

Table S6. Rank abundance of 1179 taxa.

Genus	Species	HOT	Status	Clones	Rank	Rank Abundance	Relative Abundance
<i>Veillonella</i>	<i>parvula</i>	161	Named	2304	1	0.066296	1.00000
<i>Streptococcus</i>	<i>mitis</i>	677	Named	1968	2	0.056628	0.85417
<i>Streptococcus</i>	<i>mutans</i>	686	Named	1499	3	0.043133	0.65061
<i>Streptococcus</i>	<i>sanguinis</i>	758	Named	851	4	0.024487	0.36936
<i>Veillonella</i>	<i>dispar</i>	160	Named	848	5	0.024401	0.36806
<i>Streptococcus</i>	<i>anginosus</i>	543	Named	561	6	0.016142	0.24349
<i>Gemella</i>	<i>morbilorum</i>	46	Named	544	7	0.015653	0.23611
<i>Gemella</i>	<i>haemolyans</i>	626	Named	536	8	0.015423	0.23264
<i>Granulicatella</i>	<i>adiacens</i>	534	Named	513	9	0.014761	0.22266
<i>Streptococcus</i>	<i>gordonii</i>	622	Named	478	10	0.013754	0.20747
<i>Fusobacterium</i>	<i>nucleatum subsp. vincentii</i>	200	Named	469	11	0.013495	0.20356
<i>Streptococcus</i>	sp.	64	PhyloTYPE	460	12	0.013236	0.19965
<i>Dialister</i>	<i>invisus</i>	118	Named	459	13	0.013207	0.19922

<i>Streptococcus</i>	<i>mitis</i> bv 2	398	Named	410	14	0.011798	0.17795
<i>Streptococcus</i>	<i>parasanguinis II</i>	411	Named	365	15	0.010503	0.15842
<i>Campylobacter</i>	<i>gracilis</i>	623	Named	362	16	0.010416	0.15712
<i>Selenomonas</i>	<i>noxia</i>	130	Named	360	17	0.010359	0.15625
<i>Capnocytophaga</i>	<i>granulosa</i>	325	Named	344	18	0.009898	0.14931
<i>Streptococcus</i>	<i>intermedius</i>	644	Named	334	19	0.009611	0.14497
<i>Streptococcus</i>	<i>cristatus</i>	578	Named	325	20	0.009352	0.14106
<i>Selenomonas</i>	<i>sputigena</i>	151	Named	305	21	0.008776	0.13238
<i>Enterococcus</i>	<i>casseliflava</i>	801	Named	282	22	0.008114	0.12240
<i>Haemophilus</i>	<i>parainfluenzae</i>	826	Named	280	23	0.008057	0.12153
<i>Bacillus</i>	<i>fusiformis</i>	614	Named	279	24	0.008028	0.12109
<i>Fusobacterium</i>	<i>nucleatum</i> subsp. <i>animalis</i>	420	Named	255	25	0.007337	0.11068
<i>Streptococcus</i>	sp.	58	Unnamed	251	26	0.007222	0.10894
<i>Parvimonas</i>	<i>micra</i>	111	Named	237	27	0.006820	0.10286
<i>Abiotrophia</i>	<i>defectiva</i>	389	Named	236	28	0.006791	0.10243
<i>Bacteroidaceae [G-I]</i>	sp.	272	Unnamed	210	29	0.006043	0.09115

<i>Veillonella</i>	<i>atypica</i>	524	Named	208	T-30	0.005985	0.09028
<i>Streptococcus</i>	<i>infantis</i>	638	Named	208	T-30	0.005985	0.09028
<i>Lactococcus</i>	<i>lactis</i>	804	Named	208	T-30	0.005985	0.09028
<i>Rothia</i>	<i>dentocariosa</i>	587	Named	207	33	0.005956	0.08984
<i>Capnocytophaga</i>	<i>gingivalis</i>	337	Named	202	34	0.005812	0.08767
<i>Pseudoramibacter</i>	<i>alactolyticus</i>	538	Named	197	35	0.005669	0.08550
<i>Peptostreptococcus</i>	<i>stomatis</i>	112	Named	191	36	0.005496	0.08290
<i>Streptococcus</i>	sp.	71	Unnamed	189	37	0.005438	0.08203
<i>Dialister</i>	<i>pneumosintes</i>	736	Named	185	38	0.005323	0.08030
<i>Streptococcus</i>	<i>australis</i>	73	Named	178	T-39	0.005122	0.07726
<i>Micrococcus</i>	sp.	F54	Phylotype	178	T-39	0.005122	0.07726
<i>Streptococcus</i>	sp.	C65	Phylotype	176	41	0.005064	0.07639
<i>Escherichia</i>	<i>coli</i>	574	Named	171	42	0.004920	0.07422
<i>Corynebacterium</i>	<i>matruchotii</i>	666	Named	169	43	0.004863	0.07335
<i>Mitsuokella</i>	sp.	131	Phylotype	167	44	0.004805	0.07248
<i>Atopobium</i>	<i>rimae</i>	750	Named	164	45	0.004719	0.07118

<i>Propionibacterium</i>	<i>acidifaciens</i> .	191	Unnamed	159	46	0.004575	0.06901
<i>Streptococcus</i>	<i>constellatus</i>	576	Named	158	47	0.004546	0.06858
<i>Fusobacterium</i>	<i>nucleatum subsp. polymorphum</i>	202	Named	156	48	0.004489	0.06771
<i>Anaeroglobus</i>	<i>geminatus</i>	121	Named	154	49	0.004431	0.06684
<i>Prevotella</i>	<i>oris</i>	311	Named	153	50	0.004402	0.06641
<i>Lactobacillus</i>	<i>rhamnosus</i>	B85	Named	142	51	0.004086	0.06163
<i>Streptococcus</i>	sp.	431	Phylotype	140	52	0.004028	0.06076
<i>Lautropia</i>	<i>mirabilis</i>	22	Named	138	T-53	0.003971	0.05990
<i>Enterococcus</i>	<i>faecalis</i>	604	Named	138	T-53	0.003971	0.05990
<i>Gramulicatella</i>	<i>elegans</i>	596	Named	135	T-55	0.003885	0.05859
<i>Enterobacter</i>	<i>hormaechei</i>	634	Named	135	T-55	0.003885	0.05859
<i>Pseudomonas</i>	<i>aeruginosa</i>	536	Named	133	57	0.003827	0.05773
<i>Streptococcus</i>	sp.	70	Unnamed	131	T-58	0.003769	0.05686
<i>Streptococcus</i>	sp.	C56	Phylotype	131	T-58	0.003769	0.05686
<i>Eubacterium [XII] [G-5]</i>	<i>saphenum</i>	759	Named	130	60	0.003741	0.05642

<i>Streptococcus</i>	sp.	C08	Phylotype	129	61	0.003712	0.05599
<i>Streptococcus</i>	sp.	66	Unnamed	127	62	0.003654	0.05512
<i>Streptococcus</i>	<i>salivarius</i>	755	Named	121	63	0.003482	0.05252
<i>Veillonellaceae [G-I]</i>	sp.	155	Unnamed	115	T-64	0.003309	0.04991
<i>Neisseria</i>	<i>flavescens</i>	610	Named	115	T-64	0.003309	0.04991
<i>Neisseria</i>	<i>mucosa</i>	682	Named	115	T-64	0.003309	0.04991
<i>Megasphaera</i>	<i>micronuciformis</i>	122	Named	114	T-67	0.003280	0.04948
<i>Micrococcus</i>	<i>luteus</i>	C78	Named	114	T-67	0.003280	0.04948
<i>Selenomonas</i>	<i>infelix</i>	639	Named	112	69	0.003223	0.04861
<i>Haemophilus</i>	sp.	36	Unnamed	110	T-70	0.003165	0.04774
<i>Neisseria</i>	<i>flava</i>	609	Named	110	T-70	0.003165	0.04774
<i>Eubacterium [XIVa][G-I]</i>	<i>saburreum</i>	494	Named	109	72	0.003136	0.04731
<i>Filifactor</i>	<i>alocis</i>	539	Named	108	73	0.003108	0.04688
<i>Actinomyces</i>	sp.	448	Phylotype	107	T-74	0.003079	0.04644
<i>Actinomyces</i>	<i>israelii</i>	645	Named	107	T-74	0.003079	0.04644
<i>Propionibacterium</i>	<i>propionicum</i>	739	Named	107	T-74	0.003079	0.04644

<i>Selenomonas</i>	<i>dianae</i>	139	Named	105	T-77	0.003021	0.04557
<i>Acinetobacter</i>	sp.	C51	Phylotype	105	T-77	0.003021	0.04557
<i>Lactobacillus</i>	<i>paracasei</i>	716	Named	104	79	0.002993	0.04514
<i>Atopobium</i>	<i>parvulum</i>	723	Named	103	80	0.002964	0.04470
<i>Lactobacillus</i>	sp.	461	Phylotype	102	T-81	0.002935	0.04427
<i>Streptococcus</i>	sp.	C14	Phylotype	102	T-81	0.002935	0.04427
<i>Atopobium</i>	sp.	416	Phylotype	101	83	0.002906	0.04384
<i>Haemophilus</i>	<i>influenzae</i>	641	Named	100	84	0.002877	0.04340
<i>Neisseria</i>	<i>sicca</i>	764	Named	98	85	0.002820	0.04253
<i>Mogibacterium</i>	<i>timidum</i>	42	Named	94	86	0.002705	0.04080
<i>Catonella</i>	<i>morbi</i>	165	Named	92	87	0.002647	0.03993
<i>Lactobacillus</i>	<i>plantarum</i>	D15	Named	91	88	0.002618	0.03950
<i>Porphyromonas</i>	<i>gingivalis</i>	619	Named	89	89	0.002561	0.03863
<i>Tannerella</i>	<i>forsythia</i>	613	Named	88	T-90	0.002532	0.03819
<i>Kingella</i>	<i>oralis</i>	706	Named	88	T-90	0.002532	0.03819
<i>Gemella</i>	<i>bergeri</i>	555	Named	87	92	0.002503	0.03776

<i>Lactobacillus</i>	<i>vaginalis</i>	51	Named	86	93	0.002475	0.03733
<i>Enterococcus</i>	sp.	A43	Phylotype	85	94	0.002446	0.03689
<i>Porphyromonas</i>	<i>endodontalis</i>	273	Named	83	95	0.002388	0.03602
<i>Xanthomonas</i>	sp.	37	Phylotype	82	96	0.002360	0.03559
<i>Neisseria</i>	<i>pharyngis</i>	729	Named	81	97	0.002331	0.03516
<i>Actinomyces</i>	<i>naeslundii</i>	176	Named	79	98	0.002273	0.03429
<i>Prevotella</i>	<i>melaninogenica</i>	469	Named	78	99	0.002244	0.03385
<i>Treponema</i>	<i>denticola</i>	584	Named	77	100	0.002216	0.03342
<i>Shuttleworthia</i>	<i>satelles</i>	95	Named	76	T-101	0.002187	0.03299
<i>Prevotella</i>	<i>nigrescens</i>	693	Named	76	T-101	0.002187	0.03299
<i>Treponema</i>	<i>socranskii</i> subsp. <i>socranskii</i>	769	Named	75	103	0.002158	0.03255
<i>Actinomyces</i>	sp.	171	Unnamed	73	T-104	0.002101	0.03168
<i>Streptococcus</i>	<i>sobrinus</i>	768	Named	73	T-104	0.002101	0.03168
<i>Selenomonas</i>	sp.	140	Unnamed	70	T-106	0.002014	0.03038
<i>Propionibacterium</i>	<i>acnes</i>	530	Named	70	T-106	0.002014	0.03038
<i>Bacillus</i>	sp.	B77	Phylotype	70	T-106	0.002014	0.03038

<i>Campylobacter</i>	<i>concisus</i>	575	Named	69	T-109	0.001985	0.02995
<i>Campylobacter</i>	<i>showae</i>	763	Named	69	T-109	0.001985	0.02995
<i>Synergistetes [G-3]</i>	sp.	360	Phylotype	66	111	0.001899	0.02865
<i>Prevotella</i>	sp.	317	Unnamed	65	112	0.001870	0.02821
<i>Clostridiales [F-2][G-1]</i>	sp.	75	Phylotype	64	T-113	0.001842	0.02778
<i>Prevotella</i>	<i>tannerae</i>	466	Named	64	T-113	0.001842	0.02778
<i>Streptococcus</i>	<i>parasanguinis I</i>	721	Named	64	T-113	0.001842	0.02778
<i>Burkholderia</i>	sp.	C87	Phylotype	63	116	0.001813	0.02734
<i>Streptococcus</i>	<i>pneumoniae</i>	734	Named	62	117	0.001784	0.02691
<i>Streptococcus</i>	<i>vestibularis</i>	21	Named	61	T-118	0.001755	0.02648
<i>Actinomyces</i>	sp.	169	Unnamed	61	T-118	0.001755	0.02648
<i>Prevotella</i>	<i>denticola</i>	291	Named	61	T-118	0.001755	0.02648
<i>Lactobacillus</i>	<i>fermentum</i>	608	Named	60	T-121	0.001726	0.02604
<i>Neisseria</i>	<i>meningitidis</i>	669	Named	60	T-121	0.001726	0.02604
<i>Neisseria</i>	sp.	14	Phylotype	59	123	0.001698	0.02561
<i>Lachnospiraceae [G-2]</i>	sp.	100	Phylotype	58	124	0.001669	0.02517

