteromonas haloplanktis str. TAC125 CR9542
AY040230.1 gg_id:21879 Pseudoalteromonas translucida str. KMM520 AY0402
AF316144.1 gg_id:52879 Pseudoalteromonas issachenkonii str. KMM 3549; F13 AF3161
EU880522.1 gg_id:351145 Pseudoalteromonas elyakovii str. P18 EU8805
AF316142.1 gg_id:37455 Pseudoalteromonas issachenkonii str. KMM 3558; F2 AF3161
FJ457238.1 gg_id:568943 Pseudoalteromonas luteoviolacea str. S4060 FJ4572
FJ952795.1 gg_id:446774 Pseudoalteromonas sp. str. 1tc11 FJ9527
FJ200646.1 gg_id:308744 Pseudoalteromonas haloplanktis str. I 6 FJ2006
EF584051.1 gg_id:228654 Pseudoalteromonas sp. str. GAS9 EF5840
EF584064.1 gg_id:227631 Pseudoalteromonas sp. str. LS3 EF5840

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DQ531952.1 gg_id:161487 Pseudoalteromonas sp. str. MJ5322
X82147.1 gg_id:8972 Pseudoalteromonas rubra str. ATCC 29570T
FJ457188.1 gg_id:539701 Pseudoalteromonas rubra str. S2676
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AF529062.1 gg_id:60988 Pseudoalteromonas citrea str. CIP 105339
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FJ200640.1 gg_id:341432 Pseudoalteromonas agarivorans str. C 36
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AF539773.1 gg_id:84384 Pseudoalteromonas sp. str. E36
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X82136.1 gg_id:8908 Pseudoalteromonas carrageenovora str. ATCC 12662T
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AY864628.1 gg_id:111346 Pseudoalteromonas sp. str. U50
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AF313497.1 gg_id:111481 Pseudoalteromonas sp. str. A3
DQ923439.1 gg_id:187538 Pseudoalteromonas sp. W-5
AB257336.2 gg_id:182468 Pseudoalteromonas mariniglutinosa str. Do-52
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AJ507251.1 gg_id:59526 Pseudoalteromonas mariniglutinosa str. KMM 3635
FJ172090.1 gg_id:348422 Pseudoalteromonas sp. str. SBC7

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