

Supplementary Table S1. The full list of identified using 16S rRNA metabarcoding pathogens

OTUs Abundance, %	Case 1	Case 2	Case 3	Case 4	Case 5	Case 6	Case 7	Case 8	Case 9	Case 10
OTU_4 (unc_Roseateles)	9.0	0.8	0	2.7	15	6.7	5.8	2.4	15	19
OTU_2 (unc_Sphingomonas)	6.0	0.6	0	2.5	4.0	3.3	5.5	1.7	11	22
OTU_3 (unc_Enterococcus)	0.1	8.3	100	0	0	0	0	1.6E-03	0	22
OTU_6 (Pseudomonas_stutzeri)	27	1.5	1.0E-02	5.5	11	19	19	12	16	2.5E-03
OTU_1 (unc_Bacillus)	0	67	0	0	0	0	0	0	0	0
OTU_5 (unc_Streptococcus)	0	0.2	0	0	1.3	1.3	13	47	0	0
OTU_7 (Ralstonia_pickettii)	0	5.4	0.3	55	0	1.7E-03	0.1	0	0	0.1
OTU_19 (unc_Acinetobacter)	14	0.9	4.8E-03	4.0	6.7	8.4	4.0	4.5	13	5.0E-03
OTU_18 (unc_Pseudomonas)	8.0	1.1	9.5E-03	3.1	4.5	8.0	6.3	3.8	4.7	0.2
OTU_9 (unc_Sediminibacterium)	0	0.2	4.8E-03	0	4.1	0.8	0.9	0.8	0.4	4.8
OTU_8 (Sphingobium_limneticum)	4.1	0.5	0	2.4	1.0	9.6	3.8	2.2	9.7	0
OTU_12 (unc_Agrobacterium)	5.7	0.3	9.5E-03	0.7	0.8	6.4	10	2.3	3.2	5.0E-03
OTU_10 (unc_Reyranella)	1.3	0.1	0	0.7	6.1	0.6	0.1	1.7	1.6	6.8
OTU_11 (Taonella_mepensis)	1.7	0.2	4.8E-03	0.6	3.0	1.5	2.3	1.7	0.5	3.3
OTU_26 (unc_Betaproteobacteria)	0.8	0.1	0	0.3	3.4	0.9	1.0	0.3	2.8	2.1
OTU_16 (unc_Bradyrhizobium)	0.2	0.3	2.0E-02	1.5	1.4	0.4	1.4	0.6	3.2	4.2
OTU_28 (Corynebacterium_kroppenstedtii)	0.6	2.0E-02	4.8E-03	3.8	2.7	0.4	1.1	0.6	2.5	0.3
OTU_21 (unc_Phyllobacterium)	0	1.3	0.1	1.0	0	2.0	9.7	0.6	0.4	1.0E-02
OTU_17 (unc_Methylobacterium)	0.3	0.1	0	0.3	0.3	2.8	0.3	1.8	0.5	0.7
OTU_20 (unc_Brevundimonas)	4.4	0.3	0	1.0	0.9	1.1	2.0	1.7	1.4	0
OTU_15 (unc_Staphylococcus)	0.8	0.3	0	0.5	1.5	2.8	0.7	0.2	2.8	1.3
OTU_23 (unc_Dietzia)	0	0	0	0	1.9	1.2	0.2	0.8	0.1	5.0E-03

OTU_30 (<i>Aquabacterium</i> _parvum)	2.0	0.3	0	0.7	1.8	0.6	1.5	0.7	1.3	0
OTU_25 (unc_Chitinophagaceae)	0	7.1E-03	0	0	1.7	0.1	0.1	0.2	2.3	0
OTU_29 (unc_Acinetobacter)	2.5	0.1	0	0.3	1.6	0.8	0.1	1.3	0	0
OTU_35 (unc_Micrococcus)	0.7	0.1	0	0.6	0.6	0	0.8	0.9	0	0
OTU_36 (unc_Delftia)	0.6	0.1	0	0.6	1.2	2.1	0.3	0.5	0.3	0
OTU_27 (unc_Stenotrophobacter)	0	0	0	0	0	0.1	0	0	0	5.4
OTU_39 (unc_Streptococcus)	0.3	0.2	0	0.0	0.1	0.4	0.1	1.0E-02	0.2	0
OTU_32 (unc_Sphingopyxis)	0.7	0.2	0	0.8	1.6	0	0.5	0.5	0	0