<i>Eubacterium [XI][G-1]</i>	<i>infirmum</i>	105	Named	57	125	0.001640	0.02474
<i>Selenomonas</i>	<i>artemidis</i>	124	Named	56	T-126	0.001611	0.02431
<i>Gemella</i>	<i>sanguinis</i>	757	Named	56	T-126	0.001611	0.02431
<i>Bifidobacterium</i>	<i>dentium</i>	588	Named	55	T-128	0.001583	0.02387
<i>Prevotella</i>	<i>intermedia</i>	643	Named	55	T-128	0.001583	0.02387
<i>Neisseria</i>	<i>bacilliformis</i>	13	Named	54	T-130	0.001554	0.02344
<i>Lachnospiraceae [G-1]</i>	sp.	107	Unnamed	54	T-130	0.001554	0.02344
<i>Parvimonas</i>	sp.	110	Unnamed	54	T-130	0.001554	0.02344
<i>Bacteroidetes [G-3]</i>	sp.	281	PhyloTYPE	54	T-130	0.001554	0.02344
<i>Bacilli [G]</i>	sp.	C43	PhyloTYPE	54	T-130	0.001554	0.02344
<i>Streptococcus</i>	sp.	74	Unnamed	53	T-135	0.001525	0.02300
<i>Fusobacterium</i>	<i>periodonticum</i>	201	Named	53	T-135	0.001525	0.02300
<i>Selenomonas</i>	sp.	G00	PhyloTYPE	53	T-135	0.001525	0.02300
<i>Actinomyces</i>	sp.	175	Unnamed	52	138	0.001496	0.02257
<i>Bacteroidales [G-2]</i>	sp.	274	Unnamed	51	T-139	0.001467	0.02214
<i>Neisseria</i>	<i>subflava</i>	476	Named	51	T-139	0.001467	0.02214

<i>Paracoccus</i>	<i>marcusii</i>	D41	Named	51	T-139	0.001467	0.02214
<i>Leptotrichia</i>	<i>wadei</i>	222	Named	50	142	0.001439	0.02170
<i>Streptococcus</i>	sp.	61	Unnamed	49	T-143	0.001410	0.02127
<i>Selenomonas</i>	sp.	149	Unnamed	49	T-143	0.001410	0.02127
<i>Streptococcus</i>	<i>downei</i>	594	Named	49	T-143	0.001410	0.02127
<i>Oribacterium</i>	<i>sinus</i>	457	Named	48	T-146	0.001381	0.02083
<i>Prevotella</i>	sp.	G57	Phylotype	48	T-146	0.001381	0.02083
<i>Fusobacteria [G-1]</i>	sp.	220	Phylotype	47	T-148	0.001352	0.02040
<i>Prevotella</i>	sp.	298	Unnamed	47	T-148	0.001352	0.02040
<i>Cardiobacterium</i>	<i>hominis</i>	633	Named	47	T-148	0.001352	0.02040
<i>Enterococcus</i>	<i>saccharominimus</i>	803	Named	47	T-148	0.001352	0.02040
<i>Fusobacterium</i>	sp.	A11	Phylotype	47	T-148	0.001352	0.02040
<i>Synergistetes [G-3]</i>	sp.	363	Phylotype	46	T-153	0.001324	0.01997
<i>Oribacterium</i>	sp.	372	Unnamed	46	T-153	0.001324	0.01997
<i>Bacillus</i>	<i>subtilis</i>	468	Named	45	T-155	0.001295	0.01953
<i>Corynebacterium</i>	sp.	B00	Phylotype	45	T-155	0.001295	0.01953

<i>Lactobacillus</i>	<i>rhamnosus</i>	749	Named	44	157	0.001266	0.01910
<i>Megasphaera</i>	sp.	123	Phylotype	43	158	0.001237	0.01866
<i>Treponema</i>	sp.	237	Phylotype	42	T-159	0.001209	0.01823
<i>Porphyromonas</i>	sp.	279	Unnamed	42	T-159	0.001209	0.01823
<i>Prevotella</i>	<i>oralis</i>	705	Named	42	T-159	0.001209	0.01823
<i>Eubacterium [XII] [G-7]</i>	<i>yurii</i>	377	Named	40	T-162	0.001151	0.01736
<i>Neisseria</i>	<i>elongata</i>	598	Named	40	T-162	0.001151	0.01736
<i>Veillonella</i>	sp.	158	Unnamed	39	T-164	0.001122	0.01693
<i>Capnocytophaga</i>	sp.	329	Unnamed	39	T-164	0.001122	0.01693
<i>Capnocytophaga</i>	<i>sputigena</i>	775	Named	38	166	0.001093	0.01649
<i>Actinomyces</i>	sp.	170	Unnamed	36	T-167	0.001036	0.01563
<i>Selenomonas</i>	sp.	479	Unnamed	36	T-167	0.001036	0.01563
<i>Actinomyces</i>	<i>gerencseriae</i>	618	Named	36	T-167	0.001036	0.01563
<i>Burkholderia</i>	sp.	406	Phylotype	33	T-170	0.000950	0.01432
<i>Lachnospiraceae [G-4]</i>	sp.	419	Unnamed	33	T-170	0.000950	0.01432
<i>Rothia</i>	<i>mucilaginosa</i>	681	Named	33	T-170	0.000950	0.01432

<i>Lactobacillus</i>	<i>salivarius</i>	756	Named	33	T-170	0.000950	0.01432
<i>Streptococcus</i>	<i>sinensis</i>	767	Named	33	T-170	0.000950	0.01432
<i>Olsenella</i>	<i>profusa</i>	806	Named	33	T-170	0.000950	0.01432
<i>Desulfobulbus</i>	sp.	41	Phylotype	32	T-176	0.000921	0.01389
<i>Streptococcus</i>	sp.	68	Unnamed	32	T-176	0.000921	0.01389
<i>Fusobacterium</i>	sp.	370	Unnamed	32	T-176	0.000921	0.01389
<i>Prevotella</i>	sp.	473	Unnamed	31	T-179	0.000892	0.01345
<i>Cryptobacterium</i>	<i>curtum</i>	579	Named	31	T-179	0.000892	0.01345
<i>Streptococcus</i>	sp.	56	Unnamed	30	T-181	0.000863	0.01302
<i>Veillonellaceae [G-I]</i>	sp.	132	Phylotype	30	T-181	0.000863	0.01302
<i>Veillonellaceae [G-I]</i>	sp.	150	Phylotype	30	T-181	0.000863	0.01302
<i>Achromobacter</i>	<i>xylooxidans</i>	343	Named	30	T-181	0.000863	0.01302
<i>Eikenella</i>	<i>corrodens</i>	577	Named	30	T-181	0.000863	0.01302
<i>Kingella</i>	<i>denitrificans</i>	582	Named	30	T-181	0.000863	0.01302
<i>Proteus</i>	<i>mirabilis</i>	676	Named	30	T-181	0.000863	0.01302
<i>Fusobacterium</i>	sp.	203	Phylotype	29	T-188	0.000834	0.01259

<i>Dialister</i>	sp.	A97	Phylotype	29	T-188	0.000834	0.01259
<i>Veillonellaceae [G-1]</i>	sp.	145	Phylotype	28	T-190	0.000806	0.01215
<i>Treponema</i>	sp.	232	Phylotype	28	T-190	0.000806	0.01215
<i>Corynebacterium</i>	<i>durum</i>	595	Named	28	T-190	0.000806	0.01215
<i>Stenotrophomonas</i>	<i>maltophilia</i>	663	Named	28	T-190	0.000806	0.01215
<i>Veillonella</i>	sp.	780	Unnamed	28	T-190	0.000806	0.01215
<i>Sphingomonas</i>	sp.	A66	Phylotype	28	T-190	0.000806	0.01215
<i>Paenibacillus</i>	sp.	E29	Phylotype	28	T-190	0.000806	0.01215
<i>Firmicutes [G]</i>	sp.	G23	Phylotype	28	T-190	0.000806	0.01215
<i>Treponema</i>	sp.	247	Phylotype	27	T-198	0.000777	0.01172
<i>Synergistetes [G-3]</i>	sp.	358	Phylotype	27	T-198	0.000777	0.01172
<i>Acinetobacter</i>	<i>baumannii</i>	554	Named	27	T-198	0.000777	0.01172
<i>Lactobacillus</i>	<i>oris</i>	709	Named	27	T-198	0.000777	0.01172
<i>Treponema</i>	sp.	228	Phylotype	25	T-202	0.000719	0.01085
<i>Selenomonas</i>	sp.	E83	Phylotype	25	T-202	0.000719	0.01085
<i>Clostridiales [F-2][G-2]</i>	sp.	85	Phylotype	24	T-204	0.000691	0.01042

<i>Peptostreptococcaceae</i> [XI][G-4]	sp.		369	Phylotype	24	T-204	0.000691	0.01042
<i>Pseudomonas</i>	sp.		B99	Phylotype	24	T-204	0.000691	0.01042
<i>Selenomonas</i>	sp.		F19	Phylotype	24	T-204	0.000691	0.01042
<i>Streptococcus</i>	sp.		65	Unnamed	23	T-208	0.000662	0.00998
<i>Selenomonas flueggei</i>			125	Named	23	T-208	0.000662	0.00998
<i>Scardovia</i>	sp.		195	Unnamed	23	T-208	0.000662	0.00998
<i>Capnocytophaga</i>	sp.		338	Unnamed	23	T-208	0.000662	0.00998
<i>TM7 [G-1]</i>	sp.		346	Phylotype	23	T-208	0.000662	0.00998
<i>Actinomyces</i>	sp.		B78	Phylotype	23	T-208	0.000662	0.00998
<i>Pseudomonas</i>	sp.		D11	Phylotype	23	T-208	0.000662	0.00998
<i>Selenomonas</i>	sp.		H64	Phylotype	23	T-208	0.000662	0.00998
<i>Paracoccus</i>	sp.		H67	Phylotype	23	T-208	0.000662	0.00998
<i>Streptococcus</i>	sp.		57	Unnamed	22	T-217	0.000633	0.00955
<i>Oribacterium</i>	sp.		108	Unnamed	22	T-217	0.000633	0.00955
<i>Actinomyces</i>	sp.		180	Unnamed	22	T-217	0.000633	0.00955

<i>Peptoniphilus</i>	sp.	836	Unnamed	22	T-217	0.000633	0.00955
<i>Actinobaculum</i>	sp.	183	Unnamed	21	T-221	0.000604	0.00911
<i>Rothia</i>	sp.	188	Unnamed	21	T-221	0.000604	0.00911
<i>Treponema</i>	sp.	231	Phylotype	21	T-221	0.000604	0.00911
<i>Prevotella</i>	<i>pleuritidis</i>	303	Named	21	T-221	0.000604	0.00911
<i>Pyramidobacter</i>	<i>piscolens</i>	357	Named	21	T-221	0.000604	0.00911
<i>Pseudomonas</i>	sp.	C61	Phylotype	21	T-221	0.000604	0.00911
<i>Streptococcus</i>	sp.	E98	Phylotype	21	T-221	0.000604	0.00911
<i>Lachnospiraceae [G-3]</i>	sp.	96	Unnamed	20	T-228	0.000575	0.00868
<i>Selenomonas</i>	sp.	138	Unnamed	20	T-228	0.000575	0.00868
<i>Bacteroidetes [G-3]</i>	sp.	365	Phylotype	20	T-228	0.000575	0.00868
<i>Eubacterium [XI] [G-1]</i>	<i>sulci</i>	467	Named	20	T-228	0.000575	0.00868
<i>Solobacterium</i>	<i>moorei</i>	678	Named	20	T-228	0.000575	0.00868
<i>Streptococcus</i>	<i>oralis</i>	707	Named	20	T-228	0.000575	0.00868
<i>Streptococcus</i>	<i>peroris</i>	728	Named	20	T-228	0.000575	0.00868
<i>Lachnospiraceae [G]</i>	sp.	A17	Phylotype	20	T-228	0.000575	0.00868

<i>Corynebacterium</i>	<i>durum</i>	A65	Named	20	T-228	0.000575	0.00868
<i>Selenomonas</i>	sp.	137	Unnamed	19	T-237	0.000547	0.00825
<i>Prevotella</i>	sp.	313	PhyloTYPE	19	T-237	0.000547	0.00825
<i>Peptostreptococcaceae</i> [XI][G-1]	sp.	383	PhyloTYPE	19	T-237	0.000547	0.00825
<i>Mitsuokella</i>	sp.	521	PhyloTYPE	19	T-237	0.000547	0.00825
<i>Ochrobactrum</i>	<i>anthropi</i>	544	Named	19	T-237	0.000547	0.00825
<i>Eubacterium</i> [XI][G-6]	<i>minutum</i>	673	Named	19	T-237	0.000547	0.00825
<i>Capnocytophaga</i>	sp.	B29	PhyloTYPE	19	T-237	0.000547	0.00825
<i>Selenomonas</i>	sp.	F21	PhyloTYPE	19	T-237	0.000547	0.00825
<i>Selenomonas</i>	sp.	H23	PhyloTYPE	19	T-237	0.000547	0.00825
<i>Peptostreptococcaceae</i>	sp.	113	PhyloTYPE	18	T-246	0.000518	0.00781
[XIII][G-1]							
<i>Veillonellaceae</i> [G-1]	sp.	129	Unnamed	18	T-246	0.000518	0.00781
<i>Prevotella</i>	<i>histicola</i>	289	Named	18	T-246	0.000518	0.00781
<i>Prevotella</i>	sp.	300	PhyloTYPE	18	T-246	0.000518	0.00781

<i>TM7 [G-1]</i>	sp.	347	Phylotype	18	T-246	0.000518	0.00781
<i>Prevotella</i>	sp.	376	Unnamed	18	T-246	0.000518	0.00781
<i>Fusobacterium</i>	<i>nucleatum subsp. nucleatum</i>	698	Named	18	T-246	0.000518	0.00781
<i>Haemophilus</i>	<i>parainfluenzae</i>	718	Named	18	T-246	0.000518	0.00781
<i>Paracoccus</i>	sp.	A23	Phylotype	18	T-246	0.000518	0.00781
<i>Bacillus</i>	sp.	F28	Phylotype	18	T-246	0.000518	0.00781
<i>Selenomonas</i>	sp.	G55	Phylotype	18	T-246	0.000518	0.00781
<i>Fusobacterium</i>	sp.	H27	Phylotype	18	T-246	0.000518	0.00781
<i>Peptococcus</i>	sp.	167	Unnamed	17	T-258	0.000489	0.00738
<i>Leptotrichia</i>	sp.	215	Phylotype	17	T-258	0.000489	0.00738
<i>Leptotrichia</i>	<i>hofstadii</i>	224	Named	17	T-258	0.000489	0.00738
<i>Treponema</i>	sp.	257	Unnamed	17	T-258	0.000489	0.00738
<i>Lactobacillus</i>	sp.	418	Unnamed	17	T-258	0.000489	0.00738
<i>Parascardovia</i>	<i>denticolens</i>	586	Named	17	T-258	0.000489	0.00738
<i>Lautropia</i>	sp.	A94	Phylotype	17	T-258	0.000489	0.00738
<i>Porphyromonas</i>	sp.	B47	Phylotype	17	T-258	0.000489	0.00738

