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AJ867253.1 gg_id:111057 Roseobacter gallaeciensis X6 AJ8672
DQ514304.1 gg_id:161029 Roseobacter sp. str. BSs20188 DQ514
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DQ915619.1 gg_id:311058 Phaeobacter gallaeciensis str. NBRC16654 DQ915
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GQ348550.1 gg_id:533795 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
FJ203339.1 gg_id:314415 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2033
FJ202225.1 gg_id:350338 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2022
FJ716879.1 gg_id:554782 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7168
FJ203234.1 gg_id:305694 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
DQ811850.1 gg_id:172275 mangrove soil clone MSB-3ax5 DQ811
EU799430.1 gg_id:308804 It's all ranking aquatic Newport Harbour RI clone 1C227029 EU799
FJ203312.1 gg_id:327709 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2033
FJ664800.1 gg_id:554245 Quantitative dynamics cells plankton-fed microbial fuel cell clone plankton_D11 FJ6648
EU642854.1 gg_id:279275 Pseudoruegeria sp. JC2496 str. JC2496; IMSNU 14150 EU642
EU246829.1 gg_id:248590 polyphasic description Pocillopora meandrina Palmyra Atoll Calcinus obscurus abdon EU246
AF384204.1 gg_id:23935 Rhodobaca bogoriensis str. CL1 AF384
EF554833.1 gg_id:224271 Rhodobaca barguzinensis str. VKM B-2406 EF5548
AF513929.1 gg_id:62508 Rhodobacter group clone LA1-B42N AF513
AJ605746.2 gg_id:148330 Rhodobacter sp. str. EL-50 AJ6057
AF454295.1 gg_id:63841 hypersaline Mono Lake clone ML812M-6 AF454
AJ132383.1 gg_id:73883 Natronohydrobacter thiooxidans str. AHO 1 sediments ly alkaline soda lakes AJ1323
DQ659236.1 gg_id:170558 Roseinatronobacter monicus str. Ros35 DQ659
AB369166.1 gg_id:243619 Microbiological Assessment Circulation Mud Fluids During First Operation Drilling De AB369
DQ432310.1 gg_id:156883 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake Hamra sediment clon DQ432
AF454292.1 gg_id:63844 hypersaline Mono Lake clone ML602M-13 AF454
DQ432219.1 gg_id:156640 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake UmRisha sediment cl DQ432
DQ432393.1 gg_id:156929 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake Hamra water clone W DQ432

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EU703461.1 gg_id:377032 lakes on Eastern Tin Plateau freshwater lake altitude 4279 masl clone XZZLH1 EU703
AB355088.1 gg_id:240664 Monitoring and archaeal two heavy polluted areas lake Egypt based on sediment Ma AB355
AY212706.1 gg_id:102583 water 10 m downstream manure clone 257ds10 AY212
EU580696.1 gg_id:268039 Rhodobacter sp. str. NP25b EU580
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EU104326.1 gg_id:321409 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
GU257946.1 gg_id:566382 Rhodobacter changlensis str. HXG-D9 GU257
DQ342322.1 gg_id:214765 Rhodobacter blasticus str. ATCC 33485; KC2138 DQ342
AB077986.1 gg_id:67721 Rhodobacter gluconicum AB077
DQ676414.1 gg_id:225039 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
AM690348.2 gg_id:225526 Rhodobacter katedanii str. JA234 AM690
EU652478.1 gg_id:277835 Catellibacterium sp. str. JPB-2.07 EU652
GU454898.1 gg_id:520640 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
EF192900.1 gg_id:201726 sediments temperate artificial lake Rapel reservoir sediment clone Rap2_6C EF1929
AF234745.1 gg_id:35565 sludge clone A26 AF2347
FJ382199.1 gg_id:512277 Potentially pathogenic and air stem cell transplant summer hospital shower water BN FJ3821
AF035433.1 gg_id:5793 Rhodobacter apigmentum AF0354
EU464681.1 gg_id:301082 Hartmann's mountain zebra feces clone AFZEB_aaj69a12 EU4646
AY957924.1 gg_id:114895 drinking water biofilm clone B3NR69D9 AY9579
DQ337005.1 gg_id:147896 subsurface water clone EV818CFSSAHH73 DQ337
EU979475.1 gg_id:307279 Rhodobacter sp. str. CR07-8 EU9794
AB122032.1 gg_id:98295 Rhodobacter sp. str. Jip03 AB1220
D16429.1 gg_id:5800 Rhodobacter blasticus str. ATCC33485 D16429
AF368183.1 gg_id:58379 non-EBPR sludge clone SBRT155 AF3681
FJ196030.1 gg_id:333434 Rhodobacter sp. str. ZS3-33 FJ1960
AM696303.1 gg_id:217363 Rhodovulum marinum str. JA219 AM696
D16420.1 gg_id:5840 Rhodovulum sp str. MB260 D16420
U55277.1 gg_id:5835 Rhodovulum sulfidophilum str. W-1S=FERM P-15320 U55277
GU370080.1 gg_id:519755 Rhodovulum sulfidophilum str. P210(4) GU370
EU770422.1 gg_id:306266 Rhodovulum sulfidophilum str. sy59 EU7704

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AM921780.1 gg_id:291146 Rhodovulum puniceilacus str. JA322 AM921
NR_025222.1 gg_id:555568 Albidovulum inexpectatum str. FRR-10 NR_02
AB079635.2 gg_id:62339 Rhodovulum strictum str. JCM 9221 AB079
NR_025608.1 gg_id:568416 Roseisalinus antarcticus str. EL-88; DSM 11466 NR_02
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FJ897782.1 gg_id:543161 Roseivivax sp. str. BH87090 FJ8977
AB166989.1 gg_id:102398 Roseivivax sp. str. NT N51 AB166
GQ243710.1 gg_id:578525 consortia immobilizing nitrifying bioreactors designed penaeid and non-penaeid larv GQ243
EF512126.1 gg_id:220146 Roseobacter denitrificans str. CSQ-3 EF5121
NC_008209.1 gg_id:163757 Roseobacter denitrificans str. OCh 114 NC_00
GQ249575.1 gg_id:513025 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 12-14 cm layer cl GQ249
AY654757.1 gg_id:104921 Oculina mucus isolate 112 AY654
AY177714.1 gg_id:83398 Rhodobacteraceae T11 AY177
DQ446103.2 gg_id:164817 Surface Mucopolysaccharide Layer and Band Microbial Mat Black Band-Diseased Sid DQ446
AJ844283.1 gg_id:160252 Rubellimicrobium thermophilum str. E-R2A-8a AJ8442
FM874293.1 gg_id:360432 occupant source house mattress dust clone MC04B03 FM874
EU338486.1 gg_id:242423 Pararubellimicrobium aerilata str. 5715S-9 EU338
EF547368.1 gg_id:218626 Rubellimicrobium sp. str. MSL 20 EF5473
DQ532142.1 gg_id:167156 Geographical Variation Microbial Burden and Spacecraft Assembly Jet Propulsion Lal DQ532
DQ532274.1 gg_id:172438 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space Ce DQ532
EF682991.1 gg_id:243852 Eastern Mediterranean atmosphere clone F13_3E_FL EF6829
AJ968649.1 gg_id:144280 Ruegeria atlantica str. N5III AJ9686
DQ915630.1 gg_id:330249 Silicibacter lacuscaerulensis str. DSM11314 DQ915
DQ915631.1 gg_id:306399 Silicibacter pomeroyi str. DSM15171 DQ915
EF092256.1 gg_id:264007 Persistent Axinella corrugata Roseobacter clone MAx29_D12 EF0922
AM709700.1 gg_id:254802 symbiotic Ruegeria strains cuttlefish Sepia officinalis egg yolk clone B4SOVto5 AM709
AM709703.1 gg_id:254660 symbiotic Ruegeria strains cuttlefish Sepia officinalis shell sacs clone 2SOSC91 AM709
AM930441.1 gg_id:268418 involved Tripneustes gratilla (Echinoidea) cloning and FISH host body wall lesions cl AM930
GU118794.1 gg_id:533437 Threatened Corals Provide Microbial Habitats clone Mfra_N19 GU118
FJ203171.1 gg_id:354074 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
GU118756.1 gg_id:593469 Threatened Corals Provide Microbial Habitats clone Mfra_C19 GU118
FJ203011.1 gg_id:350278 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
AM709701.1 gg_id:248371 symbiotic Ruegeria strains cuttlefish Sepia officinalis embryo clone B4SOEmb23 AM709
FJ403069.1 gg_id:357398 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4030
AJ633996.1 gg_id:114809 host's egg clone IEGG2 AJ6339
DQ357745.1 gg_id:149148 Marine Hatchery: Balance Between Pathogenic and Potentially Probiotic Strains mo DQ357
FJ716889.1 gg_id:549008 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7168
EF629830.1 gg_id:274739 Changes marine Mycale laxissima on transfer aqua sponge tissue EF6298

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D88526.1 gg_id:5714 Ruegeria atlantica str. IAM14463 D8852
FJ403081.1 gg_id:369403 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4030
EU195952.1 gg_id:251321 Roseobacter sp. str. P123 EU195
DQ917807.1 gg_id:220204 Microbial profiles bleached and wild-type Muricea elongata Oleiphilus clone BME10 DQ917
FJ203260.1 gg_id:346584 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
AM709702.1 gg_id:256402 symbiotic Ruegeria strains cuttlefish Sepia officinalis renal appendage clone B4SOAr AM709
FJ937894.1 gg_id:534136 Ruegeria sp. str. LS36 FJ9378
DQ917855.1 gg_id:223353 Microbial profiles bleached and wild-type Muricea elongata Ruegeria clone ME108 DQ917
GQ391987.1 gg_id:522785 Silicibacter sp. str. PaD1.36a GQ391
FJ203147.1 gg_id:315763 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
AM709704.1 gg_id:235710 symbiotic Ruegeria strains cuttlefish Sepia officinalis shell sacs clone B6SOSCV22_36 AM709
EU907925.1 gg_id:328931 Ruegeria sp. str. SKUK MB1010 EU907
U58356.1 gg_id:5685 Sagittula stellata U5835
AJ557871.1 gg_id:84036 Arctic deep sea Isolation common chemoorganotrophic oxygen-respiring polar current AJ5578
FJ425632.1 gg_id:362713 communnities healthy and diseased colonies Caribbean coral Montastrea faveolata S FJ4256
FJ203292.1 gg_id:321452 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
DQ915633.1 gg_id:351778 Sulfitobacter brevis str. DSM11443 DQ915
DQ915634.1 gg_id:345214 Sulfitobacter delicatus str. LMG20554 DQ915
AM990658.1 gg_id:278280 Sulfitobacter sp. MOLA 381 AM990
DQ357749.1 gg_id:146001 Sulfitobacter sp. G5-5b DQ357
DQ357751.1 gg_id:150184 Sulfitobacter sp. Cl-H6 DQ357
AF007254.2 gg_id:5638 Sulfitobacter sp. str. EE-36 AF007
NZ_AALV01000004.1 gg_id:149068 Sulfitobacter sp. str. EE-36 NZ_AA
AY794211.1 gg_id:109147 soil clone F457 AY7942
GU061076.1 gg_id:592471 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
GU061073.1 gg_id:590614 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
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EU823295.1 gg_id:310853 Sulfitobacter sp. str. P64 EU823
AF007260.2 gg_id:5681 Sulfitobacter sp. str. GAI-37 AF007
AF173972.1 gg_id:71580 Roseobacter sp. str. KT0917 AF173
GU061114.1 gg_id:579705 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone YAF- GU061
GU061035.1 gg_id:537289 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone Mix GU061
AJ534237.1 gg_id:75228 Sulfitobacter sp. LM-16 AJ5342
GU061104.1 gg_id:592909 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
GU061041.1 gg_id:571091 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone Mix- GU061
AY162107.1 gg_id:135250 Cultivating Sargasso Sea clone GMD16F09 AY162
GU061096.1 gg_id:572670 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
AY771778.1 gg_id:144788 Arctic Sulfitobacter brevis clone SE96 AY7717
EU919845.1 gg_id:315930 Kongsfjorden western Spitsbergen Svalbard ocean water clone s81 EU919
GU061088.1 gg_id:593911 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061

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DQ659417.1 gg_id:166927 Roseobacter sp. str. SO3 DQ659
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AY690671.1 gg_id:107730 Sulfitobacter sp. str. GC02 AY6906
AB121782.1 gg_id:104397 Jannaschia cystaugens str. CFPB-A9 AB1217
DQ915638.1 gg_id:316408 Thalassobacter stenotrophicus str. CECT5294 DQ915
AB121783.1 gg_id:104544 Jannaschia cystaugens str. CFPB-A5 AB1217
AJ631302.1 gg_id:109582 Thalassobacter stenotrophicus str. CECT 5294 = 5SM22 AJ6313
DQ915639.1 gg_id:315323 Thalassobius gelatinovorus str. NBRC15761 DQ915
AJ878874.1 gg_id:142415 Thalassobius mediterraneus str. CECT 5383 = XSM19 AJ8788
FJ202550.1 gg_id:341164 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
AJ968650.1 gg_id:143564 Roseovarius pelophilus str. G5II AJ9686
AF114484.2 gg_id:5704 CVSP isolate str. CV919-312 AF1144
EU236364.1 gg_id:297563 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg91D8 EU2363
FJ203176.1 gg_id:337783 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
FJ203129.1 gg_id:341596 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
DQ376573.1 gg_id:152175 Microbial ecology removing phosphate wastewater EBPR process aerobic sequencin DQ376
EF540436.1 gg_id:220130 oil-shale semi-coke solid waste clone PK_V EF5404
DQ640665.1 gg_id:233567 full-scale EBPR sludge clone Skagenf58 DQ640
EU979473.1 gg_id:344892 Rhodobacter sp. str. CR07-5 EU9794
AM403212.1 gg_id:163121 Nitrate-reducing and Denitrifying Marine Aqua Biofilter marine aqua AM403
EF205262.1 gg_id:188150 Paracoccus sp. 'CJSPY1 (P-I)' str. CJSPY1 (P-I) EF2052
FJ872532.2 gg_id:593437 Rhodobacter sp. str. ZH15 FJ8725
AY656719.1 gg_id:99632 Thioclava pacifica str. TL2 AY6567
AM745438.1 gg_id:243786 Rhodobacter maris str. JA276 AM745
AY914074.1 gg_id:114095 Rhodobacter sp. R-8 AY9140
GQ355283.1 gg_id:593578 Thioclava sp. str. GT-5 GQ355
FJ542826.1 gg_id:530503 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
AY172658.1 gg_id:77954 Rhodobacter sp. str. B-7005 AY1726
AM408117.1 gg_id:186127 Rhodobacter vinaykumarii str. JA123 AM408
EF061947.1 gg_id:193660 Xiamen China mangrove sediment clone XME33 EF0619
EU491911.1 gg_id:264288 Abundance and microbial life ocean crust deep seawater East Pacific Rise clone EPR4 EU4919
FJ202842.1 gg_id:306401 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2028
FJ497541.1 gg_id:586757 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4975
GU220746.1 gg_id:591725 Microbial mineralization and involved microbial Basin field temperature hydrotherm GU220
EU287008.1 gg_id:239638 Pacific arctic surface sediment clone B78-44 EU2870
FJ905620.1 gg_id:568278 s Tonga-Kermadec iron oxide chimney-like structure on Volcano 19 Tonga Arc clone V FJ9056
AF468367.2 gg_id:80906 Arctic sea ice ARK10115 AF4683
EF573178.1 gg_id:231130 Coco's Island site 23 marine clone S23_1277 EF5731
FJ905703.1 gg_id:552641 s Tonga-Kermadec iron oxide sediments Volcano 1 Tonga Arc clone V1F23b FJ9057

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

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DQ395796.1 gg_id:153730 deep-sea octacoral clone ctg_CGOF245 DQ395
FJ152917.1 gg_id:310932 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1529
FJ203141.1 gg_id:347865 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
FM242431.1 gg_id:346708 sequences sediment clone 17 T12d-oil FM242
AF353234.1 gg_id:45295 marine clone Arctic96A-8 AF3532
FJ889529.1 gg_id:560186 Sulfitobacter sp. str. KS8-4 FJ8895
FJ202949.1 gg_id:340342 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2029
GU061063.1 gg_id:543382 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
FM958454.1 gg_id:535414 situ detection (Hippocampus guttulatus) intestinal content and mucus farmed seahc FM958
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DQ915629.1 gg_id:305904 Salipiger mucosus str. LMG22090 DQ915
EU491858.1 gg_id:272287 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918
FJ202508.1 gg_id:341377 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
AJ002565.1 gg_id:5632 Marine (isolate SRF3) clone AJ0025
FJ202423.1 gg_id:345716 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2024
DQ103627.1 gg_id:135997 hypersaline endoevaporitic microbial mat clone E4aC05 DQ103
AJ002563.1 gg_id:25112 Marine (isolate SRF1) clone AJ0025
AF365993.1 gg_id:29620 marine str. BBAT2 AF3659
AF098494.1 gg_id:5645 Sulfitobacter sp. str. GAI-109 AF0984
EF202614.1 gg_id:202204 Sulfitobacter donghicola str. DSW-25 EF2026
EU573104.1 gg_id:277995 and Dependent Microbial Oil Reservoir North Sea Ekofisk oil field Norwegian sector c EU5731
AY654818.1 gg_id:99671 Oculina mucus isolate 80 AY6548
FJ202330.1 gg_id:348420 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2023
AB266065.2 gg_id:170549 Citreicella sp. str. 2-2A AB2660
GU061006.1 gg_id:564192 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone SF79 GU061
AF513400.1 gg_id:69921 Rhodobacter sp. str. 1-5 AF5134
FJ203661.1 gg_id:307998 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
EU544713.1 gg_id:277244 free-living particle- aggregate- archaeal and Laptev Sea 0.22-1.0 um size fraction (fre EU5447
AY162098.1 gg_id:135249 Cultivating Sargasso Sea clone GMD21H11 AY1620
DQ517132.1 gg_id:159870 Cultivation-intensive study ephemeral hypereutrophic aquatic microbial Mojave Des DQ517
EU236329.1 gg_id:301814 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a1F11 EU2363
AY007684.1 gg_id:41631 marine isolate JP88.1 AY0076
DQ856513.1 gg_id:245016 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C1T DQ856
AY770942.1 gg_id:108310 water injection well Dagang oilfield clone W2 AY7709
AY922214.1 gg_id:112728 whalefall clone 131693 AY9222
AF468376.1 gg_id:79810 Arctic sea ice ARK10233 AF4683
GU061139.1 gg_id:584007 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone SF33 GU061
AB491855.1 gg_id:562026 affiliation (Litopenaeus vannamei) Banyuwangi East Java microbial flocs floating brac AB4918
GU118249.1 gg_id:592387 Threatened Corals Provide Microbial Habitats clone Dstr_B07 GU118

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GQ232737.2 gg_id:548120 Phaeobacter sp. str. G4 GQ232
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AAYC01000011.1 gg_id:215094 Roseobacter sp. str. SK209-2-6 AAYC0
FJ202158.1 gg_id:306945 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2021
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DQ015795.1 gg_id:140619 Antarctic lake water clone ELB16-005 DQ015
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AY682197.1 gg_id:103418 Sulfitobacter sp. str. KMM 3457 AY682
EF646118.1 gg_id:240900 Potential interactions particle- anammox and archaeal partners namibian Namibian u EF6461
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EF123343.1 gg_id:249977 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1233
AJ968647.1 gg_id:144272 Roseobacter sp. str. N05I AJ9686
AY701455.1 gg_id:108540 dinoflagellate symbiont clone GCHU11_G AY7014
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FJ516790.1 gg_id:575915 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5167
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AF235129.1 gg_id:39391 North Sea clone KTc0993 AF2351
EU464692.1 gg_id:296206 Hartmann's mountain zebra feces clone AFZEB_aaj69c08 EU4646
EF659442.1 gg_id:247277 within Poole Harbour seawater clone PH-A6 EF6594
AM930445.1 gg_id:268773 involved Tripneustes gratilla (Echinoidea) cloning and FISH host body wall lesions clo AM930
AF507828.1 gg_id:62731 Mono Lake at depth 35 m station 6 July 2000 clone ML635J-2 proteobacterium AF5078
AM697339.1 gg_id:220866 sequencing libraries indoor dust clone BF0002C036 AM697
FJ437982.1 gg_id:542595 Benthic okenone production Fayetteville New York Green Lake surface sediments 16. FJ4379
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AJ534243.1 gg_id:76030 Sulfitobacter sp. PIC-76 AJ5342
DQ915443.3 gg_id:168588 Maritimibacter alkaliphilus str. HTCC2654 DQ915
AY862758.2 gg_id:161659 prokaryotic genetic Salar de Atacama Northern Chile athalassohaline lake water clon AY8627
CU467448.1 gg_id:313991 Prokaryotic Tunisian saltern Saltern solar multipond clone CU467
GU117994.1 gg_id:591140 Threatened Corals Provide Microbial Habitats clone Acer_I13 GU117
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FJ202782.1 gg_id:315498 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2027
FJ716900.1 gg_id:575317 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7169
FJ202529.1 gg_id:318405 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
FJ202470.1 gg_id:321216 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2024
FJ222605.1 gg_id:350516 Albidovulum sp. str. S1K1 FJ2226
GQ433928.1 gg_id:560840 Subseafloor ocean crust microbial observatories: (F-through Osmo Colonization Syst GQ433
GU118103.1 gg_id:532669 Threatened Corals Provide Microbial Habitats clone Apal_O05 GU118
EF123406.1 gg_id:251365 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1234
Y15011.1 gg_id:5826 Rhodovulum iodosum str. N1 Y15011

