

Table 2: OTUs enriched in stover amended soil relative to control
BLAST against Living Tree Project

OTU ID	Day of Response ^a	Top BLAST hits	BLAST %ID
OTU.3	12, 82	<i>Arthrobacter spp.</i>	100.0
OTU.4	12	<i>Massilia tieshanensis</i> , <i>Massilia aerilata</i>	100.0
OTU.6	12, 82	<i>Bacillus megaterium</i> , <i>Bacillus flexus</i>	100.0
OTU.9	12	<i>Sphingobacterium siyangense</i>	100.0
OTU.10	12, 82	<i>Terrabacter terrae</i> , <i>Terrabacter tumescens</i> , <i>Terrabacter sp. ON10</i> , <i>Terrabacter lapilli</i> , <i>Terrabacter sp. PY2</i>	100.0
OTU.18	12, 82	<i>Buttiauxella warmboldiae</i> , <i>Buttiauxella izardii</i> , <i>Buttiauxella agrestis</i> , <i>Pantoea rwandensis</i> , <i>Pantoea rodasii</i> , <i>Enterobacter amnigenus</i> , <i>Buttiauxella noackiae</i>	100.0
OTU.28	12, 82	<i>Stenotrophomonas chelatiphaga</i>	98.67
OTU.33	82	<i>Roseomonas aquatica</i>	98.13
OTU.39	12, 82	<i>Thermomonas dokdonensis</i>	99.47
OTU.45	82	<i>Rudaea cellulosilytica</i>	97.06
OTU.57	12, 82	<i>Methylobacterium radiotolerans</i> , <i>Methylobacterium longum</i>	100.0
OTU.58	12, 82	<i>Aquamonas fontana</i>	99.73
OTU.62	12, 82	<i>Streptomyces spp.</i>	100.0
OTU.78	12	<i>Oxalicibacterium flavum</i>	98.92
OTU.82	12, 82	<i>Rhizobium endophyticum</i> , <i>Rhizobium tubonense</i> , <i>Rhizobium tibeticum</i>	100.0
OTU.83	82	<i>Terrimonas sp. M-8</i> , <i>Terrimonas sp. RIB1-6</i>	95.95
OTU.84	12, 82	<i>Tumebacillus permanentifrigoris</i>	99.46
OTU.88	12	<i>Chryseobacterium sp. THMBM1</i>	99.73
OTU.89	82	<i>Arenimonas sp. CH15-1</i>	99.2
OTU.91	12	<i>Chryseobacterium jejuense</i> , <i>Chryseobacterium sp. RBA2-6</i>	98.92
OTU.93	12	<i>Methylocystis echinoides</i>	96.51
OTU.100	12	<i>Novosphingobium naphthalenivorans</i> , <i>Novosphingobium mathurense</i> , <i>Novosphingobium pentaromativorans</i> , <i>Novosphingobium resinovorum</i>	99.46
OTU.112	12, 82	<i>Pseudomonas fulva</i> , <i>Pseudomonas flavescens</i> , <i>Pseudomonas benzenivorans</i> , <i>Pseudomonas argentinensis</i>	100.0
OTU.114	12, 82	<i>Microvirga quangxiensis</i>	99.46

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OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.118	12, 82	<i>Flavobacterium beibuense</i> , <i>Flavobacterium sp. FCS-5</i>	98.11
OTU.120	82	<i>Devosia insulae</i>	99.46
OTU.132	82	<i>Ilumatobacter fluminis</i>	97.87
OTU.133	12	<i>Deinococcus gobiensis</i> , <i>Deinococcus navajonensis</i>	95.41
OTU.139	12, 82	<i>Burkholderia glathei</i>	100.0
OTU.140	82	<i>Rhizobiales bacterium Mfc52</i>	92.74
OTU.141	12, 82	<i>Achromobacter spanius</i> , <i>Achromobacter insolitus</i>	100.0
OTU.152	82	<i>Chitinophaga arvensicola</i>	94.32
OTU.154	12, 82	<i>Comamonas thiooxydans</i> , <i>Comamonas testosteroni</i>	99.73
OTU.160	82	<i>Hydrotalea flava</i>	95.41
OTU.166	82	<i>Niastella sp. JCN-23</i>	96.49
OTU.176	82	No hits of at least 90% identity	86.74
OTU.177	12, 82	<i>Microvirga aerilata</i>	98.66
OTU.184	12, 82	<i>Adhaeribacter terreus</i>	98.11
OTU.185	12, 82	<i>Erythromicrobium ramosum</i> , <i>Porphyrobacter tepidarius</i>	99.46
OTU.190	12	<i>Lysobacter sp. RCML-52</i>	99.73
OTU.191	12	<i>Geothrix fermentans</i>	96.26
OTU.193	82	No hits of at least 90% identity	84.88
OTU.218	12, 82	<i>Rhizobium cellulosilyticum</i>	100.0
OTU.229	12	<i>Flavisolibacter ginsengisoli</i>	96.49
OTU.265	12	<i>Segetibacter koreensis</i>	99.19
OTU.269	82	<i>Aminobacter aminovorans</i> , <i>Mesorhizobium loti</i> , <i>Mesorhizobium australicum WSM2073</i> , <i>Mesorhizobium shangrilense</i>	100.0
OTU.271	12, 82	<i>Bosea sp. R-46060</i>	100.0
OTU.287	12, 82	<i>Flavisolibacter ginsengisoli</i>	95.14
OTU.311	82	<i>Burkholderia sordidicola</i>	92.78
OTU.316	12, 82	<i>Caulobacter henricii</i>	99.46
OTU.319	82	<i>Dokdonella sp. KIS28-6</i>	100.0
OTU.333	12, 82	No hits of at least 90% identity	89.97
OTU.339	12, 82	<i>Brevundimonas halotolerans</i>	99.73
OTU.343	12, 82	<i>Pedobacter rhizosphaerae</i>	100.0

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OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.349	82	No hits of at least 90% identity	88.13
OTU.352	82	<i>Sphingobacteria bacterium RYG</i>	93.28
OTU.357	12, 82	<i>Ochrobactrum pseudogrignonense</i>	100.0
OTU.361	12, 82	<i>Streptosporangium amethystogenes</i> subsp. <i>amethystogenes</i> , <i>Streptosporangium longisporum</i> , <i>Streptosporangium album</i> , <i>Streptosporangium roseum</i> , <i>Streptosporangium oxazolinicum</i> , <i>Streptosporangium amethystogenes</i> subsp. <i>fukuense</i> , <i>Streptosporangium canum</i>	100.0
OTU.373	12, 82	<i>Paenibacillus pabuli</i> , <i>Paenibacillus tundrae</i> , <i>Paenibacillus taichungensis</i> , <i>Paenibacillus xylanexedens</i> , <i>Paenibacillus xylanilyticus</i>	100.0
OTU.379	82	No hits of at least 90% identity	86.77
OTU.391	12	<i>Flavisolibacter ginsengisoli</i>	95.68
OTU.395	12, 82	<i>Solibius ginsengiterrae</i>	95.41
OTU.399	82	<i>Flavobacterium</i> sp. <i>DK69</i>	96.49
OTU.402	12	<i>Brevundimonas vesicularis</i> , <i>Brevundimonas nasdae</i>	100.0
OTU.412	82	<i>Sphingobacteria bacterium RYG</i>	92.43
OTU.413	12, 82	<i>Cellulomonas aerilata</i> , <i>Cellulomonas terrae</i> , <i>Cellulomonas humilata</i> , <i>Cellulomonas soli</i>	100.0
OTU.442	12, 82	<i>Rhodococcus wratislaviensis</i>	100.0
OTU.444	82	No hits of at least 90% identity	84.95
OTU.448	12, 82	<i>Brevundimonas alba</i>	99.19
OTU.451	12	<i>Polaromonas aquatica</i> , <i>Polaromonas jejuensis</i> , <i>Polaromonas vacuolata</i>	99.46
OTU.454	12, 82	<i>Nocardioides hwasunensis</i>	100.0
OTU.455	12, 82	<i>Methylobacterium rhodesianum</i> , <i>Methylobacterium populi</i> , <i>Methylobacterium zatmanii</i>	100.0
OTU.464	82	No hits of at least 90% identity	82.18
OTU.468	82	<i>Steroidobacter denitrificans</i>	91.18
OTU.472	82	<i>Steroidobacter denitrificans</i>	98.4
OTU.477	82	<i>Ferruginibacter alkalilentus</i> , <i>Ferruginibacter lapsinensis</i>	95.95
OTU.484	82	<i>Reyranella massiliensis</i>	97.85
OTU.503	82	<i>Opitutus terrae</i>	90.91