<i>Burkholderiaceae [G]</i>	sp.	D14	Phylotype	17	T-258	0.000489	0.00738
<i>Oribacterium</i>	sp.	102	Phylotype	16	T-267	0.000460	0.00694
<i>Tannerella</i>	sp.	286	Phylotype	16	T-267	0.000460	0.00694
<i>Burkholderia</i>	<i>cepacia</i>	571	Named	16	T-267	0.000460	0.00694
<i>Bacillus</i>	sp.	C44	Phylotype	16	T-267	0.000460	0.00694
<i>Manihot</i>	<i>esculenta</i>	C60	Named	16	T-267	0.000460	0.00694
<i>Streptococcus</i>	sp.	H24	Phylotype	16	T-267	0.000460	0.00694
<i>Lachnospiraceae [G-1]</i>	sp.	82	Unnamed	15	T-273	0.000432	0.00651
<i>Dialister</i>	sp.	119	Phylotype	15	T-273	0.000432	0.00651
<i>Leptotrichia</i>	sp.	212	Phylotype	15	T-273	0.000432	0.00651
<i>Prevotella</i>	sp.	314	Unnamed	15	T-273	0.000432	0.00651
<i>Lachnospiraceae [G-4]</i>	sp.	373	Phylotype	15	T-273	0.000432	0.00651
<i>Prevotella</i>	sp.	472	Unnamed	15	T-273	0.000432	0.00651
<i>Leptotrichia</i>	sp.	498	Unnamed	15	T-273	0.000432	0.00651
<i>Staphylococcus</i>	<i>aureus</i>	550	Named	15	T-273	0.000432	0.00651
<i>Treponema</i>	<i>maltophilum</i>	664	Named	15	T-273	0.000432	0.00651

<i>Actinomyces</i>	sp.		C25	Phylotype	15	T-273	0.000432	0.00651
<i>Brevundimonas</i>	<i>diminuta</i>		D38	Named	15	T-273	0.000432	0.00651
<i>Burkholderia</i>	sp.		D43	Phylotype	15	T-273	0.000432	0.00651
<i>Actinomyces</i>	sp.		E75	Phylotype	15	T-273	0.000432	0.00651
<i>Rhodocyclus</i>	sp.		28	Unnamed	14	T-286	0.000403	0.00608
<i>Olsenella</i>	<i>uli</i>		38	Named	14	T-286	0.000403	0.00608
<i>Selenomonas</i>	sp.		126	Phylotype	14	T-286	0.000403	0.00608
<i>Treponema</i>	sp.		251	Phylotype	14	T-286	0.000403	0.00608
<i>Porphyromonas</i>	<i>catoniae</i>		283	Named	14	T-286	0.000403	0.00608
<i>Bergeyella</i>	sp.		422	Phylotype	14	T-286	0.000403	0.00608
<i>Selenomonas</i>	sp.		481	Unnamed	14	T-286	0.000403	0.00608
<i>Deftuvibacter</i>	<i>lusatiensis</i>		660	Named	14	T-286	0.000403	0.00608
<i>Treponema</i>	sp.		A31	Phylotype	14	T-286	0.000403	0.00608
<i>Selenomonas</i>	sp.		E50	Phylotype	14	T-286	0.000403	0.00608
<i>Actinomyces</i>	sp.		172	Unnamed	13	T-296	0.000374	0.00564
<i>Treponema</i>	sp.		230	Phylotype	13	T-296	0.000374	0.00564

<i>Treponema</i>	sp.	265	Phylotype	13	T-296	0.000374	0.00564
<i>Prevotella</i>	<i>oulorum</i>	288	Named	13	T-296	0.000374	0.00564
<i>Prevotella</i>	sp.	292	Phylotype	13	T-296	0.000374	0.00564
<i>Actinomyces</i>	sp.	525	Phylotype	13	T-296	0.000374	0.00564
<i>Cardiobacterium</i>	<i>valvulum</i>	540	Named	13	T-296	0.000374	0.00564
<i>Johnsonella</i>	<i>ignava</i>	635	Named	13	T-296	0.000374	0.00564
<i>Treponema</i>	<i>lecithinolyticum</i>	653	Named	13	T-296	0.000374	0.00564
<i>Aggregatibacter</i>	<i>segnis</i>	762	Named	13	T-296	0.000374	0.00564
<i>Dialister</i>	<i>micraerophilus</i>	843	Named	13	T-296	0.000374	0.00564
<i>Streptococcus</i>	sp.	B59	Phylotype	13	T-296	0.000374	0.00564
<i>Bacillus</i>	<i>clausii</i>	45	Named	12	T-308	0.000345	0.00521
<i>Actinomyces</i>	sp.	178	Unnamed	12	T-308	0.000345	0.00521
<i>Leptotrichia</i>	sp.	213	Phylotype	12	T-308	0.000345	0.00521
<i>Synergistetes [G-3]</i>	sp.	359	Phylotype	12	T-308	0.000345	0.00521
<i>Synergistetes [G-3]</i>	sp.	362	Phylotype	12	T-308	0.000345	0.00521
<i>TM7 [G-5]</i>	sp.	437	Phylotype	12	T-308	0.000345	0.00521

<i>Actinomyces</i>	sp.	446	Phylotype	12	T-308	0.000345	0.00521
<i>Aggregatibacter</i>	sp.	458	Unnamed	12	T-308	0.000345	0.00521
<i>Leptotrichia</i>	<i>buccalis</i>	563	Named	12	T-308	0.000345	0.00521
<i>Pseudomonas</i>	<i>fluorescens</i>	612	Named	12	T-308	0.000345	0.00521
<i>Campylobacter</i>	<i>rectus</i>	748	Named	12	T-308	0.000345	0.00521
<i>Prevotella</i>	<i>multisaccharivorax</i>	794	Named	12	T-308	0.000345	0.00521
<i>Paracoccus</i>	<i>marinus</i>	A13	Named	12	T-308	0.000345	0.00521
<i>Haemophilus</i>	sp.	C82	Phylotype	12	T-308	0.000345	0.00521
<i>Actinomyces</i>	<i>massiliensis</i>	C93	Named	12	T-308	0.000345	0.00521
<i>Selenomonas</i>	sp.	E39	Phylotype	12	T-308	0.000345	0.00521
<i>Actinomyces</i>	sp.	E63	Phylotype	12	T-308	0.000345	0.00521
<i>Prevotella</i>	sp.	G70	Phylotype	12	T-308	0.000345	0.00521
<i>Mogibacterium</i>	<i>vescum</i>	8	Named	11	T-326	0.000317	0.00477
<i>Leptothrix</i>	sp.	25	Phylotype	11	T-326	0.000317	0.00477
<i>Haemophilus</i>	sp.	35	Phylotype	11	T-326	0.000317	0.00477
<i>Peptostreptococcaceae [XII][G-</i>	sp.	103	Phylotype	11	T-326	0.000317	0.00477

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<i>Selenomonas</i>	sp.			134	Unnamed	11	T-326	0.000317	0.000317	0.00477			
<i>Johnsonella</i>	sp.			166	Phylotype	11	T-326	0.000317	0.000317	0.00477			
<i>Actinomyces</i>	sp.			181	Unnamed	11	T-326	0.000317	0.000317	0.00477			
<i>Treponema</i>	sp.			239	Phylotype	11	T-326	0.000317	0.000317	0.00477			
<i>Prevotella</i>	sp.			306	Unnamed	11	T-326	0.000317	0.000317	0.00477			
<i>Lactobacillus</i>	sp.			424	Unnamed	11	T-326	0.000317	0.000317	0.00477			
<i>Neisseria</i>	sp.			499	Phylotype	11	T-326	0.000317	0.000317	0.00477			
<i>Bulleidia</i>	<i>extructa</i>			603	Named	11	T-326	0.000317	0.000317	0.00477			
<i>Actinomyces</i>	<i>odontolyticus</i>			701	Named	11	T-326	0.000317	0.000317	0.00477			
<i>Clostridiales [G]</i>	sp.			C16	Phylotype	11	T-326	0.000317	0.000317	0.00477			
<i>Selenomonas</i>	sp.			C23	Phylotype	11	T-326	0.000317	0.000317	0.00477			
<i>Parvimonas</i>	sp.			D94	Phylotype	11	T-326	0.000317	0.000317	0.00477			
<i>Lactobacillus</i>	sp.			H25	Phylotype	11	T-326	0.000317	0.000317	0.00477			
<i>Kingella</i>	sp.			12	Phylotype	10	T-343	0.000288	0.000288	0.00434			
<i>Clostridiales [F-I][G-I]</i>	sp.			93	Phylotype	10	T-343	0.000288	0.000288	0.00434			

<i>Lachnospiraceae [G-4]</i>	sp.	97	Phylotype	10	T-343	0.000288	0.00434
<i>Selenomonas</i>	sp.	133	Unnamed	10	T-343	0.000288	0.00434
<i>Veillonellaceae [G-1]</i>	sp.	148	Phylotype	10	T-343	0.000288	0.00434
<i>Catonella</i>	sp.	164	Phylotype	10	T-343	0.000288	0.00434
<i>Actinomyces</i>	sp.	177	Unnamed	10	T-343	0.000288	0.00434
<i>Treponema</i>	sp.	234	Phylotype	10	T-343	0.000288	0.00434
<i>Bacteroidetes [G-3]</i>	sp.	280	Phylotype	10	T-343	0.000288	0.00434
<i>Prevotella</i>	sp.	308	Phylotype	10	T-343	0.000288	0.00434
<i>Agrobacterium</i>	<i>tumefaciens</i>	485	Named	10	T-343	0.000288	0.00434
<i>Dialister</i>	sp.	502	Phylotype	10	T-343	0.000288	0.00434
<i>Peptostreptococcus</i>	<i>anaerobius</i>	542	Named	10	T-343	0.000288	0.00434
<i>Lactobacillus</i>	<i>crispatus</i>	817	Named	10	T-343	0.000288	0.00434
<i>Solobacterium</i>	sp.	A05	Phylotype	10	T-343	0.000288	0.00434
<i>Diaphorobacter</i>	<i>nitroreducens</i>	A07	Named	10	T-343	0.000288	0.00434
<i>Lachnospiraceae [G]</i>	sp.	A61	Phylotype	10	T-343	0.000288	0.00434
<i>Streptococcus</i>	sp.	E12	Phylotype	10	T-343	0.000288	0.00434

<i>Selenomonas</i>	sp.	E20	Phylotype	10	T-343	0.000288	0.00434
<i>Deinococcus</i>	sp.	F57	Phylotype	10	T-343	0.000288	0.00434
<i>Selenomonas</i>	sp.	F85	Phylotype	10	T-343	0.000288	0.00434
<i>Leptothrix</i>	sp.	26	Phylotype	9	T-364	0.000259	0.00391
<i>Paenibacillus</i>	sp.	48	Phylotype	9	T-364	0.000259	0.00391
<i>Veillonellaceae [G-1]</i>	sp.	135	Phylotype	9	T-364	0.000259	0.00391
<i>Leptotrichia</i>	sp.	219	Phylotype	9	T-364	0.000259	0.00391
<i>Treponema</i>	sp.	238	Phylotype	9	T-364	0.000259	0.00391
<i>Treponema</i>	sp.	254	Phylotype	9	T-364	0.000259	0.00391
<i>Bergeyella</i>	sp.	319	Phylotype	9	T-364	0.000259	0.00391
<i>Dietzia</i>	sp.	368	Unnamed	9	T-364	0.000259	0.00391
<i>Leptotrichia</i>	sp.	463	Phylotype	9	T-364	0.000259	0.00391
<i>Prevotella</i>	sp.	526	Phylotype	9	T-364	0.000259	0.00391
<i>Aggregatibacter</i>	<i>actinomycetemcomitans</i>	531	Named	9	T-364	0.000259	0.00391
<i>Prevotella</i>	<i>buccae</i>	560	Named	9	T-364	0.000259	0.00391
<i>Lactobacillus [17]</i>	<i>cateniformis</i>	569	Named	9	T-364	0.000259	0.00391

<i>Prevotella</i>	<i>pallens</i>	714	Named	9	T-364	0.000259	0.00391
<i>Centipeda</i>	<i>periodontii</i>	726	Named	9	T-364	0.000259	0.00391
<i>Burkholderiales [G]</i>	sp.	A57	Phylotype	9	T-364	0.000259	0.00391
<i>Streptococcus</i>	sp.	B66	Phylotype	9	T-364	0.000259	0.00391
<i>Prevotella</i>	<i>salivae</i>	B81	Named	9	T-364	0.000259	0.00391
<i>Pseudomonas</i>	sp.	C84	Phylotype	9	T-364	0.000259	0.00391
<i>Peptostreptococcaceae [XII][G]</i>	sp.	D17	Phylotype	9	T-364	0.000259	0.00391
<i>Streptococcus</i>	sp.	E43	Phylotype	9	T-364	0.000259	0.00391
<i>Selenomonas</i>	sp.	F80	Phylotype	9	T-364	0.000259	0.00391
<i>Selenomonas</i>	sp.	F81	Phylotype	9	T-364	0.000259	0.00391
<i>Eubacterium [XII][G]</i>	sp.	G32	Phylotype	9	T-364	0.000259	0.00391
<i>Campylobacter</i>	sp.	G43	Phylotype	9	T-364	0.000259	0.00391
<i>Treponema</i>	sp.	G85	Phylotype	9	T-364	0.000259	0.00391
<i>Pseudomonas</i>	sp.	32	Phylotype	8	T-390	0.000230	0.00347
<i>Leptotrichia</i>	<i>shahii</i>	214	Named	8	T-390	0.000230	0.00347
<i>Treponema</i>	sp.	236	Phylotype	8	T-390	0.000230	0.00347