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GU061120.1 gg_id:532672 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone YAF- GU061
AJ294354.1 gg_id:46595 Roseobacter isolate 667-4 clone AJ2943
EU246237.1 gg_id:257864 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P5-F08 EU246
DQ234221.2 gg_id:142932 determined library mangrove clone DS139 DQ234
EU236288.1 gg_id:297968 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone HgCo15 EU236
EF648144.1 gg_id:239526 dynamics during produced water treatment aerobic activated sludge clone XJ91 EF6481
DQ372842.1 gg_id:153431 Pirellula and OM43 are among dominant lineages identified Oregon coast diatom bl DQ372
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GU061099.1 gg_id:584882 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
GU118238.1 gg_id:538100 Threatened Corals Provide Microbial Habitats clone Dstr_A04 GU118
DQ860070.1 gg_id:165645 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone L16 DQ860
DQ659411.1 gg_id:168003 Roseobacter sp. str. B11 DQ659
FJ516820.1 gg_id:548219 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5168
FJ203293.1 gg_id:333321 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
DQ856523.1 gg_id:257325 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C2L DQ856
EF646123.1 gg_id:251424 Potential interactions particle- anammox and archaeal partners namibian Namibian u EF6461
EU573108.1 gg_id:279211 and Dependent Microbial Oil Reservoir North Sea Ekofisk oil field Norwegian sector c EU573
FJ203568.1 gg_id:352275 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
AY926462.1 gg_id:113266 Roseobacterium hispanicus str. B-33 AY9264
FJ202177.1 gg_id:355194 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2021
FJ976449.2 gg_id:548736 Shimia sp. str. SW-6 FJ9764
AB476254.1 gg_id:564346 Compositional and functional transition epibiotic microbial hydrothermal vent galath AB476
DQ811851.1 gg_id:165140 mangrove soil clone MSB-3B6 DQ811
EU336981.1 gg_id:240704 Saemankumia lutimaris str. SMK-117 EU336
EF632664.1 gg_id:245725 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) sedir EF6326
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DQ009316.1 gg_id:123474 bacterioplankton clone SPOTSAPR01_5m244 DQ009
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FJ203659.1 gg_id:336186 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
AY794174.1 gg_id:109121 soil clone F3C91 AY794
FJ973603.1 gg_id:397618 Periodic sulfide irruptions impact microbial structure and hypersaline lake water colu FJ9736
NR_026472.1 gg_id:567431 Sulfitobacter mediterraneus str. CH-B427; DSM 12244 NR_02
GU118899.1 gg_id:553131 Threatened Corals Provide Microbial Habitats clone Past_G06 GU118
DQ357735.1 gg_id:145906 Marine Hatchery: Balance Between Pathogenic and Potentially Probiotic Strains sea DQ357
FM180527.1 gg_id:323927 Roseovarius aestuarii str. B13 FM180
EU799662.1 gg_id:310648 It's all ranking aquatic Newport Harbour RI clone 1C227285 EU799
FJ716861.1 gg_id:542437 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7168
AM990873.1 gg_id:278099 Structure and functions coastal NW Mediterranean ecosystem sea water AM990
GQ359327.1 gg_id:556328 Oceanicola sp. str. YCWB25 GQ359

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EU488233.1 gg_id:556227 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
DQ146982.1 gg_id:150257 Vibrio sp. str. V261 DQ146
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EF186075.1 gg_id:194746 Rhodobacter sp. str. DQ12-45T EF1860
AM691099.1 gg_id:221022 Rhodobacteraceae clone EG11 AM691
FJ973583.1 gg_id:427025 Periodic sulfide irruptions impact microbial structure and hypersaline lake water colu FJ9735
AB294959.1 gg_id:222231 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma AB2949
FJ203203.1 gg_id:348315 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
FJ623308.1 gg_id:583673 Efficient polyhydroxyalkanoates production waste-activated submitted process seque FJ6233
GQ451279.1 gg_id:548853 Fecal Yunnan snub-nosed monkey feces clone J274 GQ451
DQ269086.1 gg_id:189161 marine macro-alga surface clone DPC189 DQ269
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GU118141.1 gg_id:576886 Threatened Corals Provide Microbial Habitats clone Apal_B15 GU118
DQ103616.1 gg_id:131552 hypersaline endoevaporitic microbial mat clone E4aB11 DQ103
EF205473.1 gg_id:210036 among geographically regions central Tibet geothermal spring mat clone YCB80 EF2054
DQ535898.1 gg_id:159839 Thalassobius aestuarii str. TF-212 DQ535
FJ403106.1 gg_id:362198 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4031
AM930417.1 gg_id:275915 involved Tripneustes gratilla (Echinoidea) cloning and FISH host body wall lesions clc AM930
EF659446.1 gg_id:240011 within Poole Harbour seawater clone PH-B1 EF6594
FJ516806.1 gg_id:513963 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5168
FJ203288.1 gg_id:322400 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
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AM696698.1 gg_id:217239 Rhodovulum adriaticum str. JA245 AM696
FJ202541.1 gg_id:311333 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
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EF092221.1 gg_id:262333 Persistent Axinella corrugata Roseobacter clone B255_A1h EF0922
AB473971.1 gg_id:568966 sequences microorganisms Alvaneu Bad sulfur containing freshwater spring clone AS AB4739
GQ441231.1 gg_id:577508 and dynamic mat present vs. active nitrogen-fixing microorganisms marine microbia GQ441
DQ521519.1 gg_id:184737 Antarctic lake ice cover clone ANTLV2_E06 DQ521
DQ432231.1 gg_id:156862 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake UmRisha sediment cl DQ432
AY214344.1 gg_id:82177 Rhodovulum sp. A_wp02224 AY2143
EU342372.1 gg_id:261363 Thalassobacter sp. str. GA2-M15 EU3423
DQ836762.1 gg_id:169231 attached biomass denitrifying bioreactor clone DB-89 DQ836
EU978825.1 gg_id:591654 and metabolic potential metagenome glacier ice clone glb337c EU9788
FJ425647.1 gg_id:364960 communnities healthy and diseased colonies Caribbean coral Montastrea faveolata R FJ4256
FJ592545.1 gg_id:549666 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
DQ811852.1 gg_id:163280 mangrove soil clone MSB-3E12 DQ811
FJ202586.1 gg_id:310049 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
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AM990868.1 gg_id:279616 Structure and functions coastal NW Mediterranean ecosystem sea water AM990

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EU287297.1 gg_id:255176 Pacific arctic surface sediment clone S11-114 EU287
EU581832.1 gg_id:546740 Oceanicola sp. str. JLT1210 EU581
AF445668.1 gg_id:21491 travertine hot spring clone SM1C11 AF445
DQ836751.1 gg_id:170685 attached biomass denitrifying bioreactor clone DB-9 DQ836
DQ223020.1 gg_id:140881 Roseobacter sp. str. UAzPsK-5 DQ223
DQ153145.1 gg_id:139458 Structure Biofilms on Artificial Surfaces and Free-living Estuarine Seawater surface b DQ153
FJ516819.1 gg_id:559891 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5168
GU118190.1 gg_id:537679 Threatened Corals Provide Microbial Habitats clone Dstr_M20 GU118
DQ071102.1 gg_id:135986 marine surface water clone SJC1.11 DQ071
FJ210801.1 gg_id:354233 Roseobacter sp. str. QDHT-05 FJ2108
AM945586.1 gg_id:263307 Sulfitobacter sp. MOLA:356 AM945
AB491831.1 gg_id:572325 affiliation (Litopenaeus vannamei) Banyuwangi East Java microbial flocs floating brac AB491
AY563033.1 gg_id:100625 Rhodobacter litoralis str. Mano9 AY563
AB100002.1 gg_id:79260 inactive deep-sea hydrothermal vent chimneys clone IndB2-5 AB100
EU799262.1 gg_id:318713 It's all ranking aquatic Newport Harbour RI clone 1C226841 EU799
AY162081.1 gg_id:135247 Cultivating Sargasso Sea clone GMD37D2 AY162
DQ860059.1 gg_id:165826 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone C3 DQ860
EU246226.1 gg_id:253444 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P5-E03 EU246
EF123390.1 gg_id:247890 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1233
AM779569.1 gg_id:254618 Oceanicola sp. str. DongtaiB-3030 AM779
DQ015815.1 gg_id:140242 Antarctic lake water clone ELB16-059 DQ015
AJ440751.1 gg_id:66953 Gram-negative MM 1 AJ4407
EU907924.1 gg_id:335974 Roseobacter sp. str. SKUK MB1009 EU907
DQ070788.1 gg_id:146111 Prokaryotic Structures Seafloor Basalts Varying Age and Juan de Fuca Ridge deep sea DQ070
AF468352.1 gg_id:80289 Arctic sea ice ARK10013 AF4683
FJ542926.1 gg_id:515949 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
EF123341.1 gg_id:239882 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1233
GQ500845.1 gg_id:533002 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
EU143346.1 gg_id:240707 Loktanella sp. str. J57 EU143
DQ446109.2 gg_id:173763 Surface Mucopolysaccharide Layer and Band Microbial Mat Black Band-Diseased Sid DQ446
GU362938.1 gg_id:548305 and archaeal biogeochemical cycling genes within Qiongdongnan basin Sea: pilot stu GU362
FJ403096.1 gg_id:365829 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4030
FJ825922.1 gg_id:569378 Succession during spring Sea filtered surface sea water period diatom bloom Yel Sea; FJ8259
FJ516795.1 gg_id:545503 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5167
FJ203262.1 gg_id:333817 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
AY922228.1 gg_id:113238 whalefall clone 131735 AY9222
FJ152875.1 gg_id:307203 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1528
GQ349657.1 gg_id:580330 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
AF473915.1 gg_id:46143 template extracted Diploria strigosa with BBD clone 128-19L proteobacterium AF4739
EU246804.1 gg_id:251325 polyphasic description Pocillopora meandrina Palmyra Atoll Calcinus obscurus abdon EU2468
GU061266.1 gg_id:548114 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061

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FJ716898.1 gg_id:511895 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7168
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EU907918.1 gg_id:345550 Roseobacter sp. str. SKUK MB1003 EU907
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FM872825.1 gg_id:365967 occupant source house floor dust clone FB03D09 FM872
FJ664812.1 gg_id:548236 Quantitative dynamics cells plankton-fed microbial fuel cell clone plankton_H05 FJ6648
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GU117983.1 gg_id:555923 Threatened Corals Provide Microbial Habitats clone Acer_G14 GU117
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EU624444.1 gg_id:286855 Ruegeria atlantica str. SS-05 EU624
FJ543062.1 gg_id:516831 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5430
EF632666.1 gg_id:255072 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) sedir EF6326
DQ153140.1 gg_id:138819 Structure Biofilms on Artificial Surfaces and Free-living Estuarine Seawater surface b DQ153
GU118199.1 gg_id:559757 Threatened Corals Provide Microbial Habitats clone Dstr_E23 GU118
DQ421597.2 gg_id:210820 Salt Marsh Prokaryotes salt marsh clone SIMO-4232 DQ421
FJ202464.1 gg_id:323413 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2024
FJ202775.1 gg_id:343203 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2027
AB355069.1 gg_id:257450 Monitoring and archaeal two heavy polluted areas lake Egypt based on sediment Ma AB355
EU835420.1 gg_id:349125 structure and quorum sensing reverse osmosis RO membrane biofilm clone 1B15 EU835
DQ421651.2 gg_id:211048 Salt Marsh Prokaryotes salt marsh clone SIMO-4286 DQ421
DQ659415.1 gg_id:166922 Roseobacter sp. str. COL2P DQ659
AY167261.1 gg_id:81559 Roseobacter sp. str. ANT9274 AY167
EU196325.1 gg_id:243714 Roseobacter sp. NP30 EU196
EF123331.1 gg_id:239119 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1233
FJ230841.1 gg_id:304753 Phaeobacter sp. str. G8 FJ2308
DQ856532.1 gg_id:244293 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C2N DQ856
EU835468.1 gg_id:317397 structure and quorum sensing reverse osmosis RO membrane biofilm clone 3M06 EU835
DQ234217.2 gg_id:141577 determined library mangrove clone DS134 DQ234
DQ412059.1 gg_id:154395 Roseobacter sp. str. LOB-8 DQ412
DQ667965.1 gg_id:170315 Orientalimare eburneum str. SW-277 DQ667
EF207559.1 gg_id:215772 mangrove system sediment EF2075
EF471656.1 gg_id:270937 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay (EF4716
FJ203039.1 gg_id:307415 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030

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DQ412074.1 gg_id:155494 Sulfitobacter sp. str. SPB-4	DQ412
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AY639888.1 gg_id:99168 Citricella thiooxidans str. CHLG 16	AY6398
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AF170736.1 gg_id:5781 Rhodobacter sp. str. QSSC1-20	AF1707
AY344379.1 gg_id:92016 Lake Kauhako surface clone K2-S-3	AY3443
DQ659414.1 gg_id:167392 Roseobacter sp. str. BS90	DQ659
GQ396945.1 gg_id:593441 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s	GQ396
EU236337.1 gg_id:295747 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a2D1	EU2363
AF076897.1 gg_id:5649 d isolate western Sargasso Sea clone C	AF0768
FJ202870.1 gg_id:332743 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k	FJ2028
AY922243.1 gg_id:113817 whalefall clone 131821	AY9222
EU360293.1 gg_id:271908 Pyrene-Degrading Consortium Deep-Sea Sediment West Pacific and its Key Member	EU3602
EU799036.1 gg_id:305874 It's all ranking aquatic Newport Harbour RI clone 1C226589	EU7990
DQ153126.1 gg_id:138759 Structure Biofilms on Artificial Surfaces and Free-living Estuarine Seawater surface b	DQ153
GU118282.1 gg_id:587759 Threatened Corals Provide Microbial Habitats clone Dstr_I12	GU118
EU799596.1 gg_id:350463 It's all ranking aquatic Newport Harbour RI clone 1C227214	EU7995
DQ412069.1 gg_id:156195 Sulfitobacter sp. str. KBB-2	DQ412
FJ202786.1 gg_id:352366 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k	FJ2027
EU564840.1 gg_id:259482 Pseudoruegeria sp. str. CC-GIMAT-2	EU5648
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AY845243.1 gg_id:110751 sponge clone 34P35	AY8452
EU799556.1 gg_id:355768 It's all ranking aquatic Newport Harbour RI clone 1C227168	EU7995
FJ202880.1 gg_id:319087 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k	FJ2028
EU652592.1 gg_id:277670 Seasonal variation microbial Yel Sea sediment clone B8S-134	EU6525
FJ202552.1 gg_id:331090 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k	FJ2025
AY548765.1 gg_id:103722 Sulfitobacter sp. str. PM03	AY5487
FJ872535.1 gg_id:567410 Ruegeria sp. str. ZH17	FJ8725

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EU799442.1 gg_id:354617 It's all ranking aquatic Newport Harbour RI clone 1C227041 EU7994
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FJ232451.1 gg_id:368703 Phaeobacter sp. str. NH52F FJ2324
EF659432.1 gg_id:239375 within Poole Harbour seawater clone PH-B6 EF6594
AY845237.1 gg_id:111058 sponge clone 310P5 AY8452
GQ349664.1 gg_id:579002 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
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EU419924.1 gg_id:273372 Roseobacter sp. str. RW37 EU4199
GU396991.1 gg_id:516703 Haematobacter sp. str. BC14248 GU396
EF573183.1 gg_id:234741 Coco's Island site 23 marine clone S23_1282 EF5731
EU771820.1 gg_id:337345 African elephant feces clone AFEL_aai29h10 EU7718
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AM176846.1 gg_id:160274 Recovery mangrove sediment clone SZB5 AM176
GU061078.1 gg_id:580648 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
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FJ157232.1 gg_id:318111 Acetobacter indonesiensis str. KS9 FJ1572
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FJ157228.1 gg_id:324416 Acetobacter lovaniensis str. KS1 FJ1572
AB052706.1 gg_id:25818 Acetobacter orientalis str. 21F-2 AB0527
AB470917.3 gg_id:535203 Acetobacter orientalis KL13-3 AB4709
AB117965.1 gg_id:92224 Acetobacter orientalis SF18-1 AB1179
AJ419845.1 gg_id:58362 Acetobacter orleanensis str. LMG 1583 AJ4198
AB264094.1 gg_id:187581 Acetobacter syzygii str. SKU19 AB2640
AB052713.1 gg_id:22265 Acetobacter syzygii str. 22F-1B AB0527
AB032354.1 gg_id:34202 Acetobacter tropicalis str. NRIC 0312 No.3 AB0323
AB032355.1 gg_id:34622 Acetobacter tropicalis str. NRIC 0312 No.39 AB0323
AJ419836.1 gg_id:58365 Acetobacter peroxydans str. LMG 1633 AJ4198
EU341169.1 gg_id:245064 Evaluation Rapid Technologies Estimate Microbial Burden and Commercial Airline Ca EU3411
AF127399.1 gg_id:4453 Acetobacter aceti SRI1321 AF1273
NR_025512.1 gg_id:561758 Acetobacter cerevisiae str. LMG 1625 NR_025
GQ240641.1 gg_id:551528 Acetobacter pasteurianus str. DSM 2324 GQ240
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DQ906056.1 gg_id:227409 nickel resistance genes metagenome Tinto River rhizosphere clone ERF-1C1 DQ906
AY495956.1 gg_id:101146 Acidiphilium sp. str. WJ52 AY4959
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D30776.1 gg_id:4484 Acidiphilium rubrum str. ATCC 35905 D30776
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GQ264418.1 gg_id:581747 Structure Simulated Level Waste Site simulated level waste site clone WW3_107 GQ264
AM749786.1 gg_id:243689 Acidicaldus sp. T163 AM749
AM749741.1 gg_id:253578 Isolation ing candidate division soils New Zealand steam-affected geothermal soil cl AM749
AY882839.2 gg_id:112235 hot spring clone SK290 AY8828
DQ906080.1 gg_id:223846 nickel resistance genes metagenome Tinto River rhizosphere clone ERF-G1 DQ906
AB362219.2 gg_id:240252 Acidocella aluminiidurans str. AL46 AB3622
D30771.1 gg_id:4491 Acidocella aminolytica str. 101 D30771
EU360486.1 gg_id:271114 pilot biological ferrous iron oxidation mine water treatment plant clone W-Z-59 EU3604
X77468.1 gg_id:4470 Acidomonas methanolica str. LMG 1668 X77468
AB056321.1 gg_id:67745 Kozakia baliensis str. Yo-3 AB0563
AB025928.1 gg_id:52420 Asaia bogorensis str. 71 AB0259
AB230993.2 gg_id:145122 Asaia bogorensis str. BCC 15703 AB2309
AB292238.1 gg_id:247489 Asaia bogorensis AG60 AB2922
EF635915.1 gg_id:255006 Asaia sp. str. Gb16 EF6359
AB025930.1 gg_id:33826 Asaia bogorensis str. 87 AB0259
AB292240.1 gg_id:246483 Asaia krungthepensis AE76 AB2922
AB292241.1 gg_id:254045 Asaia krungthepensis AE68 AB2922
AB292242.1 gg_id:235750 Asaia krungthepensis AE73 AB2922
AB102955.1 gg_id:95258 Asaia krungthepensis str. AA06 AB1029
AB292239.1 gg_id:239498 Asaia siamensis AE97 AB2922
AB230990.2 gg_id:149795 Asaia siamensis str. BCC 15670 AB2309
CP001189.1 gg_id:306076 Gluconacetobacter diazotrophicus 5 str. PAl 5; ATCC 49037 CP0011
DQ387435.1 gg_id:220341 Gluconacetobacter diazotrophicus str. 4L DQ387
AF127392.1 gg_id:4462 Gluconacetobacter diazotrophicus SRI1990 AF1273
AF127397.1 gg_id:4461 Gluconacetobacter diazotrophicus SRI1212 AF1273
X75620.2 gg_id:4423 Gluconacetobacter hansenii str. NCIB8746 LMG1527 X75620
AB166736.1 gg_id:100534 Gluconacetobacter hansenii str. NBRC 14820 AB1667

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EU096236.1 gg_id:277233 Gluconacetobacter intermedius str. EW-i EU096
AF127395.1 gg_id:4441 Gluconacetobacter liquefaciens SRI1994 AF1273
AF127413.1 gg_id:4430 Gluconacetobacter sacchari IF9645 AF1274
AF127406.1 gg_id:4437 Gluconacetobacter sacchari SRI1216 AF1274
AJ012698.1 gg_id:4467 Gluconacetobacter europaeus str. ZIM B028 V3 AJ0126
X75619.1 gg_id:70289 Gluconacetobacter xylinus str. NCIB11664 LMG1515 X75619
AF111841.1 gg_id:23292 Gluconacetobacter johannae str. CFN-Cf55 AF1118
AF192761.1 gg_id:36976 Gluconacetobacter azotocaptans str. CFN-Ca54 AF1927
AY958232.3 gg_id:124373 Gluconacetobacter azotocaptans DS1 AY9582
AB016865.1 gg_id:4420 Acetobacter sp AB0168
AB095100.1 gg_id:71559 Gluconacetobacter persimmonensis str. KJ145 AB0951
AF127401.1 gg_id:4460 Gluconacetobacter diazotrophicus SRI1206 AF1274
AB166741.1 gg_id:104836 Gluconacetobacter saccharivorans str. LMG 1584 AB1667
AB205220.1 gg_id:114307 Gluconacetobacter europaeus str. DSM 6160 AB2052
EU000272.1 gg_id:250024 Gluconacetobacter xylinus EU000
X80775.2 gg_id:4456 Gluconobacter cerinus str. ATCC 19441 IFO 3267 = LMG 1368 X80775
AB308276.1 gg_id:231954 Gluconobacter cerinus AB3082
DQ523496.1 gg_id:159924 Acetobacter syzygii str. A265 DQ523
AB308275.1 gg_id:231476 Gluconobacter oxydans str. LD51-2 AB3082
AB178430.1 gg_id:108207 Gluconobacter oxydans str. NBRC 12118 AB1784
AB128051.1 gg_id:98968 Gluconobacter thailandicus str. F142-1 AB1280
AB178392.1 gg_id:108166 Gluconobacter oxydans str. NBRC 3250 AB1783
AB308278.1 gg_id:229359 Gluconobacter frateurii str. CR16-1 AB3082
AB470921.2 gg_id:533805 Gluconobacter frateurii GB222-1 AB4709
EU096234.1 gg_id:274069 Gluconobacter sp. str. EW707 EU096
AB470922.2 gg_id:533831 Gluconobacter japonicus GB132-1 AB4709
X82290.1 gg_id:4464 Gluconobacter frateurii str. IFO 3264 LMG 1365 X82290
AB308277.1 gg_id:234315 Gluconobacter thailandicus str. ZN22-2 AB3082
AF173825.1 gg_id:4404 Antarctic clone LB3-94 AF1738
AY682679.1 gg_id:144291 Microbial biofilms corroding systems heating system clone 24 AY6826
EU012448.1 gg_id:255991 Roseococcus suduntuyensis str. SHET EU0124
AY150047.1 gg_id:75850 Roseomonas cervicalis str. ATCC 49957 AY1500
EU742165.1 gg_id:314674 Roseomonas cervicalis str. XTD510 EU7421
GQ047159.1 gg_id:446551 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw1009d GQ047
AF227831.1 gg_id:26143 str. 47076 AF2278
GQ043367.1 gg_id:395886 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw96 GQ043
GQ043897.1 gg_id:458437 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw97 GQ043
GQ038481.1 gg_id:399250 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ038
GQ045958.1 gg_id:387425 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw9 GQ045
GQ041000.1 gg_id:378028 Topographical and Temporal Human Skin Microbiome skin inguinal crease clone nbv GQ041