OTU_51 (unc_Priestia)	0	0	0	4.1	2.0E-02	0	0	0	0	0
OTU_40 (unc_Corynebacterium)	0	0.2	0	0	0.7	0.5	0.7	0	0	0
OTU_37 (unc_Paracoccus)	0.4	5.3E-03	0	0.1	0.3	0.0	0	0	0.3	0
OTU_31 (unc_Bosea)	0.1	0.1	0	5.3E-03	0.1	0.1	0	0.1	0	2.9
OTU_33 (unc_Methylobacterium)	1.1	0	0	0	0.8	1.8	0	0	0.1	0
OTU_34 (unc_Pedomicrobium)	0	0	0	0	0	0	0.1	8.1E-03	0.1	1.3
OTU_38 (<i>Lawsonella</i> _clevelandensis)	0	0	0	0.6	0.7	1.7	0	0	0	0
OTU_71 (unc_Blastomonas)	1.4	0.2	4.8E-03	0.2	0.5	0.1	0.1	0.4	0	0
OTU_22 (unc_Enterobacteriaceae)	0	2.8	0	1.0E-02	0	0	0	0.1	0	0
OTU_64 (<i>Sphingobacterium</i> _psychroa quaticum)	0.7	0.1	0	0.3	0	0	1.7	0	0	0
OTU_61 (unc_Sphingomonas)	0	0.2	0	0.4	0.3	0.9	0	0	0.2	0
OTU_42 (unc_Acidibacter)	0.1	0	0	0	1.1	0	0	0.4	1.0E-02	0.4
OTU_41 (unc_Acidovorax)	4.6E-03	0.1	0	0.1	1.1	1.0	0.1	0	0	0
OTU_44 (unc_Brachybacterium)	0	0.2	0	0.2	0	1.8	0	0.1	0	0
OTU_55 (unc_Kocuria)	0	0.2	0	0	0	0.2	0	0	0	1.7
OTU_50 (unc_Pseudomonas)	0	0	0	0	0	1.2	0.2	0.2	0.2	0
OTU_46 (unc_Corynebacterium)	0.1	0	0	0	0	0	0.6	1.1	0	0
OTU_63 (unc_Paraburkholderia)	0.7	0	0	0.1	0.1	0.2	0.3	0.3	0	0
OTU_45 (unc_Nevskiaceae)	0	0	0	0.1	1.2	0	0	1.0E-02	0	0
OTU_74 (<i>Cutibacterium</i> _acnes)	4.6E-03	0	0	1.0E-02	0.1	0.0	0	0	0	0
OTU_60 (unc_Brevibacterium)	0	0	0	0.1	0	0.7	0	0	0	0

OTU_68 (Truepera_radiovictrix)	0	0	0	0	0	0	0	0	1.5	0
OTU_47 (unc_Phenylobacterium)	0	0	0	0	0	0	0.0	0.2	0.6	0.6
OTU_57 (unc_Sphaerobacteraceae)	0.7	1.8E-03	0	0	0	0.6	0	0	0.2	0
OTU_79 (unc_Stenotrophomonas)	0.2	0.3	0	0	0.5	0.2	1.0E-02	0.2	0	0
OTU_53 (unc_Planococcus)	0.1	0	0	0	0	1.3	0	0	0	0
OTU_92 (unc_Brevibacterium)	0	0	0	0.3	0.2	0	0.5	0	0.1	0.4
OTU_69 (unc_Melaminivora)	0	0	0	0	0	0	0.0	0	0.1	0
OTU_78 (unc_Caballeronia)	0	0	0	0	0	0	0	0	0	0
OTU_67 (unc_Neisseriaceae)	0	0	0	0.4	0.7	0	0	0	0	0.2
OTU_109 (unc_Streptomyces)	0	0.0	0	1.2	0	0	0	0	0	0
OTU_48 (unc_Caulobacter)	0.1	0	0	0	1.1	0	0	0	0	0
OTU_227 (unc_Acinetobacter)	0.4	0	0	0	0	2.0E-02	0.5	0.2	0.1	0
OTU_75 (unc_Rubrobacter)	0	3.5E-03	0	0	0.0	0	0.2	0	0	1.0
OTU_76 (unc_Acinetobacter)	0	0	0	0	0.7	0	0	0	0.6	0
OTU_85 (unc_Prevotellaceae)	0	0	0	0	0.7	0	0	0	0	0
OTU_108 (unc_Streptophyta)	0	3.5E-03	0	0	0.1	0	0.9	0	0.2	0