<i>Treponema</i>	sp.		268	Unnamed	8	T-390	0.000230	0.00347
<i>Prevotella</i>	sp.		299	Unnamed	8	T-390	0.000230	0.00347
<i>Peptoniphilus</i>	sp.		375	Unnamed	8	T-390	0.000230	0.00347
<i>Parvimonas</i>	sp.		393	Unnamed	8	T-390	0.000230	0.00347
<i>Leptotrichia</i>	sp.		417	Phylotype	8	T-390	0.000230	0.00347
<i>Leptotrichia</i>	sp.		462	Phylotype	8	T-390	0.000230	0.00347
<i>Prevotella</i>	sp.		781	Unnamed	8	T-390	0.000230	0.00347
<i>Olsenella</i>	sp.		807	Unnamed	8	T-390	0.000230	0.00347
<i>Fusobacteria [G-I]</i>	sp.		A71	Phylotype	8	T-390	0.000230	0.00347
<i>Comamonas</i>	<i>testosteroni</i>		A85	Named	8	T-390	0.000230	0.00347
<i>Streptococcus</i>	sp.		A95	Phylotype	8	T-390	0.000230	0.00347
<i>Ralstonia</i>	sp.		B67	Phylotype	8	T-390	0.000230	0.00347
<i>Fusobacterium</i>	sp.		C10	Phylotype	8	T-390	0.000230	0.00347
<i>Selenomonas</i>	sp.		F20	Phylotype	8	T-390	0.000230	0.00347
<i>Streptococcus</i>	sp.		G62	Phylotype	8	T-390	0.000230	0.00347
<i>Selenomonas</i>	sp.		146	Unnamed	7	T-408	0.000201	0.00304

<i>Leptotrichia</i>	sp.		225	Unnamed	7	T-408	0.000201	0.00304
<i>Treponema</i>	sp.		264	Phylotype	7	T-408	0.000201	0.00304
<i>Prevotella</i>	<i>salivae</i>		307	Named	7	T-408	0.000201	0.00304
<i>Capnocytophaga</i>	sp.		326	Unnamed	7	T-408	0.000201	0.00304
<i>Capnocytophaga</i>	sp.		336	Unnamed	7	T-408	0.000201	0.00304
<i>SR1 [G-1]</i>	sp.		345	Phylotype	7	T-408	0.000201	0.00304
<i>TM7 [G-1]</i>	sp.		349	Phylotype	7	T-408	0.000201	0.00304
<i>Synergistetes [G-3]</i>	sp.		361	Phylotype	7	T-408	0.000201	0.00304
<i>Capnocytophaga</i>	sp.		412	Unnamed	7	T-408	0.000201	0.00304
<i>Actinomyces</i>	sp.		414	Phylotype	7	T-408	0.000201	0.00304
<i>Variovorax</i>	<i>paradoxus</i>		717	Named	7	T-408	0.000201	0.00304
<i>Enterococcus</i>	sp.		A78	Phylotype	7	T-408	0.000201	0.00304
<i>Lactobacillus</i>	sp.		A89	Phylotype	7	T-408	0.000201	0.00304
<i>Ralstonia</i>	<i>pickettii</i>		C02	Named	7	T-408	0.000201	0.00304
<i>Clostridiales [G]</i>	sp.		C86	Phylotype	7	T-408	0.000201	0.00304
<i>Lactobacillus</i>	sp.		D06	Phylotype	7	T-408	0.000201	0.00304

<i>Veillonella</i>	sp.		D93	Phylotype	7	T-408	0.000201	0.00304
<i>Selenomonas</i>	sp.		D96	Phylotype	7	T-408	0.000201	0.00304
<i>Streptococcus</i>	sp.		E81	Phylotype	7	T-408	0.000201	0.00304
<i>Alphaproteobacteria [G]</i>	sp.		F61	Phylotype	7	T-408	0.000201	0.00304
<i>Gemella</i>	sp.		G07	Phylotype	7	T-408	0.000201	0.00304
<i>Selenomonas</i>	sp.		G67	Phylotype	7	T-408	0.000201	0.00304
<i>Bacteroidetes [G-3]</i>	sp.		G83	Phylotype	7	T-408	0.000201	0.00304
<i>Campylobacter</i>	sp.		44	Phylotype	6	T-432	0.000173	0.00260
<i>TM7 [G-1]</i>	sp.		352	Phylotype	6	T-432	0.000173	0.00260
<i>Prevotella</i>	<i>micans</i>		378	Named	6	T-432	0.000173	0.00260
<i>Selenomonas</i>	sp.		442	Phylotype	6	T-432	0.000173	0.00260
<i>Selenomonas</i>	sp.		478	Unnamed	6	T-432	0.000173	0.00260
<i>Bacteroidetes [G-5]</i>	sp.		507	Phylotype	6	T-432	0.000173	0.00260
<i>Campylobacter</i>	<i>curvus</i>		580	Named	6	T-432	0.000173	0.00260
<i>Scardovia</i>	<i>inopinata</i>		642	Named	6	T-432	0.000173	0.00260
<i>Treponema</i>	<i>medium</i>		667	Named	6	T-432	0.000173	0.00260

<i>Aggregatibacter</i>	<i>paraphrophilus</i>	720	Named	6	T-432	0.000173	0.00260
<i>Bacillus</i>	sp.	A03	Phylotype	6	T-432	0.000173	0.00260
<i>Bacillus</i>	sp.	A08	Phylotype	6	T-432	0.000173	0.00260
<i>Actinomyces</i>	sp.	A50	Phylotype	6	T-432	0.000173	0.00260
<i>Acinetobacter</i>	sp.	A58	Phylotype	6	T-432	0.000173	0.00260
<i>Kytococcus</i>	<i>sedentarius</i>	A81	Named	6	T-432	0.000173	0.00260
<i>Cupriavidus</i>	<i>basilensis</i>	A84	Named	6	T-432	0.000173	0.00260
<i>Actinomyces</i>	<i>dentalis</i>	B28	Named	6	T-432	0.000173	0.00260
<i>Porphyromonas</i>	sp.	C34	Phylotype	6	T-432	0.000173	0.00260
<i>Olsenella</i>	sp.	C46	Phylotype	6	T-432	0.000173	0.00260
<i>Actinomyces</i>	sp.	C55	Phylotype	6	T-432	0.000173	0.00260
<i>Firmicutes [G]</i>	sp.	C68	Phylotype	6	T-432	0.000173	0.00260
<i>Bacillaceae [G]</i>	sp.	D26	Phylotype	6	T-432	0.000173	0.00260
<i>Marinomonas</i>	sp.	D33	Phylotype	6	T-432	0.000173	0.00260
<i>Cellulomonas</i>	<i>cellulans</i>	D36	Named	6	T-432	0.000173	0.00260
<i>Peptococcus</i>	sp.	D92	Phylotype	6	T-432	0.000173	0.00260

<i>Selenomonas</i>	sp.		E44	Phylotype	6	T-432	0.000173	0.00260
<i>Capnocytophaga</i>	sp.		E54	Phylotype	6	T-432	0.000173	0.00260
<i>Streptococcus</i>	sp.		E78	Phylotype	6	T-432	0.000173	0.00260
<i>Selenomonas</i>	sp.		F22	Phylotype	6	T-432	0.000173	0.00260
<i>Acinetobacter</i>	sp.		F58	Phylotype	6	T-432	0.000173	0.00260
<i>Sphingomonas</i>	sp.		F71	Phylotype	6	T-432	0.000173	0.00260
<i>Selenomonas</i>	sp.		F72	Phylotype	6	T-432	0.000173	0.00260
<i>Aggregatibacter</i>	sp.		G24	Phylotype	6	T-432	0.000173	0.00260
<i>Selenomonas</i>	sp.		G51	Phylotype	6	T-432	0.000173	0.00260
<i>Selenomonas</i>	sp.		H66	Phylotype	6	T-432	0.000173	0.00260
<i>Neisseria</i>	sp.		18	Phylotype	5	T-467	0.000144	0.00217
<i>Neisseria</i>	sp.		20	Unnamed	5	T-467	0.000144	0.00217
<i>Treponema</i>	<i>vincentii</i>		29	Named	5	T-467	0.000144	0.00217
<i>Oribacterium</i>	sp.		78	Unnamed	5	T-467	0.000144	0.00217
<i>Peptostreptococcaceae</i> [XII][G-2]	sp.		91	Phylotype	5	T-467	0.000144	0.00217

<i>Treponema</i>	sp.		253	Phylotype	5	T-467	0.000144	0.00217
<i>Treponema</i>	sp.		256	Phylotype	5	T-467	0.000144	0.00217
<i>Treponema</i>	sp.		263	Phylotype	5	T-467	0.000144	0.00217
<i>Porphyromonas</i>	sp.		278	Phylotype	5	T-467	0.000144	0.00217
<i>Porphyromonas</i>	sp.		284	Unnamed	5	T-467	0.000144	0.00217
<i>Prevotella</i>	sp.		304	Phylotype	5	T-467	0.000144	0.00217
<i>TM7 [G-1]</i>	sp.		353	Phylotype	5	T-467	0.000144	0.00217
<i>Leptotrichia</i>	sp.		392	Phylotype	5	T-467	0.000144	0.00217
<i>Pseudomonas</i>	<i>stutzeri</i>		477	Named	5	T-467	0.000144	0.00217
<i>Lachnospiraceae [G-8]</i>	sp.		500	Phylotype	5	T-467	0.000144	0.00217
<i>Aggregatibacter</i>	sp.		512	Unnamed	5	T-467	0.000144	0.00217
<i>Treponema</i>	sp.		517	Phylotype	5	T-467	0.000144	0.00217
<i>Treponema</i>	sp.		518	Phylotype	5	T-467	0.000144	0.00217
<i>Haemophilus</i>	<i>aegyptius</i>		535	Named	5	T-467	0.000144	0.00217
<i>Prevotella</i>	<i>baroniae</i>		553	Named	5	T-467	0.000144	0.00217
<i>Mycoplasma</i>	<i>faucium</i>		606	Named	5	T-467	0.000144	0.00217

<i>Simonsiella</i>	<i>muelleri</i>	683	Named	5	T-467	0.000144	0.00217
<i>Jonquetella</i>	<i>anthropi</i>	777	Named	5	T-467	0.000144	0.00217
<i>Anaerococcus</i>	<i>lactolyticus</i>	A64	Named	5	T-467	0.000144	0.00217
<i>Brachybacterium</i>	sp.	A83	Phylotype	5	T-467	0.000144	0.00217
<i>Methylobacterium</i>	sp.	B53	Phylotype	5	T-467	0.000144	0.00217
<i>Kocuria</i>	sp.	B56	Phylotype	5	T-467	0.000144	0.00217
<i>Alphaproteobacteria [G]</i>	sp.	B69	Phylotype	5	T-467	0.000144	0.00217
<i>Actinomycetales [G]</i>	sp.	C05	Phylotype	5	T-467	0.000144	0.00217
<i>Finegoldia</i>	sp.	D24	Phylotype	5	T-467	0.000144	0.00217
<i>Actinomyces</i>	sp.	E91	Phylotype	5	T-467	0.000144	0.00217
<i>Bacillus</i>	sp.	F14	Phylotype	5	T-467	0.000144	0.00217
<i>Selenomonas</i>	sp.	F30	Phylotype	5	T-467	0.000144	0.00217
<i>Lachnospiraceae [G]</i>	sp.	F93	Phylotype	5	T-467	0.000144	0.00217
<i>Selenomonas</i>	sp.	F96	Phylotype	5	T-467	0.000144	0.00217
<i>Selenomonas</i>	sp.	F98	Phylotype	5	T-467	0.000144	0.00217
<i>Terrahaemophilus</i>	sp.	G25	Phylotype	5	T-467	0.000144	0.00217

<i>Veillonella</i>	sp.		G61	Phylotype	5	T-467	0.000144	0.00217
<i>Eubacterium [XII][G]</i>	sp.		G75	Phylotype	5	T-467	0.000144	0.00217
<i>Selenomonas</i>	sp.		H63	Phylotype	5	T-467	0.000144	0.00217
<i>Delftia</i>	<i>acidovorans</i>		23	Named	4	T-507	0.000115	0.00174
<i>Lachnospiraceae [G-5]</i>	sp.		80	Phylotype	4	T-507	0.000115	0.00174
<i>Lachnospiraceae [G-7]</i>	sp.		86	Phylotype	4	T-507	0.000115	0.00174
<i>Leptotrichia</i>	sp.		218	Phylotype	4	T-507	0.000115	0.00174
<i>Prevotella</i>	sp.		296	Phylotype	4	T-507	0.000115	0.00174
<i>Bergeyella</i>	sp.		322	Phylotype	4	T-507	0.000115	0.00174
<i>TM7 [G-1]</i>	sp.		348	Phylotype	4	T-507	0.000115	0.00174
<i>Peptoniphilus</i>	sp.		386	Unnamed	4	T-507	0.000115	0.00174
<i>Clostridiales [F-1][G-2]</i>	sp.		402	Phylotype	4	T-507	0.000115	0.00174
<i>Catonella</i>	sp.		451	Phylotype	4	T-507	0.000115	0.00174
<i>Prevotella</i>	sp.		474	Phylotype	4	T-507	0.000115	0.00174
<i>Bacteroidetes [G-5]</i>	sp.		511	Phylotype	4	T-507	0.000115	0.00174
<i>Aggregatibacter</i>	sp.		513	Phylotype	4	T-507	0.000115	0.00174