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EU104101.1 gg_id:309755 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
EU182887.1 gg_id:236056 Roseomonas sp. str. MH59 EU1828
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DQ815048.1 gg_id:163698 zebrafish digestive tract clone aab49c08 DQ815
DQ990962.2 gg_id:197174 Microbial Rock Varnish Yungay Atacama Desert Chile rock varnish clone AYRV2-159 DQ990
DQ129635.1 gg_id:194558 microarray level air two cities Texas urban aerosol clone AKIW418 DQ129
FJ479560.1 gg_id:573663 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4795
FM874262.1 gg_id:358386 occupant source house mattress dust clone MC03F05 FM874
GQ097319.1 gg_id:444359 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw453e0 GQ097
NR_028984.1 gg_id:587551 Roseomonas rosea str. 173-96; DSM 14916 NR_02
AJ488504.1 gg_id:64742 Teichococcus ludipueritiae str. 170-96 AJ4885
EF682998.1 gg_id:246217 Eastern Mediterranean atmosphere clone F14_5H_FL EF6829
AB114611.1 gg_id:102348 rice plant clone ER-17 AB114
DQ129249.1 gg_id:151603 microarray level air two cities Texas urban aerosol clone AKIW963 DQ129
AJ871428.1 gg_id:110099 Paracraurococcus ruber CP2C AJ8714
AF513983.1 gg_id:64695 Leahibacter roseus str. LCR1 AF5139
GQ097329.1 gg_id:458521 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw453f08 GQ097
AB009013.1 gg_id:4416 Paracraurococcus ruber str. NS102 AB009
AY150045.1 gg_id:75313 Roseomonas gilardii str. ATCC 49956 AY150
EU540200.1 gg_id:280239 profile human microbiota antecubital fossa (inner elbow) skin clone nbt247a02 EU540
AF170742.1 gg_id:4413 Paracraurococcus sp. str. QSSC5-3 AF1707

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AB445099.1 gg_id:315688 Swaminathania salitolerans str. NBRC 104436 AB4450
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EF446204.1 gg_id:208105 Prokaryotic and Ecology Floating Macroscopic Filaments Extreme Acidic River Rio Tint EF4462
EF516844.1 gg_id:220042 grassland soil clone FCPN646 EF5168
L22481.1 gg_id:8030 Heliothis virescens testis endosymbiont L22481
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AM162403.1 gg_id:154314 and situ identification acidic sphagnum peat bog AM162
Y07647.2 gg_id:4411 Drentse grassland soil clone vii Y07647
EU409601.1 gg_id:276553 Gluconobacter sp. str. EW707 EU4096
DQ906066.1 gg_id:223618 nickel resistance genes metagenome Tinto River rhizosphere clone ERF-C4 DQ906
AM697086.1 gg_id:221365 sequencing libraries indoor dust clone BF0002B036 AM697
AY395443.1 gg_id:102882 pasture soil clone EB1124 AY3954
EU360476.1 gg_id:259611 pilot biological ferrous iron oxidation mine water treatment plant clone FS-Z1-34 EU3604
AJ292597.1 gg_id:53994 PCB-polluted soil clone WD238 AJ2925

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AJ292602.1 gg_id:32348 PCB-polluted soil clone WD271 AJ2926
EF520426.1 gg_id:230396 structure lakes: what controls ? acid-impacted lake clone ADK-SGh02-10 EF5204
FJ465991.1 gg_id:509891 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4659
GQ059256.1 gg_id:458919 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw: GQ059
AJ292598.1 gg_id:43037 PCB-polluted soil clone WD248 AJ2925
AM162450.1 gg_id:151048 and situ identification acidic sphagnum peat bog clone B78 AM162
EU250247.1 gg_id:250255 acid mine drainage clone GXDC-33 EU250
AB381935.1 gg_id:263857 Rhodovastum atsumiense str. G2-11 AB381
GQ402643.1 gg_id:585535 Accessing microbial biodiversity and metabolic potential tropical -based approach pe GQ402
AM691109.1 gg_id:222002 Acetobacteraceae clone BF15 AM691
EU445203.1 gg_id:554674 Isolation Weight Forest Topsoil Metagenomic tropical forest topsoil clone R16 EU445
DQ378187.1 gg_id:152683 oil-polluted soil clone F23_Pitesti DQ378
DQ378196.1 gg_id:155293 oil-polluted soil clone F33_Pitesti DQ378
AM696849.1 gg_id:226512 sequencing libraries indoor dust clone BF0002A034 AM696
AJ867914.1 gg_id:141085 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone C2 AJ8679
FJ466191.1 gg_id:572984 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4661
EU861954.1 gg_id:319672 effects chronic fertilization on alpine tundra microbial implications carbon and cyclin EU861
EU250257.1 gg_id:253895 acid mine drainage clone GXDC-43 EU250
FJ228277.1 gg_id:315855 sulfate-reducing enriched during bioremediation experiment sediments (Brandenbur FJ2282
DQ532333.1 gg_id:171696 Geographical Variation Microbial Burden and Spacecraft Assembly Kennedy Space C DQ532
AF465654.1 gg_id:21795 geothermal soil clone YNPRH71B AF4656
GQ495410.1 gg_id:591379 terrestrial volcanic rocks Iceland Hnausahraun lava f clone Bas-7-52 GQ495
GQ402607.1 gg_id:509496 Accessing microbial biodiversity and metabolic potential tropical -based approach pe GQ402
AJ292611.1 gg_id:44239 PCB-polluted soil clone WD2108 AJ2926
GQ054463.1 gg_id:451765 Topographical and Temporal Human Skin Microbiome skin back clone nbw1089b08c GQ054
EF516502.1 gg_id:223286 grassland soil clone FCPT506 EF5165
FJ774959.1 gg_id:509511 Symbionts Brown Planthopper (Homoptera: Delphacidae) symbiont Nilaparvata luger FJ7749
DQ659205.1 gg_id:220971 acid mine drainage clone DBS-Clone30 DQ659
DQ833767.1 gg_id:172972 Geochemical Controls on Microbial Rainbow and Springs Yellowstone National Park DQ833
AY192273.1 gg_id:81029 Rocky Mountain alpine soil clone W2b-8C AY192
FM874399.1 gg_id:364038 occupant source house mattress dust clone MD01H12 FM874
AB286050.1 gg_id:229374 Asaia lannaensis str. BCC 15733 AB286
AB297501.1 gg_id:250834 Roseomonas sp. str. K-20 (=JCM 14634) AB297
GQ055435.1 gg_id:432032 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw1115c GQ055
AM934721.1 gg_id:545175 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
AY594267.2 gg_id:122816 Photorhabdus luminescens AY594
AY741146.1 gg_id:163171 Azospirillum amazonense str. 21R AY741
GU256437.1 gg_id:580136 Azospirillum amazonense str. DSM 2787 GU256
DQ288688.1 gg_id:151705 Azospirillum brasilense str. MTCC4037 DQ288
GQ284588.2 gg_id:542408 Azospirillum sp. str. ptl3 GQ284

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DQ304794.1 gg_id:151417 studies on and nifH Lasiurus sindicus Sewan grass Thar Desert India root clone C3 DQ304
AY344369.2 gg_id:91010 Lake Kauhako 30 m clone K2-30-36 AY3443
AB542395.1 gg_id:546390 Azospirillum sp. str. TSA36t AB5423
AY283791.1 gg_id:86765 Azospirillum sp. str. LOD4 AY2837
AB508899.1 gg_id:568722 Azospirillum sp. str. TSH98 AB5088
FJ871055.1 gg_id:565380 Azospirillum lipoferum str. ATCC 29707 FJ8710
DQ288690.1 gg_id:156178 Azospirillum brasilense str. MTCC4039 DQ288
AY563022.1 gg_id:104372 Azospirillum sp. WAI-19 AY5630
AM922283.1 gg_id:248382 Azospirillum picis str. IMMIB Tar-3 AM922
AF521650.1 gg_id:58733 Azospirillum sp. str. Arm2-2 AF5216
DQ676344.1 gg_id:220178 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
AB307662.1 gg_id:260920 clay wall used Japanese traditional buildings clone R63 AB3076
AJ238567.1 gg_id:4572 Azospirillum doebereinerae str. 63f AJ2385
DQ104848.1 gg_id:131955 Azospirillum brasilense str. A154 DQ104
DQ787330.1 gg_id:171362 Azospirillum lipoferum str. B2 DQ787
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EU264075.1 gg_id:344876 Azospirillum sp. str. CA6 EU2640
GU256443.1 gg_id:590104 Azospirillum oryzae str. JCM 21588 GU256
AB208736.1 gg_id:115023 AEOC021 clone AB2087
DQ682470.1 gg_id:163594 Azospirillum oryzae str. N7 DQ682
AY692040.1 gg_id:107382 UASB reactor clone I43 AY6920
AY660962.1 gg_id:101185 Azospirillum lipoferum str. CN1 AY6609
AF170353.1 gg_id:4586 Azospirillum sp. str. TTI AF1703
AY118224.1 gg_id:63341 Azospirillum sp. str. DA2-3-1 AY1182
DQ288686.1 gg_id:152173 Azospirillum brasilense str. MTCC4035 DQ288
AB508878.1 gg_id:573964 Azospirillum sp. str. TSH5 AB5088
AB487643.1 gg_id:594099 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4876
AB508897.1 gg_id:562150 Azospirillum sp. str. TSH96 AB5088
FJ970032.1 gg_id:439656 Inquilinus sp. str. CCBAU 01442 FJ9700
AM162580.1 gg_id:153285 and role capsules Sepia officinalis host egg clone 3SOMBO87 AM162
GU118836.1 gg_id:572584 Threatened Corals Provide Microbial Habitats clone Mfra_A15 GU118
AJ244806.1 gg_id:25832 Sepia officinalis symbiont AJ2448
AM162572.1 gg_id:150158 and role capsules Sepia officinalis cuttlefish egg capsule clone 3iSOMBO24 AM162
GU118235.1 gg_id:537977 Threatened Corals Provide Microbial Habitats clone Dstr_L14 GU118
GU118741.1 gg_id:555264 Threatened Corals Provide Microbial Habitats clone Mfra_E20 GU118
AJ244796.1 gg_id:34409 Sepia officinalis symbiont AJ2447
GU118225.1 gg_id:587505 Threatened Corals Provide Microbial Habitats clone Dstr_G01 GU118

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GU118960.1 gg_id:582926 Threatened Corals Provide Microbial Habitats clone Past_N22 GU118
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FJ203583.1 gg_id:329326 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
AM709743.1 gg_id:256953 symbiotic Ruegeria strains cuttlefish Sepia officinalis renal appendages clone B4SOA AM709
EF612768.1 gg_id:242309 Aquaspirillum peregrinum subsp. integrum str. LMG 5407 EF6127
AM901294.1 gg_id:328525 Phaeospirillum cystidoformans str. JA317 AM901
NR_027605.1 gg_id:543918 Magnetospirillum gryphiswaldense MSR-1 str. MSR-1; DSM 6361 NR_02
AJ488096.1 gg_id:66169 chlorobenzene-degrading consortium clone IIIA-9 AJ4880
D14433.1 gg_id:4535 Phaeospirillum fulvum str. NCIMB11762 D14433
AY171615.1 gg_id:78316 Dechlorospirillum sp. str. SN1 AY1716
AJ617837.1 gg_id:110792 oxic-anoxic interphase flooded paddy soil clone D14426 AJ6178
AB250624.1 gg_id:155751 Phaeospirillum sp. str. TUT3101 AB2506
AB307663.1 gg_id:276270 clay wall used Japanese traditional buildings clone B21 AB3076
Y17390.1 gg_id:4531 Magnetospirillum sp. str. MSM-4 Y17390
AB378599.1 gg_id:259049 dominate denitrification nitrate- succinate-ammended stable istope probing 13C fra AB3785
FJ230905.1 gg_id:306797 under Level Antibiotic Selection Pressures river water clone F56 FJ2309
AB487549.1 gg_id:580150 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4875
AJ292593.1 gg_id:37406 PCB-polluted soil clone WD229 AJ2925
AB111108.1 gg_id:200462 Electrochemically assisted iron respiration grow previously iron-reducing clone LS-2 AB111
DQ659219.1 gg_id:217481 acid mine drainage clone DBS-Clone44 DQ659
AM990911.1 gg_id:279352 Nisaea sp. MOLA 137 AM990
FJ202804.1 gg_id:310439 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2028
DQ665839.1 gg_id:162790 Nisaea denitrificans str. DR41_18 DQ665
AB074520.1 gg_id:66304 Aquaspirillum itersonii subsp. nipponicum str. IAM 14945 subsp. AB0745
DQ234645.1 gg_id:143736 subsurface water clone DR546BH1103001SAD24 DQ234
EU835446.1 gg_id:313151 structure and quorum sensing reverse osmosis RO membrane biofilm clone 2B21 EU8354
AF293000.1 gg_id:50973 ferromanganous micronodule clone MNG3 AF2930
NR_028855.1 gg_id:574444 Rhodocista pekingensis str. 3-p; JCM 11669 NR_02
FM177580.1 gg_id:313312 Rhodocista pekingensis str. JA405 FM177
AM072288.1 gg_id:138132 Rhodocista xerospirillum AM072
NC_011420.2 gg_id:326714 Rhodospirillum centenum SW str. SW; ATCC 51521 NC_01
CP000230.1 gg_id:143149 Rhodospirillum rubrum str. ATCC 11170 CP0002
D30777.1 gg_id:4516 Rhodospirillum photometricum str. E-11 D30777
D14431.1 gg_id:6144 Rhodothalassium salexigens str. ATCC35888 D14431
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FJ152947.1 gg_id:356548 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1529
GU447293.1 gg_id:524419 Rhodovibrio salinarum str. LYG30 GU447
DQ432379.1 gg_id:156957 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake Hamra water clone W DQ432
DQ103606.1 gg_id:136158 hypersaline endoevaporitic microbial mat clone E4aB02 DQ103
DQ103605.1 gg_id:133137 hypersaline endoevaporitic microbial mat clone E4aF06 DQ103

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AY987846.1 gg_id:136616 Rhodovibrio sp. 2Mb1 AY9878
AM284389.1 gg_id:181835 Roseospira goa str. JA133 AM284
DQ103603.1 gg_id:132484 hypersaline endoevaporitic microbial mat clone E6aD01 DQ103
AM691104.1 gg_id:225998 Rhodobacteraceae clone EG16 AM691
DQ672568.1 gg_id:172363 Skermanella sp. str. 5416T-32 DQ672
GU195651.1 gg_id:584833 Skermanella sp. str. Py-2-1 GU195
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FJ823930.1 gg_id:525260 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8239
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AB071665.1 gg_id:67722 Tistrella mobilis AB071
FN556563.1 gg_id:514434 Tistrella sp. str. JW16.1a FN556
AM259857.1 gg_id:212680 Spatial sponge- Mediterranean Tethya aurantium sponge cortex clone TAA-10-13 AM259
EF682992.1 gg_id:235586 Eastern Mediterranean atmosphere clone F14_1C_FL EF6829
AM743175.1 gg_id:252146 Azospirillum sp. str. AP-500 AM743
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DQ167235.1 gg_id:138716 Brucella sp. str. HJ114 DQ167
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FJ746303.1 gg_id:568242 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7463
GQ350189.1 gg_id:545318 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU881096.1 gg_id:331647 Karst region primary forest soil clone KPF200711-101 EU881
EF032744.1 gg_id:198274 cyanobacterial mat in lava cave clone HAVOmat83 EF0327
AY598790.1 gg_id:102860 Germany: Lake Staffelsee clone SS31 AY5987
AF391979.1 gg_id:45882 thermal soil clone YNPFFP3 AF3919
EF494345.1 gg_id:225101 Nunnock River granitic landscape clone NR.1.078 EF4943
AM293856.1 gg_id:206331 Subaequorebacter tamlense str. HST3-12 AM293
GQ264472.1 gg_id:588466 Structure Simulated Level Waste Site simulated level waste site clone WW3_179 GQ264
EU050764.1 gg_id:241084 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_04_78 EU050
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EF471705.1 gg_id:259136 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay EF4717
EU919770.1 gg_id:341359 Kongsfjorden western Spitsbergen Svalbard ocean water clone b31 EU9197
EU048669.1 gg_id:250376 and archaeal surface marine sediment Slope South China Sea clone MD2896-B9 EU0486
EU236336.1 gg_id:292058 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a2C6 EU2363
DQ001629.1 gg_id:157721 Microbial seeps asphalt seep clone E91-35 DQ001
AB193945.1 gg_id:108907 deep-sea vent clone Sc-EB05 AB1939
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EU491336.1 gg_id:261444 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
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EU133518.1 gg_id:236426 tallgrass prarie soil clone FFCH17724 EU1335
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FJ746316.1 gg_id:540101 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7463
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EU834769.1 gg_id:545652 methods extraction nucleic acid activated lab scale EBPR-activated sludge clone L24 EU8347
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DQ015800.1 gg_id:139714 Antarctic lake water clone ELB16-121 DQ015
DQ070828.1 gg_id:147136 Prokaryotic Structures Seafloor Basalts Varying Age East Pacific Rise and Ridge basal DQ070
GQ337284.1 gg_id:572538 Hydrography shapes biogeography deep Arctic Ocean water depth 996m clone NB0 GQ337
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AJ519623.1 gg_id:75071 Uranium mill tailings soil sample clone Sh765B-TzT-31 proteobacterium AJ5196

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

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AY627380.1 gg_id:106889 marine clone AY627380 AY6273
AY913376.1 gg_id:112806 forest soil clone DUNssu169 (-7A) (OTU#134) AY9133
AF104473.1 gg_id:4539 Olavius loisae endosymbiont 2 AF1044
CU923620.1 gg_id:545476 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU923
EU386000.1 gg_id:275575 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
EU652588.1 gg_id:278603 Seasonal variation microbial Yel Sea sediment clone D8S-126 EU6525
EU800562.1 gg_id:307071 It's all ranking aquatic Delaware Bay NJ clone 2C228703 EU8005
EU369121.1 gg_id:304307 libraries oyster shell clone MBIOS-05 EU3691
AM935823.1 gg_id:552916 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
AY913278.1 gg_id:114037 forest soil clone DUNssu056 (-2C) (OTU#127) AY9132
EU245313.1 gg_id:244972 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H5-H03 EU2453
GQ346810.1 gg_id:585746 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
EU656113.1 gg_id:282361 Thalassobaculum sp. P24 EU6561
AY654771.1 gg_id:105164 Oculina mucus isolate 38 AY6547
AM997954.1 gg_id:574275 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
FJ592569.1 gg_id:574659 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
GQ060714.1 gg_id:419408 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw04e09 GQ060
GQ263876.1 gg_id:538639 Structure Simulated Level Waste Site simulated level waste site clone WC2_121 GQ263
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DQ395408.1 gg_id:154449 deep-sea octacoral clone ctg_CGOAA92 DQ395
EU589252.1 gg_id:262768 China Ningxia rice paddy field soil clone 1_A7 EU5892
EU675666.1 gg_id:300711 Magnetospirillum sp. str. QH-2 EU6756
GQ246307.1 gg_id:513629 Temporal and spatial China North Yel Sea sediments clone Sd1-26 GQ246
EF632901.1 gg_id:244241 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6329
EF018460.1 gg_id:210561 aspen rhizosphere clone Amb_16S_1086 EF0184
GQ337197.1 gg_id:581302 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
AF407727.1 gg_id:60574 Great Artesian Basin clone B79 AF4077
Z29637.1 gg_id:4555 Azospirillum species Z29637
GQ346973.1 gg_id:580534 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
DQ984631.1 gg_id:253590 YK oil-contaminated soil clone YK_106 DQ984
GQ348477.1 gg_id:583127 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU861390.1 gg_id:318694 marine magnetite-containing magnetic spirillum str. MMS-1 EU8613
GQ349430.1 gg_id:542625 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
DQ906721.1 gg_id:171043 and marine microorganisms Antarctic sea water 20 m clone AntCL1D10 DQ906
EU202861.1 gg_id:351776 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2028
EF515275.1 gg_id:222372 Electricigen Enrichment MFC upf microbial fuel cell anode clone 22c07 EF5152
AY660963.1 gg_id:101335 Azospirillum irakense str. Rsb1 AY6609
EF018935.1 gg_id:211514 aspen rhizosphere clone Amb_16S_1444 EF0189
DQ396323.1 gg_id:152292 deep-sea octacoral clone ctg_NISA029 DQ396
EU449569.1 gg_id:561462 Soil are Changed Agricultural Practices agricultural soil clone Plot4-A11 EU4495