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OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.507	12, 82	<i>Methylophilus methylotrophus</i> , <i>Methylothera versatilis</i> 301	99.2
OTU.518	82	<i>Amaricoccus kaplicensis</i>	96.51
OTU.519	12	<i>Acinetobacter nosocomialis</i> , <i>Acinetobacter calcoaceticus</i> , <i>Acinetobacter pittii</i>	100.0
OTU.523	82	<i>Phaselicystis flava</i>	92.25
OTU.524	82	<i>Aeromonas spp.</i>	100.0
OTU.529	82	No hits of at least 90% identity	88.38
OTU.534	12, 82	<i>Methylobacterium aquaticum</i>	100.0
OTU.535	12, 82	<i>Cupriavidus necator</i> , <i>Wautersia numazuensis</i> , <i>Cupriavidus basilensis</i>	100.0
OTU.536	82	<i>Ferruginibacter alkalilentus</i> , <i>Ferruginibacter lapsinensis</i>	94.07
OTU.541	12	<i>Flavobacterium sp. MK3</i>	99.46
OTU.548	82	No hits of at least 90% identity	89.3
OTU.550	82	<i>Ohtaekwangia koreensis</i>	92.72
OTU.564	82	<i>Solitalea koreensis</i>	99.73
OTU.565	82	<i>Marinobacter salicampi</i>	92.51
OTU.567	12	<i>Dyella koreensis</i> , <i>Dyella soli</i>	100.0
OTU.571	82	<i>Rhodopseudomonas sp. JA576</i> , <i>Rhodopseudomonas faecalis</i> , <i>Rhodopseudomonas palustris</i> , <i>Rhodopseudomonas sp. R-45977</i>	98.12
OTU.582	82	No hits of at least 90% identity	88.3
OTU.589	12, 82	<i>Sphingomonas trueperi</i> , <i>Sphingomonas pituitosa</i>	98.66
OTU.590	12, 82	<i>Chryseobacterium piscium</i> , <i>Chryseobacterium balustinum</i>	99.19
OTU.599	82	<i>Polynucleobacter acidiphobus</i>	90.43
OTU.604	12, 82	<i>Pseudochrobactrum sp. KSS 7.8</i> , <i>Pseudochrobactrum saccharolyticum</i> , <i>Pseudochrobactrum asaccharolyticum</i>	97.85
OTU.609	82	No hits of at least 90% identity	89.65
OTU.612	12, 82	<i>Burkholderia phenoliruptrix</i> , <i>Burkholderia graminis</i>	100.0
OTU.614	82	<i>Singulisphaera acidiphila</i>	99.73
OTU.619	12, 82	<i>Nocardioides hankookensis</i>	98.67
OTU.628	82	<i>Spirochaeta aurantia subsp. aurantia</i>	95.17

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OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.629	82	No hits of at least 90% identity	83.99
OTU.635	12	<i>Sphingomonadaceae bacterium KMM 6042</i>	99.2
OTU.636	12	<i>Rummeliibacillus pycnus</i>	99.2
OTU.639	12, 82	<i>Pedobacter ginsengisoli</i>	100.0
OTU.640	12, 82	<i>Rhodococcus jostii</i>	99.73
OTU.642	82	No hits of at least 90% identity	88.03
OTU.645	12, 82	<i>Kaistia sp. 5YN7-3, Kaistia sp. B6-12</i>	100.0
OTU.649	12	<i>Kribbella antibiotica, Kribbella flavida, Kribbella sp. PIP 118</i>	100.0
OTU.655	12	<i>Bacillus gibsonii</i>	100.0
OTU.673	12	<i>Ramlibacter tataouinensis</i>	98.66
OTU.678	12, 82	<i>Shinella granuli, Shinella zoogloeoides</i>	100.0
OTU.680	82	No hits of at least 90% identity	88.0
OTU.688	82	<i>Sphingobacteria bacterium RYG</i>	96.22
OTU.689	82	No hits of at least 90% identity	87.17
OTU.697	12, 82	<i>Dokdonella koreensis</i>	95.19
OTU.702	82	No hits of at least 90% identity	83.89
OTU.704	12, 82	<i>Microbacterium fluvii, Microbacterium pumilum, Microbacterium deminutum, Microbacterium terricola</i>	100.0
OTU.706	82	<i>Cellvibrio fulvus</i>	100.0
OTU.707	82	<i>Paracoccus sp. NB88</i>	99.46
OTU.715	82	No hits of at least 90% identity	86.74
OTU.720	82	<i>Luteolibacter sp. CCTCC AB 2010415</i>	99.2
OTU.734	12, 82	<i>Sphingomonas changbaiensis</i>	98.66
OTU.746	82	No hits of at least 90% identity	88.5
OTU.767	82	No hits of at least 90% identity	89.57
OTU.774	82	<i>Sphingobacteria bacterium RYG</i>	90.84
OTU.779	82	<i>Phenylobacterium falsum</i>	95.43
OTU.791	82	No hits of at least 90% identity	85.83
OTU.814	82	<i>Pirellula staleyi DSM 6068</i>	91.4
OTU.816	12	<i>Terriglobus saanensis SP1PR4</i>	98.13
OTU.827	82	<i>Roseomonas terrae, Roseomonas lacus</i>	98.12
OTU.831	82	<i>Acidiphilium acidophilum</i>	94.65
OTU.835	82	No hits of at least 90% identity	89.81
OTU.837	82	No hits of at least 90% identity	83.29

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OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.838	12	No hits of at least 90% identity	85.41
OTU.840	82	<i>Dongia mobilis</i>	96.78
OTU.862	12, 82	<i>Mucilaginibacter sp. BR-18</i>	99.19
OTU.863	82	<i>Singulisphaera rosea</i>	92.8
OTU.873	12	<i>Hymenobacter algoricola</i>	100.0
OTU.876	12, 82	<i>Shimazuella kribbensis</i>	99.19
OTU.887	82	<i>Gemmatimonas aurantiaca</i>	92.74
OTU.897	82	<i>Bryobacter aggregatus</i>	92.47
OTU.900	12, 82	<i>Paenibacillus pocheonensis</i>	100.0
OTU.908	82	No hits of at least 90% identity	82.67
OTU.909	82	No hits of at least 90% identity	89.25
OTU.923	82	<i>Chitinimonas taiwanensis</i> , <i>Burkholderia soli</i>	90.45
OTU.924	82	No hits of at least 90% identity	88.56
OTU.939	12, 82	<i>Dyadobacter beijingensis</i>	97.57
OTU.940	82	<i>Asticcacaulis taihuensis</i>	96.26
OTU.942	12	<i>Hymenobacter rigui</i>	99.46
OTU.964	12	<i>Pelosinus fermentans</i> , <i>Pelosinus propionicus</i>	99.2
OTU.967	82	No hits of at least 90% identity	85.75
OTU.971	82	<i>Sphingobacteria bacterium RYG</i>	92.45
OTU.976	12, 82	<i>Paenibacillus borealis</i>	99.73
OTU.993	12, 82	<i>Fimbriimonas ginsengisoli Gsoil 348</i>	92.55
OTU.1008	82	<i>Phaselicystis flava</i>	92.25
OTU.1013	82	<i>Ilumatobacter fluminis</i>	94.15
OTU.1016	82	<i>Sphingomonas changbaiensis</i>	96.25
OTU.1022	12, 82	No hits of at least 90% identity	85.87
OTU.1031	82	<i>Bryobacter aggregatus</i>	90.86
OTU.1035	82	<i>Spirochaeta aurantia subsp. aurantia</i>	94.37
OTU.1045	12, 82	<i>Rhodanobacter sp. DCY45</i> , <i>Rhodanobacter fulvus</i>	100.0
OTU.1046	12	<i>Clostridium saccharoperbutylacetonicum</i>	99.73
OTU.1057	12	<i>Vampirovibrio chlorellavorus</i>	94.32
OTU.1062	82	<i>Ohtaekwangia koreensis</i>	93.51
OTU.1083	82	<i>Aciditerrimonas ferrireducens</i>	92.8
OTU.1085	12, 82	<i>Sphingomonas endophytica</i> , <i>Sphingomonas phyllosphaerae</i>	100.0
OTU.1090	82	<i>Ohtaekwangia koreensis</i>	94.61