<i>Lactobacillus</i>	<i>acidophilus</i>	529	Named	4	T-507	0.000115	0.00174
<i>Treponema</i>	<i>amylovorum</i>	541	Named	4	T-507	0.000115	0.00174
<i>Afipia</i>	<i>sp. genomospecies 4</i>	636	Phylotype	4	T-507	0.000115	0.00174
<i>Prevotella</i>	<i>marshii</i>	665	Named	4	T-507	0.000115	0.00174
<i>Moraxella</i>	<i>osloensis</i>	711	Named	4	T-507	0.000115	0.00174
<i>Klebsiella</i>	<i>pneumoniae</i>	731	Named	4	T-507	0.000115	0.00174
<i>Pseudomonas</i>	<i>otitidis</i>	834	Named	4	T-507	0.000115	0.00174
<i>Lactobacillus</i>	<i>iners</i>	838	Named	4	T-507	0.000115	0.00174
<i>Peptoniphilus</i>	<i>indolicus</i>	840	Named	4	T-507	0.000115	0.00174
<i>Streptococcus</i>	<i>sp.</i>	A04	Phylotype	4	T-507	0.000115	0.00174
<i>Brevundimonas</i>	<i>vesicularis</i>	A14	Named	4	T-507	0.000115	0.00174
<i>Corynebacterium</i>	<i>sp.</i>	A16	Phylotype	4	T-507	0.000115	0.00174
<i>Sphingomonas</i>	<i>koreensis</i>	A37	Named	4	T-507	0.000115	0.00174
<i>Comomonadaceae [G]</i>	<i>sp.</i>	A82	Phylotype	4	T-507	0.000115	0.00174
<i>Micrococcus</i>	<i>sp.</i>	B64	Phylotype	4	T-507	0.000115	0.00174
<i>Alphaproteobacteria [G]</i>	<i>sp.</i>	B70	Phylotype	4	T-507	0.000115	0.00174

<i>Anoxybacillus</i>	<i>pushchinoensis</i>	B72	Named	4	T-507	0.000115	0.00174
<i>Betaproteobacteria [G]</i>	sp.	B96	Phylotype	4	T-507	0.000115	0.00174
<i>Dialister</i>	sp.	C18	Phylotype	4	T-507	0.000115	0.00174
<i>Bacilli [G]</i>	sp.	C35	Phylotype	4	T-507	0.000115	0.00174
<i>Streptococcus</i>	sp.	C97	Phylotype	4	T-507	0.000115	0.00174
<i>Acinetobacter</i>	sp.	C99	Phylotype	4	T-507	0.000115	0.00174
<i>Centipeda</i>	sp.	D18	Phylotype	4	T-507	0.000115	0.00174
<i>Aerococcus</i>	<i>christensenii</i>	D44	Named	4	T-507	0.000115	0.00174
<i>Centipeda</i>	sp.	D51	Phylotype	4	T-507	0.000115	0.00174
<i>Neisseria</i>	sp.	D61	Phylotype	4	T-507	0.000115	0.00174
<i>Streptococcus</i>	sp.	D84	Phylotype	4	T-507	0.000115	0.00174
<i>Streptococcus</i>	sp.	E18	Phylotype	4	T-507	0.000115	0.00174
<i>Porphyromonas</i>	sp.	E21	Phylotype	4	T-507	0.000115	0.00174
<i>Fusobacterium</i>	sp.	E22	Phylotype	4	T-507	0.000115	0.00174
<i>Kingella</i>	sp.	E73	Phylotype	4	T-507	0.000115	0.00174
<i>Selenomonas</i>	sp.	F33	Phylotype	4	T-507	0.000115	0.00174

<i>Paenibacillus</i>	sp.		F45	Phylotype	4	T-507	0.000115	0.00174
<i>Selenomonas</i>	sp.		F82	Phylotype	4	T-507	0.000115	0.00174
<i>Capnocytophaga</i>	sp.		F95	Phylotype	4	T-507	0.000115	0.00174
<i>Veillonella</i>	sp.		G30	Phylotype	4	T-507	0.000115	0.00174
<i>Veillonella</i>	sp.		G38	Phylotype	4	T-507	0.000115	0.00174
<i>Treponema</i>	sp.		G53	Phylotype	4	T-507	0.000115	0.00174
<i>Streptococcus</i>	sp.		G63	Phylotype	4	T-507	0.000115	0.00174
<i>Mitsuokella</i>	sp.		G68	Phylotype	4	T-507	0.000115	0.00174
<i>Selenomonas</i>	sp.		H30	Phylotype	4	T-507	0.000115	0.00174
<i>Selenomonas</i>	sp.		H65	Phylotype	4	T-507	0.000115	0.00174
<i>Desulfovibrio</i>	sp.		40	Phylotype	3	T-562	0.000086	0.00130
<i>Staphylococcus</i>	<i>warneri</i>		76	Named	3	T-562	0.000086	0.00130
<i>Peptostreptococcaceae [XI][G-7]</i>	sp.		81	Phylotype	3	T-562	0.000086	0.00130
<i>Lachnospiraceae [G-1]</i>	sp.		83	Phylotype	3	T-562	0.000086	0.00130
<i>Selenomonas</i>	sp.		136	Unnamed	3	T-562	0.000086	0.00130

<i>Treponema</i>	sp.		235	Phylotype	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.		262	Phylotype	3	T-562	0.000086	0.00130
<i>Flavobacteriales [G-I]</i>	sp.		321	Phylotype	3	T-562	0.000086	0.00130
<i>Selenomonas</i>	sp.		388	Phylotype	3	T-562	0.000086	0.00130
<i>Syntrophomonadaceae [VIII][G-I]</i>	sp.		435	Phylotype	3	T-562	0.000086	0.00130
<i>Streptococcus</i>	sp.		486	Unnamed	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.		490	Phylotype	3	T-562	0.000086	0.00130
<i>Bacteroidetes [G-5]</i>	sp.		505	Phylotype	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.		508	Phylotype	3	T-562	0.000086	0.00130
<i>Aggregatibacter</i>	<i>aphrophilus</i>		545	Named	3	T-562	0.000086	0.00130
<i>Staphylococcus</i>	<i>caprae</i>		567	Named	3	T-562	0.000086	0.00130
<i>Staphylococcus</i>	<i>epidermidis</i>		601	Named	3	T-562	0.000086	0.00130
<i>Rhizobium</i>	<i>loti</i>		659	Named	3	T-562	0.000086	0.00130
<i>Mogibacterium</i>	<i>neglectum</i>		691	Named	3	T-562	0.000086	0.00130
<i>Mogibacterium</i>	<i>pumilum</i>		742	Named	3	T-562	0.000086	0.00130

<i>Mycoplasma</i>	<i>salivarium</i>	754	Named	3	T-562	0.000086	0.00130
<i>Lactobacillus</i>	<i>colehominis</i>	816	Named	3	T-562	0.000086	0.00130
<i>Rhodobacter</i>	<i>capsulatus</i>	A00	Named	3	T-562	0.000086	0.00130
<i>Paracoccus</i>	<i>marcusii</i>	A21	Named	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.	A33	Phylotype	3	T-562	0.000086	0.00130
<i>Bacillus</i>	sp.	A51	Phylotype	3	T-562	0.000086	0.00130
<i>Leuconostoc</i>	<i>pseudomesenteroides</i>	A60	Named	3	T-562	0.000086	0.00130
<i>Stenotrophomonas</i>	sp.	A62	Phylotype	3	T-562	0.000086	0.00130
<i>Alphaproteobacteria [G]</i>	sp.	A67	Phylotype	3	T-562	0.000086	0.00130
<i>Actinomyces</i>	<i>cardiffensis</i>	A75	Named	3	T-562	0.000086	0.00130
<i>Pseudomonas</i>	<i>mosselii</i>	A88	Named	3	T-562	0.000086	0.00130
<i>Parvimonas</i>	sp.	A91	Phylotype	3	T-562	0.000086	0.00130
<i>Peptostreptococcaceae</i>	sp.	B17	Phylotype	3	T-562	0.000086	0.00130
[XIII][G]							
<i>Lachnospiraceae</i>	sp.	B32	Phylotype	3	T-562	0.000086	0.00130
<i>Atopostipes</i>	sp.	B55	Phylotype	3	T-562	0.000086	0.00130

<i>Brevibacterium</i>	<i>casei</i>	B88	Named	3	T-562	0.000086	0.00130
<i>Porphyromonas</i>	sp.	B94	Phylotype	3	T-562	0.000086	0.00130
<i>Pseudomonas</i>	<i>anguilliseptica</i>	C06	Named	3	T-562	0.000086	0.00130
<i>Clostridiales [G]</i>	sp.	C07	Phylotype	3	T-562	0.000086	0.00130
<i>Microbacterium</i>	sp.	C11	Phylotype	3	T-562	0.000086	0.00130
<i>Betaproteobacteria [G]</i>	sp.	C12	Phylotype	3	T-562	0.000086	0.00130
<i>Propionivibrio</i>	sp.	C33	Phylotype	3	T-562	0.000086	0.00130
<i>Rhizobium</i>	sp.	C41	Phylotype	3	T-562	0.000086	0.00130
<i>Erysipelotrichales [G]</i>	sp.	C62	Phylotype	3	T-562	0.000086	0.00130
<i>Dialister</i>	sp.	C66	Phylotype	3	T-562	0.000086	0.00130
<i>Arsenicococcus</i>	sp.	C74	Phylotype	3	T-562	0.000086	0.00130
<i>Ralstonia</i>	sp.	C76	Phylotype	3	T-562	0.000086	0.00130
<i>Methylobacterium</i>	<i>rhodesianum</i> ;	C94	Named	3	T-562	0.000086	0.00130
<i>Enterobacter</i>	sp.	C98	Phylotype	3	T-562	0.000086	0.00130
<i>Janibacter</i>	sp.	D01	Phylotype	3	T-562	0.000086	0.00130
<i>Granulicatella</i>	sp.	D03	Phylotype	3	T-562	0.000086	0.00130

<i>Gemella</i>	sp.		D42	Phylotype	3	T-562	0.000086	0.00130
<i>Acinetobacter</i>	sp.		D46	Phylotype	3	T-562	0.000086	0.00130
<i>Kingella</i>	sp.		D49	Phylotype	3	T-562	0.000086	0.00130
<i>Filifactor</i>	sp.		D70	Phylotype	3	T-562	0.000086	0.00130
<i>Selenomonas</i>	sp.		D75	Phylotype	3	T-562	0.000086	0.00130
<i>Veillonella</i>	sp.		D88	Phylotype	3	T-562	0.000086	0.00130
<i>Streptococcus</i>	sp.		E19	Phylotype	3	T-562	0.000086	0.00130
<i>Clostridiales [G]</i>	sp.		E35	Phylotype	3	T-562	0.000086	0.00130
<i>Peptostreptococcaceae</i> [XIII][G]	sp.		E45	Phylotype	3	T-562	0.000086	0.00130
<i>Kingella</i>	sp.		E51	Phylotype	3	T-562	0.000086	0.00130
<i>Clostridia [G]</i>	sp.		E60	Phylotype	3	T-562	0.000086	0.00130
<i>Neisseria</i>	sp.		E61	Phylotype	3	T-562	0.000086	0.00130
<i>Selenomonas</i>	sp.		E80	Phylotype	3	T-562	0.000086	0.00130
<i>Selenomonas</i>	sp.		F09	Phylotype	3	T-562	0.000086	0.00130
<i>Bacillus</i>	sp.		F16	Phylotype	3	T-562	0.000086	0.00130

<i>Bacteroidetes [G]</i>	sp.		F31	Phylotype	3	T-562	0.000086	0.00130
<i>Selenomonas</i>	sp.		F83	Phylotype	3	T-562	0.000086	0.00130
<i>Selenomonas</i>	sp.		F84	Phylotype	3	T-562	0.000086	0.00130
<i>Streptococcus</i>	sp.		F97	Phylotype	3	T-562	0.000086	0.00130
<i>Lachnospiraceae [XIVa][G]</i>	sp.		G33	Phylotype	3	T-562	0.000086	0.00130
<i>Capnocytophaga</i>	sp.		G34	Phylotype	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.		G41	Phylotype	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.		G49	Phylotype	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.		G52	Phylotype	3	T-562	0.000086	0.00130
<i>Prevotella</i>	sp.		G54	Phylotype	3	T-562	0.000086	0.00130
<i>Prevotella</i>	sp.		G64	Phylotype	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.		G86	Phylotype	3	T-562	0.000086	0.00130
<i>Treponema</i>	sp.		G87	Phylotype	3	T-562	0.000086	0.00130
<i>Parvimonas</i>	sp.		H07	Phylotype	3	T-562	0.000086	0.00130
<i>Capnocytophaga</i>	sp.		H18	Phylotype	3	T-562	0.000086	0.00130
<i>Cardiobacterium</i>	sp.		H38	Phylotype	3	T-562	0.000086	0.00130

<i>Micrococcus</i>	<i>luteus</i>	H68	Named	3	T-562	0.000086	0.00130
<i>Caulobacter</i>	sp.	2	Phylotype	2	T-645	0.000058	0.00087
<i>Sphingomonas</i>	sp.	3	Phylotype	2	T-645	0.000058	0.00087
<i>Sphingomonas</i>	sp.	7	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus</i>	sp.	55	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus</i>	sp.	67	Phylotype	2	T-645	0.000058	0.00087
<i>Lachnospiraceae [G-1]</i>	sp.	89	Phylotype	2	T-645	0.000058	0.00087
<i>Butyrivibrio</i>	sp.	94	Phylotype	2	T-645	0.000058	0.00087
<i>Peptostreptococcaceae [XII][G-7]</i>	sp.	106	Unnamed	2	T-645	0.000058	0.00087
<i>Lachnospiraceae [G-7]</i>	sp.	163	Phylotype	2	T-645	0.000058	0.00087
<i>Peptococcus</i>	sp.	168	Phylotype	2	T-645	0.000058	0.00087
<i>Microbacterium</i>	sp.	185	Phylotype	2	T-645	0.000058	0.00087
<i>Kocuria</i>	sp.	189	Phylotype	2	T-645	0.000058	0.00087
<i>Leptotrichia</i>	sp.	217	Phylotype	2	T-645	0.000058	0.00087
<i>Leptotrichia</i>	sp.	223	Phylotype	2	T-645	0.000058	0.00087