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

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

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GU118834.1 gg_id:578664 Threatened Corals Provide Microbial Habitats clone Mfra_G08 GU118
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EF157232.1 gg_id:257142 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
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AJ633959.1 gg_id:114556 host's accessory nidamental gland clone 30ANG6 AJ6339
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DQ451508.1 gg_id:156725 forest soil clone FAC69 DQ451
EU491777.1 gg_id:266165 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4917
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EU592393.1 gg_id:548310 Seasonal changes Salton Sea hypersaline water clone SSW14N EU5923
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EF516219.1 gg_id:217798 grassland soil clone FCPP413 EF5162
FJ746286.1 gg_id:571876 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7462
EU491571.1 gg_id:273715 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR40 EU4915
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GQ264041.1 gg_id:541154 Structure Simulated Level Waste Site simulated level waste site clone WC3_135 GQ264
EF572964.1 gg_id:231481 Coco's Island site 23 marine clone S23_1063 EF5729
EU528234.1 gg_id:284226 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU5282
DQ070813.1 gg_id:147895 Prokaryotic Structures Seafloor Basalts Varying Age and Juan de Fuca Ridge basalt gl DQ070
EU491885.1 gg_id:271221 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918

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FJ902662.1 gg_id:578073 phreatic limestone sinkholes northeastern Mexico biofilm 1m deep hydrothermal ver FJ9026
EU799997.1 gg_id:334995 It's all ranking aquatic Newport Harbour RI clone 1C227677 EU799
FJ755769.1 gg_id:509529 vertical archaeal water and Lake Taihu sediment clone LT-SB-B68 FJ7557
DQ378260.1 gg_id:155886 oil-polluted soil clone M43_Pitesti DQ378
EU801583.1 gg_id:322226 It's all ranking aquatic Chesapeake Bay MD clone 3C002913 EU801
GQ347647.1 gg_id:577589 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
DQ906739.1 gg_id:172572 and marine microorganisms Antarctic sea water 20 m clone AntCL1H5 DQ906
EU133575.1 gg_id:256291 tallgrass prarie soil clone FFCH4971 EU133
EU386027.1 gg_id:273866 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
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EF018202.1 gg_id:210884 aspen rhizosphere clone Amb_16S_548 EF0182
X99671.1 gg_id:4509 Rhodospira trueperi X99671
EU050757.1 gg_id:247141 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_03_94 EU050
FJ624900.1 gg_id:514040 Influence lead contamination on boreal pine forest soil clone HF_C_27 FJ6249
DQ129334.1 gg_id:138266 microarray level air two cities Texas urban aerosol clone AKIW400 DQ129
AM936707.1 gg_id:544803 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
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AB054178.1 gg_id:22043 marine sponge clone HOC44 AB054
FM242428.1 gg_id:309504 sequences sediment clone 38 T12d-oil FM242
FJ826483.1 gg_id:557922 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8264
AM935865.1 gg_id:548884 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
EF573465.1 gg_id:229444 Coco's Island site 23 marine clone S23_1564 EF5734
AF143822.1 gg_id:72005 hydrocarbon-degrading consortium clone 4-Org1-14 AF1438
AM997828.1 gg_id:547681 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EU799295.1 gg_id:343368 It's all ranking aquatic Newport Harbour RI clone 1C226876 EU799
AJ581592.1 gg_id:94143 soil near uranium mill tailings clone KCM-B-56 AJ5815
EU800342.1 gg_id:351874 It's all ranking aquatic Delaware Bay NJ clone 2C228398 EU800
EU799038.1 gg_id:352113 It's all ranking aquatic Newport Harbour RI clone 1C226591 EU799
EU287307.1 gg_id:256038 Pacific arctic surface sediment clone S26-7 EU287
GQ350021.1 gg_id:545223 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU804599.1 gg_id:355413 It's all ranking aquatic 250 miles Panama City clone 6C232548 EU804
FM242426.1 gg_id:326326 sequences sediment clone 62 T12d-oil FM242
GU119311.1 gg_id:560169 Threatened Corals Provide Microbial Habitats reef water clone Reef_B20 GU119
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GU118274.1 gg_id:583340 Threatened Corals Provide Microbial Habitats clone Dstr_J21 GU118
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EU925871.1 gg_id:314210 sediments sediment station DBS1 northern Bering Sea clone 074F57 EU9258
EU799960.1 gg_id:342027 It's all ranking aquatic Newport Harbour RI clone 1C227638 EU799
FM242370.1 gg_id:336124 sequences sediment clone 37 T9d+oil FM242

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AY033313.1 gg_id:68981 marine bacterioplankton clone MB12C06 AY033
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DQ071034.1 gg_id:131466 marine surface water clone Chl1.16 DQ071
EU799958.1 gg_id:320279 It's all ranking aquatic Newport Harbour RI clone 1C227636 EU799
EU802957.1 gg_id:349161 It's all ranking aquatic Northeast Colon Panama clone 4C230356 EU802
FM242449.1 gg_id:354033 sequences sediment clone 34 T12d+oil FM242
FJ826476.1 gg_id:558348 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8264
AB054154.1 gg_id:36213 marine sponge clone HOC20 AB054
U70679.2 gg_id:4573 Cape Hatteras picoplankton clone OM38 U7067
FM242318.1 gg_id:307417 sequences sediment clone 51 T9d-oil FM242
EU802384.1 gg_id:308776 It's all ranking aquatic Northeast Colon Panama clone 4C229705 EU802
GQ350321.1 gg_id:586003 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU803029.1 gg_id:303610 It's all ranking aquatic Northeast Colon Panama clone 4C230433 EU803
GQ347926.1 gg_id:577083 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
AY744399.1 gg_id:108446 EBAC2C11 complete clone EBAC2C11 AY744
EF573783.1 gg_id:233886 site S25 near Coco's Island marine clone S25_127 EF5737
AJ240913.1 gg_id:72555 5' clone CHAB-III-8 AJ2409
EU010169.1 gg_id:252561 Biogeography Singapore seawaters seawater clone B50 EU010
FM242430.1 gg_id:330363 sequences sediment clone 64 T12d-oil FM242
FM242317.1 gg_id:329580 sequences sediment clone 2 T9d-oil FM242
EU800341.1 gg_id:336625 It's all ranking aquatic Delaware Bay NJ clone 2C228397 EU800
GU061713.1 gg_id:555701 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
EF540391.1 gg_id:226996 oil-shale semi-coke solid waste clone MJK32_MK EF5403
EU799440.1 gg_id:311029 It's all ranking aquatic Newport Harbour RI clone 1C227039 EU799
EF100695.1 gg_id:175841 Caenimicrobium bisanense str. K93 EF1006
AB110414.1 gg_id:87741 Blastochloris sulfoviridis str. RN1 AB110
FJ169957.1 gg_id:346128 Anaplasma bovis str. Zhongxian FJ1699
FJ226454.1 gg_id:350108 Anaplasma marginale str. Ishigaki-2007 FJ2264
AF318945.1 gg_id:50430 Anaplasma ovis 1/11/2/480 sheep 3573/7 AF318
AJ633050.1 gg_id:99202 Anaplasma ovis Yuzhong AJ6330
AJ633051.1 gg_id:99352 Anaplasma ovis Zhangjiachuan AJ6330
AY530806.1 gg_id:100745 Anaplasma platys AY5308
AF287153.1 gg_id:18302 Anaplasma platys AF287

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GU118640.1 gg_id:590270 Threatened Corals Provide Microbial Habitats clone Mfav_F04 GU118
AJ566640.2 gg_id:136326 Ixodes ricinus endosymbiont 1 (IricES1) AJ5666
FJ389575.1 gg_id:365478 cloning and Anaplasma sp. goat Chongqing blood clone szg-2 FJ3895
AY969011.1 gg_id:141855 Anaplasma phagocytophilum J4 AY9690
EU978823.1 gg_id:536819 and metabolic potential metagenome glacier ice clone glb335c EU9788
EU491849.1 gg_id:264506 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918
GU118498.1 gg_id:587318 Threatened Corals Provide Microbial Habitats clone Gven_P15 GU118
AF069962.1 gg_id:5951 Acanthamoeba endosymbiont clone UWC36 AF0699
AF283007.1 gg_id:25427 Anaplasma centrale AF2830
EF667897.1 gg_id:256410 Long-term maintenance species-specific microbiota basal metazoan Hydra epithelium EF6678
AY527214.1 gg_id:112067 Anaplasma phagocytophilum str. Strong AY5272
FJ981659.1 gg_id:563266 Flea Vectors Plague clone XC1 FJ9816
AY969014.1 gg_id:143969 Anaplasma phagocytophilum J4 AY9690
FJ478621.1 gg_id:513411 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
AF497583.1 gg_id:82176 detected Haemaphysalis wellingtoni removed chicken Thailand clone Hw124 AF4975
AM997528.1 gg_id:562891 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EF667921.1 gg_id:236767 Long-term maintenance species-specific microbiota basal metazoan Hydra epithelium EF6679
EU799164.1 gg_id:355833 It's all ranking aquatic Newport Harbour RI clone 1C226733 EU7991
AB196721.1 gg_id:152987 Anaplasma phagocytophilum str. SS33P-L AB1967
AY969013.1 gg_id:143770 Anaplasma phagocytophilum J4 AY9690
GU118630.1 gg_id:582550 Threatened Corals Provide Microbial Habitats clone Mfav_B15 GU118
GU118616.1 gg_id:569314 Threatened Corals Provide Microbial Habitats clone Mfav_P11 GU118
AF156784.1 gg_id:6042 Anaplasma platys str. Gzh981 AF1567
AF493952.1 gg_id:66159 tick-borne pathogen order Rickettsiales East: evidence ticks humans identical 2 Homo AF4939
EF633744.1 gg_id:278455 Neoehrlichia lotori str. RAC413 EF6337
AY135531.1 gg_id:59553 Candidatus Neoehrlichia mikurensis str. Rattus strain mikurensis AY1355
DQ915970.1 gg_id:172589 Ehrlichia canis DQ915
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NC_007354.1 gg_id:139026 Ehrlichia canis str. Jake NC_00
U86664.1 gg_id:6016 Ehrlichia chaffeensis Jax U86664
AB013009.1 gg_id:6031 Ehrlichia muris str. NA1 AB0130
AF318021.1 gg_id:16712 Ehrlichia ruminantium str. Porto Henrique AF3180
M73227.1 gg_id:6026 Ehrlichia ewingii M7322
NC_007798.1 gg_id:153370 Neorickettsia sennetsu str. Miyayama NC_00
AF037210.1 gg_id:6005 Neorickettsia risticii str. SHSN-1 AF0372
AF206298.1 gg_id:17901 Ehrlichia sp. 'trout isolate' trout AF2062
AF037211.1 gg_id:6006 Neorickettsia risticii str. SHSN-2 AF0372

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

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

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

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FJ826203.1 gg_id:537705 Succession during spring Sea filtered surface sea water decay period after diatom blo FJ8262
GQ351049.1 gg_id:579779 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ351
GQ348133.1 gg_id:563097 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU802469.1 gg_id:338687 It's all ranking aquatic Northeast Colon Panama clone 4C229801 EU802
GQ350531.1 gg_id:538597 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
DQ071141.1 gg_id:136072 marine surface water clone Surf1.16 DQ071
EU800481.1 gg_id:326350 It's all ranking aquatic Delaware Bay NJ clone 2C228595 EU800
EU804453.1 gg_id:348575 It's all ranking aquatic 250 miles Panama City clone 6C232382 EU804
EU805349.1 gg_id:346932 It's all ranking aquatic 250 miles Panama City clone 6C233353 EU805
EU804887.1 gg_id:339824 It's all ranking aquatic 250 miles Panama City clone 6C232859 EU804
EU800477.1 gg_id:331844 It's all ranking aquatic Delaware Bay NJ clone 2C228590 EU800
GQ350584.1 gg_id:538959 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU802582.1 gg_id:316095 It's all ranking aquatic Northeast Colon Panama clone 4C229931 EU802
EU800272.1 gg_id:327151 It's all ranking aquatic Delaware Bay NJ clone 2C228312 EU800
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DQ655710.1 gg_id:161812 ocean) brackish zone shal coastal lagoon (South Atlantic Ocean) clone 8 DQ655
DQ071057.1 gg_id:136068 marine surface water clone Chl1.44 DQ071
EF573476.1 gg_id:228126 Coco's Island site 23 marine clone S23_1575 EF5734
EF572494.1 gg_id:232394 Coco's Island site 23 marine clone S23_593 EF5724
EU805294.1 gg_id:309191 It's all ranking aquatic 250 miles Panama City clone 6C233294 EU805
EU801916.1 gg_id:339935 It's all ranking aquatic Chesapeake Bay MD clone 3C003295 EU801
EU805290.1 gg_id:336652 It's all ranking aquatic 250 miles Panama City clone 6C233290 EU805
EU592342.1 gg_id:563209 Seasonal changes Salton Sea hypersaline water clone SSW26Ap EU5923
EF574158.1 gg_id:231586 site S25 near Coco's Island marine clone S25_502 EF5741
EF572576.1 gg_id:234584 Coco's Island site 23 marine clone S23_675 EF5725
GQ348209.1 gg_id:568564 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU804711.1 gg_id:311847 It's all ranking aquatic 250 miles Panama City clone 6C232669 EU804
EU805387.1 gg_id:316104 It's all ranking aquatic 250 miles Panama City clone 6C233396 EU805
EU805281.1 gg_id:329803 It's all ranking aquatic 250 miles Panama City clone 6C233281 EU805
GU061691.1 gg_id:586444 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
GU061692.1 gg_id:591579 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
EU005773.1 gg_id:276591 Major activity inside and outside natural iron-fertilized phytoplankton bloom 3 micro EU005
EU800693.1 gg_id:345826 It's all ranking aquatic Delaware Bay NJ clone 2C228861 EU800
DQ071050.1 gg_id:132375 marine surface water clone Chl1.38 DQ071
EU801209.1 gg_id:353280 It's all ranking aquatic Chesapeake Bay MD clone 3C002456 EU801
EU268085.1 gg_id:240606 summer Dalian and Liaodong bay determined sea water clone L2104 EU268
EF573262.1 gg_id:231495 Coco's Island site 23 marine clone S23_1361 EF5732
EU805181.1 gg_id:334084 It's all ranking aquatic 250 miles Panama City clone 6C233174 EU805
GU230228.1 gg_id:535781 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
EF572832.1 gg_id:229346 Coco's Island site 23 marine clone S23_931 EF5728
EF573364.1 gg_id:232550 Coco's Island site 23 marine clone S23_1463 EF5733

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EF572751.1 gg_id:234005 Coco's Island site 23 marine clone S23_850 EF5727
EU805034.1 gg_id:325083 It's all ranking aquatic 250 miles Panama City clone 6C233016 EU805
EU805307.1 gg_id:349966 It's all ranking aquatic 250 miles Panama City clone 6C233309 EU805
GU230237.1 gg_id:589643 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
L10935.1 gg_id:5908 marine clone FL11 L10935
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EU802266.1 gg_id:305764 It's all ranking aquatic Northeast Colon Panama clone 4C229568 EU802
EF574501.1 gg_id:229394 site S25 near Coco's Island marine clone S25_845 EF5745
GQ349360.1 gg_id:562690 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
EU800659.1 gg_id:353974 It's all ranking aquatic Delaware Bay NJ clone 2C228821 EU800
EU804959.1 gg_id:342655 It's all ranking aquatic 250 miles Panama City clone 6C232937 EU804
EU803180.1 gg_id:336030 It's all ranking aquatic Northeast Colon Panama clone 4C230617 EU803
EU804468.1 gg_id:326529 It's all ranking aquatic 250 miles Panama City clone 6C232399 EU804
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GU230215.1 gg_id:545228 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
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EU804553.1 gg_id:317436 It's all ranking aquatic 250 miles Panama City clone 6C232500 EU804
EF571925.1 gg_id:232961 Coco's Island site 23 marine clone S23_24 EF5719
AB186970.1 gg_id:105858 Mariana trough hydrothermal vent water 0.2micro-m filterable fraction clone MT-NE AB186
GU061794.1 gg_id:586928 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
EU638843.1 gg_id:329172 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU638
GU061451.1 gg_id:562584 and function microbial mesoscale eddy perturbations South China Sea clone CE1-DC GU061
GU061695.1 gg_id:581190 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
EU802853.1 gg_id:324621 It's all ranking aquatic Northeast Colon Panama clone 4C230237 EU802
EU801758.1 gg_id:345184 It's all ranking aquatic Chesapeake Bay MD clone 3C003127 EU801
FJ744964.1 gg_id:558120 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7449
GQ351154.1 gg_id:554776 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
EU800851.1 gg_id:336634 It's all ranking aquatic Delaware Bay NJ clone 2C229079 EU800
FJ745165.1 gg_id:561678 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7451
EU805345.1 gg_id:342433 It's all ranking aquatic 250 miles Panama City clone 6C233349 EU805
EF573040.1 gg_id:234332 Coco's Island site 23 marine clone S23_1139 EF5730
DQ071171.1 gg_id:130137 marine surface water clone Surf1.8 DQ071
GQ348102.1 gg_id:590363 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
GQ348021.1 gg_id:590277 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
GU061790.1 gg_id:571620 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
GU230229.1 gg_id:575529 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
EU804396.1 gg_id:344788 It's all ranking aquatic 250 miles Panama City clone 6C232311 EU804
GU119384.1 gg_id:576863 Threatened Corals Provide Microbial Habitats reef water clone Reef_H22 GU119
EU803151.1 gg_id:308243 It's all ranking aquatic Northeast Colon Panama clone 4C230583 EU803
GU230235.1 gg_id:546752 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230

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GU061320.1 gg_id:581682 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061
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EU799608.1 gg_id:338826 It's all ranking aquatic Newport Harbour RI clone 1C227227 EU799
EU803146.1 gg_id:356204 It's all ranking aquatic Northeast Colon Panama clone 4C230578 EU803
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GU230234.1 gg_id:582241 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
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EU800264.1 gg_id:331734 It's all ranking aquatic Delaware Bay NJ clone 2C228304 EU8002
EU804878.1 gg_id:325227 It's all ranking aquatic 250 miles Panama City clone 6C232849 EU8048
EU592389.1 gg_id:577168 Seasonal changes Salton Sea hypersaline water clone SSW78Ap EU5923
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GQ348097.1 gg_id:548057 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
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EF575211.1 gg_id:229546 site S25 near Coco's Island marine clone S25_1555 EF5752
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o__Rickettsiales	f__Pelagibacteraceae

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GU119458.1 gg_id:538125 Threatened Corals Provide Microbial Habitats reef water clone Reef_O05 GU119
GU061332.1 gg_id:569420 and function microbial mesoscale eddy perturbations South China Sea clone CE1-5m GU061
GQ346735.1 gg_id:589317 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
AY907725.1 gg_id:136383 identification picoplankton contrasting waters Arabian Sea water clone A312018 AY907
EU801049.1 gg_id:331948 It's all ranking aquatic Delaware Bay NJ clone 2C229348 EU801
EU799135.1 gg_id:317800 It's all ranking aquatic Newport Harbour RI clone 1C226699 EU799
EU804369.1 gg_id:341897 It's all ranking aquatic 250 miles Panama City clone 6C232281 EU804
GU061709.1 gg_id:534335 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
GU061714.1 gg_id:582991 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
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GQ349905.1 gg_id:554098 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
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EU802428.1 gg_id:333447 It's all ranking aquatic Northeast Colon Panama clone 4C229755 EU802
GU061338.1 gg_id:564801 and function microbial mesoscale eddy perturbations South China Sea clone CE1-5m GU061
GU061507.1 gg_id:571143 and function microbial mesoscale eddy perturbations South China Sea clone CE2-5m GU061
GQ348152.1 gg_id:543428 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
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EF572654.1 gg_id:227884 Coco's Island site 23 marine clone S23_753 EF5726
GQ349030.1 gg_id:592949 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
EU800753.1 gg_id:330543 It's all ranking aquatic Delaware Bay NJ clone 2C228941 EU800
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EF572230.1 gg_id:233341 Coco's Island site 23 marine clone S23_329 EF5722
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FJ826405.1 gg_id:551570 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8264
GU119456.1 gg_id:561610 Threatened Corals Provide Microbial Habitats reef water clone Reef_A07 GU119
GU119421.1 gg_id:586285 Threatened Corals Provide Microbial Habitats reef water clone Reef_I03 GU119
EU805029.1 gg_id:322492 It's all ranking aquatic 250 miles Panama City clone 6C233011 EU805
GU061701.1 gg_id:573467 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
EF572907.1 gg_id:230670 Coco's Island site 23 marine clone S23_1006 EF5729
DQ071055.1 gg_id:133857 marine surface water clone Chl1.42 DQ071
EF572304.1 gg_id:234195 Coco's Island site 23 marine clone S23_403 EF5723
EU800672.1 gg_id:356107 It's all ranking aquatic Delaware Bay NJ clone 2C228836 EU800
EU268069.1 gg_id:239902 summer Dalian and Liaodong bay determined sea water clone D4758 EU268
EF572266.1 gg_id:231018 Coco's Island site 23 marine clone S23_365 EF5722
GU230227.1 gg_id:575839 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230

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EU237291.1 gg_id:252418 Genes transcripts Eastern and Western Equatorial North Atlantic Ocean microbial as: EU237
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EU800228.1 gg_id:347716 It's all ranking aquatic Delaware Bay NJ clone 2C228250 EU8002
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EU805370.1 gg_id:355973 It's all ranking aquatic 250 miles Panama City clone 6C233377 EU8053
GQ348121.1 gg_id:574611 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU800573.1 gg_id:336168 It's all ranking aquatic Delaware Bay NJ clone 2C228719 EU8005
GQ347984.1 gg_id:535016 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
FJ826463.1 gg_id:552472 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8264
EU805242.1 gg_id:334464 It's all ranking aquatic 250 miles Panama City clone 6C233239 EU8052
GU230241.1 gg_id:548888 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
GU061807.1 gg_id:554292 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
EF572202.1 gg_id:228993 Coco's Island site 23 marine clone S23_301 EF5722
EF571907.1 gg_id:233932 Coco's Island site 23 marine clone S23_6 EF5719
EU804502.1 gg_id:348475 It's all ranking aquatic 250 miles Panama City clone 6C232440 EU8045
EU800005.1 gg_id:322498 It's all ranking aquatic Newport Harbour RI clone 1C227685 EU8000
EF572463.1 gg_id:232144 Coco's Island site 23 marine clone S23_562 EF5724
EF572449.1 gg_id:232239 Coco's Island site 23 marine clone S23_548 EF5724
GU230236.1 gg_id:555992 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
GQ347181.1 gg_id:591181 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
EU804475.1 gg_id:348011 It's all ranking aquatic 250 miles Panama City clone 6C232409 EU8044
EF572912.1 gg_id:233011 Coco's Island site 23 marine clone S23_1011 EF5729
EU802138.1 gg_id:312524 It's all ranking aquatic Chesapeake Bay MD clone 3C003551 EU8021
EU394584.1 gg_id:343762 Coccolithophore bloom dynamics shape (NE-Atl) northern Bay Biscay pelagic particle EU3945
GU061702.1 gg_id:536611 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061