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OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.1102	82	<i>Geobacter grbiciae</i> , <i>Desulfuromonas michiganensis</i> , <i>Geobacter metallireducens</i>	90.27
OTU.1104	12	No hits of at least 90% identity	87.5
OTU.1106	82	<i>Adhaeribacter aerophilus</i>	91.83
OTU.1107	82	<i>Lysobacter sp. DCY21T</i>	99.2
OTU.1108	82	<i>Pirellula staleyi DSM 6068</i>	95.16
OTU.1119	12, 82	<i>Devosia crocina</i> , <i>Devosia riboflavina</i>	98.66
OTU.1120	82	<i>Chitinimonas taiwanensis</i>	91.47
OTU.1123	82	<i>Porticoccus litoralis</i>	92.49
OTU.1133	12, 82	<i>Pedobacter insulae</i>	98.11
OTU.1134	82	<i>Sporocytophaga myxococcoides</i>	98.65
OTU.1145	82	<i>Cohaesibacter gelatinilyticus</i>	90.45
OTU.1155	12	No hits of at least 90% identity	80.58
OTU.1166	82	<i>Planctomyces limnophilus</i>	91.47
OTU.1182	12	<i>Pseudomonas psychrotolerans</i>	99.46
OTU.1192	82	<i>Rhodococcus triatomae</i>	97.87
OTU.1195	12	<i>Segetibacter koreensis</i>	97.3
OTU.1200	82	<i>Frigoribacterium mesophilum</i>	99.73
OTU.1208	12	<i>Bacteriovorax stolpii</i>	98.93
OTU.1217	12	<i>Chitinophaga pinensis</i>	98.65
OTU.1222	12	<i>Bacillus aerophilus</i> , <i>Bacillus pumilus</i> , <i>Bacillus safensis</i> , <i>Bacillus stratosphericus</i> , <i>Bacillus altitudinis</i>	100.0
OTU.1226	12, 82	<i>Nocardiosis alba</i>	100.0
OTU.1234	82	<i>Schlesneria paludicola</i>	97.31
OTU.1237	82	<i>Terrimonas ferruginea</i>	92.45
OTU.1240	12	<i>Hymenobacter sp. A2-91</i>	98.11
OTU.1252	82	<i>Singulisphaera rosea</i>	94.93
OTU.1259	82	No hits of at least 90% identity	80.11
OTU.1264	82	<i>Terrimonas lutea</i>	93.55
OTU.1272	12	No hits of at least 90% identity	85.11
OTU.1276	82	<i>Turneriella parva</i>	99.47
OTU.1288	82	No hits of at least 90% identity	86.15
OTU.1289	12	<i>Clostridium pascui</i> , <i>Clostridium peptidivorans</i>	95.72
OTU.1295	12, 82	<i>Leifsonia poae</i>	99.2
OTU.1301	82	<i>Burkholderia sp. ATSB16</i>	94.39

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OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.1304	82	<i>Methylobacillus glycogenes</i> , <i>Methylobacillus sp. Lap</i>	97.33
OTU.1321	82	<i>Sphingobacteria bacterium RYG</i>	94.09
OTU.1323	82	<i>Cystobacter badius</i> , <i>Cystobacter velatus</i> , <i>Cystobacter miniatus</i>	94.93
OTU.1324	12, 82	<i>Sphingomonas japonica</i>	98.37
OTU.1329	82	No hits of at least 90% identity	89.04
OTU.1344	12, 82	<i>Hymenobacter ocellatus</i>	97.3
OTU.1347	82	<i>Friedmanniella spumicola</i> , <i>Friedmanniella okinawensis</i> , <i>Friedmanniella antarctica</i> , <i>Friedmanniella sp. W6</i>	100.0
OTU.1351	82	No hits of at least 90% identity	89.89
OTU.1366	12, 82	<i>Agromyces humatus</i>	99.73
OTU.1373	12, 82	<i>Dyella marensis</i> , <i>Dyella terrae</i> , <i>Fulvimonas soli</i> , <i>Dokdonella sp. LM 2-5</i>	95.45
OTU.1378	12	<i>Chitinophaga arvensicola</i> , <i>Chitinophaga niastensis</i>	93.24
OTU.1380	82	No hits of at least 90% identity	89.36
OTU.1384	12, 82	<i>Sphingomonas trueperi</i> , <i>Sphingomonas pituitosa</i>	96.78
OTU.1385	12, 82	<i>Sphingopyxis panaciterrae</i> , <i>Sphingopyxis chilensis</i>	100.0
OTU.1389	12	<i>Arthrobacter crystallopoietes</i>	98.13
OTU.1406	82	<i>Geminicoccus roseus</i>	92.49
OTU.1425	12	No hits of at least 90% identity	89.54
OTU.1428	82	<i>Chitinophaga ginsengisegetis</i>	92.18
OTU.1436	12	No hits of at least 90% identity	88.47
OTU.1439	82	<i>Chitinophaga sp. CS5-B1</i>	99.73
OTU.1490	82	No hits of at least 90% identity	81.42
OTU.1495	82	<i>Haliea mediterranea</i>	92.53
OTU.1496	12, 82	<i>Bdellovibrio bacteriovorus</i>	93.32
OTU.1506	12	<i>Paenibacillus turicensis</i>	98.66
OTU.1518	82	<i>Steroidobacter denitrificans</i>	96.79
OTU.1522	82	<i>Haliangium ochraceum</i>	91.03
OTU.1524	82	<i>Roseococcus suduntuyensis</i>	98.39
OTU.1535	82	<i>Nocardioides sp. OS4</i> , <i>Nocardioides fonticola</i>	98.13
OTU.1544	82	<i>Armatimonas rosea</i>	92.0
OTU.1550	12, 82	<i>Roseomonas ludipueritiae</i>	98.39