<i>Treponema</i>	sp.		227	Phylotype	2	T-645	0.000058	0.00087
<i>Treponema</i>	sp.		252	Phylotype	2	T-645	0.000058	0.00087
<i>Treponema</i>	sp.		270	Phylotype	2	T-645	0.000058	0.00087
<i>Treponema</i>	sp.		271	Phylotype	2	T-645	0.000058	0.00087
<i>Porphyromonas</i>	sp.		275	Phylotype	2	T-645	0.000058	0.00087
<i>Porphyromonas</i>	sp.		277	Phylotype	2	T-645	0.000058	0.00087
<i>Prevotella</i>	sp.		302	Unnamed	2	T-645	0.000058	0.00087
<i>Prevotella</i>	sp.		315	Phylotype	2	T-645	0.000058	0.00087
<i>Flavobacteriales [G-1]</i>	sp.		318	Phylotype	2	T-645	0.000058	0.00087
<i>Flavobacteriales [G-2]</i>	sp.		320	Phylotype	2	T-645	0.000058	0.00087
<i>Capnocytophaga</i>	sp.		332	Unnamed	2	T-645	0.000058	0.00087
<i>Capnocytophaga</i>	sp.		335	Unnamed	2	T-645	0.000058	0.00087
<i>TM7 [G-2]</i>	sp.		350	Phylotype	2	T-645	0.000058	0.00087
<i>Clostridiales [F-2][G-3]</i>	sp.		366	Phylotype	2	T-645	0.000058	0.00087
<i>Clostridiales [F-2][G-3]</i>	sp.		381	Phylotype	2	T-645	0.000058	0.00087
<i>Porphyromonas</i>	sp.		395	Phylotype	2	T-645	0.000058	0.00087

<i>Acinetobacter</i>	sp.	408	Phylotype	2	T-645	0.000058	0.00087
<i>Actinomyces</i>	sp.	449	Phylotype	2	T-645	0.000058	0.00087
<i>Veillonellaceae [G-1]</i>	sp.	483	Phylotype	2	T-645	0.000058	0.00087
<i>Peptostreptococcaceae [XI][G-3]</i>	sp.	495	Phylotype	2	T-645	0.000058	0.00087
<i>Selenomonas</i>	sp.	501	Phylotype	2	T-645	0.000058	0.00087
<i>Bacteroidetes [G-6]</i>	sp.	516	Phylotype	2	T-645	0.000058	0.00087
<i>Neisseria</i>	sp.	523	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus agalactiae</i>		537	Named	2	T-645	0.000058	0.00087
<i>Propionibacterium avidum</i>		552	Named	2	T-645	0.000058	0.00087
<i>Afipia broomeae</i>		559	Named	2	T-645	0.000058	0.00087
<i>Enterobacter cancerogenus</i>		565	Named	2	T-645	0.000058	0.00087
<i>Prevotella dentalis</i>		583	Named	2	T-645	0.000058	0.00087
<i>Mogibacterium diversum</i>		593	Named	2	T-645	0.000058	0.00087
<i>Slackia exigua</i>		602	Named	2	T-645	0.000058	0.00087
<i>Bacteroides heparinolyticus</i>		630	Named	2	T-645	0.000058	0.00087

<i>Prevotella</i>	<i>multiformis</i>	685	Named	2	T-645	0.000058	0.00087
<i>Eubacterium [XII][G-6]</i>	<i>nodatum</i>	694	Named	2	T-645	0.000058	0.00087
<i>Desulfomicrobium</i>	<i>orale</i>	703	Named	2	T-645	0.000058	0.00087
<i>Erythromicrobium</i>	<i>ramosum</i>	747	Named	2	T-645	0.000058	0.00087
<i>Clostridiales [G]</i>	sp.	A10	Phylotype	2	T-645	0.000058	0.00087
<i>Stenotrophomonas</i>	sp.	A24	Phylotype	2	T-645	0.000058	0.00087
<i>Herbaspirillum</i>	sp.	A32	Phylotype	2	T-645	0.000058	0.00087
<i>Corynebacterium</i>	<i>simulans</i>	A35	Named	2	T-645	0.000058	0.00087
<i>Firmicutes [G]</i>	sp.	A55	Phylotype	2	T-645	0.000058	0.00087
<i>Delftia</i>	sp.	A59	Phylotype	2	T-645	0.000058	0.00087
<i>Pseudomonas</i>	<i>stutzeri</i>	A68	Named	2	T-645	0.000058	0.00087
<i>Burkholderia</i>	sp.	A72	Phylotype	2	T-645	0.000058	0.00087
<i>Moraxella</i>	sp.	B07	Phylotype	2	T-645	0.000058	0.00087
<i>Bifidobacterium</i>	<i>longum</i>	B54	Named	2	T-645	0.000058	0.00087
<i>Eubacterium [XII][G]</i>	sp.	B60	Phylotype	2	T-645	0.000058	0.00087
<i>Nocardioopsis</i>	sp.	B75	Phylotype	2	T-645	0.000058	0.00087

<i>Comamonadaceae [G]</i>	sp.		B76	Phylotype	2	T-645	0.000058	0.00087
<i>Comonadaceae [G]</i>	sp.		B87	Phylotype	2	T-645	0.000058	0.00087
<i>Microbacterium</i>	<i>hominis</i>		C00	Named	2	T-645	0.000058	0.00087
<i>Bacillus</i>	sp.		C03	Phylotype	2	T-645	0.000058	0.00087
<i>Clostridiales [G]</i>	sp.		C20	Phylotype	2	T-645	0.000058	0.00087
<i>Rhodobacter</i>	sp.		C30	Phylotype	2	T-645	0.000058	0.00087
<i>Deinococcus</i>	sp.		C45	Phylotype	2	T-645	0.000058	0.00087
<i>Clostridiales [G]</i>	sp.		C47	Phylotype	2	T-645	0.000058	0.00087
<i>Bacillus</i>	sp.		C52	Phylotype	2	T-645	0.000058	0.00087
<i>Brevundimonas</i>	<i>vesicularis</i>		C59	Named	2	T-645	0.000058	0.00087
<i>Methylobacterium</i>	sp.		C71	Phylotype	2	T-645	0.000058	0.00087
<i>Pseudomonas</i>	sp.		C85	Phylotype	2	T-645	0.000058	0.00087
<i>Alphaproteobacteria [G]</i>	sp.		D00	Phylotype	2	T-645	0.000058	0.00087
<i>Leptothrix</i>	sp.		D13	Phylotype	2	T-645	0.000058	0.00087
<i>Rubrobacter</i>	<i>xylanophilus</i>		D32	Named	2	T-645	0.000058	0.00087
<i>Rhizobium</i>	<i>rhizogenes</i>		D34	Named	2	T-645	0.000058	0.00087

<i>Sanguibacter</i>	<i>keddieii</i>	D35	Named	2	T-645	0.000058	0.00087
<i>Synergistetes [G]</i>	sp.	D48	Phylotype	2	T-645	0.000058	0.00087
<i>Kingella</i>	sp.	D55	Phylotype	2	T-645	0.000058	0.00087
<i>Neisseria</i>	sp.	D56	Phylotype	2	T-645	0.000058	0.00087
<i>Veillonella</i>	sp.	D89	Phylotype	2	T-645	0.000058	0.00087
<i>Porphyromonas</i>	sp.	D90	Phylotype	2	T-645	0.000058	0.00087
<i>Granulicatella</i>	sp.	E03	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus</i>	sp.	E06	Phylotype	2	T-645	0.000058	0.00087
<i>Prevotella</i>	sp.	E13	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus</i>	sp.	E28	Phylotype	2	T-645	0.000058	0.00087
<i>Actinomyces</i>	sp.	E33	Phylotype	2	T-645	0.000058	0.00087
<i>Megasphaera</i>	sp.	E34	Phylotype	2	T-645	0.000058	0.00087
<i>Veillonellaceae [G-I]</i>	sp.	E37	Phylotype	2	T-645	0.000058	0.00087
<i>Dialister</i>	sp.	E47	Phylotype	2	T-645	0.000058	0.00087
<i>Aggregatibacter</i>	sp.	E49	Phylotype	2	T-645	0.000058	0.00087
<i>Selenomonas</i>	sp.	E57	Phylotype	2	T-645	0.000058	0.00087

<i>Clostridiales [F-I][G]</i>	sp.		E58	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus</i>	sp.		E66	Phylotype	2	T-645	0.000058	0.00087
<i>Selenomonas</i>	sp.		E76	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus</i>	sp.		E85	Phylotype	2	T-645	0.000058	0.00087
<i>Selenomonas</i>	sp.		E97	Phylotype	2	T-645	0.000058	0.00087
<i>Sphingomonas</i>	sp.		F02	Phylotype	2	T-645	0.000058	0.00087
<i>Tessaracoccus</i>	sp.		F04	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus</i>	sp.		F10	Phylotype	2	T-645	0.000058	0.00087
<i>Alphaproteobacteria [G]</i>	sp.		F12	Phylotype	2	T-645	0.000058	0.00087
<i>Veillonellaceae [G-I]</i>	sp.		F24	Phylotype	2	T-645	0.000058	0.00087
<i>Atopobium</i>	sp.		F34	Phylotype	2	T-645	0.000058	0.00087
<i>Paracoccus</i>	<i>zeaxanthinifaciens</i>		F59	Named	2	T-645	0.000058	0.00087
<i>Bacteroidetes [G]</i>	sp.		F60	Phylotype	2	T-645	0.000058	0.00087
<i>Paracoccus</i>	sp.		F63	Phylotype	2	T-645	0.000058	0.00087
<i>Synergistetes [G]</i>	sp.		F67	Phylotype	2	T-645	0.000058	0.00087
<i>Bacillus</i>	sp.		F79	Phylotype	2	T-645	0.000058	0.00087

<i>Peptococcus</i>	sp.		F94	Phylotype	2	T-645	0.000058	0.00087
<i>Aggregatibacter</i>	sp.		G01	Phylotype	2	T-645	0.000058	0.00087
<i>Haemophilus</i>	sp.		G13	Phylotype	2	T-645	0.000058	0.00087
<i>TM7 [G-1]</i>	sp.		G29	Phylotype	2	T-645	0.000058	0.00087
<i>Synergistetes [G-3]</i>	sp.		G36	Phylotype	2	T-645	0.000058	0.00087
<i>Treponema</i>	sp.		G48	Phylotype	2	T-645	0.000058	0.00087
<i>Streptococcus</i>	sp.		G59	Phylotype	2	T-645	0.000058	0.00087
<i>Clostridiales [G]</i>	sp.		G74	Phylotype	2	T-645	0.000058	0.00087
<i>Treponema</i>	sp.		G76	Phylotype	2	T-645	0.000058	0.00087
<i>Selenomonas</i>	sp.		G78	Phylotype	2	T-645	0.000058	0.00087
<i>Campylobacter</i>	sp.		G90	Phylotype	2	T-645	0.000058	0.00087
<i>Lactobacillus</i>	sp.		G98	Phylotype	2	T-645	0.000058	0.00087
<i>Veillonella</i>	sp.		H02	Phylotype	2	T-645	0.000058	0.00087
<i>Parvimonas</i>	sp.		H16	Phylotype	2	T-645	0.000058	0.00087
<i>Prevotella</i>	sp.		H59	Phylotype	2	T-645	0.000058	0.00087
<i>Sphingomonas</i>	sp.		4	Phylotype	1	T-769	0.000029	0.00043

<i>Sphingomonas</i>	sp.	6	Phylotype	1	T-769	0.000029	0.00043
<i>Eikenella</i>	sp.	11	Unnamed	1	T-769	0.000029	0.00043
<i>Bdellovibrio</i>	sp.	39	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	69	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G-2]</i>	sp.	88	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G-6]</i>	sp.	90	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.	143	Phylotype	1	T-769	0.000029	0.00043
<i>Microbacterium</i>	sp.	186	Unnamed	1	T-769	0.000029	0.00043
<i>Propionibacterium</i>	sp.	193	Unnamed	1	T-769	0.000029	0.00043
<i>Propionibacterium</i>	sp.	194	Phylotype	1	T-769	0.000029	0.00043
<i>Atopobium</i>	sp.	199	Unnamed	1	T-769	0.000029	0.00043
<i>Fusobacterium</i>	sp.	205	Phylotype	1	T-769	0.000029	0.00043
<i>Fusobacteria [G-1]</i>	sp.	210	Phylotype	1	T-769	0.000029	0.00043
<i>Leptotrichia</i>	sp.	221	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	226	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	242	Phylotype	1	T-769	0.000029	0.00043

<i>Treponema</i>	sp.		246	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		249	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		250	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		255	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		258	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		260	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		293	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		301	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		305	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		309	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		310	Phylotype	1	T-769	0.000029	0.00043
<i>Capnocytophaga</i>	sp.		324	Unnamed	1	T-769	0.000029	0.00043
<i>Capnocytophaga</i>	sp.		334	Phylotype	1	T-769	0.000029	0.00043
<i>TM7 [G-3]</i>	sp.		351	Phylotype	1	T-769	0.000029	0.00043
<i>TM7 [G-4]</i>	sp.		355	Phylotype	1	T-769	0.000029	0.00043
<i>TM7 [G-5]</i>	sp.		356	Phylotype	1	T-769	0.000029	0.00043