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EU800405.1 gg_id:347665 It's all ranking aquatic Delaware Bay NJ clone 2C228495 EU800
AB193898.1 gg_id:108822 deep-sea vent clone MT797-NB06 AB193
GQ347809.1 gg_id:581936 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
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EU802293.1 gg_id:335921 It's all ranking aquatic Northeast Colon Panama clone 4C229602 EU802
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GQ348074.1 gg_id:557720 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
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EU237270.1 gg_id:255537 Genes transcripts Eastern and Western Equatorial North Atlantic Ocean microbial as EU237
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EU010187.1 gg_id:253778 Biogeography Singapore seawaters seawater clone C36 EU010
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EU010177.1 gg_id:252942 Biogeography Singapore seawaters seawater clone C19 EU010
EU394549.1 gg_id:336437 Coccolithophore bloom dynamics shape (NE-Atl) northern Bay Biscay pelagic free-livi EU394
GQ351239.1 gg_id:547409 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
AF406553.1 gg_id:60623 bacterioplankton clone AEGEAN_248 AF406
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AY274843.1 gg_id:82782 Delaware River estuary clone 3G9 AY274
GQ340185.1 gg_id:581266 Prokaryotic Marathonas Reservoir Greece water column clone K07-79-BAC GQ340
EU802303.1 gg_id:306827 It's all ranking aquatic Northeast Colon Panama clone 4C229613 EU802
EU259796.1 gg_id:242575 summer sea Dalian and Liaodong determined bay water clone D2893 EU259
EU237308.1 gg_id:241412 Genes transcripts Eastern and Western Equatorial North Atlantic Ocean microbial as EU237
GQ337312.1 gg_id:553107 Hydrography shapes biogeography deep Arctic Ocean water depth 252m clone NB2 GQ337
AACY01057101.1 gg_id:106333 Sargasso Sea AACY0
GQ337233.1 gg_id:534129 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
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GU119433.1 gg_id:579477 Threatened Corals Provide Microbial Habitats reef water clone Reef_C06 GU119
AY921813.1 gg_id:113622 farm soil clone AKYH1091 AY9218
EU802363.1 gg_id:354980 It's all ranking aquatic Northeast Colon Panama clone 4C229680 EU802
EF573485.1 gg_id:231006 Coco's Island site 23 marine clone S23_1584 EF5734
AB193924.1 gg_id:108885 deep-sea vent clone Sc-NB08 AB1939
EU802142.1 gg_id:330438 It's all ranking aquatic Chesapeake Bay MD clone 3C003555 EU802
EU802938.1 gg_id:312639 It's all ranking aquatic Northeast Colon Panama clone 4C230334 EU802
FJ545616.1 gg_id:560900 Diveristy Seawater and Sediment Northern North Yel Sea seawater clone W2-33 FJ5456
DQ906712.1 gg_id:171482 and marine microorganisms Antarctic sea water 20 m clone AntCL1B7 DQ906
FJ612305.1 gg_id:510517 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP7.5.89 FJ6123
EF573216.1 gg_id:230136 Coco's Island site 23 marine clone S23_1315 EF5732
AB193906.1 gg_id:108863 deep-sea vent clone Mb-NB10 AB1939
GQ348295.1 gg_id:582171 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
GQ347833.1 gg_id:579810 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
AF353209.1 gg_id:40986 marine clone Arctic96B-3 AF3532
EU802246.1 gg_id:337290 It's all ranking aquatic Northeast Colon Panama clone 4C229547 EU802

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EU802309.1 gg_id:328726 It's all ranking aquatic Northeast Colon Panama clone 4C229620 EU802
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GU061818.1 gg_id:582665 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
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GQ351188.1 gg_id:542603 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
GQ337278.1 gg_id:551848 Hydrography shapes biogeography deep Arctic Ocean water depth 996m clone NB0 GQ337
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AF534425.1 gg_id:85388 Toolik Lake main station at 3 m depth clone TLM01/TLMdgge14 proteobacterium AF5344
GQ349786.1 gg_id:593932 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
EU803037.1 gg_id:343789 It's all ranking aquatic Northeast Colon Panama clone 4C230445 EU803
EU010144.1 gg_id:245665 Biogeography Singapore seawaters seawater clone B55 EU010
EU802276.1 gg_id:354196 It's all ranking aquatic Northeast Colon Panama clone 4C229581 EU802
GQ351241.1 gg_id:585642 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
EU802321.1 gg_id:346262 It's all ranking aquatic Northeast Colon Panama clone 4C229633 EU802
EF572632.1 gg_id:227854 Coco's Island site 23 marine clone S23_731 EF5726
EU802028.1 gg_id:317489 It's all ranking aquatic Chesapeake Bay MD clone 3C003417 EU802
EU800327.1 gg_id:326805 It's all ranking aquatic Delaware Bay NJ clone 2C228380 EU800
U75256.1 gg_id:5902 marine planktonic clone SAR211 U7525
GQ346815.1 gg_id:551062 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
DQ071172.1 gg_id:128312 marine surface water clone Surf1.9 DQ071
EU703194.1 gg_id:572065 taxon richness -altitude lakes eastern tin plateau predominance Bacteroidetes and S EU703
GQ337168.1 gg_id:544967 Hydrography shapes biogeography deep Arctic Ocean water depth 410m clone CB1 GQ337
EU804826.1 gg_id:307486 It's all ranking aquatic 250 miles Panama City clone 6C232791 EU804
EU805084.1 gg_id:309659 It's all ranking aquatic 250 miles Panama City clone 6C233069 EU805
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GQ351217.1 gg_id:593777 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
GQ337299.1 gg_id:555674 Hydrography shapes biogeography deep Arctic Ocean water depth 252m clone NB2 GQ337
EU802668.1 gg_id:308130 It's all ranking aquatic Northeast Colon Panama clone 4C230026 EU802
EU802856.1 gg_id:316482 It's all ranking aquatic Northeast Colon Panama clone 4C230240 EU802
EU801709.1 gg_id:307941 It's all ranking aquatic Chesapeake Bay MD clone 3C003069 EU801
AB187007.1 gg_id:106132 Suiyo seamount hydrothermal vent water 0.2micro-m filterable fraction clone SSM-N AB187
EU804121.1 gg_id:323752 It's all ranking aquatic 250 miles Panama City clone 6C232013 EU804
GQ337208.1 gg_id:571518 Hydrography shapes biogeography deep Arctic Ocean water depth 400m clone CB1 GQ337
GQ337194.1 gg_id:557396 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
GQ337176.1 gg_id:568404 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
EU802580.1 gg_id:332667 It's all ranking aquatic Northeast Colon Panama clone 4C229928 EU802
GQ347006.1 gg_id:538870 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
EU802269.1 gg_id:318647 It's all ranking aquatic Northeast Colon Panama clone 4C229571 EU802
EU801648.1 gg_id:348015 It's all ranking aquatic Chesapeake Bay MD clone 3C002994 EU801
GU061225.1 gg_id:587129 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061
EU803197.1 gg_id:355010 It's all ranking aquatic Northeast Colon Panama clone 4C230639 EU803
GU061335.1 gg_id:545379 and function microbial mesoscale eddy perturbations South China Sea clone CE1-5m GU061
EU802749.1 gg_id:323099 It's all ranking aquatic Northeast Colon Panama clone 4C230121 EU802
EF572257.1 gg_id:234083 Coco's Island site 23 marine clone S23_356 EF5722
EU802486.1 gg_id:346488 It's all ranking aquatic Northeast Colon Panama clone 4C229820 EU802
U75255.1 gg_id:5901 marine planktonic clone SAR203 U7525
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GU061700.1 gg_id:583930 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
GQ337179.1 gg_id:543492 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
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GQ347972.1 gg_id:585433 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
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GQ351145.1 gg_id:588235 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
EU802447.1 gg_id:346851 It's all ranking aquatic Northeast Colon Panama clone 4C229777 EU802
DQ071038.1 gg_id:136067 marine surface water clone Chl1.20 DQ071
EU803946.1 gg_id:326593 It's all ranking aquatic Lake Gatun clone 5C231612 EU803
GQ337081.1 gg_id:552977 Hydrography shapes biogeography deep Arctic Ocean water depth 1191m clone NB GQ337
GQ337211.1 gg_id:588959 Hydrography shapes biogeography deep Arctic Ocean water depth 400m clone CB1 GQ337
GU061319.1 gg_id:541800 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061
EF573562.1 gg_id:228042 Coco's Island site 23 marine clone S23_1661 EF5735
EU800521.1 gg_id:322179 It's all ranking aquatic Delaware Bay NJ clone 2C228652 EU800
EU803139.1 gg_id:321861 It's all ranking aquatic Northeast Colon Panama clone 4C230570 EU803

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EF572364.1 gg_id:233869 Coco's Island site 23 marine clone S23_463 EF5723
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U17644.1 gg_id:5947 Rickettsia australis str. PHS U17644
L36101.1 gg_id:5923 Rickettsia australis str. Phillips L36101
U25042.1 gg_id:5927 Rickettsia sp str. S unnamed isolate spotted fever group U2504
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L36215.1 gg_id:5973 Rickettsia montana L36215
DQ150682.1 gg_id:138092 Rickettsia rickettsii str. 1989CO1 DQ150
AY737685.1 gg_id:107037 Rickettsia marmionii str. KB AY7376
U12462.1 gg_id:5956 Rickettsia sibirica str. 246 U12462
L36221.1 gg_id:5932 Rickettsia typhi str. Wilmington L36221
AF060705.2 gg_id:5958 Rickettsia honei str. RB AF0607
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U15162.1 gg_id:5968 Rickettsia canadensis str. McKeil U15162
EU036982.1 gg_id:345789 Rickettsia raoultii str. Elanda-23/95 EU0369
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U12461.1 gg_id:5953 Rickettsia parkeri str. Maculatum-20 U12461
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L36213.1 gg_id:5948 Rickettsia japonica str. YM L36213
AF172942.1 gg_id:112194 Rickettsia hulinii str. HL-93 AF1729
DQ100164.1 gg_id:145296 Rickettsia monacensis str. IrR/Munich DQ100
CU466797.1 gg_id:352699 Evry wastewater treatment plant anoxic basin clone CU466
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AJ628417.1 gg_id:112673 Wolbachia pipientis AJ6284
AJ548799.1 gg_id:102098 Wolbachia pipientis AJ5487
AJ548802.1 gg_id:106063 Wolbachia pipientis AJ5488
L02884.1 gg_id:6105 Trichogramma deion symbiont L02884

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s__Wolbachia endosymbiont of Brugia malayi	72067
s__Wolbachia endosymbiont of Chorthippus parallelus	72856
s__Wolbachia endosymbiont of Culex quinquefasciatus	72911
s__Wolbachia endosymbiont of Dirofilaria immitis	72016
s__Wolbachia endosymbiont of Dirofilaria immitis	72397
s__unclassified	61261
s__Erythrobacter aquimaris	60202
s__Erythrobacter citreus	60282
s__Erythrobacter flavus	61226
s__Erythrobacter longus	60643
s__Erythrobacter longus	60648
s__Erythrobacter vulgaris	61220
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s__Erythromicrobium ramosum	60457
s__Erythromicrobium ramosum	60593
s__Erythromicrobium ramosum	61039
s__Erythromicrobium ramosum	61375
s__unclassified	60419

AJ548800.1 gg_id:105759 Wolbachia pipientis AJ5488
AJ548798.1 gg_id:101940 Wolbachia pipientis AJ5487
U83091.1 gg_id:6095 Wolbachia sp U8309
L02888.1 gg_id:6104 Trichogramma deion symbiont L02888
AJ628416.1 gg_id:112509 Wolbachia pipientis AJ6284
FJ152094.1 gg_id:308082 Inherited Fungal Endosymbionts Parasitic Wasp and Its Cockroach Host clone b6 FJ1520
L02882.1 gg_id:6082 Muscifidurax uniraptor symbiont L02882
EU833482.1 gg_id:434746 symbiont EU8334
U83092.1 gg_id:6096 Wolbachia sp U8309
FJ152096.1 gg_id:310108 Inherited Fungal Endosymbionts Parasitic Wasp and Its Cockroach Host clone cb5 FJ1520
AF035160.1 gg_id:6093 Wolbachia sp AF035
AJ010275.1 gg_id:6124 Wolbachia endosymbiont of Brugia malayi AJ0102
AB475124.1 gg_id:525041 Wolbachia pipientis str. TUA AB475
AM999887.1 gg_id:308581 Wolbachia pipientis str. wPip AM999
AF304445.1 gg_id:43030 Wolbachia sp. Okinawa-RS32 AF3044
AJ276500.1 gg_id:23566 filarial nematode Dirofilaria repens symbiont of AJ2765
AM990908.1 gg_id:279145 Citromicrobium sp. MOLA 134 AM990
AY461442.1 gg_id:106575 Erythrobacter aquimaris str. SW-116 AY4614
AM990822.1 gg_id:279411 Erythrobacter sp. MOLA 47 AM990
AF500005.1 gg_id:64083 Erythrobacter flavus str. SW-52 AF5000
L01786.1 gg_id:6333 Erythrobacter longus str. OCh101 L01786
AF465835.1 gg_id:107785 Erythrobacter longus str. DSM 6997 AF4658
AY706937.1 gg_id:99838 Erythrobacter vulgaris str. 022 2-12 AY7069
AF432339.1 gg_id:31853 template extracted sediment oil-contaminated nutrient amended beach microcosms; AF4323
EU935322.1 gg_id:539587 str. daSW.23 EU9353
FJ843590.1 gg_id:550126 Sphingomonas phyllosphaerae str. FA2 FJ8435
AM990791.1 gg_id:277841 Erythrobacter sp. MOLA 15 AM990
EU440989.1 gg_id:326335 Erythrobacter flavus str. 2PR56-3 EU4409
EU488295.1 gg_id:575919 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4882
DQ985049.1 gg_id:179170 Erythrobacter sp. str. JL993 DQ985
DQ480144.1 gg_id:157107 Erythrobacter sp. D3043 DQ480
FJ607439.1 gg_id:534887 Erythrobacter citreus str. 107 FJ6074
EF512708.1 gg_id:224600 Erythrobacter vulgaris str. JL271 EF5127
FN431786.1 gg_id:562499 Erythrobacter gaetbuli str. AMV3 AMV17 FN4317
FM180516.1 gg_id:335238 Erythrobacter sp. ZZ-2008 str. B2 FM180
FJ999661.1 gg_id:415733 Sphingomonas phyllosphaerae str. FA2 FJ9996
FN668376.1 gg_id:519549 Erythromicrobium sp. str. BF8 FN6683
GQ284449.1 gg_id:541199 Erythromicrobium ramosum str. THWCS10 GQ284
AB013355.1 gg_id:6348 Erythromicrobium ramosum str. DSM8510T AB0133
GQ472116.1 gg_id:563240 Microcystis aeruginosa freshwater lake clone C-71 GQ472
GU321359.1 gg_id:546220 Porphyrobacter sp. str. NMC22 GU321

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DQ001633.1 gg_id:158572 Microbial seeps asphalt seep clone E91-43 DQ001
FJ997597.1 gg_id:421125 Altererythrobacter sp. str. KYW147 FJ9975
GQ264058.1 gg_id:588310 Structure Simulated Level Waste Site simulated level waste site clone WC3_156 GQ264
DQ676427.1 gg_id:219175 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
DQ811848.1 gg_id:162725 mangrove soil clone MSB-3A6 DQ811
EU440968.1 gg_id:312240 Lutibacterium anuloederans str. PR54-13 EU440
NR_025816.1 gg_id:536261 Porphyrobacter donghaensis str. SW-132; KCTC 12229 NR_02
AB033328.1 gg_id:6344 Porphyrobacter tepidarius str. DSM 10594 AB0333
AF465839.1 gg_id:99433 Porphyrobacter tepidarius str. DSM 10594 AF4658
AM691106.1 gg_id:221235 Erythrobacter longus str. EG6 AM691
EU770257.1 gg_id:346240 Porphyrobacter sp. str. A2 EU770
FJ516793.1 gg_id:539522 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5167
EU276570.1 gg_id:362799 -DGGE and Approaches Analyze Effect Agricultural Practices on Soil agricultural soil c EU276
GQ016328.1 gg_id:493339 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ016
DQ399262.2 gg_id:156567 Erythrobacter indica str. YPM1 DQ399
DQ448852.1 gg_id:164787 Novosphingobium nitrogenifigens str. Y88 DQ448
AF465836.1 gg_id:79367 Erythrobacter litoralis str. DSM 8509 AF4658
EF424398.1 gg_id:205122 Erythrobacter sp. str. SM115 EF4243
DQ985038.1 gg_id:181578 Erythrobacter sp. str. JL1020 DQ985
FJ674974.1 gg_id:516226 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6749
AF407713.1 gg_id:60586 Great Artesian Basin clone B10 AF407
AM176884.1 gg_id:161637 Recovery mangrove sediment clone SZB32 AM176
GQ476825.2 gg_id:534168 Altererythrobacter sp. str. R83-1 GQ476
FJ672518.1 gg_id:531645 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6725
EU118985.1 gg_id:243523 Novosphingobium yangbajingensis str. YBSY-12-2 EU118
EF520398.1 gg_id:231088 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-25 EF5203
DQ520195.1 gg_id:160939 shal China sequences hypertrophic freshwater lake clone ML-9-79.2 DQ520
AY576736.1 gg_id:139435 Erythrobacter sp. str. 5IX/A01/140 AY576
DQ304436.1 gg_id:147665 Alterierythrobacter epoxidivorans str. JCS350 DQ304
DQ532296.1 gg_id:171201 Geographical Variation Microbial Burden and Spacecraft Assembly Kennedy Space C DQ532
AM936339.1 gg_id:547624 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
AY922042.1 gg_id:114156 farm soil clone AKYG623 AY922
AM746688.1 gg_id:256823 etiology rosy discoloration ancient wall paintings rosy-discoloured mural clone A10 AM746
EU328097.1 gg_id:278897 dynamics during bioremediation crude oil contaminated moderate saline soil clone Y EU328
EF157081.1 gg_id:256321 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF157
GQ048646.1 gg_id:445570 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ048
FJ675449.1 gg_id:518998 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6754
AB012062.1 gg_id:6337 Erythrobacter sp. str. MBIC3019 D1215 AB012
EU104298.1 gg_id:345528 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
CU914890.1 gg_id:445733 and dynamics adapted microbiota during degradation Tunisian zarzatine enriched se CU914

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AJ000920.2 gg_id:37092 Novosphingobium sp. str. K16 AJ0009
DQ416556.1 gg_id:154250 s1cb26 DQ416
AY921774.1 gg_id:113287 farm soil clone AKYG900 AY9217
EF433552.1 gg_id:211658 Sphingomonas sp. str. WW11 EF4335
DQ396158.1 gg_id:151573 deep-sea octacoral clone ctg_NISA006 DQ396
EF157121.1 gg_id:257302 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1571
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EU571151.1 gg_id:266620 eubacterium sp. str. 7-22 EU5711
DQ011529.1 gg_id:122133 Porphyrobacter dokdonensis str. DSW-74 DQ011
AF361178.2 gg_id:31470 Sphingomonas sp. Pee Shoal AF3611
DQ860069.1 gg_id:166645 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone L30 DQ860
AF407721.1 gg_id:56491 Great Artesian Basin clone B37 AF4077
DQ856560.1 gg_id:240257 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C3X DQ856
EU784670.1 gg_id:341576 Sphingomonas sp. str. FG03 EU7846
CU466791.1 gg_id:328857 Evry wastewater treatment plant anoxic basin clone CU4667
EU131005.1 gg_id:251580 Sphingomonas sp. BAC151 EU1310
DQ860058.1 gg_id:164190 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone C1 DQ860
EF516627.1 gg_id:225426 grassland soil clone FCPS560 EF5166
DQ985034.1 gg_id:182499 Erythrobacter sp. str. JL990 DQ985
DQ898300.1 gg_id:162144 Sphingomonas sp. str. MTR-71 DQ898
EU360289.1 gg_id:263034 Pyrene-Degrading Consortium Deep-Sea Sediment West Pacific and its Key Member EU3602
AY792292.1 gg_id:123759 Humic Lake clone CrystalBog5F2 AY7922
DQ454121.1 gg_id:156628 Porphyrobacter sp. str. J3-AN66 DQ454
FN397674.1 gg_id:460604 Erythromicrobium sp. str. AMV10 FN3976
EU283461.1 gg_id:277643 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A553 EU2834
EU327998.1 gg_id:277975 dynamics during bioremediation crude oil contaminated moderate saline soil clone B EU3279
GQ060166.1 gg_id:459373 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ060
Y10678.1 gg_id:6365 Sandaracinobacter sibiricus Y10678
CU920729.1 gg_id:533603 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU9207
FJ444668.1 gg_id:569696 Cotton Rhizosphere cotton rhizosphere clone 2y-17 FJ4446
AB094461.1 gg_id:78562 Erythrobacter sp. str. 2216.25.25 AB0944
EF173348.1 gg_id:191991 Options situ remediation contaminated mixture chlorinated hydrocarbons hydrocarb EF1733
FM873313.1 gg_id:367391 occupant source house floor dust clone FD01E08 FM873
GQ169059.1 gg_id:543056 Novosphingobium sp. str. CTDB1 GQ169
AB033327.1 gg_id:6350 Porphyrobacter neustonensis str. DSM 9434 AB0333
AF468357.1 gg_id:79915 Arctic sea ice ARK10031 AF4683
AF468353.1 gg_id:79611 Arctic sea ice ARK10016 AF4683
AB363004.1 gg_id:269458 Erythrobacter ishigakiensis str. NITE-AP48 AB3630
DQ453142.1 gg_id:157567 Erythrobacter sp. str. CNU001 DQ453