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OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.1552	82	No hits of at least 90% identity	85.56
OTU.1553	82	<i>Singulisphaera rosea</i>	91.2
OTU.1556	12	<i>Paenibacillus agarexedens</i> , <i>Paenibacillus sputi</i>	98.12
OTU.1564	82	<i>Niastella yeongjuensis</i>	99.73
OTU.1566	82	<i>Panacagrimonas perspica</i>	98.12
OTU.1570	82	No hits of at least 90% identity	78.88
OTU.1574	12	<i>Dyadobacter ginsengisoli</i>	98.38
OTU.1575	12	<i>Exiguobacterium indicum</i> , <i>Exiguobacterium acetylicum</i>	100.0
OTU.1576	12, 82	<i>Delftia tsuruhatensis</i> , <i>Delftia lacustris</i>	98.66
OTU.1577	82	<i>Isosphaera-like str. OJF2</i>	94.16
OTU.1587	82	No hits of at least 90% identity	85.68
OTU.1588	82	<i>Pseudolabrys taiwanensis</i>	96.51
OTU.1593	82	No hits of at least 90% identity	89.87
OTU.1594	82	<i>Schlesneria paludicola</i>	94.1
OTU.1599	82	No hits of at least 90% identity	86.98
OTU.1613	82	<i>Phaselicystis flava</i>	91.47
OTU.1633	82	No hits of at least 90% identity	89.46
OTU.1636	82	<i>Phenylobacterium falsum</i>	97.04
OTU.1640	82	<i>Nocardioides halotolerans</i>	98.13
OTU.1659	12	<i>Paenibacillus daejeonensis</i>	100.0
OTU.1667	82	No hits of at least 90% identity	84.17
OTU.1670	82	No hits of at least 90% identity	89.84
OTU.1675	82	No hits of at least 90% identity	86.54
OTU.1680	82	<i>Pseudomonas alcaligenes</i>	98.1
OTU.1682	12	<i>Collimonas fungivorans</i> , <i>Oxalicibacterium faecigallinarum</i> , <i>Oxalicibacterium horti</i>	97.85
OTU.1684	12, 82	<i>Serratia marcescens</i> , <i>Serratia nematodiphila</i>	100.0
OTU.1714	82	No hits of at least 90% identity	83.33
OTU.1717	12	<i>Paenibacillus sacheonensis</i>	100.0
OTU.1724	82	No hits of at least 90% identity	84.1
OTU.1726	82	No hits of at least 90% identity	88.77
OTU.1727	12	<i>Salinirepens amamiensis</i>	94.35
OTU.1737	82	<i>Ferrimicrobium acidiphilum</i>	91.76
OTU.1741	82	<i>Cytophaga hutchinsonii ATCC 33406</i>	99.19

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.1743	12, 82	<i>Clostridium saccharobutylicum</i>	96.25
OTU.1786	82	No hits of at least 90% identity	84.57
OTU.1796	12	<i>Bdellovibrio bacteriovorus</i>	93.32
OTU.1818	82	<i>Roseivirga spongicola</i>	91.35
OTU.1824	82	<i>Solibius ginsengiterrae</i>	91.85
OTU.1832	12, 82	<i>Phyllobacterium bourgognense</i> , <i>Phyllobacterium brassicacearum</i> , <i>Phyllobacterium sp. PEPV15</i> , <i>Phyllobacterium trifolii</i> , <i>Phyllobacterium myrsinacearum</i>	99.73
OTU.1836	82	No hits of at least 90% identity	89.12
OTU.1838	82	<i>Singulisphaera rosea</i>	90.43
OTU.1840	82	No hits of at least 90% identity	88.68
OTU.1842	82	<i>Adhaeribacter aquaticus</i>	90.76
OTU.1857	82	<i>Roseivirga spongicola</i>	93.48
OTU.1873	82	No hits of at least 90% identity	80.06
OTU.1877	12	<i>Stenotrophomonas terrae</i>	100.0
OTU.1892	82	<i>Halioglobus pacificus</i>	91.2
OTU.1897	82	<i>Fluviicola taffensis</i>	93.82
OTU.1919	82	No hits of at least 90% identity	89.13
OTU.1929	82	<i>Chitinophaga niastensis</i> , <i>Chitinophaga niabensis</i>	96.76
OTU.1945	82	<i>Ohtaekwangia koreensis</i>	90.59
OTU.1951	82	<i>Prostheco bacter de j on ge ii</i> , <i>Prostheco bacter de b on t ii</i>	96.52
OTU.1952	82	No hits of at least 90% identity	87.47
OTU.1956	82	No hits of at least 90% identity	88.8
OTU.1958	82	<i>Legionella pneumophila subsp. pneumophila str. Philadelphia 1</i> , <i>Legionella anisa</i> , <i>Legionella pneumophila subsp. pascullei</i>	98.92
OTU.1974	82	<i>Solitalea canadensis</i>	100.0
OTU.1975	12	<i>Flavobacterium sp. THG 01</i>	94.61
OTU.1982	82	<i>Bryobacter aggregatus</i>	93.28
OTU.1986	12, 82	<i>Isosphaera-like str. OJF2</i>	93.9
OTU.1992	12, 82	<i>Streptomyces viridosporus</i>	99.2
OTU.1995	12	<i>Pseudoduganella violaceinigra</i>	99.46
OTU.1999	12, 82	<i>Aurantimonas sp. L9-753</i>	99.2
OTU.2018	12, 82	<i>Cellvibrio gandavensis</i>	98.12

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.2026	12	No hits of at least 90% identity	82.31
OTU.2028	82	No hits of at least 90% identity	89.49
OTU.2067	82	<i>Sphingobacteria bacterium RYG</i>	92.45
OTU.2069	82	<i>Ferruginibacter alkalilentus</i>	94.91
OTU.2094	12, 82	No hits of at least 90% identity	88.86
OTU.2098	12	<i>Paenibacillus larvae subsp. larvae</i>	94.86
OTU.2100	12	<i>Sphingobacterium composti Ten et al. 2007</i>	94.07
OTU.2101	82	No hits of at least 90% identity	89.84
OTU.2132	82	<i>Ferruginibacter lapsinanis</i>	96.5
OTU.2138	12, 82	<i>Fluviicola taffensis</i>	98.92
OTU.2144	82	<i>Asticcacaulis benevestitus</i>	98.92
OTU.2148	12, 82	<i>Solibius ginsengiterrae</i>	91.6
OTU.2152	12	No hits of at least 90% identity	88.47
OTU.2153	82	No hits of at least 90% identity	77.21
OTU.2157	12, 82	<i>Sphingobium rhizovicinum</i>	100.0
OTU.2166	82	No hits of at least 90% identity	86.02
OTU.2200	82	No hits of at least 90% identity	89.84
OTU.2208	12, 82	<i>Mucilaginibacter sp. ANJLI2</i>	99.46
OTU.2213	82	<i>Singulisphaera rosea</i>	92.0
OTU.2216	12, 82	<i>Chitinophaga niabensis</i>	91.01
OTU.2220	82	No hits of at least 90% identity	89.54
OTU.2222	82	<i>Gemmata obscuriglobus</i>	94.39
OTU.2239	82	<i>Adhaeribacter aerophilus</i>	93.19
OTU.2276	82	<i>Singulisphaera rosea</i>	93.87
OTU.2277	12, 82	<i>Camelimonas lactis</i>	100.0
OTU.2279	82	No hits of at least 90% identity	79.2
OTU.2281	82	<i>Byssovorax cruenta</i>	94.12
OTU.2282	82	<i>Caedibacter caryophilus</i>	92.18
OTU.2287	12	No hits of at least 90% identity	88.0
OTU.2288	82	<i>Phenylobacterium lituiforme</i>	96.24
OTU.2294	82	<i>Dongia mobilis</i>	97.86
OTU.2301	82	<i>Pseudoxanthomonas mexicana</i>	100.0
OTU.2329	82	<i>Mucilaginibacter paludis</i>	99.19
OTU.2331	82	<i>Lysinimicrobium mangrovi</i> , <i>Demequina aestuarii</i>	98.93