OTU_49 (unc_Brevundimonas)	0	0.2	0	0.4	0.5	0	0	0	0	2.5E-03
OTU_105 (unc_Glutamicibacter)	0.8	0	0	0.3	0	0	0	0	0	0
OTU_119 (Haemophilus_parainfluenzae)	0	0	0	0	0.7	0	0	0	0	0
OTU_43 (unc_Brevundimonas)	0	0	0	0	0	1.0	0	0	0	0
OTU_95 (Anaerococcus_octavius)	0	0	0	2.0E-02	0	0	0	0	0	0
OTU_94 (unc_Cupriavidus)	0	0	0	0	0	0.9	0	0	0	0
OTU_62 (unc_Mammaliicoccus)	0.1	0	0	0	0.8	0	0	0	0	0
OTU_59 (unc_Faecalibacterium)	0	0	0	0	0	0	0	0.8	0.1	0
OTU_54 (Peptoniphilus_duerdenii)	0	0	0	0	0.8	0	0	0	0	0
OTU_99 (unc_Gammaproteobacteria)	0	0	0	0	0	0.8	0	0	0	0
OTU_104 (unc_Thiopseudomonas)	0	0	0	0	0	0	0	0	0	0
OTU_179 (unc_Veillonella)	0.1	0.1	0	0	0	0	0.6	0	0	0
OTU_158 (Pedobacter_nutrimenti)	0	0	0	0	0	0	0.7	0	0	0
OTU_52 (unc_Hyphomicrobiales)	0	0	0	0	0	0.7	0	0	0	0

OTU_96 (<i>Moraxella_osloensis</i>)	0.1	0.0	0	0.1	0.2	0	0.3	0.1	0	0
OTU_90 (<i>unc_Proteiniphilum</i>)	0	0	0	0	0	0	0	0	0	0
OTU_70 (<i>unc_Janibacter</i>)	0	0	0	0	0.4	0	0	0.3	0	0
OTU_81 (<i>unc_Corynebacterium</i>)	0	0	0	0	0	0	0	0	0	0
OTU_244 (<i>Haemophilus_haemolyticus</i>)	0	0	0	0	0.6	0	0	0	0	0
OTU_84 (<i>Granulicatella_elegans</i>)	0	0	0	0	0.6	0	0	0	0	0
OTU_118 (<i>unc_Haemophilus</i>)	0	0	0	0	0.6	0	0	0	0	0
OTU_91 (<i>unc_Barrientosiimonas</i>)	0	0	0	0	0.6	0	0	0	0	0
OTU_97 (<i>unc_Neisseria</i>)	0	1.8E-03	0	0	0.6	0	0	0	0	0
OTU_72 (<i>Anaerococcus_nagya</i>)	0	0	0	0	0.6	0	0	0	0	0
OTU_87 (<i>unc_Neisseriaceae</i>)	0	0	0	0.1	0	0.3	0	0.2	0	0
OTU_262 (<i>Abiotrophia_defectiva</i>)	0.6	0	0	0	0	0	0	0	0	0
OTU_100 (<i>unc_Anaerobacillus</i>)	0	3.5E-03	0	0	0	0.5	0.1	0	0	0
OTU_24 (<i>unc_Escherichia/Shigella</i>)	0	0.6	0	0	0	0	0	0	0	0
OTU_206 (<i>Tepidiphilus_margaritifer</i>)	0	0	0	0	0	0	0.5	0	0	0
OTU_132 (<i>Cloacibacterium_normanense</i>)	0	1.8E-03	0	0	0	0	0	0	0.5	0
OTU_190 (<i>unc_Bacillaceae</i>)	0.5	3.5E-03	0	0	0	0	0	0	0	0
OTU_185 (<i>unc_Parazoarcus</i>)	0	0	0	0	0	0	0	0	0.5	0
OTU_89 (<i>unc_Brevibacillus</i>)	0	0	0	0	0	0	0	0.5	0	0
OTU_98 (<i>unc_Bacillus</i>)	0	0.1	0	0	0.5	0	0	0	0	0
OTU_140 (<i>unc_Microbacterium</i>)	0.1	0	0	0	8.1E-03	0	0	0	2.0E-02	0
OTU_175 (<i>unc_Pseudoxanthomonas</i>)	0	0	0	0	0	0	0.4	0	0.2	0
OTU_82 (<i>unc_Azospirillaceae</i>)	0	0.1	0	0.3	0	0.2	0	0	0	0