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U90216.1 gg_id:6294 Sphingomonas sp. 2MPII U9021
EU682685.1 gg_id:342749 Sphingomonas sp. str. V2M44 EU682
GQ020378.1 gg_id:489291 Topographical and Temporal Human Skin Microbiome skin glabella clone nbu268d11 GQ020
U20775.2 gg_id:44107 Novosphingobium stygium str. SMCC B0712 U2077
EU335142.1 gg_id:257791 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
DQ906835.1 gg_id:256780 Oman subsurface soil clone 7B-8 DQ906
GU166344.1 gg_id:582973 Altererythrobacter sp. str. JM27 GU166
EU707558.1 gg_id:352352 Novosphingobium sp. str. YC6720 EU707
AJ746092.1 gg_id:113876 Novosphingobium sp. MG35 AJ746
EU726272.1 gg_id:349727 Alterierythrobacter sp. str. H32 EU726
FJ542890.1 gg_id:525932 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
DQ497759.1 gg_id:161477 Sphingomonas sp. str. M80C3 DQ497
DQ295907.1 gg_id:147648 Cohn's Crenothrix filamentous oxidizer unusual methane monooxygenase physical e DQ295
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AM940945.1 gg_id:282135 Sphingomonas sp. ECN-2008 AM940
D13737.1 gg_id:6255 Sphingomonas suberifaciens str. IFO 15211 D13737
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EF516033.1 gg_id:222080 grassland soil clone FCPS756 EF5160
EF580938.1 gg_id:278358 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM4 EF5809
GQ008488.1 gg_id:501177 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ008
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DQ248234.1 gg_id:143386 carbon tetrachloride contaminated soil clone UE1 DQ248
EU223953.1 gg_id:249385 -RFLP microbes Qinghai-Tin rail soil clone 120 EU2239
EU193103.1 gg_id:367223 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU193
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EU193107.1 gg_id:364127 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU193
AB258386.1 gg_id:159768 Kaistobacter terrae str. KCTC12630 AB2583
EU202894.1 gg_id:344045 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2028
AY769084.1 gg_id:108378 Kaistobacter koreensis str. PB180 AY7690
EU202882.1 gg_id:340856 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2028
AM935913.1 gg_id:563989 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
EU133392.1 gg_id:242675 tallgrass prarie soil clone FFCH13278 EU133
GQ051344.1 gg_id:414905 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ051
FJ266331.1 gg_id:549411 Novosphingobium capsulatum str. RFNB21 FJ2663
AB196250.1 gg_id:115182 Sphingomonas sp. str. IC081 AB196

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AB275605.1 gg_id:214353 Sphingomonas bosoensis str. WF3 AB275
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AB088760.1 gg_id:78821 Novosphingobium sp. str. J30 AB088
EU304250.1 gg_id:248028 Novosphingobium str. Hg4-04 EU304
AB453304.1 gg_id:514645 Sphingomonas sp. str. SO11 AB453
GQ388869.1 gg_id:511687 quality deterioration drinking system during serious red water outbreak clone Z28 GQ388
EU730906.1 gg_id:332390 Novosphingobium stygium str. 159 EU730
DQ137853.2 gg_id:138108 Novosphingobium sp. str. Geo25 DQ137
GQ388914.1 gg_id:580916 quality deterioration drinking system during serious red water outbreak clone Z76 GQ388
FJ425737.1 gg_id:364924 Novosphingobium sp. str. CC-TPE-1 FJ4257
DQ337059.1 gg_id:149880 subsurface water clone EV818SWSAP5 DQ337
GQ264150.1 gg_id:579694 Structure Simulated Level Waste Site simulated level waste site clone WC3_91 GQ264
GQ023414.1 gg_id:486255 Topographical and Temporal Human Skin Microbiome skin buttock clone nbu318b1 GQ023
U20755.2 gg_id:73612 Novosphingobium aromaticivorans str. SMCC B0695 U20755
AB362778.1 gg_id:256401 Sphingomonas sp. str. YT0136 AB362
GQ103626.1 gg_id:438411 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ103
GQ069267.1 gg_id:374280 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw241e0 GQ069
AY571829.1 gg_id:102884 Sphingomonas sp. 44/40 AY5718
AY792290.1 gg_id:127870 Humic Lake clone CrystalBog2A11 AY792
FJ377543.1 gg_id:369876 Novosphingobium sp. str. F2 FJ3775
GQ038636.1 gg_id:412772 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ038
GQ020999.1 gg_id:488670 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu2 GQ020
EU937904.1 gg_id:334944 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
GQ021683.1 gg_id:487986 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ021
GQ472779.1 gg_id:576436 Biodiversity planktonic Northern surface water northern Bering Sea clone u3 GQ472
FJ459983.1 gg_id:453127 Novosphingobium subterraneum str. C1-3-2; IMER-A3-1 FJ4599
GQ038752.1 gg_id:378163 Topographical and Temporal Human Skin Microbiome skin buttock clone nbu522f03 GQ038
AB331237.1 gg_id:239753 Novosphingobium aromaticivorans str. FM-1 AB331
AJ617857.1 gg_id:111304 oxic-anoxic interphase flooded paddy soil clone D28217 AJ6178
EU937862.1 gg_id:313934 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
AY921998.1 gg_id:113391 farm soil clone AKYG1872 AY921
EU644200.1 gg_id:281742 and Samoylov Island Lena active layer -centred polygon near surface horizon; Rim_T EU644
AJ416411.1 gg_id:38548 Novosphingobium hassiacum W-51 AJ4164
AB177224.1 gg_id:133463 seafloor methane hydrate clone ODP1230B9.15 AB177
EF127604.1 gg_id:208477 Fossil Genes and Microbes Entombed Oldest Ice on Earth ice clone EME109 EF1276
GQ083294.1 gg_id:403709 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ083
AF411072.1 gg_id:39863 Sphingomonas sp. str. ACM-3962 AF411
GQ038964.1 gg_id:371390 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ038
GQ102705.1 gg_id:382549 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ102

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GQ083414.1 gg_id:402526 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ083
DQ327674.1 gg_id:165838 Biological filtration limits carbon availability affects downstream biofilm formation a DQ327
GQ023478.1 gg_id:486191 Topographical and Temporal Human Skin Microbiome skin buttock clone nbu319b0(GQ023
GQ092194.1 gg_id:449436 Topographical and Temporal Human Skin Microbiome skin palm clone nbw409f03c1 GQ092
EF549586.1 gg_id:252275 Novosphingobium indicum str. H25 EF5495
AY328849.1 gg_id:90554 drinking water MB27 AY3288
FJ801185.1 gg_id:513803 Investigation Water -DGGE and Clone Library Zoige Alpine Wetland clone ZWB2-14 FJ8011
GQ099344.1 gg_id:439803 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw5(GQ099
EU431800.1 gg_id:562657 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU4318
AB240479.1 gg_id:142028 microbial structure rhizosphere biofilm Sapporo root-tip (0 40 mm) Phragmites Sosei AB2404
DQ463204.1 gg_id:157220 Acid Drainage Tong Lushan Copper Mine China acid mine drainage clone tls3-9 DQ463
GQ022145.1 gg_id:487522 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu3 GQ022
AF025352.1 gg_id:6293 Sphingomonas sp. str. D16 AF0253
EF029110.1 gg_id:221739 Novosphingobium resinovorum str. NCIMB 8767 EF0291
DQ450761.1 gg_id:160417 Microbial Alpine Tundra Wet Meadow Soil: Chloroflexi Cold Water-Saturated saturat DQ450
EU639137.1 gg_id:323096 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU6391
DQ501315.1 gg_id:189621 Seasonal dynamics free-living and particle- four lakes northeastern Germany freshw DQ501
FJ801188.1 gg_id:542132 Investigation Water -DGGE and Clone Library Zoige Alpine Wetland clone ZWB3-1 FJ8011
GQ020347.1 gg_id:489319 Topographical and Temporal Human Skin Microbiome skin glabella clone nbu267h11 GQ020
U20756.2 gg_id:136117 Novosphingobium aromaticivorans str. SMCC F199 U20756
DQ676432.1 gg_id:224427 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
FM872861.1 gg_id:360253 occupant source house floor dust clone FB03H05 FM872
FM178802.1 gg_id:311289 Monitoring abundance and expression 'dehalococcoides' species chloroethene-redu FM178
AY792291.1 gg_id:150609 Temporal Variation and Humic Lake clone CrystalBog022E8 AY7922
DQ207361.1 gg_id:139541 Sphingomonas sp. str. PF-I DQ207
EU104173.1 gg_id:341018 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1041
CU917745.1 gg_id:533625 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU9177
EU827494.1 gg_id:352634 Sphingomonas sp. tfd44 str. TFD44 EU8274
AF097802.2 gg_id:21583 activated sludge clone 1958 AF0978
AY212578.1 gg_id:99539 water 10 m downstream manure clone 127ds10 AY2125
GQ050992.1 gg_id:435628 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ050
AB286635.1 gg_id:214915 Reducing production and domination comamonadaceae reducing oxygen supply was AB2866
AY922004.1 gg_id:113584 farm soil clone AKYG395 AY9220
FJ177534.2 gg_id:341537 Novosphingobium sp. str. HU1-AH51 FJ1775
GQ069243.1 gg_id:397437 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw241b0 GQ069
GQ037883.1 gg_id:464605 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ037
AJ292613.1 gg_id:150636 Combined use and study polychlorinated biphenyl-polluted soil clone WD2119 AJ2926
Z93995.1 gg_id:6292 activated sludge clone T28 Z93995
U20774.2 gg_id:70401 Novosphingobium aromaticivorans str. SMCC B0522 U20774
AB237654.1 gg_id:156847 Novosphingobium subarcticum str. T7b AB2376
AB177883.1 gg_id:103401 Novosphingobium sp. str. TUT562 AB1778

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AF418947.1 gg_id:19427 USA: Colorado Fort collins Horsetooth Reservoir clone HTB9 AF418
GQ093042.1 gg_id:426566 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw408d11 GQ093
GQ022398.1 gg_id:487271 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbu31 GQ022
GU208446.1 gg_id:563756 Study on Sediment Dongping Lake sediment clone Fr2-8 GU208
GQ008813.1 gg_id:500852 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ008
AY328817.1 gg_id:90016 drinking water Y14 AY328
DQ337064.1 gg_id:146133 subsurface water clone EV818SWSAP16 DQ337
EU937930.1 gg_id:348236 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
AY500142.1 gg_id:104274 Novosphingobium sp. T3-B9 str. T3-B9; AS 1.3432 AY500
EU907884.1 gg_id:320881 Identification hexahydro-135-trinitro-135-triazine (RDX)-degrading microorganisms v EU907
EU431820.1 gg_id:554799 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU431
EU984513.1 gg_id:306710 Novosphingobium sp. str. Rr 2-17 EU984
GQ035853.1 gg_id:418017 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbu37 GQ035
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AB074191.1 gg_id:24166 Sphingomonas sp. str. HL7 AB074
EU937960.1 gg_id:342517 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
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AB491315.1 gg_id:546882 Sphingobium fuliginis str. TIK-1 AB491
AB042233.1 gg_id:18499 Sphingobium herbicidovorans AB042
EF065102.1 gg_id:179619 Sphingobium herbicidovorans str. FL EF0651
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DQ885954.1 gg_id:162032 Sphingomonas sp. str. BSL8 DQ885
AY574367.1 gg_id:106220 Sphingobium yanoikuyae str. DD109 AY5743
AB331239.1 gg_id:240760 Sphingobium yanoikuyae str. FM-3 AB3312
Y12803.1 gg_id:38158 Sphingomonas agrestis str. HV3 Y12803
DQ374096.1 gg_id:156568 Sphingomonas agrestis str. MP5 DQ374
AB025530.1 gg_id:6164 Sphingomonas sp. str. MBIC3990 Lo251 AB0255
AM489507.1 gg_id:203347 Sphingomonas olei str. IMMIB HF-1 AM489
GQ025136.1 gg_id:415594 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbw1 GQ025
EU744623.1 gg_id:349718 Sphingobium sp. str. W1 EU744
AF157490.1 gg_id:6196 Sphingomonas sp. str. JS5 AF1574
EU050705.1 gg_id:238200 Microbial waters production water Daqing oil field clone DQ328-85 EU050
EU286535.1 gg_id:239979 Sphingobium sp. 2F5-2 EU2865
AB112465.1 gg_id:88144 Suiyo Seamount hydrothermal plume clone SUP27-27 AB1124
AY771799.1 gg_id:108045 Sphingomonas sp. 12-7 AY7717
GQ096356.1 gg_id:423731 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw4 GQ096
DQ202286.1 gg_id:139757 Sphingomonas sp. str. PF-K DQ202
AB000106.1 gg_id:6183 Sphingomonas sp. str. VT1 AB000

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EU440973.1 gg_id:306928 Sphingobium sp. str. PR52-27 EU440
AB299582.1 gg_id:254239 Sphingomonas sp. str. M30-VN10-1W AB299
FJ192049.1 gg_id:315748 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1920
EU159273.1 gg_id:354715 Sphingobium sp. str. YBL3 EU159
D13728.1 gg_id:6148 Sphingobium yanoikuyae str. IFO15102 D13728
EU009209.1 gg_id:255180 Sphingobium yanoikuyae str. WP01 EU009
X94098.1 gg_id:6200 Sphingomonas xenophaga str. BN6 X94098
GQ086522.1 gg_id:467619 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw29 GQ086
AY367025.1 gg_id:93363 Sphingomonas sp. As1-3 AY367
X94100.1 gg_id:6163 Sphingomonas paucimobilis str. EPA 505 X94100
AF177917.1 gg_id:6193 Sphingomonas sp. str. DhA-95 AF177
AF191022.1 gg_id:6167 Sphingomonas sp. str. CD AF191
EU183233.1 gg_id:251876 Sphingobium yanoikuyae str. XJ EU183
EU307932.1 gg_id:239648 Sphingobium yanoikuyae str. BF-18 EU307
FJ191324.1 gg_id:310721 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1913
EU375008.1 gg_id:275989 Sphingomonas sp. str. PhS5B EU375
AM998824.1 gg_id:552210 Sphingomonas sp. VC-230 str. VC-230=CCUG 55809 AM998
X85023.1 gg_id:6161 Sphingobium yanoikuyae str. PAH-degrading X85023
AJ011505.1 gg_id:6162 Sphingomonas sp. str. DhA-33 AJ0115
FJ191470.1 gg_id:332159 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1914
AJ000918.2 gg_id:40332 Sphingomonas sp. str. K6 AJ0009
FJ191669.1 gg_id:329197 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1916
EU240401.1 gg_id:252134 Sphingomonas xenophaga str. B107 EU240
AY367026.1 gg_id:93744 Sphingomonas sp. As1-5 AY367
FJ966197.1 gg_id:589286 Sphingobium sp. str. P21 FJ9661
FJ193541.1 gg_id:322175 Genetic inventory 'LMA-MTF' spacecraft assembly clean room during Phoenix missior FJ1935
U37341.1 gg_id:6178 Sphingomonas paucimobilis U3734
AY635898.1 gg_id:101815 Sphingomonas sp. str. AMS7 AY6358
EU616621.1 gg_id:271841 Sphingomonas sp. str. JK1 soil EU616
U37525.1 gg_id:6160 Sphingobium yanoikuyae str. Q1 U37525
EU159274.1 gg_id:313442 Sphingobium sp. str. YBL1 EU159
D16145.1 gg_id:6171 Sphingobium yanoikuyae str. GIFU9882 D16145
GQ019477.1 gg_id:490192 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbu190d GQ019
GQ101175.1 gg_id:370761 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw49 GQ101
AY151393.1 gg_id:73215 Sphingomonas cloacae LH227 AY1513
AB299577.1 gg_id:251303 Sphingomonas sp. str. M5-VN5-4W AB299
AB214867.1 gg_id:131055 Sphingomonas sp. str. ED8 AB2148
FJ191966.1 gg_id:306789 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1919
GQ087461.1 gg_id:416479 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw357g0 GQ087
GQ023300.1 gg_id:486369 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu2 GQ023

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EU057882.1 gg_id:253243 Sphingomonas sp. CH30-8 EU0578
AB040739.1 gg_id:41591 Sphingomonas cloacae str. S-3 AB0407
EU095328.1 gg_id:250814 Sphingobium sp. str. x23 EU0953
AJ009708.2 gg_id:51138 Sphingobium sp. K40 AJ0097
AB120764.1 gg_id:94883 Sphingobium yanoikuyae str. EC-S01 AB1207
EU219973.1 gg_id:251683 Sphingomonas sp. str. KMC001 EU2199
AF038030.1 gg_id:6195 Sphingomonas sp. str. Lep1 AF0380
FJ191658.1 gg_id:343666 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1916
GQ093336.1 gg_id:377426 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw42 GQ093
GQ036338.1 gg_id:409030 Topographical and Temporal Human Skin Microbiome skin palm clone nbu363b03c1 GQ036
AY921715.1 gg_id:113008 farm soil clone AKYH784 AY9217
EU240418.1 gg_id:257571 Sphingobium yanoikuyae str. B166 EU2404
AY151392.1 gg_id:55376 Sphingomonas chungbukensis VM0440 AY1513
FJ193550.1 gg_id:307264 Genetic inventory 'LMA-MTF' spacecraft assembly clean room during Phoenix missior FJ1935
AB214868.1 gg_id:131403 Sphingomonas sp. str. ED9 AB2148
AF494538.1 gg_id:67345 Sphingomonas sp. str. BPC4 AF4945
AY328789.1 gg_id:90697 drinking water system simulator clone DSSD92 AY3287
AY519129.1 gg_id:99094 Sphingomonas paucimobilis str. B90A AY5191
AB288318.1 gg_id:212603 Sphingomonas sp. str. SY-17 AB2883
AY921943.1 gg_id:113573 farm soil clone AKYG1014 AY9219
AY026948.1 gg_id:32571 Sphingomonas sp. str. ML1 AY0269
GQ087010.1 gg_id:432007 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbw3 GQ087
EU580697.1 gg_id:261294 Sphingobium sp. str. NP22b EU5806
DQ238831.1 gg_id:142891 Sphingobium sp. VM2 DQ238
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GQ022171.1 gg_id:487497 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu3 GQ022
FJ192016.1 gg_id:324267 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1920
FJ192036.1 gg_id:350560 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1920
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FJ796422.1 gg_id:462836 Sphingomonas sp. str. S-II-13 FJ7964
AJ007800.1 gg_id:6234 Asticcacaulis excentricus str. ACM 1263 (T) AJ0078
AF331660.1 gg_id:136175 Caulobacter leidyia AF3316
FN421869.1 gg_id:593077 proteogenomics physiology phyllosphere clover clone 2_E01 FN4218
AY571825.1 gg_id:102263 hydrocarbon-contaminated soil clone 433E AY5718
FJ605269.1 gg_id:560773 Sphingomonas sp. str. DCY44 FJ6052
AM229669.1 gg_id:152429 Sphingomonas mucosissima str. CP173-2 AM229
AF409011.1 gg_id:102915 Sphingomonas sp. str. Ellin169 AF4090
GU377113.1 gg_id:523159 Sphingomonas sp. str. sptzw24 GU377
FN421984.1 gg_id:585288 proteogenomics physiology phyllosphere clover clone 5_F05 FN4219

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DQ917254.1 gg_id:168642 and functional genes 24-D degrading clone 57-39	DQ917
DQ532256.1 gg_id:165325 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space Ce	DQ532
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GQ031784.1 gg_id:423523 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n	GQ031
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DQ129612.1 gg_id:137871 microarray level air two cities Texas urban aerosol clone AKIW723	DQ129
EU753665.1 gg_id:316463 modern freshwater stromatolites Ruidera Pools Natural Park Spain dry stromatolite	EU753
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GQ340201.1 gg_id:541929 Prokaryotic Marathonas Reservoir Greece water column clone VE07-03-BAC	GQ340
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EU337119.1 gg_id:510578 Sphingomonas sp. str. 3B3	EU337
DQ532255.1 gg_id:173459 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space Ce	DQ532
EU707561.1 gg_id:319666 Sphingomonas sp. str. YC6723	EU707
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EU400651.1 gg_id:263151 Sphingomonas aquatilis LBS10 EU4006
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FJ834325.1 gg_id:561245 Sphingomonas sp. str. BH3 FJ8343
EU099873.1 gg_id:243743 Sphingomonas sp. str. CJ70 EU0998

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FJ382671.1 gg_id:590918 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3826
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FJ193525.1 gg_id:347975 Genetic inventory 'LMA-MTF' spacecraft assembly clean room during Phoenix missior FJ1935
EU071479.1 gg_id:251041 and resistance microorganisms European clean room ESTEC HYDRA facility clone EHF EU0714
DQ532204.1 gg_id:162721 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space Ce DQ532
AM697187.1 gg_id:226138 sequencing libraries indoor dust clone BF0001C028 AM697