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.2354	82	<i>Methylovorus mays</i> , <i>Methylovorus glucosotrophus</i> , <i>Methylovorus menthalis</i>	98.66
OTU.2381	82	<i>Bryobacter aggregatus</i>	93.28
OTU.2383	82	No hits of at least 90% identity	89.54
OTU.2393	82	<i>Alkanibacter difficilis</i>	93.09
OTU.2405	82	No hits of at least 90% identity	85.11
OTU.2411	12, 82	<i>Dyadobacter sp. BZ26</i>	100.0
OTU.2420	82	No hits of at least 90% identity	89.7
OTU.2446	82	No hits of at least 90% identity	85.95
OTU.2450	82	No hits of at least 90% identity	86.9
OTU.2463	82	<i>Roseomonas sp. enrichment culture clone 03SU 06-B1</i> <i>Roseomonas stagni</i>	96.51
OTU.2478	82	<i>Prostheco bacter fluviatilis</i>	99.47
OTU.2482	12	<i>Paenibacillus castaneae</i>	99.46
OTU.2489	82	No hits of at least 90% identity	85.07
OTU.2497	82	No hits of at least 90% identity	84.76
OTU.2519	82	<i>Anaeromyxobacter dehalogenans</i>	90.19
OTU.2524	12	<i>Acinetobacter johnsonii</i>	100.0
OTU.2526	82	<i>Terrimonas sp. M-8</i> , <i>Terrimonas sp. RIB1-6</i>	97.3
OTU.2533	82	<i>Legionella sp. LegA</i>	97.59
OTU.2539	82	<i>Planctomyces maris</i>	90.32
OTU.2541	82	No hits of at least 90% identity	88.27
OTU.2548	82	No hits of at least 90% identity	86.84
OTU.2553	82	No hits of at least 90% identity	88.0
OTU.2565	82	<i>Caedibacter caryophilus</i>	91.11
OTU.2576	82	No hits of at least 90% identity	88.27
OTU.2584	82	<i>Flaviumibacter petaseus</i>	100.0
OTU.2590	82	No hits of at least 90% identity	83.52
OTU.2594	82	<i>Vampirovibrio chlorellavorus</i>	93.78
OTU.2611	82	No hits of at least 90% identity	85.91
OTU.2619	82	<i>Isosphaera-like str. OJF2</i>	92.59
OTU.2622	82	No hits of at least 90% identity	79.51
OTU.2623	12, 82	<i>Sphingomonas insulae</i>	97.86
OTU.2624	82	No hits of at least 90% identity	87.16
OTU.2654	82	<i>Kaistia sp. 5YN7-3</i> , <i>Kaistia sp. B6-12</i>	96.77

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.2656	82	No hits of at least 90% identity	89.04
OTU.2658	82	<i>Thiohalobacter thiocyanaticus</i>	92.78
OTU.2660	82	No hits of at least 90% identity	89.59
OTU.2669	82	<i>Singulisphaera acidiphila</i>	93.33
OTU.2671	82	No hits of at least 90% identity	82.62
OTU.2675	82	<i>Sphingomonas trueperi</i> , <i>Sphingomonas pituitosa</i> , <i>Sphingomonas sp. C16y</i>	96.26
OTU.2701	12, 82	<i>Paenibacillus agaridevorans</i>	98.93
OTU.2708	82	No hits of at least 90% identity	87.8
OTU.2714	12, 82	<i>Flavobacterium pectinovorum</i> , <i>Flavobacterium chungnamense</i>	97.27
OTU.2737	12, 82	<i>Agromyces sp. NIO-1018</i> , <i>Agromyces sp. MJ21</i>	100.0
OTU.2738	12	<i>Pseudomonas rhizosphaerae</i> , <i>Pseudomonas abietaniphila</i>	100.0
OTU.2739	82	<i>Zavarzinella formosa</i>	92.49
OTU.2741	82	No hits of at least 90% identity	79.84
OTU.2759	82	No hits of at least 90% identity	83.73
OTU.2769	82	No hits of at least 90% identity	84.88
OTU.2770	82	No hits of at least 90% identity	86.06
OTU.2778	82	No hits of at least 90% identity	88.8
OTU.2780	82	<i>Fimbriimonas ginsengisoli Gsoil 348</i>	96.27
OTU.2788	82	<i>Singulisphaera rosea</i>	94.13
OTU.2791	82	No hits of at least 90% identity	87.5
OTU.2794	82	<i>Ohtaekwangia koreensis</i>	95.41
OTU.2821	82	No hits of at least 90% identity	88.95
OTU.2828	82	<i>Sphingopyxis taejonensis</i>	100.0
OTU.2833	82	<i>Rhizomicrobium palustre</i> , <i>Rhizobiales bacterium Mfc52</i>	91.69
OTU.2865	82	<i>Rhizomicrobium palustre</i>	93.01
OTU.2885	12	<i>Chitinophaga sancti</i>	94.28
OTU.2895	82	<i>Verrucomicrobiaceae bacterium DC2a-G7</i>	99.46
OTU.2926	82	No hits of at least 90% identity	88.62
OTU.2932	82	<i>Sediminibacterium salmoneum</i>	91.6
OTU.2943	82	<i>Afpia massiliensis</i>	97.58
OTU.2948	12, 82	<i>Variovorax paradoxus</i> , <i>Xenophilus aerolatus</i>	98.92
OTU.2968	82	<i>Opitutus terrae</i>	92.78

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.2978	82	No hits of at least 90% identity	86.15
OTU.2996	12, 82	<i>Luteolibacter</i> sp. CCTCC AB 2010415	97.06
OTU.3033	82	<i>Nannocystis pusilla</i>	100.0
OTU.3040	12	<i>Enterococcus saccharolyticus</i> , <i>Enterococcus casseliflavus</i> , <i>Enterococcus gallinarum</i>	100.0
OTU.3050	82	<i>Sulfuricella denitrificans</i>	92.72
OTU.3061	12, 82	<i>Pigmentiphaga litoralis</i>	100.0
OTU.3068	12, 82	<i>Promicromonospora thailandica</i>	99.47
OTU.3071	82	No hits of at least 90% identity	88.5
OTU.3072	82	<i>Chitinophaga niabensis</i>	90.49
OTU.3081	82	<i>Hymenobacter</i> sp. OR362-8	97.03
OTU.3082	12	<i>Kineococcus radiotolerans</i> , <i>Kineococcus</i> sp. RP-B16	100.0
OTU.3094	12, 82	<i>Azospirillum melinis</i>	99.73
OTU.3105	82	<i>Terrimonas lutea</i>	94.86
OTU.3106	12, 82	<i>Mucilaginibacter</i> sp. BDR-9	98.92
OTU.3116	82	<i>Chelatococcus daeguensis</i> , <i>Rhizobium</i> sp. HT4	94.35
OTU.3119	82	No hits of at least 90% identity	87.47
OTU.3121	82	No hits of at least 90% identity	86.12
OTU.3133	12	<i>Brevibacillus ginsengisoli</i>	100.0
OTU.3135	12, 82	<i>Bdellovibrio bacteriovorus</i>	93.6
OTU.3147	82	<i>Legionella shakespearei</i>	95.72
OTU.3148	12	<i>Epilithonimonas lactis</i>	98.37
OTU.3149	82	No hits of at least 90% identity	81.96
OTU.3160	82	No hits of at least 90% identity	89.87
OTU.3208	12	<i>Novosphingobium</i> sp. LL02, <i>Novosphingobium hassiacum</i> , <i>Novosphingobium aromaticivorans</i> DSM 12444	97.59
OTU.3226	12	<i>Sphingobacterium bambusae</i>	98.92
OTU.3230	82	<i>Isosphaera</i> -like str. OJF2	92.59
OTU.3255	82	<i>Verrucomicrobium spinosum</i>	97.86
OTU.3274	82	No hits of at least 90% identity	81.12
OTU.3279	82	<i>Phaselicystis flava</i>	90.11
OTU.3282	82	<i>Sediminibacterium salmoneum</i>	95.14
OTU.3303	82	<i>Luteolibacter</i> sp. CCTCC AB 2010415	91.76
OTU.3313	82	<i>Methylococcus capsulatus</i>	90.13