OTU_123 (<i>unc_Bacteria</i>)	0.5	0	0	0	0	0	0	0	0	0
OTU_88 (<i>unc_WPS-2_genera_incertae_sedis</i>)	0	0	0	0.1	0	0	0	0	0	0
OTU_130 (<i>unc_Peribacillus</i>)	0	0.2	0	0.3	0	0	0	0	0	0
OTU_151 (<i>unc_Rhizobiaceae</i>)	0	0	0	0	0	0.5	0	0	0	0
OTU_117 (<i>Comamonas_aquatis</i>)	0	0	0	0.4	0.0	0	0	0	0.0	0
OTU_65 (<i>Corynebacterium_otitidis</i>)	0	0.5	0	0	0	0	0	0	0	0

OTU_80 (unc_Gallionellaceae)	0	0	0	0	0	0	0	0.5	0	0
OTU_83 (unc_Jeotgalicoccus)	0	0	0	0	0	0	0	0.5	0	0
OTU_66 (unc_Anaerococcus)	0	0	0	0	0	0	0	0.5	0	0
OTU_13 (Limosilactobacillus_fermentum)	0	0.4	0	0	0	0	0	0	0	0
OTU_115 (unc_Chryseobacterium)	0	0	0	0	0	0	0	0	0	0
OTU_191 (unc_Acinetobacter)	0	0	0	0	0	0	0.4	0	0	0
OTU_93 (Porphyromonas_pasteri)	0	0	0	0	0.4	0	0	0	0	0
OTU_73 (Fusobacterium_necrophorum)	0	0	0	0	0.4	0	0	0	0	0
OTU_201 (Weizmannia_coagulans)	0	0	0	0	0	0	0.4	0	0	0
OTU_106 (unc_Bacillaceae)	0	0	0	0	0	0	0	0	0	0
OTU_146 (Bacteroides_uniformis)	0	0.4	0	0	0	0	0	0	0	0
OTU_102 (Alloiococcus_otitis)	0	0.4	0	0	0	0	0	0	0	0
OTU_164 (Deinococcus_budaensis)	0	0	0	0	0	0	0	0	0.4	0
OTU_178 (unc_Comamonadaceae)	0	0	0	0	4.0E-03	0.0	0	0.0	0.3	0
OTU_139 (unc_Rubellimicrobium)	0	0	0	0	0	0	0	0.3	0	0
OTU_111 (Tissierella_creatinophila)	0	0	0	0.3	0	0	0	0	0	0
OTU_116 (Hoylesella_nanceiensis)	0	0	0	0	0.3	0	0	0	0	0
OTU_122 (unc_Helcococcus)	0	0	0	0	0	0.3	0	0	0	0
OTU_101 (unc_Aliterella)	0	0	0	0	0	0.3	0	0	0	0
OTU_112 (unc_Bacillus)	0	0.1	0	0	0	0	0	0.2	0	0
OTU_110 (unc_Isoptericola)	0	2.0E-02	0	0	0	0	0	0.3	0	0
OTU_184 (Luteimonas_terrae)	0	0	0	0	0	0.1	0	0	0.2	0
OTU_202 (Polynucleobacter_difficilis)	0	0	0	0	0	0	0.3	0	0	0
OTU_86 (Terriglobus_roseus)	0	0	0	0	0	0	0	0.3	0	0
OTU_148 (unc_Coleofasciculales)	0.3	0	0	0	0	0	0	0	0	0
OTU_103 (Ezakiella_coagulans)	0	0	0	0	0.3	0	0	0	0	0
OTU_113 (Mesorhizobium_terrae)	0	0	0	0	0	0	0	0	0	0.3
OTU_168 (unc_Aquabacterium)	0.2	0	0	0	0	0	0.0	0.0	0	0
OTU_136 (Chryseobacterium_hominis)	0	0	0	0.3	0	0	0	0	0	0

OTU_114 (unc_Curvibacter)	0.1	1.8E-03	0	2.0E-02	0.0	0	0	0.2	0	0
OTU_142 (unc_Ornithinimicrobium)	0	0	0	0	0	0	0.0	0.2	0	0
OTU_147 (unc_Herbaspirillum)	0	0	0	0	0	0.3	0	0	0	0
OTU_125 (Finegoldia_magna)	0	0.1	0	0	0.2	0	0	0	0	0
OTU_188 (Flaviflexus_salsibiostraticola)	0	0	0	0	0	0.0	0	0	0.2	0