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EU464953.1 gg_id:292322 African elephant feces clone AFYEL_aaj65g04 EU4649
AJ871278.1 gg_id:114482 Sphingomonas sp. str. DS3-1 AJ8712
EU816422.1 gg_id:346880 Sphingopyxis sp. str. Geo48 EU8164
DQ112242.1 gg_id:130329 Sphingopyxis sp. str. LH21 DQ112
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AY212804.1 gg_id:81691 Sphingomonas sp. str. EVA17 AY2128
EU075217.1 gg_id:242041 Sphingopyxis soli DCY34 EU0752
FJ382554.1 gg_id:514081 Potentially pathogenic water and air stem cell transplant summer hospital shower ae FJ3825
EU143356.1 gg_id:248554 Sphingopyxis sp. str. J24 EU1433
FJ382356.1 gg_id:539068 Potentially pathogenic water and air stem cell transplant summer hospital shower ae FJ3823
FJ382353.1 gg_id:510825 Potentially pathogenic water and air stem cell transplant summer hospital shower ae FJ3823
Z73631.1 gg_id:6327 Sphingomonas sp str. RB2256 Z73631
AB245343.1 gg_id:147540 Sphingopyxis ginsengisoli str. Gsoil 250 AB2453
AB219940.1 gg_id:132488 Sphingosinicella microcystinivorans str. MDB2 AB2199
EU703332.1 gg_id:384499 lakes on Eastern Tin Plateau freshwater lake altitude 4298 masl clone XZTSH2 EU7033
EU431722.1 gg_id:559317 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU4317
DQ087403.1 gg_id:128144 Alisphingobium solum str. KSL-125 DQ087
AJ620201.1 gg_id:99158 Sphingomonas sp. str. PlS5 AJ6202
EU403958.1 gg_id:305626 Microbial septic tank effluent and soil biozone infiltrative surface clean water clone (EU4039
AJ303009.1 gg_id:52593 Novosphingobium lentum str. MT1 AJ3030
EU016180.1 gg_id:239608 Sphingomonas sp. str. S11-S-9 EU0161
DQ366009.1 gg_id:152371 Dominant soils Marble Point and Wright Valley Victoria Land Antarctica soil clone 20 DQ366
EU937972.1 gg_id:311613 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
AY608604.1 gg_id:100015 Sphingomonas baekryungensis str. SW-150 AY6086
GQ389106.1 gg_id:565345 quality deterioration drinking system during serious red water outbreak clone S60 GQ389
FJ675209.1 gg_id:525875 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6752
DQ823172.1 gg_id:171409 Techniques Microbial Studies Oregon Caves National Monument Quad 3 on trail clo DQ823
FJ478653.1 gg_id:582012 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
GQ389029.1 gg_id:570614 quality deterioration drinking system during serious red water outbreak clone J76 GQ389
AF170744.1 gg_id:6263 Sphingomonas sp. str. QSSC5-6 AF1707
FN397673.1 gg_id:426622 Blastomonas sp. str. AMV5 FN3976
D16149.1 gg_id:6203 Sphingomonas sp str. SYK6 D16149
FJ801196.1 gg_id:555964 Investigation Water -DGGE and Clone Library Zoige Alpine Wetland clone ZWB3-13 FJ8011

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EF688392.1 gg_id:238429 soil clone F6-154 EF6883
DQ781320.1 gg_id:166587 Sphingopyxis sp. str. FR1087 DQ781
EU104232.1 gg_id:349315 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
EU276573.1 gg_id:361840 -DGGE and Approaches Analyze Effect Agricultural Practices on Soil agricultural soil c EU276
FN421536.1 gg_id:549682 proteogenomics physiology phyllosphere Arabidopsis thaliana clone 14_A04 FN421
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AY167827.2 gg_id:77547 Sphingomonas sp. str. SAFR-042 AY167
GQ263416.1 gg_id:564561 Structure Simulated Level Waste Site simulated level waste site clone FW2_09B GQ263
GQ111232.1 gg_id:436152 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ111
EF522271.1 gg_id:222183 Endolithic Microbial Ecosystems Rocky Mountain endolithic sandstone clone FQSS05 EF5222
EU016177.1 gg_id:236085 Sphingomonas sp. str. S11-S-6 EU016
D13727.1 gg_id:6320 Sphingopyxis terrae str. IFO15098 D13727
AY921978.1 gg_id:113153 farm soil clone AKYG1729 AY921
GQ263560.1 gg_id:586875 Structure Simulated Level Waste Site simulated level waste site clone FW3_112C GQ263
GU479372.1 gg_id:527848 Sphingomonas sp. str. Sphing1 GU479
GQ500764.1 gg_id:577484 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
DQ269104.1 gg_id:190640 marine macro-alga surface clone DPC150 DQ269
AM990821.1 gg_id:277792 Structure and functions coastal NW Mediterranean ecosystem sea water AM990
AM697063.1 gg_id:220477 sequencing libraries indoor dust clone BF0002B013 AM697
EU137618.1 gg_id:267439 Bartonella-positive fleas: and assembly patterns Oropsylla hirsuta (prairie dog flea) C EU137
EU196323.1 gg_id:238220 Sphingomonas sp. NP32 EU196
FJ889532.1 gg_id:564413 Sphingopyxis sp. str. SC022 FJ8895
EU573097.1 gg_id:278193 and Dependent Microbial Oil Reservoir North Sea Ekofisk oil field Norwegian sector c EU573
GQ103823.1 gg_id:397122 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ103
DQ228401.1 gg_id:141916 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
GQ264328.1 gg_id:582134 Structure Simulated Level Waste Site simulated level waste site clone WW2_159 GQ264
EU564841.1 gg_id:261978 Sphingosinicella sp. str. CC-TBT-3 EU5648
DQ129622.1 gg_id:138630 microarray level air two cities Texas urban aerosol clone AKIW980 DQ129
D84626.2 gg_id:557168 Novosphingobium sp. str. S23435 D84626
EU133364.1 gg_id:256653 tallgrass prarie soil clone FFCH11613 EU133
EU404009.1 gg_id:336188 Microbial septic tank effluent and soil biozone infiltrative surface clean water clone C EU404
AM697055.1 gg_id:217733 sequencing libraries indoor dust clone BF0002B005 AM697
EU642848.1 gg_id:278075 Sandarakinorhabdus sp. JC2488 str. JC2488; IMSNU 14142 EU6428
AF281033.1 gg_id:39861 Zymomonas mobilis AF281
AF498741.1 gg_id:58089 soil isolate Ellin359 AF4987
GQ264264.1 gg_id:555926 Structure Simulated Level Waste Site simulated level waste site clone WW1_b69 GQ264
AY913375.1 gg_id:113174 forest soil clone DUNssu168 (-7A) (OTU#169) AY9133
AB081581.3 gg_id:74124 Rhizobiales str. A48 AB081

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EU939387.1 gg_id:325850 Fecal wild pygmy loris feces clone WF16S_04 EU9393
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EU472130.1 gg_id:288432 francois langur feces clone FL_1aaa01a11 EU4721
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AJ536878.1 gg_id:78776 uranium mining waste pile soil sample clone JG30-KF-AS50 AJ5368
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EU104244.1 gg_id:325439 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104

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AY957890.1 gg_id:114903 drinking water biofilm clone B1NR70D5
DQ889887.1 gg_id:166172 natural product producing octocoral Erythropodium caribaeorum clone EC154
DQ856551.1 gg_id:255188 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C3M
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EU862292.1 gg_id:353739 Achromobacter ruhlandii str. jql-9
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EU360470.1 gg_id:268783 Achromobacter xylosoxidans str. B3A
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DQ104980.1 gg_id:136390 Achromobacter xylosoxidans str. JS 1-2
AJ551119.1 gg_id:94243 Achromobacter xylosoxidans mp5
EU195874.1 gg_id:241186 Alcaligenes sp. str. SBW1
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AJ875020.1 gg_id:111930 Achromobacter xylosoxidans str. A8
EU006066.1 gg_id:246414 Achromobacter xylosoxidans str. SL06500
AF205370.1 gg_id:49793 Achromobacter ruhlandii str. QSSC3-1
EU877076.1 gg_id:311931 Achromobacter xylosoxidans str. EC3_3502

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AJ133493.2 gg_id:6859 Alcaligenes sp. CCUG 36768 AJ1334
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FJ674784.1 gg_id:518282 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6747
AB188125.1 gg_id:132621 Alcaligenes latus str. IAM 12599 AB188
EF032746.1 gg_id:185782 cyanobacterial mat in lava cave clone HAVOmat12 EF0327
AY435511.1 gg_id:107880 propane-sparged ground water well C4 clone MAFB-C4-17 AY4355
FJ535109.1 gg_id:557756 microbial colonizing walls Karstic Slovenia cave wall biofilm clone 3PJM33 FJ5351
AB487742.1 gg_id:589854 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB487
AB486168.1 gg_id:579191 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSBARC AB486
AJ867848.1 gg_id:169526 Leptothrix sp. B2 AJ8678
AY493948.1 gg_id:136596 Fingerprinting aggregates soil clone 11 AY4939
AB378584.1 gg_id:260565 dominate denitrification nitrate- succinate-ammended stable istope probing 13C fra AB3785
AY466115.1 gg_id:101912 Bordetella holmesii G6128 AY466
EU082165.1 gg_id:244327 Bordetella petrii str. AU11495 EU082
DQ132877.1 gg_id:137970 Bordetella sp. str. QJ9-6 DQ132
EU082157.1 gg_id:251205 Bordetella sp. str. AU9919 EU082
DQ256466.1 gg_id:142658 Bordetella sp. str. 1-3 DQ256
EU835434.1 gg_id:346019 structure and quorum sensing reverse osmosis RO membrane biofilm clone 2B04 EU835
EU643523.1 gg_id:280840 Bordetella bronchiseptica str. Bb5 EU6435
AB255107.1 gg_id:171796 corroded concrete clone SC-127 AB255
EU603444.1 gg_id:263781 Bordetella petrii str. PTG4-17 EU6034
AJ249861.1 gg_id:21781 Bordetella petrii str. DSM 12804 Se-1111R AJ2498
EU082151.1 gg_id:240317 Bordetella sp. str. AU7206 EU082
AB039335.1 gg_id:37958 Bordetella sp. str. M1-6 AB0393
E06073.1 gg_id:25458 Bordetella bronchiseptica E06073
AY594190.1 gg_id:126199 Bordetella ansorpii str. SMC-8986 AY5941
AF142326.1 gg_id:6889 Bordetella pertussis str. Tohama; CIP 81.32 AF1423
AF366577.1 gg_id:49946 Bordetella parapertussis str. B24 AF3665
GQ480840.1 gg_id:541437 Bordetella petrii str. B1 GQ480
EU000383.1 gg_id:250883 Bordetella sp. str. BK-52 EU000
FJ969842.3 gg_id:445459 Bordetella parapertussis str. HT200 FJ9698
AJ005447.1 gg_id:6845 Alcaligenes defragrans str. 54Pin AJ0054
AB195160.1 gg_id:108771 Alcaligenes defragrans str. PD-16 AB195
Y12639.1 gg_id:6856 Denitrobacter permanens Y12639
EF205259.1 gg_id:189889 Alcaligenes sp. 'TJOPY2 (-I)' str. TJOPY2 (-I) EF2052
AF227163.1 gg_id:71251 Oligella urethralis str. ATCC17960 AF2271
AJ251912.1 gg_id:35834 Oligella ureolytica str. LMG 6519 AJ2519
AF190911.1 gg_id:6840 Pelistega europaea str. LMG15725 AF1909
FN421876.1 gg_id:594220 proteogenomics physiology phyllosphere clover clone 2_F01 FN4218
EU583723.1 gg_id:263890 Pigmentiphaga sp. str. JSM 061001 EU5837
EU170477.1 gg_id:238547 Pigmentiphaga sp. str. Zn-d-2 EU1704
AB300989.1 gg_id:272661 Sutterella parvirubra str. YIT 11816 (= JCM 14724 = DSM 19 AB3009
GQ033291.1 gg_id:455791 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw: GQ033

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EF400872.1 gg_id:213646 human fecal clone SJTU_D_02_65 EF4008
AJ832129.1 gg_id:181656 Sutterella morbirenis str. CCUG 49335 AJ8321
EU532008.1 gg_id:271525 Predominant role host genetics controlling gut microbiota FMF remission stool clone EU532
EU009806.1 gg_id:254552 intestinal microbiota domestic and wild turkeys turkey cecum clone WTB_D10 EU0098
EU009787.1 gg_id:245510 intestinal microbiota domestic and wild turkeys turkey cecum clone DTB_T27 EU0097
AB491209.1 gg_id:547221 Parasutterella sp. str. YIT 12071 AB491
EU767287.1 gg_id:339461 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone C2 EU767
GQ011110.1 gg_id:498556 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ011
AJ566849.1 gg_id:104793 Sutterella stercoricanis str. CCUG 47620 AJ5668
GQ031994.1 gg_id:445073 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ031
EU767511.1 gg_id:322193 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone C2 EU767
EU763570.1 gg_id:319660 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone A4 EU763
EF071208.1 gg_id:176809 library survey and quantitative measurement major mucosa- IBD human colonic muc EF0712
AY981032.1 gg_id:126136 human rectum mucosal biopsy clone V241 AY9810
FJ368987.1 gg_id:358132 feces adult twins and mothers; TS4 clone TS4_a02h09 FJ3689
EU767098.1 gg_id:345647 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone C1 EU767
DQ795867.1 gg_id:196501 human fecal clone RL245_aai83f07 DQ795
NR_024915.1 gg_id:540774 Taylorella asinigenitalis str. UCD-1; ATCC 700933 NR_024
AB200410.1 gg_id:112165 Taylorella equigenitalis str. Aus4 AB2004
AJ864470.1 gg_id:109028 Tetrathiobacter kashmirae str. WT001 AJ8644
AJ864472.1 gg_id:109030 Tetrathiobacter kashmirae str. 445C AJ8644
AJ864474.1 gg_id:109032 Tetrathiobacter kashmirae str. WPT AJ8644
AJ864473.1 gg_id:109031 Tetrathiobacter kashmirae str. WGT AJ8644
AJ864471.1 gg_id:109029 Tetrathiobacter kashmirae str. 445A AJ8644
X86589.1 gg_id:6882 Alcaligenes sp str. 15-62 X86589
AY880023.1 gg_id:111634 Alcaligenes monasteriensis str. DPN7 AY8800
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AM944735.1 gg_id:323924 Advenella incenata 4 AM944
DQ469586.1 gg_id:157497 Alcaligenes monasteriensis str. Ac2-VH DQ469
X86580.1 gg_id:6833 Alcaligenes sp str. 05-36 X86580
X86590.1 gg_id:6883 Alcaligenes sp str. 30-68 X86590
AY569458.1 gg_id:107606 Advenella incenata str. R-16599 AY5694
GQ203113.1 gg_id:558099 Ochrobactrum pseudogrignonense str. 21-6PIN GQ203
AM944734.1 gg_id:348042 Advenella incenata str. CCUG 45225 AM944
EU447156.1 gg_id:273668 Tetrathiobacter kashmirensis str. CA1 EU447
AY622228.1 gg_id:106744 subsurface soil clone C-CG17 AY6222
FJ477679.1 gg_id:511551 Bordetella sp. HU-1 str. 1 FJ4776
U82826.1 gg_id:6846 Alcaligenes sp. str. NKNTAU U82826
AY792261.1 gg_id:120103 Humic Lake clone CrystalBog1G5 AY7922
EU801712.1 gg_id:309818 It's all ranking aquatic Chesapeake Bay MD clone 3C003072 EU801
EU703295.1 gg_id:466157 lakes on Eastern Tin Plateau polysaline lake altitude 3204 masl clone XZQH78 EU703

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

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FJ755942.1 gg_id:512238 Alcaligenes sp. str. RCT1 FJ7559
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GQ254283.1 gg_id:574340 Castellaniella sp. str. H13 GQ254
FJ477650.1 gg_id:584565 gut symbionts are tightly linked herbivory ants whole extracted single ant clone 96j FJ4776
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FM873876.1 gg_id:368602 occupant source house mattress dust clone MB02C10 FM873
EF471233.1 gg_id:220259 Alcaligenes sp. str. BBTR16 EF4712
X92415.1 gg_id:6857 Alcaligenes sp. str. L6 X92415
DQ802483.1 gg_id:196880 human fecal clone RL188_aah18d02 DQ802
DQ799440.1 gg_id:184082 human fecal clone RL305aal86c09 DQ799
AJ277742.1 gg_id:22985 Brackiella oedipodis str. LMG 1945 R8846 AJ2777
AF232922.1 gg_id:22530 United Kingdom: Montserrat geothermal springs clone MS8 MS8 AF2329
EU133883.1 gg_id:237924 tallgrass prarie soil clone FFCH2283 EU1338
EF602797.1 gg_id:228235 Salmonella typhimurium Exploits Inflammation Compete Intestinal Microbiota mouse EF6027
GU138383.1 gg_id:585406 Achromobacter sp. str. THG 42 GU138
DQ346507.1 gg_id:153290 during composting libraries compost clone 2B23 DQ346
AY921769.1 gg_id:114410 farm soil clone AKYG1037 AY9217
AY792262.1 gg_id:123064 Humic Lake clone CrystalBog5E7 AY7922
EU800906.1 gg_id:330902 It's all ranking aquatic Delaware Bay NJ clone 2C229148 EU8009

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GQ105915.1 gg_id:458158 Topographical and Temporal Human Skin Microbiome skin gluteal crease clone nbw GQ105

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GQ111823.1 gg_id:422183 Topographical and Temporal Human Skin Microbiome skin popliteal fossa clone nbw GQ111
GQ101656.1 gg_id:468045 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw497f11c GQ101
GQ103464.1 gg_id:459681 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ103
GQ102629.1 gg_id:396678 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ102
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GQ104473.1 gg_id:423244 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ104
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GQ104691.1 gg_id:370959 Topographical and Temporal Human Skin Microbiome skin inguinal crease clone nbv GQ104
GQ098323.1 gg_id:457430 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw40 GQ098
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o__Burkholderiales	f__Aquabacteriaceae

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GQ097174.1 gg_id:445004 Topographical and Temporal Human Skin Microbiome skin gluteal crease clone nbw GQ097
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GQ114235.1 gg_id:409114 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw59 GQ114
GQ094988.1 gg_id:397716 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ094
GQ105166.1 gg_id:419019 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ105
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GQ103013.1 gg_id:411167 Topographical and Temporal Human Skin Microbiome skin inguinal crease clone nbv GQ103
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FJ387582.1 gg_id:359267 Burkholderia andropogonis str. 6268 FJ3875
FM872738.1 gg_id:367189 occupant source house floor dust clone FB01C03 FM872
AY792252.1 gg_id:125947 Humic Lake clone CrystalBog2B1 AY792
AY792253.1 gg_id:119501 Humic Lake clone CrystalBog571B4 AY792
EU827473.1 gg_id:344718 Burkholderia sp. str. TFD4 EU827
FJ796443.1 gg_id:569235 Burkholderia terrae str. IMER-B1-41 FJ7964
Y17009.1 gg_id:7211 Burkholderia caribensis str. MWAP64 Y17009
Y17011.1 gg_id:7220 Burkholderia caribensis str. MWAP84 Y17011
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EU477824.1 gg_id:260877 and validation prototype -based taxonomic microarray members genus Burkholderia EU4778
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GQ200825.1 gg_id:589059 Burkholderia cepacia str. V1 GQ200
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AF148555.1 gg_id:72468 Burkholderia multivorans str. LMG 13010 AF1485
AF043302.1 gg_id:7249 Burkholderia vietnamiensis str. AMMD AF0433
U96932.1 gg_id:7246 Burkholderia plantarii str. LMG16020T U9693
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AY154372.1 gg_id:76898 Burkholderia phenazinium Hg 8 AY1543
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EU024161.1 gg_id:239364 Cupriavidus metallidurans str. CIP 107179 EU0241
EU024165.1 gg_id:239356 Cupriavidus pauculus str. CIP 105943 EU0241
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EU827495.1 gg_id:310258 Ralstonia eutropha str. JMP134 EU8274
GU202015.1 gg_id:576596 Cupriavidus taiwanensis str. JSG2 GU202
NC_010530.1 gg_id:293752 Cupriavidus taiwanensis str. LMG 19424 NC_010

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EF446928.1 gg_id:214206 Cupriavidus taiwanensis str. AAJ14 EF4469
EF114432.1 gg_id:547037 Cupriavidus taiwanensis MSMB31 EF1144
GQ342296.1 gg_id:538911 Cupriavidus necator str. CG-T3 GQ342
AF067833.1 gg_id:7397 Ralstonia sp. str. TFD41 AF0678
AY583652.1 gg_id:102442 Ralstonia sp. str. 1-1 AY5836
AF500586.1 gg_id:66050 Wautersia respiraculi str. AU1618 AF5005
AY860251.1 gg_id:111512 Cupriavidus taiwanensis str. AUX0106 AY8602
AY571787.1 gg_id:100095 Ralstonia sp. str. 1-1 AY5717
AY860242.1 gg_id:111723 Cupriavidus respiraculi str. AU4630 AY8602
EU447184.1 gg_id:261584 Cupriavidus basilensis str. SE15 EU4471
D87999.1 gg_id:7386 Cupriavidus necator str. DSM 2839 D87999
AB007995.1 gg_id:58406 Wautersia eutropha str. P11 AB0079
EF114429.1 gg_id:545936 Cupriavidus gilardii MSMB32 EF1144
AF251026.1 gg_id:41961 Ralstonia sp. str. LD35 AF2510
AY860237.1 gg_id:111546 Cupriavidus respiraculi str. AU3775 AY8602
D88001.1 gg_id:7387 Ralstonia sp. str. CT14 D88001
AF312022.1 gg_id:73500 Wautersia basilensis str. DSM 11853 AF3120
DQ409082.1 gg_id:154472 Ralstonia sp. DUT_AHX str. DUT_AUX DQ409
AY860227.1 gg_id:111839 Cupriavidus gilardii str. AU3713 AY8602
AY860234.1 gg_id:111890 Cupriavidus metallidurans str. AU4057 AY8602
AY860250.1 gg_id:111665 Ralstonia sp. str. AU5980 AY8602
AY303977.1 gg_id:88067 Wautersia taiwanensis str. MS1 AY3039
AB109778.1 gg_id:83846 Wautersia basilensis str. KF708 AB1097
NR_027607.1 gg_id:555836 Cupriavidus metallidurans CH34 str. CH34; ATCC 43123 NR_027
AF361022.1 gg_id:33253 Ralstonia sp. str. NAH8 AF3610
AB104447.1 gg_id:111989 Wautersia numadzuensis str. TE26 AB1044
AB271141.1 gg_id:168999 Cupriavidus gilardii str. MXC3-9 AB2711
EU827493.1 gg_id:325237 Cupriavidus respiraculi str. TFD48 EU8274
AF300325.2 gg_id:47468 Wautersia taiwanensis str. LMG 19425 AF3003
AF500587.1 gg_id:56268 Wautersia respiraculi str. AU3369 AF5005
DQ298326.1 gg_id:152771 studies on and nifH Lasiurus sindicus Sewan grass Thar Desert Rajasthan India rhizos DQ298
AY568502.2 gg_id:106836 Wautersia basilensis str. C-10 AY5685
AY860241.1 gg_id:111717 Cupriavidus respiraculi str. AU4622 AY8602
AB167180.1 gg_id:107898 Ralstonia sp. str. c310 AB1671
Y10825.1 gg_id:7382 Wautersia eutropha str. DS185 Y10825
AM905275.1 gg_id:327185 Cupriavidus taiwanensis JC4 AM905
AF312021.1 gg_id:70450 Wautersia basilensis str. ER121 AF3120
AY860235.1 gg_id:112054 Cupriavidus metallidurans str. AU6284 AY8602
AY860229.1 gg_id:112248 Cupriavidus gilardii str. AU4927 AY8602
FJ823153.1 gg_id:563047 Lautropia sp. str. TeTO FJ8231