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.3315	12	<i>Sphingobacterium kitahiroshimense</i> , <i>Sphingobacterium anhuiense</i> , <i>Sphingobacterium faecium</i>	100.0
OTU.3343	82	<i>Reyranella massiliensis</i>	96.25
OTU.3345	82	No hits of at least 90% identity	88.14
OTU.3368	82	<i>Aeromicrobium ponti</i>	98.66
OTU.3373	12, 82	<i>Skermanella aerolata</i>	99.2
OTU.3415	82	No hits of at least 90% identity	89.63
OTU.3416	12	<i>Paenibacillus vulneris</i>	96.78
OTU.3425	82	No hits of at least 90% identity	89.52
OTU.3432	12	<i>Duganella sp. T54</i> , <i>Duganella zoogloeoides</i>	98.66
OTU.3454	82	<i>Legionella quinlivanii</i>	96.52
OTU.3469	82	<i>Legionella shakespearei</i>	97.07
OTU.3479	12	<i>Chryseobacterium anthropi</i>	98.38
OTU.3481	82	<i>Frankia sp. S9-650</i>	95.2
OTU.3509	12	<i>Massilia timonae</i>	98.39
OTU.3512	82	<i>Rickettsia sibirica</i> , <i>Rickettsia rickettsii</i> , <i>Rickettsia akari</i> (agent of rickettsialpox)	90.32
OTU.3515	82	<i>Dyadobacter ginsengisoli</i>	96.76
OTU.3522	12	<i>Pantoea eucalypti</i> , <i>Pantoea wallisii</i> , <i>Pantoea vagans</i> , <i>Pantoea agglomerans</i>	97.86
OTU.3540	12	No hits of at least 90% identity	88.77
OTU.3553	12	<i>Acidovorax sp. NF1078</i>	98.36
OTU.3558	82	<i>Paucibacter toxinivorans</i>	98.39
OTU.3562	82	<i>Solibius ginsengiterrae</i>	92.74
OTU.3573	82	No hits of at least 90% identity	87.7
OTU.3579	82	<i>Terrimonas sp. M-8</i>	97.57
OTU.3601	82	<i>Gluconacetobacter johannae</i> , <i>Gluconacetobacter azotocaptans</i>	95.17
OTU.3606	82	<i>Ilumatobacter fluminis</i>	93.35
OTU.3630	12	<i>Saccharibacillus kuerlensis</i>	100.0
OTU.3666	82	<i>Legionella cincinnatiensis</i> , <i>Legionella longbeachae</i>	96.52
OTU.3676	12, 82	<i>Chitinophaga ginsengisegetis</i>	100.0
OTU.3723	12	<i>Flavobacterium sp. DS-12</i>	97.03
OTU.3745	82	No hits of at least 90% identity	84.31
OTU.3748	82	No hits of at least 90% identity	89.54
OTU.3776	12	<i>Chryseobacterium ureilyticum</i>	98.92

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.3844	82	<i>Luteolibacter</i> sp. CCTCC AB 2010415	96.27
OTU.3845	82	No hits of at least 90% identity	87.19
OTU.3852	82	<i>Arcticibacter svalbardensis</i> MN12-7	90.03
OTU.3865	82	No hits of at least 90% identity	85.09
OTU.3873	82	<i>Chondromyces robustus</i>	92.02
OTU.3882	82	<i>Legionella pneumophila</i> subsp. <i>fraseri</i>	94.12
OTU.3891	82	<i>Ferrovibrio denitrificans</i>	91.44
OTU.3898	82	<i>Legionella cincinnatiensis</i>	96.53
OTU.3910	12, 82	<i>Flavobacterium caeni</i> , <i>Flavobacterium</i> sp. R-HLS-17	95.08
OTU.3932	12, 82	<i>Herbaspirillum rhizosphaerae</i> , <i>Herbaspirillum hiltneri</i> , <i>Herbaspirillum autotrophicum</i>	97.58
OTU.3954	12, 82	<i>Paucimonas lemoignei</i>	97.58
OTU.3986	82	<i>Hyalangium minutum</i> , <i>Cystobacter gracilis</i>	97.86
OTU.3989	12	<i>Labrys methylaminiphilus</i>	99.73
OTU.4052	82	No hits of at least 90% identity	81.43
OTU.4067	82	<i>Phenylobacterium lituiforme</i>	97.55
OTU.4074	82	<i>Cytophaga hutchinsonii</i> ATCC 33406	91.64
OTU.4124	82	<i>Geminicoccus roseus</i>	97.31
OTU.4127	12	<i>Devosia subaequoris</i>	97.31
OTU.4147	12	<i>Ornithinibacter aureus</i>	99.2
OTU.4151	12, 82	<i>Singulisphaera rosea</i>	94.13
OTU.4166	82	<i>Leucobacter</i> sp. NAL101, <i>Leucobacter tardus</i> , <i>Leucobacter alluvii</i> , <i>Leucobacter chromiirensistens</i> JG 31	98.67
OTU.4167	12	<i>Paenibacillus pectinilyticus</i>	95.69
OTU.4168	82	No hits of at least 90% identity	86.35
OTU.4182	82	<i>Cystobacter armeniacae</i> , <i>Cystobacter miniatus</i>	95.14
OTU.4188	82	<i>Sphingomonas japonica</i>	95.98
OTU.4216	82	<i>Mesorhizobium albiziae</i>	98.39
OTU.4222	82	No hits of at least 90% identity	87.81
OTU.4234	82	<i>Ohtaekwangia koreensis</i>	93.51
OTU.4238	82	No hits of at least 90% identity	87.11
OTU.4249	82	<i>Cytophaga hutchinsonii</i> ATCC 33406, <i>Adhaeribacter aquaticus</i>	91.08
OTU.4271	82	<i>Afpia felis</i> (cat scratch disease bacillus), <i>Oligotropha carboxidovorans</i>	97.33

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.4275	82	<i>Hydrotalea flava</i>	91.89
OTU.4393	82	No hits of at least 90% identity	86.36
OTU.4403	82	No hits of at least 90% identity	88.24
OTU.4426	82	No hits of at least 90% identity	86.52
OTU.4435	82	<i>Sphingomonas jaspisi</i> , <i>Kaistobacter terrae</i>	96.0
OTU.4440	82	No hits of at least 90% identity	88.86
OTU.4442	12	<i>Dyadobacter fermentans</i>	97.3
OTU.4485	82	<i>Hydrotalea flava</i>	95.37
OTU.4498	82	<i>Legionella drancourtii</i> , <i>Legionella lytica</i> , <i>Legionella sp.</i>	97.87
OTU.4502	82	<i>Caulobacter fusiformis</i>	94.09
OTU.4520	82	<i>Nubsella zeaxanthinifaciens</i> , <i>Pedobacter rhizosphaerae</i>	96.76
OTU.4524	12, 82	<i>Sphingomonas asaccharolytica</i> , <i>Sphingomonas sp. S8-3</i>	98.39
OTU.4576	82	<i>Luteolibacter sp. E100</i>	95.72
OTU.4588	82	No hits of at least 90% identity	85.64
OTU.4593	82	<i>Devosia albogilva</i>	94.92
OTU.4599	12	<i>Clostridium argentinense</i>	94.39
OTU.4620	12, 82	<i>Chitinophaga arvensicola</i>	93.53
OTU.4652	82	No hits of at least 90% identity	88.58
OTU.4688	82	<i>Sphingomonas japonica</i>	95.98
OTU.4703	12, 82	<i>Sphingobium rhizovicinum</i>	96.78
OTU.4769	12, 82	No hits of at least 90% identity	89.04
OTU.4776	12	<i>Lentzea waywayandensis</i> , <i>Lentzea flaviverrucosa</i>	99.73
OTU.4782	12	<i>Paenibacillus hodogayensis</i>	97.86
OTU.4783	82	<i>Rhodoblastus acidophilus</i> , <i>Rhodoblastus sphagnicola</i>	95.97
OTU.4804	82	No hits of at least 90% identity	87.06
OTU.4855	82	<i>Acidisphaera rubrifaciens</i> , <i>Roseomonas vinacea</i>	95.17
OTU.4883	82	<i>Adhaeribacter aerolatus</i>	92.08
OTU.4905	82	<i>Ohtaekwangia kribbensis</i>	94.07
OTU.5024	82	<i>Pirellula staleyi</i> DSM 6068	90.03
OTU.5046	82	No hits of at least 90% identity	87.9
OTU.5050	82	<i>Cupriavidus sp. CPDB6</i>	94.93
OTU.5056	12, 82	<i>Pedobacter koreensis</i>	97.57