<i>Capnocytophaga</i>	sp.		380	Unnamed	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		396	Phylotype	1	T-769	0.000029	0.00043
<i>Bifidobacteriaceae [G-2]</i>	sp.		407	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		423	Unnamed	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G-3]</i>	sp.		436	Phylotype	1	T-769	0.000029	0.00043
<i>Chloroflexi [G-1]</i>	sp.		439	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		443	Phylotype	1	T-769	0.000029	0.00043
<i>Synergistetes [G-3]</i>	sp.		452	Phylotype	1	T-769	0.000029	0.00043
<i>Synergistetes [G-3]</i>	sp.		453	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G-5]</i>	sp.		455	Phylotype	1	T-769	0.000029	0.00043
<i>Kingella</i>	sp.		459	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		475	Unnamed	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		487	Phylotype	1	T-769	0.000029	0.00043
<i>TM7 [G-1]</i>	sp.		488	Phylotype	1	T-769	0.000029	0.00043
<i>Peptostreptococcaceae [XI][G-5]</i>	sp.		493	Phylotype	1	T-769	0.000029	0.00043

<i>Lachnospiraceae [G-1]</i>	sp.	496	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G-3]</i>	sp.	503	Phylotype	1	T-769	0.000029	0.00043
<i>Tenericutes [G-1]</i>	sp.	504	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G-4]</i>	sp.	509	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.	515	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	<i>brevis</i>	558	Named	1	T-769	0.000029	0.00043
<i>Prevotella</i>	<i>veroralis</i>	572	Named	1	T-769	0.000029	0.00043
<i>Actinomyces</i>	<i>georgiae</i>	617	Named	1	T-769	0.000029	0.00043
<i>Mycoplasma</i>	<i>hominis</i>	632	Named	1	T-769	0.000029	0.00043
<i>Afpia</i>	Sp.	652	Phylotype	1	T-769	0.000029	0.00043
<i>Mycobacterium</i>	<i>neoaurum</i>	692	Named	1	T-769	0.000029	0.00043
<i>Actinomyces</i>	<i>oricola</i>	708	Named	1	T-769	0.000029	0.00043
<i>Pseudomonas</i>	<i>pseudoalcaligenes</i>	740	Named	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.	782	Unnamed	1	T-769	0.000029	0.00043
<i>Anaerococcus</i>	<i>tetradius</i>	788	Named	1	T-769	0.000029	0.00043
<i>Peptostreptococcaceae</i>	sp.	790	Phylotype	1	T-769	0.000029	0.00043

[XIII][G-2]												
<i>Tannerella</i>	sp.			808	Phylotype	1	T-769	0.000029	0.000029	0.00043		
<i>Atopobium</i>	sp			810	Unnamed	1	T-769	0.000029	0.000029	0.00043		
<i>Lactobacillus</i>	<i>johnsonii</i>			819	Named	1	T-769	0.000029	0.000029	0.00043		
<i>Gardnerella</i>	<i>vaginalis</i>			829	Named	1	T-769	0.000029	0.000029	0.00043		
<i>Sphingomonas</i>	<i>aromaticivorans</i>			A01	Named	1	T-769	0.000029	0.000029	0.00043		
<i>Paenibacillus</i>	sp.			A06	Phylotype	1	T-769	0.000029	0.000029	0.00043		
<i>Sphingomonas</i>	sp.			A09	Phylotype	1	T-769	0.000029	0.000029	0.00043		
<i>Shewanella</i>	<i>algae</i>			A12	Named	1	T-769	0.000029	0.000029	0.00043		
<i>Veillonella</i>	sp.			A15	Phylotype	1	T-769	0.000029	0.000029	0.00043		
<i>Erysipelothrichaceae [G]</i>	sp.			A18	Phylotype	1	T-769	0.000029	0.000029	0.00043		
<i>Agrococcus</i>	<i>baldiri</i>			A19	Named	1	T-769	0.000029	0.000029	0.00043		
<i>Rickettsiella</i>	<i>melolonthae</i>			A20	Named	1	T-769	0.000029	0.000029	0.00043		
<i>Corynebacterium</i>	sp.			A22	Phylotype	1	T-769	0.000029	0.000029	0.00043		
<i>Betaproteobacteria [G]</i>	sp.			A25	Phylotype	1	T-769	0.000029	0.000029	0.00043		
<i>Treponema</i>	sp.			A26	Phylotype	1	T-769	0.000029	0.000029	0.00043		

<i>Alphaproteobacteria [G]</i>	sp.	A28	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	A30	Phylotype	1	T-769	0.000029	0.00043
<i>Rhizobium</i>	<i>cellulosthyticus</i>	A34	Named	1	T-769	0.000029	0.00043
<i>Klebsiella</i>	sp.	A36	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	A38	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	A39	Phylotype	1	T-769	0.000029	0.00043
<i>Bifidobacterium</i>	<i>scardovii</i>	A40	Named	1	T-769	0.000029	0.00043
<i>Fusobacterium</i>	sp.	A42	Phylotype	1	T-769	0.000029	0.00043
<i>Staphylococcus</i>	<i>pettenkoferi</i>	A44	Named	1	T-769	0.000029	0.00043
<i>Leptotrichia</i>	sp.	A45	Phylotype	1	T-769	0.000029	0.00043
<i>Corynebacterium</i>	sp.	A46	Phylotype	1	T-769	0.000029	0.00043
<i>Capnocytophaga</i>	sp.	A48	Phylotype	1	T-769	0.000029	0.00043
<i>Acidobacteriaceae [G]</i>	sp.	A52	Phylotype	1	T-769	0.000029	0.00043
<i>Alphaproteobacteria [G]</i>	sp.	A53	Phylotype	1	T-769	0.000029	0.00043
<i>Paenibacillus</i>	sp.	A54	Phylotype	1	T-769	0.000029	0.00043
<i>TM7 [G-2]</i>	sp.	A56	Phylotype	1	T-769	0.000029	0.00043

<i>Clostridiales [G]</i>	sp.	A63	Phylotype	1	T-769	0.000029	0.00043
<i>Kocuria</i>	<i>polaris</i>	A70	Named	1	T-769	0.000029	0.00043
<i>Alphaproteobacteria [G]</i>	sp.	A73	Phylotype	1	T-769	0.000029	0.00043
<i>Sphingomonas</i>	sp.	A74	Phylotype	1	T-769	0.000029	0.00043
<i>Malikia</i>	<i>granosa</i>	A79	Named	1	T-769	0.000029	0.00043
<i>Propionibacterium</i>	<i>acidipropionici</i>	A80	Named	1	T-769	0.000029	0.00043
<i>Anaerococcus</i>	sp.	A86	Phylotype	1	T-769	0.000029	0.00043
<i>Peptoniphilus</i>	sp.	A87	Phylotype	1	T-769	0.000029	0.00043
<i>Hafnia</i>	sp.	A90	Phylotype	1	T-769	0.000029	0.00043
<i>Acinetobacter</i>	sp.	A92	Phylotype	1	T-769	0.000029	0.00043
<i>Clostridiales [G]</i>	sp.	A96	Phylotype	1	T-769	0.000029	0.00043
<i>Centipeda</i>	sp.	B01	Phylotype	1	T-769	0.000029	0.00043
<i>Catonella</i>	sp.	B06	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G]</i>	sp.	B18	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonellaceae [G-1]</i>	sp.	B19	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.	B30	Phylotype	1	T-769	0.000029	0.00043

<i>Porphyromonas</i>	sp.		B43	Phylotype	1	T-769	0.000029	0.00043
<i>Granulicatella</i>	sp.		B44	Phylotype	1	T-769	0.000029	0.00043
<i>Herminiimonas</i>	<i>saxobsidens</i>		B51	Named	1	T-769	0.000029	0.00043
<i>Synergistetes [G]</i>	sp.		B52	Phylotype	1	T-769	0.000029	0.00043
<i>Leptotrichia</i>	sp.		B57	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		B58	Phylotype	1	T-769	0.000029	0.00043
<i>Peptostreptococcaceae [XI][G]</i>	sp.		B61	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		B62	Phylotype	1	T-769	0.000029	0.00043
<i>Dialister</i>	sp.		B63	Phylotype	1	T-769	0.000029	0.00043
<i>Sphingomonas</i>	sp.		B65	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G]</i>	sp.		B68	Phylotype	1	T-769	0.000029	0.00043
<i>Corynebacterium</i>	sp.		B71	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.		B73	Phylotype	1	T-769	0.000029	0.00043
<i>Guggenheimella</i>	<i>bovis</i>		B74	Named	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.		B80	Phylotype	1	T-769	0.000029	0.00043
<i>Sphingomonas</i>	sp.		B83	Phylotype	1	T-769	0.000029	0.00043

<i>Methylobacterium</i>	sp.		B84	Phylotype	1	T-769	0.000029	0.00043
<i>Kluyvera</i>	sp.		B86	Phylotype	1	T-769	0.000029	0.00043
<i>Bacillus</i>	sp.		B89	Phylotype	1	T-769	0.000029	0.00043
<i>Deftuvibacter</i>	sp.		B90	Phylotype	1	T-769	0.000029	0.00043
<i>Leptothrix</i>	sp.		B91	Phylotype	1	T-769	0.000029	0.00043
<i>Fusobacterium</i>	<i>gonidiaformans</i>		B92	Named	1	T-769	0.000029	0.00043
<i>Gemella</i>	sp.		B93	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		B95	Phylotype	1	T-769	0.000029	0.00043
<i>Burkholderia</i>	sp.		B97	Phylotype	1	T-769	0.000029	0.00043
<i>Burkholderiales [G]</i>	sp.		B98	Phylotype	1	T-769	0.000029	0.00043
<i>Devosia</i>	sp.		C01	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		C04	Phylotype	1	T-769	0.000029	0.00043
<i>Acinetobacter</i>	sp.		C13	Phylotype	1	T-769	0.000029	0.00043
<i>Parvimonas</i>	sp.		C15	Phylotype	1	T-769	0.000029	0.00043
<i>Kingella</i>	sp.		C21	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		C22	Phylotype	1	T-769	0.000029	0.00043

<i>Corynebacterium</i>	<i>urealyticum</i>	C28	Named	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.	C29	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.	C32	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroides [G]</i>	sp.	C36	Phylotype	1	T-769	0.000029	0.00043
<i>Actinomyces</i>	<i>radinae</i>	C37	Named	1	T-769	0.000029	0.00043
<i>Kocuria</i>	sp.	C39	Phylotype	1	T-769	0.000029	0.00043
<i>Pseudomonas</i>	sp.	C48	Phylotype	1	T-769	0.000029	0.00043
<i>Caulobacter</i>	sp.	C49	Phylotype	1	T-769	0.000029	0.00043
<i>Proteus</i>	sp.	C50	Phylotype	1	T-769	0.000029	0.00043
<i>Deltaproteobacteria [G]</i>	sp.	C53	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	C54	Phylotype	1	T-769	0.000029	0.00043
<i>Acinetobacter</i>	sp.	C57	Phylotype	1	T-769	0.000029	0.00043
<i>Parvimonas</i>	sp.	C63	Phylotype	1	T-769	0.000029	0.00043
<i>Megasphaera</i>	sp.	C64	Phylotype	1	T-769	0.000029	0.00043
<i>Clostridia [G]</i>	sp.	C67	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroides</i>	sp.	C69	Phylotype	1	T-769	0.000029	0.00043

<i>Burkholderia</i>	sp.	C72	Phylotype	1	T-769	0.000029	0.00043
<i>Flavoabacterium</i>	sp.	C73	Phylotype	1	T-769	0.000029	0.00043
<i>Actinomyces</i>	<i>denticolens</i>	C75	Named	1	T-769	0.000029	0.00043
<i>Cardiobacterium</i>	sp.	C77	Phylotype	1	T-769	0.000029	0.00043
<i>Brevundimonas</i>	sp.	C79	Phylotype	1	T-769	0.000029	0.00043
<i>Acinetobacter</i>	sp.	C80	Phylotype	1	T-769	0.000029	0.00043
<i>Acinetobacter</i>	sp.	C81	Phylotype	1	T-769	0.000029	0.00043
<i>Betaproteobacteria [G]</i>	sp.	C83	Phylotype	1	T-769	0.000029	0.00043
<i>Lautropia</i>	sp.	C89	Phylotype	1	T-769	0.000029	0.00043
<i>Bacillus</i>	sp.	C90	Phylotype	1	T-769	0.000029	0.00043
<i>Oxalobacteraceae [G]</i>	sp.	C91	Phylotype	1	T-769	0.000029	0.00043
<i>Microclunatus</i>	sp.	C95	Phylotype	1	T-769	0.000029	0.00043
<i>Alphaproteobacteria [G]</i>	sp.	C96	Phylotype	1	T-769	0.000029	0.00043
<i>Williamsia</i>	<i>serinedens</i>	D02	Named	1	T-769	0.000029	0.00043
<i>Actinomyces</i>	sp.	D05	Phylotype	1	T-769	0.000029	0.00043
<i>Calycanthus</i>	<i>floridus</i>	D07	Named	1	T-769	0.000029	0.00043

<i>Bacillaceae [G]</i>	sp.	D08	Phylotype	1	T-769	0.000029	0.00043
<i>Haemophilus</i>	sp.	D10	Phylotype	1	T-769	0.000029	0.00043
<i>Dietzia</i>	sp.	D12	Phylotype	1	T-769	0.000029	0.00043
<i>Microbacterium</i>	sp.	D16	Phylotype	1	T-769	0.000029	0.00043
<i>Mitsuokella</i>	sp.	D19	Phylotype	1	T-769	0.000029	0.00043
<i>Olsenella</i>	sp.	D22	Phylotype	1	T-769	0.000029	0.00043
<i>Brachybacterium</i>	sp.	D23	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroides</i>	sp.	D25	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G]</i>	sp.	D27	Phylotype	1	T-769	0.000029	0.00043
<i>Corynebacterium</i>	sp.	D28	Phylotype	1	T-769	0.000029	0.00043
<i>Acinetobacter</i>	sp.	D29	Phylotype	1	T-769	0.000029	0.00043
<i>Pinus</i>	<i>thunbergii</i>	D30	Named	1	T-769	0.000029	0.00043
<i>Triticum</i>	<i>aestivum</i>	D31	Named	1	T-769	0.000029	0.00043
<i>Gordonia</i>	<i>aichiensis</i>	D37	Named	1	T-769	0.000029	0.00043
<i>Paracoccus</i>	<i>solventivorans</i>	D39	Named	1	T-769	0.000029	0.00043
<i>Rhizobium</i>	<i>larrymoorei</i>	D45	Named	1	T-769	0.000029	0.00043