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GQ074337.1 gg_id:389328 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw161g GQ074
AB366174.1 gg_id:239543 Limnobacter sp. str. KP1-19 AB366
AY268172.1 gg_id:89390 Pandoraea apista str. CCUG 38412 AY268
Y09879.1 gg_id:7183 Pandoraea norimbergensis Y09879
AY268174.1 gg_id:88183 Pandoraea norimbergensis str. CCUG 39188 AY268
EF596910.1 gg_id:230684 Pandoraea pnomenusa str. B-356 EF5969
AY268168.1 gg_id:88090 Pandoraea pnomenusa str. 2001008157 AY268
AB244466.1 gg_id:240824 Pandoraea pulmonicola str. C10-2 AB244
DQ837222.1 gg_id:255307 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_1B_3 DQ837
AY677092.1 gg_id:139269 Pandoraea sputorum TC84 AY677
AM921627.1 gg_id:238898 Pandoraea sputorum PhyCEm-128 AM921
AB118229.1 gg_id:105307 Pandoraea sp. PAMU-2.78 AB118
AY693821.1 gg_id:107349 anaerobic sludge clone EUB64 AY6938
AF139176.1 gg_id:72789 Pandoraea sputorum str. LMG 18819 AF139
AM110110.2 gg_id:139991 Polynucleobacter sp. str. MWH-Braz-FAM3C AM110
AF534432.1 gg_id:85148 Toolik Lake main station at 3 m depth clone TLM08/TLMdgge25a proteobacterium AF5344
AF534427.1 gg_id:86341 Toolik Lake main station at 3 m depth clone TLM03/TLMdgge33 proteobacterium AF5344
EF667903.1 gg_id:240040 Long-term maintenance species-specific microbiota basal metazoan Hydra epithelium EF6679
FM208180.1 gg_id:320042 Polynucleobacter sp. MWH-PoolGreenA3 FM208
AY947963.1 gg_id:142102 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY9479
EU800885.1 gg_id:352838 It's all ranking aquatic Delaware Bay NJ clone 2C229123 EU8008
EU803597.1 gg_id:315886 It's all ranking aquatic Lake Gatun clone 5C231196 EU8035
AJ811013.1 gg_id:106508 Polynucleobacter sp. FC1 AJ8110
AJ270258.1 gg_id:48140 Ralstonia mannitolilytica str. LMG6866 AJ2702
EU024152.1 gg_id:251899 Ralstonia mannitolilytica str. CIP 107281 EU0241
DQ924950.1 gg_id:249852 Ralstonia solanacearum str. SL341 DQ924
AB024604.1 gg_id:50898 Ralstonia solanacearum MAFF 211266 AB0246
AY464966.1 gg_id:114414 Ralstonia syzygii str. T327 AY4649
FJ193945.1 gg_id:320934 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1939
DQ997838.1 gg_id:177991 Ralstonia pickettii str. HM-1 DQ997
EF562212.1 gg_id:247420 Subsurface microbial deep-granitic-fracture water colorado Henderson Molybdenum EF5622
AB024608.1 gg_id:40253 Ralstonia solanacearum MAFF 301559 AB0246
FJ193860.1 gg_id:338003 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1938
FJ194034.1 gg_id:308094 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1940
EF562231.1 gg_id:249054 Subsurface microbial deep-granitic-fracture water colorado Henderson Molybdenum EF5622
FJ191398.1 gg_id:326241 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1913
EF207564.1 gg_id:210668 mangrove system sediment EF2075
FJ193764.1 gg_id:350437 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1937
AJ583178.1 gg_id:95147 radioactive site ground water clone S15D-MN15 AJ5831
FJ193746.1 gg_id:349604 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1937

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AY043378.1 gg_id:112275 Ralstonia mannitolilytica str. AU428 AY0433
FJ191418.1 gg_id:310934 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1914
EF688395.1 gg_id:252030 soil clone F6-168 EF6883
EF562200.1 gg_id:246230 Subsurface microbial deep-granitic-fracture water colorado Henderson Molybdenum EF5622
FJ193846.1 gg_id:352582 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1938
FJ193974.1 gg_id:332274 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1939
FJ192332.1 gg_id:324427 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
NW_002196521.1 gg_id:341686 Ralstonia solanacearum str. MolK2 NW_00
FJ821603.1 gg_id:563550 Ralstonia sp. str. TPS8 FJ8216
FJ191397.1 gg_id:309765 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1913
FJ193962.1 gg_id:323373 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1939
FJ193740.1 gg_id:308235 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1937
EF554882.1 gg_id:223957 Ralstonia sp. str. EF38 EF5548
AY646884.1 gg_id:103925 Ralstonia mannitolilytica A55 AY6468
FJ193891.1 gg_id:341457 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1938
EF016361.1 gg_id:184466 Ralstonia solanacearum str. LMG 2299 EF0163
FJ192336.1 gg_id:305713 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
FJ193886.1 gg_id:311507 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1938
EF600590.1 gg_id:299883 leaf litter clone E2-41 EF6005
AB021403.1 gg_id:7408 Ralstonia syzygii str. ATCC 49543T AB0214
DQ256353.1 gg_id:143763 subsurface water clone EV818BHEB5102702SAS31 DQ256
EU234261.1 gg_id:236043 river and un techniques downstream Wang Yang River receiving penicillin G producti EU2342
FJ625332.1 gg_id:511989 Influence lead contamination on boreal pine forest soil clone HF_M_29 FJ6253
CU466909.1 gg_id:313184 Evry wastewater treatment plant anoxic basin clone CU4669
AB089482.1 gg_id:96705 Derxia gummosa str. IAM 13946 AB0894
EU304284.1 gg_id:244827 Ralstonia sp. str. 22 EU3042
FJ516875.1 gg_id:558389 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled lower s FJ5168
CU927753.1 gg_id:558915 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU9277
FJ484886.1 gg_id:512042 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4848
EU133821.1 gg_id:253342 tallgrass prarie soil clone FFCH11971 EU1338
EU799756.1 gg_id:319986 It's all ranking aquatic Newport Harbour RI clone 1C227401 EU7997
FJ167462.1 gg_id:310487 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1674
DQ450176.1 gg_id:151384 (Uruguay) Atlantic coastal lagoon clone R7C70 DQ450
AJ853520.1 gg_id:110392 landfill leachate clone GZKB25 AJ8535
EU491796.1 gg_id:260119 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4917
FJ612332.1 gg_id:545233 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.1.39 FJ6123
EF654710.1 gg_id:236325 suspended sludge anammox reactor clone KIST-JJY041 EF6547
AB487015.1 gg_id:546434 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB4870
AY218566.1 gg_id:81664 penguin droppings sediments clone KD1-79 AY2185
EU703319.1 gg_id:412714 lakes on Eastern Tin Plateau polysaline lake altitude 3204 masl clone XZQH193 EU7033

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CU466912.1 gg_id:355680 Evry wastewater treatment plant anoxic basin clone CU466
AY921755.1 gg_id:113342 farm soil clone AKYH739 AY9217
EU802154.1 gg_id:324381 It's all ranking aquatic Chesapeake Bay MD clone 3C003572 EU8021
EF397578.1 gg_id:287984 Burkholderia sp. str. ATSB16 EF3975
AB240293.1 gg_id:141532 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB2402
FJ375442.1 gg_id:558588 T-RFLP powerful tool rapid assessment cells microbial fuel cell sheet carbon cloth ano FJ3754
FJ612382.1 gg_id:567560 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.3.32 FJ6123
AY922020.1 gg_id:113843 farm soil clone AKYG464 AY9220
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GQ010467.1 gg_id:499198 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ010
FJ230924.1 gg_id:326020 under Level Antibiotic Selection Pressures river water clone F2 FJ2309
AY509453.2 gg_id:103790 freshwater clone LiUU-5-131 AY5094
FJ674610.1 gg_id:515098 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface FJ6746
EF516760.1 gg_id:225714 grassland soil clone FCPS725 EF5167
AB252925.1 gg_id:170130 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 57 AB2529
EU727457.1 gg_id:354510 Ralstonia sp. str. JOS55-1 EU7274
GQ500805.1 gg_id:570888 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
GQ389057.1 gg_id:591556 quality deterioration drinking system during serious red water outbreak clone S13 GQ389
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GQ284434.1 gg_id:593931 Acidovorax delafieldii str. PCWCS28 GQ284
EU730927.1 gg_id:315092 Acidovorax facilis str. 228 EU730
EU304287.1 gg_id:251328 Acidovorax sp. str. 75 EU3042
Y18616.1 gg_id:7043 Acidovorax defluvii str. BSB411 Y18616
AB487337.1 gg_id:553499 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB4873
EU104261.1 gg_id:346700 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1042
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GU372409.1 gg_id:575562 Acidovorax sp. str. S7 GU372
GQ106011.1 gg_id:431652 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ106

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FJ593725.1 gg_id:531432 midgut tropical caterpillars fed on diets larvae clone CLSM148 FJ5937
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AB076844.1 gg_id:28959 Acidovorax sp. str. OS-6 AB076
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GQ102686.1 gg_id:413906 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ102
FJ497649.1 gg_id:585668 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4976
GQ104106.1 gg_id:447512 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw563c0 GQ104
AF015487.1 gg_id:7077 Pseudomonas sp. str. P51 AF015
FJ658868.1 gg_id:568875 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Sumn FJ6588
FJ477624.1 gg_id:552067 gut symbionts are tightly linked herbivory ants whole extracted single ant clone UAB- FJ4776
FJ477611.1 gg_id:578828 gut symbionts are tightly linked herbivory ants whole extracted mid- and hind-gut sin FJ4776
AF508114.1 gg_id:66230 Acidovorax avenae C1 AF508
FJ657748.1 gg_id:545408 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6577

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FJ658799.1 gg_id:510766 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Sumn FJ6587
FJ658952.1 gg_id:553262 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Sumn FJ6589
FJ658390.1 gg_id:544123 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6583
AB120964.1 gg_id:99037 Aquamonas fontana str. AQ10 AB120
DQ640723.1 gg_id:227889 full-scale EBPR sludge clone Skagen135 DQ640
AF526926.2 gg_id:62982 Mars Odyssey Orbiter and encapsulation facility clone T5-4 sp. AF5269
AY133071.1 gg_id:56377 TCE-contaminated site clone ccs221 AY1330
DQ521541.1 gg_id:193826 Antarctic lake ice cover clone ANTLV7_G03 DQ521
AY662013.1 gg_id:137452 uranium-contaminated groundwater clone 015C-G04 AY6620
AF526920.2 gg_id:62991 Mars Odyssey Orbiter and encapsulation facility clone T8-7 sp. AF5269
EU150266.1 gg_id:279514 acidobacteria and soil Niwot Ridge LTER clone Ni_F8P2 EU1502
NR_028713.1 gg_id:585432 Curvibacter delicatus str. 146; LMG 4328 NR_028
D14320.1 gg_id:6938 Brachymonas denitrificans str. -P1 D14320
EU434344.1 gg_id:271636 Brachymonas denitrificans str. a265 EU4343
EU434398.1 gg_id:265311 Brachymonas denitrificans str. a310 EU4343
AB286490.1 gg_id:208338 Reducing production and domination comamonadaceae reducing oxygen supply was AB2864
AJ318129.1 gg_id:21007 Waste-gas biofilter clone BIfdi44 AJ3181
AM420229.1 gg_id:239064 Noma subgingival plaque clone 601H06(oral) AM420
FJ982927.1 gg_id:460813 Comamonas sp. str. ST18 FJ9829
AB164432.1 gg_id:103333 Comamonas badia str. IAM 14839 AB1644
AJ576357.1 gg_id:107318 hindgut homogenate Pachnoda ephippiata larva clone PeH41 AJ5763
AF234720.1 gg_id:73090 sludge clone A6b AF2347
EU346912.1 gg_id:265983 Brachymonas sp. str. AImA4 EU3469
AB286456.1 gg_id:216049 Reducing production and domination comamonadaceae reducing oxygen supply was AB2864
FJ959768.1 gg_id:584835 bottlenose dolphin upper respiratory tract blowhole clone D8A8_102 FJ9597
AB286474.1 gg_id:214136 Reducing production and domination comamonadaceae reducing oxygen supply was AB2864
EU881354.1 gg_id:306436 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
AB286506.1 gg_id:211133 Reducing production and domination comamonadaceae reducing oxygen supply was AB2865
GQ010546.1 gg_id:499118 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ010
GQ110427.1 gg_id:447775 Topographical and Temporal Human Skin Microbiome skin back clone nbw659e10c1 GQ110
AM283038.1 gg_id:159681 Caldimonas sp. str. Han85 AM283
EU024140.1 gg_id:247295 Comamonas aquatica str. CIP 107986 EU0241
EU464974.1 gg_id:295063 African elephant feces clone AFYEL_aaj66a08 EU4649
DQ854969.1 gg_id:161904 Comamonas sp. str. GPTSA5 DQ854
EU464791.1 gg_id:289604 Hartmann's mountain zebra feces clone AFZEB_aaj70h03 EU4647
AF408327.1 gg_id:36914 Comamonas sp. clone KL-11-1-9 AF4083
EU841519.1 gg_id:337127 Comamonas aquatica str. 406 EU8415
DQ824524.1 gg_id:188323 human fecal clone RL185_aaj69c02 DQ824
AJ430344.1 gg_id:73664 Comamonas aquatica str. LMG 2370 AJ4303
AF233878.1 gg_id:40166 Comamonas denitrificans str. 2.99g AF2338
AF233879.1 gg_id:74723 Comamonas denitrificans str. 5.38g AF2338

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AF233877.1 gg_id:23191 Comamonas denitrificans str. 123	AF2338
AJ430348.1 gg_id:66684 Comamonas kerstersii str. LMG 5323	AJ4303
AB021418.1 gg_id:6936 Comamonas terrigena str. IAM 12052T	AB0214
AJ430343.1 gg_id:66690 Comamonas terrigena str. LMG 1249	AJ4303
AJ420326.1 gg_id:17652 Comamonas terrigena str. DSM 7099	AJ4203
EU024142.1 gg_id:240089 Comamonas terrigena str. LMG 5520	EU0241
AM184229.1 gg_id:160747 Comamonas terrigena str. WAB1888	AM184
NR_028719.1 gg_id:545968 Comamonas terrigena str. IMI 359870; ATCC 8461	NR_028
EU337122.1 gg_id:545663 Comamonas denitrificans str. 2B7	EU3371
GU454930.1 gg_id:524752 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir	GU454
DQ824755.1 gg_id:189547 human fecal clone RL185_aaj72g08	DQ824
EU465136.1 gg_id:290700 African elephant feces clone AFYEL_aaj68e11	EU4651
FJ658617.1 gg_id:509630 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Sumn	FJ6586
EU464722.1 gg_id:300641 Hartmann's mountain zebra feces clone AFZEB_aaj69f11	EU4647
AY566581.1 gg_id:108740 Comamonas sp. str. PG-08	AY5665
GQ179675.1 gg_id:465713 Heterogeneity vaginal microbial within individuals vagina clone VE3A04	GQ179
EU465091.1 gg_id:291187 African elephant feces clone AFYEL_aaj67h12	EU4650
EU465027.1 gg_id:296452 African elephant feces clone AFYEL_aaj66h05	EU4650
EU536017.1 gg_id:286592 profile human microbiota antecubital fossa (inner elbow) skin clone nbt161h12	EU5360
EU465029.1 gg_id:292604 African elephant feces clone AFYEL_aaj66h08	EU4650
EU465048.1 gg_id:302343 African elephant feces clone AFYEL_aaj67c02	EU4650
EU465099.1 gg_id:302440 African elephant feces clone AFYEL_aaj68a11	EU4650
AY635871.1 gg_id:99471 Comamonas sp. str. BM-9_6	AY6358
EU464843.1 gg_id:295086 Hartmann's mountain zebra feces clone AFZEB_aaj71g07	EU4648
EU465006.1 gg_id:293491 African elephant feces clone AFYEL_aaj66e11	EU4650
EU771834.1 gg_id:310863 African elephant feces clone AFYEL_aaj65h12	EU7718
AB076865.1 gg_id:42211 anaerobic sludge clone OS1L-7	AB0768
EU771819.1 gg_id:310322 African elephant feces clone AFYEL_aaj68f04	EU7718
EU464881.1 gg_id:301051 Hartmann's mountain zebra feces clone AFZEB_aaj72d04	EU4648
EU465050.1 gg_id:287469 African elephant feces clone AFYEL_aaj67c06	EU4650
AY856846.1 gg_id:111198 Comamonas sp. str. 46	AY8568
EU464826.1 gg_id:287428 Hartmann's mountain zebra feces clone AFZEB_aaj71e07	EU4648
GQ389171.1 gg_id:571114 quality deterioration drinking system during serious red water outbreak clone D40	GQ389
EU771661.1 gg_id:334294 African elephant feces clone AFEL_aai29g11	EU7716
EU464941.1 gg_id:302302 African elephant feces clone AFYEL_aaj65e07	EU4649
EU465157.1 gg_id:292068 African elephant feces clone AFYEL_aaj68h04	EU4651
EU465046.1 gg_id:299793 African elephant feces clone AFYEL_aaj67b12	EU4650
EU464997.1 gg_id:294410 African elephant feces clone AFYEL_aaj66d12	EU4649
EU465004.1 gg_id:294874 African elephant feces clone AFYEL_aaj66e08	EU4650
EU465015.1 gg_id:298465 African elephant feces clone AFYEL_aaj66f12	EU4650
DQ453128.1 gg_id:156701 Comamonas sp. str. Dant 3-8	DQ453

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EF114434.1 gg_id:534604 strains recovered during surveys Burkholderia pseudomallei northern Australia rod EF1144
EF565937.1 gg_id:228582 Comamonas sp. str. NF27 EF5659
EU465126.1 gg_id:287764 African elephant feces clone AFYEL_aaj68d09 EU4651
AF501880.1 gg_id:71872 Comamonas testosteroni str. SMCC B0589 AF5018
EU467917.1 gg_id:289291 cheetah feces clone CE2_a08 EU4679
GU454975.1 gg_id:515113 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
EU468077.1 gg_id:291421 cheetah feces clone CE2_d08_3 EU4680
EU464621.1 gg_id:289827 African elephant feces clone AFEL_aai32a05 EU4646
AY291591.1 gg_id:87769 Comamonas testosteroni str. CNB1 AY2915
AY134850.1 gg_id:58604 Comamonas sp. str. SFCD1 AY1348
EU464754.1 gg_id:291460 Hartmann's mountain zebra feces clone AFZEB_aaj70c03 EU4647
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EU464969.1 gg_id:298206 African elephant feces clone AFYEL_aaj66a02 EU4649
EU464920.1 gg_id:299963 African elephant feces clone AFYEL_aaj65b11 EU4649
AY819705.1 gg_id:109404 Comamonas sp. str. JS46 AY8197
AJ606336.1 gg_id:101448 Comamonas testosteroni SB4 AJ6063
EU464815.1 gg_id:287989 Hartmann's mountain zebra feces clone AFZEB_aaj71c10 EU4648
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EU779419.1 gg_id:343489 Hartmann's Mountain Zebra feces clone AFZEB_aaj72d07 EU7794
EU464929.1 gg_id:294023 African elephant feces clone AFYEL_aaj65c11 EU4649
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GQ008024.1 gg_id:501642 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw: GQ008
EU464698.1 gg_id:298530 Hartmann's mountain zebra feces clone AFZEB_aaj69d04 EU4646
EU464986.1 gg_id:294420 African elephant feces clone AFYEL_aaj66c06 EU4649
EU537205.1 gg_id:281659 profile human microbiota antecubital fossa (inner elbow) skin clone nbt68d07 EU5372
FJ893904.1 gg_id:540808 Matriptase-deficient mice exhibit ichthyotic selective shift skin microbiota B1E; ear pu FJ8939
AF532869.1 gg_id:91652 Comamonas sp. str. D1 AF5328
EU464891.1 gg_id:300806 Hartmann's mountain zebra feces clone AFZEB_aaj72f06 EU4648
EU464935.1 gg_id:293225 African elephant feces clone AFYEL_aaj65d12 EU4649
EU464774.1 gg_id:288302 Hartmann's mountain zebra feces clone AFZEB_aaj70e07 EU4647
EU465010.1 gg_id:298312 African elephant feces clone AFYEL_aaj66f07 EU4650
EU464915.1 gg_id:291947 African elephant feces clone AFYEL_aaj65a11 EU4649
AJ251577.1 gg_id:38824 Comamonas nitrativorans str. 23310 AJ2515
AF233876.1 gg_id:18248 Comamonas denitrificans str. 110 AF2338
EU465109.1 gg_id:287376 African elephant feces clone AFYEL_aaj68b09 EU4651
EU465146.1 gg_id:297240 African elephant feces clone AFYEL_aaj68g02 EU4651
EU536192.1 gg_id:282831 profile human microbiota antecubital fossa (inner elbow) skin clone nbt12g10 EU5361
AB109889.1 gg_id:106408 Curvibacter gracilis str. 7-1 AB1098
DQ001153.1 gg_id:118400 Rhodobacter sphaeroides str. Il106 DQ001

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