OTU_134 (unc_Mycobacterium)	0	3.5E-03	0	0	0.0	0.2	0.1	0	0	0
OTU_127 (unc_Comamonas)	0	2.0E-02	0	0	0	0	0	0.2	0	0
OTU_149 (Diaphorobacter_nitroreducens)	0	0	0	0	0.2	0.1	1.0E-02	0	0	0
OTU_166 (unc_Pseudonocardia)	4.6E-03	0	0	0	0	0	0	0	0	0
OTU_135 (unc_Rubellimicrobium)	0	0	0	0	0.2	0	0	0	0	0
OTU_131 (unc_Parvimonas)	0	0	0	0	0.2	0	0	0	0	0
OTU_126 (unc_Actinomycetota)	0	0.2	0	0	0	0	0	0	0	0
OTU_137 (Cytophaga_aurantiaca)	0	0	0	0	0.2	0	0	0	0	0
OTU_195 (unc_Leuconostoc)	0.2	0	0	0	0	0	0	0	0	0
OTU_176 (Rothia_dentocariosa)	0	0	0	0.2	0	0	0	0	0	0
OTU_120 (Marmoricola_aquaticus)	0	0	0	0	0	0	0	0.2	0	0
OTU_177 (unc_Fictibacillus)	0	0	0	0	0	0.2	0	0	0	0
OTU_128 (unc_Anaerolineae)	0	0	0	0	0	0	0	0.2	0	0
OTU_183 (unc_Phenylobacterium)	0	0	0	0	0	0	0	0	0.2	0
OTU_129 (unc_Aureimonas)	0	0	0	0	0.2	0	0	0	0	0
OTU_153 (Sandaracinobacteroides_hominis)	0	0	0	0.1	0	0.0	0	0	0	0
OTU_160 (unc_Legionella)	0	0.1	0	0	0	8.3E-03	0.1	1.0E-02	0	0
OTU_233 (Pseudoneobacillus_rhizospherae)	0	0	0	0	0	0	0	0	0.2	0
OTU_133 (unc_Granulicella)	0	0	0	0	0.2	0	0	0	0	0
OTU_199 (unc_Solirubrobacterales)	0	0	0	0.2	0	0	0	0	0	0
OTU_238 (unc_Paraburkholderia)	0.2	0	0	0	0	0	0	0	0	0
OTU_172 (unc_Streptococcus)	0	0	0	0	0.1	0	0	0	0	0
OTU_269 (Acinetobacter_marinus)	0	0	0	0	0	0	0.1	0	0	0

OTU_154 (unc_Gemella)	0	0	0	0	0.1	0.1	0	0	0	0
OTU_197 (Rufibacter_ruber)	0.1	0	0	0	0	0	0	0	0	0
OTU_138 (unc_Intestinibacter)	0	0.1	0	0	0	0	0	0	0	0
OTU_141 (unc_Lachnospiraceae)	0	0.1	0	0	0	0	0	0	0	0
OTU_145 (unc_Rubrobacter)	0	8.9E-03	0	0	0	0.1	0	0	0	0
OTU_143 (unc_Lactobacillus)	0	0.1	0	0	0	0	0	0	0	0
OTU_171 (unc_Deinococcus)	0	0	0	0.1	0	0	0	0.1	0	0
OTU_229 (unc_Halomonas)	0.1	0	0	0	0	0	0	0	0	0
OTU_162 (unc_Leptolyngbyaceae)	0	0.1	0	0	0	0	0	0	0	0
OTU_174 (unc_Alphaproteobacteria)	0	0	0	0	0.1	0	0	0	0	0
OTU_152 (Porphyromonas_pasteri)	0	0	0	0	0.1	0	0	0	0	0
OTU_159 (Patulibacter_minatonensis)	0	0.1	0	0	0	0	0	0	0	0
OTU_180 (unc_Actinomadura)	0	0	0	0	0.1	0	0	0	0	0
OTU_157 (unc_Corynebacterium)	0	0.1	0	0	0	0	0	0	0	0
OTU_165 (unc_Peptoniphilus)	0	0	0	0	0.1	0	1.0E-02	0	0	0
OTU_194 (unc_Bacteria)	0	0	0	0	0.1	0	0	0	0	0
OTU_228 (unc_Dietzia)	0	0	0	0	0	0	0.1	0	0	0