<i>Actinomyces</i>	sp.		D50	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		D53	Phylotype	1	T-769	0.000029	0.00043
<i>Rothia</i>	sp.		D54	Phylotype	1	T-769	0.000029	0.00043
<i>Pseudomonas</i>	sp.		D58	Phylotype	1	T-769	0.000029	0.00043
<i>Neisseria</i>	sp.		D59	Phylotype	1	T-769	0.000029	0.00043
<i>Corynebacterium</i>	sp.		D60	Phylotype	1	T-769	0.000029	0.00043
<i>Synergistetes [G]</i>	sp.		D62	Phylotype	1	T-769	0.000029	0.00043
<i>Megasphaera</i>	sp.		D63	Phylotype	1	T-769	0.000029	0.00043
<i>Synergistetes [G]</i>	sp.		D64	Phylotype	1	T-769	0.000029	0.00043
<i>Synergistetes [G]</i>	sp.		D65	Phylotype	1	T-769	0.000029	0.00043
<i>Campylobacter</i>	sp.		D66	Phylotype	1	T-769	0.000029	0.00043
<i>Fusobacterium</i>	sp.		D68	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		D71	Phylotype	1	T-769	0.000029	0.00043
<i>Sphingomonas</i>	sp.		D72	Phylotype	1	T-769	0.000029	0.00043
<i>Firmicutes [G]</i>	sp.		D73	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G]</i>	sp.		D74	Phylotype	1	T-769	0.000029	0.00043

<i>Veillonellaceae [G-1]</i>	sp.		D76	Phylotype	1	T-769	0.000029	0.00043
<i>Atopobium</i>	sp.		D77	Phylotype	1	T-769	0.000029	0.00043
<i>Fusobacterium</i>	sp.		D78	Phylotype	1	T-769	0.000029	0.00043
<i>Actinomyces</i>	sp.		D79	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		D81	Phylotype	1	T-769	0.000029	0.00043
<i>Capnocytophaga</i>	sp.		D86	Phylotype	1	T-769	0.000029	0.00043
<i>Tannerella</i>	sp.		D91	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		D98	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		D99	Phylotype	1	T-769	0.000029	0.00043
<i>Dialister</i>	sp.		E00	Phylotype	1	T-769	0.000029	0.00043
<i>Peptococcus</i>	sp.		E01	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		E04	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		E05	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		E07	Phylotype	1	T-769	0.000029	0.00043
<i>Synergistetes [G-3]</i>	sp.		E08	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		E10	Phylotype	1	T-769	0.000029	0.00043

<i>Streptococcus</i>	sp.	E11	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E14	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E15	Phylotype	1	T-769	0.000029	0.00043
<i>Clostridiales [G]</i>	sp.	E16	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E17	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E23	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E24	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E25	Phylotype	1	T-769	0.000029	0.00043
<i>Granulicatella</i>	sp.	E26	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E27	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E30	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E38	Phylotype	1	T-769	0.000029	0.00043
<i>Actinomyces</i>	sp.	E40	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E41	Phylotype	1	T-769	0.000029	0.00043
<i>Peptostreptococcaceae [XII][G]</i>	sp.	E46	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G]</i>	sp.	E48	Phylotype	1	T-769	0.000029	0.00043

<i>Streptococcus</i>	sp.	E52	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	E53	Phylotype	1	T-769	0.000029	0.00043
<i>Cyanobacteria [G]</i>	sp.	E55	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.	E56	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G-8]</i>	sp.	E59	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.	E64	Phylotype	1	T-769	0.000029	0.00043
<i>Campylobacter</i>	sp.	E65	Phylotype	1	T-769	0.000029	0.00043
<i>Campylobacter</i>	sp.	E67	Phylotype	1	T-769	0.000029	0.00043
<i>Campylobacter</i>	sp.	E68	Phylotype	1	T-769	0.000029	0.00043
<i>Kingella</i>	sp.	E69	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	E70	Phylotype	1	T-769	0.000029	0.00043
<i>Porphyromonas</i>	sp.	E71	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E72	Phylotype	1	T-769	0.000029	0.00043
<i>Lautropia</i>	sp.	E74	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E77	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G]</i>	sp.	E79	Phylotype	1	T-769	0.000029	0.00043

<i>Corynebacterium</i>	sp.	E82	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.	E86	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E87	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	E88	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	E89	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	E90	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	E92	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	E93	Phylotype	1	T-769	0.000029	0.00043
<i>Paenibacillus</i>	sp.	E94	Phylotype	1	T-769	0.000029	0.00043
<i>Corynebacterium</i>	sp.	E95	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	E96	Phylotype	1	T-769	0.000029	0.00043
<i>Mitsuokella</i>	sp.	E99	Phylotype	1	T-769	0.000029	0.00043
<i>Cellulomonas</i>	sp.	F01	Phylotype	1	T-769	0.000029	0.00043
<i>Firmicutes [G]</i>	sp.	F03	Phylotype	1	T-769	0.000029	0.00043
<i>Lactococcus</i>	sp.	F05	Phylotype	1	T-769	0.000029	0.00043
<i>Firmicutes [G]</i>	sp.	F06	Phylotype	1	T-769	0.000029	0.00043

<i>Ralstonia</i>	sp.		F07	Phylotype	1	T-769	0.000029	0.00043
<i>Enterococcus</i>	sp.		F08	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		F11	Phylotype	1	T-769	0.000029	0.00043
<i>Gammaproteobacteria [G]</i>	sp.		F13	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G]</i>	sp.		F15	Phylotype	1	T-769	0.000029	0.00043
<i>Catonella</i>	sp.		F17	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		F18	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		F23	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		F25	Phylotype	1	T-769	0.000029	0.00043
<i>Bacillus</i>	sp.		F26	Phylotype	1	T-769	0.000029	0.00043
<i>Bacillus</i>	sp.		F27	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		F29	Phylotype	1	T-769	0.000029	0.00043
<i>Clostridiales [G]</i>	sp.		F32	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G]</i>	sp.		F44	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		F46	Phylotype	1	T-769	0.000029	0.00043
<i>Deinococcus</i>	sp.		F48	Phylotype	1	T-769	0.000029	0.00043

<i>Micrococcus</i>	sp.		F49	Phylotype	1	T-769	0.000029	0.00043
<i>Alphaproteobacteria [G]</i>	sp.		F50	Phylotype	1	T-769	0.000029	0.00043
<i>Alphaproteobacteria [G]</i>	sp.		F51	Phylotype	1	T-769	0.000029	0.00043
<i>Mogibacterium</i>	sp.		F52	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		F53	Phylotype	1	T-769	0.000029	0.00043
<i>Ornithinimicrobium</i>	sp.		F55	Phylotype	1	T-769	0.000029	0.00043
<i>Sphingomonas</i>	sp.		F56	Phylotype	1	T-769	0.000029	0.00043
<i>Erythromicrobium</i>	sp.		F62	Phylotype	1	T-769	0.000029	0.00043
<i>Lachnospiraceae [G]</i>	sp.		F64	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		F65	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		F68	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G]</i>	sp.		F69	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G]</i>	sp.		F70	Phylotype	1	T-769	0.000029	0.00043
<i>Centipeda</i>	sp.		F73	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		F74	Phylotype	1	T-769	0.000029	0.00043
<i>Rhodobacter</i>	sp.		F75	Phylotype	1	T-769	0.000029	0.00043

<i>Selenomonas</i>	sp.		F86	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		F87	Phylotype	1	T-769	0.000029	0.00043
<i>Neisseria</i>	sp.		F88	Phylotype	1	T-769	0.000029	0.00043
<i>Clostridiales [G]</i>	sp.		F89	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		F90	Phylotype	1	T-769	0.000029	0.00043
<i>Desulfovibrio</i>	sp.		F91	Phylotype	1	T-769	0.000029	0.00043
<i>Porphyromonas</i>	sp.		F92	Phylotype	1	T-769	0.000029	0.00043
<i>Capnocytophaga</i>	sp.		F99	Phylotype	1	T-769	0.000029	0.00043
<i>Capnocytophaga</i>	sp.		G02	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		G04	Phylotype	1	T-769	0.000029	0.00043
<i>Stenotrophomonas</i>	sp.		G05	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		G08	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		G09	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		G10	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		G11	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.		G12	Phylotype	1	T-769	0.000029	0.00043

<i>Neisseria</i>	sp.	G14	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.	G15	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.	G16	Phylotype	1	T-769	0.000029	0.00043
<i>Fusobacterium</i>	sp.	G17	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.	G18	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.	G19	Phylotype	1	T-769	0.000029	0.00043
<i>Haemophilus</i>	sp.	G21	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	G22	Phylotype	1	T-769	0.000029	0.00043
<i>Peptoniphilus</i>	sp.	G26	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.	G28	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	G31	Phylotype	1	T-769	0.000029	0.00043
<i>Megasphaera</i>	sp.	G37	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	G39	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonellaceae [G-1]</i>	sp.	G40	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	G42	Phylotype	1	T-769	0.000029	0.00043
<i>Bacteroidetes [G-3]</i>	sp.	G44	Phylotype	1	T-769	0.000029	0.00043

<i>Treponema</i>	sp.		G45	Phylotype	1	T-769	0.000029	0.00043
<i>Epsilonproteobacteria [G]</i>	sp.		G46	Phylotype	1	T-769	0.000029	0.00043
<i>Porphyromonas</i>	sp.		G50	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		G58	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		G60	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		G65	Phylotype	1	T-769	0.000029	0.00043
<i>Capnocytophaga</i>	sp.		G66	Phylotype	1	T-769	0.000029	0.00043
<i>Shuttleworthia</i>	sp.		G69	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		G71	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		G72	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		G73	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		G77	Phylotype	1	T-769	0.000029	0.00043
<i>Eubacterium [XII] [G]</i>	sp.		G79	Phylotype	1	T-769	0.000029	0.00043
<i>Burkholderiaceae [G]</i>	sp.		G80	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.		G81	Phylotype	1	T-769	0.000029	0.00043
<i>Eubacterium [XII] [G]</i>	sp.		G82	Phylotype	1	T-769	0.000029	0.00043

<i>Actinomyces</i>	sp.	G84	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	G88	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	G89	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	G91	Phylotype	1	T-769	0.000029	0.00043
<i>Campylobacter</i>	sp.	G92	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	G93	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.	G94	Phylotype	1	T-769	0.000029	0.00043
<i>Neisseriaceae [G]</i>	sp.	G95	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.	G96	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	G97	Phylotype	1	T-769	0.000029	0.00043
<i>Atopobium</i>	sp.	G99	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.	H00	Phylotype	1	T-769	0.000029	0.00043
<i>Scardovia</i>	sp.	H01	Phylotype	1	T-769	0.000029	0.00043
<i>Rothia</i>	sp.	H03	Phylotype	1	T-769	0.000029	0.00043
<i>Atopobium</i>	sp.	H04	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.	H05	Phylotype	1	T-769	0.000029	0.00043

<i>Parvimonas</i>	sp.		H06	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.		H08	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.		H09	Phylotype	1	T-769	0.000029	0.00043
<i>Parvimonas</i>	sp.		H10	Phylotype	1	T-769	0.000029	0.00043
<i>Atopobium</i>	sp.		H11	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.		H12	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.		H13	Phylotype	1	T-769	0.000029	0.00043
<i>Lactobacillus</i>	sp.		H14	Phylotype	1	T-769	0.000029	0.00043
<i>Campylobacter</i>	sp.		H15	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.		H17	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.		H19	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		H20	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		H21	Phylotype	1	T-769	0.000029	0.00043
<i>Leptotrichia</i>	sp.		H22	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonella</i>	sp.		H26	Phylotype	1	T-769	0.000029	0.00043
<i>Fusobacterium</i>	sp.		H28	Phylotype	1	T-769	0.000029	0.00043

<i>Aggregatibacter</i>	sp.		H29	Phylotype	1	T-769	0.000029	0.00043
<i>Mitsuokella</i>	sp.		H31	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		H32	Phylotype	1	T-769	0.000029	0.00043
<i>Mitsuokella</i>	sp.		H33	Phylotype	1	T-769	0.000029	0.00043
<i>Pseudoramibacter</i>	sp.		H34	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonellaceae [G-I]</i>	sp.		H35	Phylotype	1	T-769	0.000029	0.00043
<i>Corynebacterium</i>	sp.		H36	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		H39	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		H42	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		H43	Phylotype	1	T-769	0.000029	0.00043
<i>Selenomonas</i>	sp.		H44	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		H45	Phylotype	1	T-769	0.000029	0.00043
<i>Treponema</i>	sp.		H46	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		H49	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		H50	Phylotype	1	T-769	0.000029	0.00043
<i>Veillonellaceae [G-I]</i>	sp.		H51	Phylotype	1	T-769	0.000029	0.00043

<i>Selenomonas</i>	sp.		H52	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		H53	Phylotype	1	T-769	0.000029	0.00043
<i>Dialister</i>	sp.		H54	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		H55	Phylotype	1	T-769	0.000029	0.00043
<i>Prevotella</i>	sp.		H56	Phylotype	1	T-769	0.000029	0.00043
<i>TM7</i>	sp.		H57	Phylotype	1	T-769	0.000029	0.00043
<i>Atopobium</i>	sp.		H58	Phylotype	1	T-769	0.000029	0.00043
<i>Streptococcus</i>	sp.		H60	Phylotype	1	T-769	0.000029	0.00043
<i>Eubacterium [XII] [G]</i>	sp.		H61	Phylotype	1	T-769	0.000029	0.00043
<i>Mogibacterium</i>	sp.		H62	Phylotype	1	T-769	0.000029	0.00043
<i>Weissella</i>	<i>cibaria</i>		H69	Named	1	T-769	0.000029	0.00043
<i>Bergeyella</i>	sp.		H70	Phylotype	1	T-769	0.000029	0.00043