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.5095	82	<i>Sphingobacteria bacterium RYG</i>	94.61
OTU.5124	12	<i>Chryseobacterium daeguense</i>	98.65
OTU.5172	82	<i>Porticoccus litoralis</i>	91.71
OTU.5180	82	<i>Phenylobacterium lituiforme</i>	98.12
OTU.5188	82	No hits of at least 90% identity	89.39
OTU.5235	82	<i>Phaselicystis flava</i>	90.79
OTU.5278	12, 82	<i>Paenibacillus lautus</i>	97.57
OTU.5289	82	<i>Singulisphaera rosea</i>	95.73
OTU.5296	82	<i>Verrucomicrobium spinosum</i>	90.19
OTU.5298	82	No hits of at least 90% identity	84.97
OTU.5310	82	<i>Thiobacillus aquaesulis</i>	91.71
OTU.5333	82	<i>Kaistia sp. B1-1</i>	99.73
OTU.5441	12, 82	<i>Rhodopseudomonas sp. R-45977</i>	98.66
OTU.5476	82	<i>Rhizomicrobium palustre</i>	92.39
OTU.5485	82	<i>Singulisphaera rosea</i>	91.73
OTU.5650	12	<i>Deinococcus gobiensis</i>	98.65
OTU.5660	82	No hits of at least 90% identity	87.6
OTU.5677	12	<i>Burkholderia hospita</i>	98.39
OTU.5705	82	<i>Chitinophaga ginsengisegetis</i>	90.22
OTU.5727	12, 82	<i>Paenibacillus rigui</i> , <i>Paenibacillus vulneris</i>	97.59
OTU.5758	82	<i>Steroidobacter denitrificans</i> , <i>Nitrosococcus oceani ATCC 19707</i>	90.11
OTU.5798	82	<i>Sphingomonas sp. YC6722</i>	99.73
OTU.5801	82	<i>Sphingobacteria bacterium RYG</i> , <i>Ohtaekwangia koreensis</i>	90.57
OTU.5812	82	<i>Mesorhizobium chacoense</i>	98.12
OTU.5815	12	<i>Singulisphaera rosea</i>	93.6
OTU.5818	82	<i>Schlesneria paludicola</i>	97.85
OTU.5861	82	No hits of at least 90% identity	83.56
OTU.5881	12	<i>Microvirga guangxiensis</i>	97.55
OTU.5887	12	<i>Geodermatophilus sp. YIM 75980</i>	96.0
OTU.5906	12	<i>Paenibacillus rhizosphaerae</i> , <i>Paenibacillus favisporus</i> , <i>Paenibacillus illinoisensis</i> , <i>Paenibacillus cineris</i> , <i>Paenibacillus cellulositrophicus</i>	95.41
OTU.5920	82	<i>Legionella pneumophila subsp. pneumophila str. Philadelphia 1</i> , <i>Legionella pneumophila subsp. pasculei</i>	97.83

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.5922	12	<i>Hymenobacter sp. DCY57</i>	98.38
OTU.5942	82	<i>Actinoplanes liguriensis</i>	99.73
OTU.5951	82	<i>Chitinophaga arvensicola</i>	92.7
OTU.6004	82	<i>Devosia limi</i>	98.92
OTU.6008	12	<i>Delftia tsuruhatensis</i> , <i>Delftia lacustris</i>	98.12
OTU.6028	82	No hits of at least 90% identity	87.13
OTU.6097	82	<i>Chitinophaga niabensis</i>	97.03
OTU.6141	12	<i>Hymenobacter elongatus</i>	98.11
OTU.6176	82	<i>Erythrobacter gaetbuli</i>	97.86
OTU.6180	12, 82	<i>Duganella sp. Sac-41</i>	97.85
OTU.6190	12	<i>Oxalicibacterium horti</i>	99.18
OTU.6238	82	<i>Ilumatobacter fluminis</i>	93.07
OTU.6263	82	<i>Isosphaera-like str. OJF2</i>	93.12
OTU.6279	82	<i>Roseomonas sp. enrichment culture clone 03SU95-71</i>	95.71
OTU.6328	12, 82	<i>Pseudomonas brassicacearum</i> subsp. <i>neoaurantiaca</i> , <i>Pseudomonas frederiksbergensis</i>	99.73
OTU.6339	82	<i>Mesorhizobium amorphae</i> , <i>Mesorhizobium opportunistum</i> WSM2075, <i>Mesorhizobium silamurunense</i>	97.55
OTU.6391	82	No hits of at least 90% identity	88.95
OTU.6395	82	<i>Nocardioides islandensis</i>	98.1
OTU.6410	82	<i>Adhaeribacter terreus</i>	90.44
OTU.6427	82	<i>Singulisphaera rosea</i>	93.73
OTU.6431	12	<i>Massilia sp. D5</i>	97.31
OTU.6436	12	<i>Microclunatus ginsengisoli</i>	99.46
OTU.6446	12, 82	<i>Diaphorobacter nitroreducens</i> , <i>Comamonas terrigena</i> , <i>Acidovorax caeni</i>	97.58
OTU.6477	12	<i>Edaphobacter aggregans</i>	97.59
OTU.6478	82	<i>Schlesneria paludicola</i>	98.01
OTU.6487	12	<i>Glaciimonas sp. A2-57</i> , <i>Glaciimonas immobilis</i> , <i>Oxalicibacterium faecigallinarum</i> , <i>Oxalicibacterium horti</i>	97.31
OTU.6495	12, 82	<i>Rhizobiales bacterium</i> WSM3557	98.92
OTU.6508	82	<i>Niastella populi</i>	96.49
OTU.6510	12, 82	<i>Rhizobium mesosinicum</i> , <i>Rhizobium alamii</i> , <i>Arthrobacter viscosus</i>	98.37
OTU.6514	82	No hits of at least 90% identity	88.92