OTU_212 (Sphingobacterium_daejeonense)	0	0	0	0	0	0	0	0	0.1	0
OTU_182 (unc_Bacteria)	0	0.1	0	0	0	0	0	0	0	0
OTU_268 (unc_Dyadobacter)	0	0	0	0	0	0	0.1	0	0	0
OTU_241 (Cumulibacter_manganitolera ns)	0	0	0	0	0	0	0	0	0	0
OTU_217 (Nakamurella_aerolata)	0	0	0	5.3E-03	0	0	0	0	0	0
OTU_231 (unc_Streptococcus)	0	0	0	0	0.1	0	0	0	0	0
OTU_259 (unc_Conexibacter)	0	0	0	0	0	0	0	0	0.1	0
OTU_14 (unc_Listeria)	0	0.1	0	0	0	0	0	0	0	0
OTU_207 (unc_Acetobacteraceae)	0	0.1	0	0	0	0	0	0	0	0
OTU_193 (unc_Spartobacteria_genera_incertae_sedis)	0	0	0	0	0.1	0	0	0	0	0
OTU_196 (Prevotella_amnii)	0	0	0	0	0	0	0	0	0	0.1
OTU_163 (unc_Aeromonas)	0	0	0	0	0	0	0.1	0	0	0
OTU_167 (unc_Bacteria)	0	0	0	0	0	0	0	0.1	0	0
OTU_169 (Akkermansia_muciniphila)	0	0.1	0	0	0	0	0	0	0	0

OTU_261 (Atopostipes_suicloacalis)	0	0	0	5.3E-03	0	0	0	0	0.1	0
OTU_225 (unc_Sphingopyxis)	0	2.0E-02	0	0	0.0	0	0	0	0	0
OTU_189 (unc_Blastococcus)	1.0E-02	0	0	0	0	0	0	0.0	0	0
OTU_208 (unc_Gemmiger)	0	0.1	0	0	0	0	0	0	0	0
OTU_232 (unc_Nocardioides)	0	0.1	0	0	0	0	0	0	0	0
OTU_187 (unc_Cytophagales)	0	0.1	0	0	0	0	0	0	0	0
OTU_203 (Phocaeicola_plebeius)	0	0.1	0	0	0	0	0	0	0	0
OTU_243 (unc_Bacillaceae)	0	0.1	0	0	0	0	0	0	0	0
OTU_230 (unc_Mobiluncus)	0	0	0	0	0.0	0	0	0	0	0
OTU_219 (unc_Prevotella)	0	0	0	0	0.0	0	0	0	0	0
OTU_216 (unc_Collinsella)	0	0.0	0	0	0	0	0	0	0	0
OTU_242 (unc_Alkalibacterium)	0	0	0	0	0	0.0	0	0	0	0
OTU_223 (unc_Muribaculaceae)	0	0	0	0	0.0	0	0	0	0	0
OTU_209 (Asticcacaulis_excentricus)	0	0	0	0	0.0	0	0	0	0	0
OTU_186 (unc_Rhodococcus)	0	0	0	0	0	0	0	0	5.8E-03	0.0
OTU_220 (unc_Brevibacterium)	0	0	0	0	0	0	0	0.0	0	0
OTU_264 (Zea_mays)	0	0	0	0.0	0	0	0	0	0	0
OTU_234 (unc_Rheinheimera)	0	0	0	0.0	0	0	0	0	0	0
OTU_249 (unc_Gp3)	0	0	0	0	0	0	0	0	2.0E-02	7.5E-03
OTU_237 (unc_Acidobacteriaceae)	0	0	0	0	0.0	0	0	0	0	0
OTU_248 (Sutterella_massiliensis)	0	2.0E-02	0	0	0	0	0	0	0	0
OTU_258 (unc_Actinobacteria)	0	0	0	0	0	0	2.0E-02	0	0	0
OTU_252 (unc_Ruminococcus)	0	2.0E-02	0	0	0	0	0	0	0	0
OTU_257 (Lachnospira_eligens)	0	2.0E-02	0	0	0	0	0	0	0	0
OTU_255 (unc_Bacteria)	0	0	0	0	0	0	0	0	2.0E-02	0
OTU_222 (unc_Porphyromonas)	0	0	0	0	1.0E-02	0	0	0	0	0
OTU_254 (unc_Paenisporosarcina)	0	7.1E-03	0	0	0	0	0	0	0	0