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.6542	82	No hits of at least 90% identity	88.12
OTU.6585	12, 82	<i>Streptomyces</i> spp.	98.38
OTU.6596	82	No hits of at least 90% identity	83.11
OTU.6599	12	<i>Methylophilus methylotrophus</i> , <i>Methylothera versatilis</i> 301, <i>Methylothera mobilis</i>	98.13
OTU.6633	12, 82	<i>Burkholderia zhejiangensis</i>	97.86
OTU.6640	12, 82	<i>Adhaeribacter terreus</i>	98.05
OTU.6704	12	<i>Chryseobacterium</i> sp. THG 15, <i>Chryseobacterium formosense</i>	98.37
OTU.6735	82	<i>Brevundimonas staleyii</i>	98.12
OTU.6749	82	<i>Sediminibacterium salmoneum</i>	95.95
OTU.6751	12	<i>Kaistobacter</i> sp. Gsoil 634	99.17
OTU.6764	12, 82	<i>Paenibacillus aestuarii</i>	96.51
OTU.6801	12	<i>Bacillus bataviensis</i> , <i>Bacillus novalis</i> , <i>Bacillus drementensis</i> , <i>Bacillus vireti</i> , <i>Bhargavaea ginsengi</i> , <i>Bacillus soli</i>	97.33
OTU.6822	12	<i>Sphingomonas changbaiensis</i>	99.16
OTU.6826	12	<i>Telluria mixta</i>	97.58
OTU.6838	12	No hits of at least 90% identity	82.56
OTU.6848	12, 82	<i>Rhizobium skienewicense</i>	98.12
OTU.6852	12	<i>Chryseobacterium jejuense</i>	96.45
OTU.6868	12, 82	<i>Caulobacter vibrioides</i> , <i>Caulobacter segnis</i>	98.66
OTU.6876	82	<i>Thiohalophilus thiocyanatolydans</i>	92.18
OTU.6898	82	<i>Arenimonas malthae</i>	96.77
OTU.6900	12	<i>Alicyclobacillus disulfidooxidans</i> , <i>Alicyclobacillus contaminans</i>	94.89
OTU.6947	12	<i>Gaiella occulta</i>	92.29
OTU.6979	12	<i>Rhizobium larrymoorei</i>	97.28
OTU.6993	82	No hits of at least 90% identity	82.8
OTU.7027	82	No hits of at least 90% identity	86.91
OTU.7041	12, 82	<i>Altererythrobacter</i> sp. H32, <i>Altererythrobacter</i> sp. MSW-14	94.09
OTU.7064	12, 82	<i>Stenotrophomonas pavanii</i> , <i>Stenotrophomonas maltophilia</i>	97.3
OTU.7074	12	No hits of at least 90% identity	89.67
OTU.7077	12	<i>Streptomyces mutomycini</i> , <i>Streptomyces atroolivaceus</i> , <i>Streptomyces finlayi</i> , <i>Streptomyces clavifer</i>	97.86

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.7083	82	No hits of at least 90% identity	87.06
OTU.7092	12	<i>Novosphingobium hassiacum</i>	97.58
OTU.7106	82	<i>Devosia neptuniae</i>	99.19
OTU.7112	12	<i>Yokenella</i> spp., <i>Pantoea</i> spp., <i>Kluyvera</i> spp., <i>Klebsiella</i> spp., <i>Erwinia</i> spp., <i>Enterobacter</i> spp., <i>Citrobacter</i> spp., <i>Cedecea</i> spp.	94.39
OTU.7117	82	No hits of at least 90% identity	88.53
OTU.7143	12	<i>Thermoleophilum minutum</i> , <i>Pseudomonas thivervalensis</i> , <i>Pseudomonas lini</i>	98.38
OTU.7167	12	<i>Sphingomonas</i> sp. BH3	95.69
OTU.7208	12	<i>Nubsella zeaxanthinifaciens</i>	98.33
OTU.7299	12	<i>Dyella</i> sp. ATSB10, <i>Dyella ginsengisoli</i>	98.38
OTU.7309	82	<i>Niastella</i> sp. JCN-23	95.1
OTU.7313	12	<i>Rhodococcus qingshengii</i> , <i>Rhodococcus erythropolis</i> , <i>Rhodococcus</i> sp. djl-6-2, <i>Nocardia coeliaca</i>	98.38
OTU.7319	82	<i>Chitinophaga niabensis</i>	95.41
OTU.7377	82	No hits of at least 90% identity	83.82
OTU.7431	82	<i>Luteimonas marina</i> , <i>Thermomonas dokdonensis</i> , <i>Luteimonas lutimaris</i>	97.33
OTU.7453	12	No hits of at least 90% identity	83.6
OTU.7454	12, 82	<i>Pedobacter boryungensis</i>	99.19
OTU.7457	82	<i>Reyranella massiliensis</i>	95.41
OTU.7461	82	<i>Sphingomonas fennica</i>	98.39
OTU.7463	82	<i>Hydrotalea flava</i> , <i>Chitinophaga arvensicola</i>	92.33
OTU.7538	82	<i>Aquabacterium parvum</i> , <i>Aquabacterium commune</i>	97.29
OTU.7609	82	<i>Niastella</i> sp. JCN-23, <i>Niastella populi</i>	95.68
OTU.7610	12, 82	<i>Schlesneria paludicola</i>	93.3
OTU.7616	12, 82	<i>Sphingomonas yunnanensis</i>	97.86
OTU.7631	82	<i>Rhizomicrobium palustre</i>	92.2
OTU.7646	12, 82	<i>Novosphingobium hassiacum</i>	97.32
OTU.7650	82	No hits of at least 90% identity	86.4
OTU.7652	82	<i>Porphyrobacter tepidarius</i>	96.51
OTU.7664	82	<i>Sphingobacteria bacterium RYG</i>	95.42
OTU.7701	12	<i>Rhodoplanes elegans</i>	96.19

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
OTU.7731	82	<i>Amnibacterium kyonggiense</i>	98.13
OTU.7735	12	<i>Paenibacillus</i> sp. WPCB018	99.46
OTU.7764	82	<i>Sporichthya polymorpha</i>	95.73
OTU.7779	12, 82	<i>Luteimonas</i> sp. KMM 9005	98.07
OTU.7786	12, 82	<i>Stenotrophomonas rhizophila</i>	98.65
OTU.7839	82	<i>Opitutus terrae</i>	91.64
OTU.7922	82	No hits of at least 90% identity	88.56
OTU.7943	12, 82	<i>Pseudomonas pseudoalcaligenes</i> , <i>Pseudomonas oleovorans</i> subsp. <i>lubricantis</i> , <i>Pseudomonas alcaliphila</i> , <i>Pseudomonas composti</i> , <i>Pseudomonas toytomiensis</i>	98.93
OTU.7950	12	<i>Massilia namucuoensis</i>	98.1
OTU.7978	12	<i>Methylobacterium goesingense</i> , <i>Methylobacterium marchantiae</i> , <i>Methylobacterium</i> sp. F3.2	98.91
OTU.7995	12, 82	<i>Roseomonas aquatica</i> , <i>Belnapia moabensis</i>	96.25
OTU.8028	12	No hits of at least 90% identity	88.77
OTU.8077	12, 82	<i>Adhaeribacter terreus</i>	97.03
OTU.8085	12	<i>Jeotgalibacillus</i> sp. WS 4628	98.13
OTU.8117	12	<i>Luteimonas lutimaris</i>	96.53
OTU.8228	82	<i>Sphingobacteria bacterium RYG</i>	92.2
OTU.8238	82	<i>Sphingobacteria bacterium RYG</i>	91.94
OTU.8259	82	<i>Rhodoplanes roseus</i>	96.2
OTU.8266	12, 82	<i>Arthrobacter koreensis</i> , <i>Arthrobacter citreus</i> , <i>Arthrobacter luteolus</i>	93.1
OTU.8270	82	No hits of at least 90% identity	86.68
OTU.8278	12, 82	<i>Isoptricola nanjingensis</i> , <i>Isoptricola hypogeus</i> , <i>Isoptricola variabilis</i>	99.73
OTU.8409	12, 82	<i>Acidovorax temperans</i>	98.91
OTU.8484	82	<i>Streptomyces aomiensis</i>	96.67
OTU.8620	12, 82	No hits of at least 90% identity	88.68
OTU.8633	82	<i>Caulobacter henricii</i> , <i>Brevundimonas</i> -like sp. LMG 11050, <i>Caulobacter</i> sp.	94.62
OTU.8661	12, 82	<i>Mucilaginibacter jinjuensis</i>	98.38
OTU.8678	82	No hits of at least 90% identity	85.03
OTU.8700	82	No hits of at least 90% identity	86.13
OTU.8710	12	<i>Pedobacter koreensis</i>	97.3

Table 2 – continued from previous page

OTU ID	Day of Response ^b	Top BLAST hits	BLAST %ID
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^a \log_2 fold change greater than 1, adjusted P-value less than 0.10.