		Week 2		VCt n=10		CCt n=10		VLr n=9		CLr n=11		Pvalues of each 2by2 comparison of taxa abundance					
Phylum	**Family**	**Genus**		**Abundance**	**Occurrence**	**Abundance**	**Occurrence**	**Abundance**	**Occurrence**	**Abundance**	**Occurrence**	**CLr vs. CCt**	**VCt vs. CCt**	**VLr vs. CCt**	**CLr vs. VCt**	**CLr vs. VLr**	**VCt vs. VLr**
Actinobacteria			P.Actinobacteria	44.5%	100%	0.5%	60%	36.0%	100%	25.4%	91%	0.015	0.008	0.005	0.426	0.683	0.720
Actinobacteria	Actinomycetaceae		F.Actinomycetaceae	0.0%	0%	0.0%	10%	0.0%	11%	0.0%	0%	0.476	1.000	0.737	1.000	0.450	0.474
Actinobacteria	Actinomycetaceae	Actinomyces	G.Actinomyces	0.0%	0%	0.0%	10%	0.0%	11%	0.0%	0%	0.476	1.000	1.000	1.000	0.450	0.474
Actinobacteria	Actinomycetaceae	Varibaculum	G.Varibaculum	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Actinobacteria	Bifidobacteriaceae		F.Bifidobacteriaceae	29.5%	80%	0.0%	50%	36.0%	100%	25.4%	91%	0.006	0.039	0.005	0.863	0.795	0.949
Actinobacteria	Bifidobacteriaceae	Aeriscardovia	G.Aeriscardovia	0.0%	10%	0.0%	0%	0.0%	11%	0.0%	18%	0.476	1.000	0.474	0.862	0.711	1.000
Actinobacteria	Bifidobacteriaceae	Bifidobacterium	G.Bifidobacterium	28.6%	80%	0.0%	50%	35.1%	100%	24.8%	91%	0.008	0.039	0.005	0.863	0.795	0.949
Actinobacteria	Bifidobacteriaceae	f_Bifidobacteriaceae_g_Unc	G.f_Bifidobacteriaceae_g_Unc	0.8%	70%	0.0%	40%	1.0%	78%	0.6%	73%	0.019	0.051	0.017	0.768	0.309	0.395
Actinobacteria	c_Actinobacteria_f_Unc		F.c_Actinobacteria_f_Unc	0.1%	60%	0.0%	10%	0.1%	56%	0.0%	9%	0.738	0.050	0.099	0.032	0.044	0.759
Actinobacteria	c_Actinobacteria_f_Unc	c_Actinobacteria_g_Unc	G.c_Actinobacteria_g_Unc	0.1%	60%	0.0%	10%	0.1%	56%	0.0%	9%	0.738	0.050	0.099	0.032	0.044	0.759
Actinobacteria	Coriobacteriaceae		F.Coriobacteriaceae	0.0%	40%	0.0%	30%	0.0%	33%	0.0%	9%	0.325	0.354	0.777	0.058	0.189	0.666
Actinobacteria	Coriobacteriaceae	Asaccharobacter	G.Asaccharobacter	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Actinobacteria	Coriobacteriaceae	Atopobium	G.Atopobium	0.0%	0%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	1.000	1.000	1.000
Actinobacteria	Coriobacteriaceae	Collinsella	G.Collinsella	0.0%	40%	0.0%	10%	0.0%	33%	0.0%	0%	0.476	0.087	0.195	0.035	0.074	0.666
Actinobacteria	Coriobacteriaceae	Eggerthella	G.Eggerthella	0.0%	10%	0.0%	0%	0.0%	11%	0.0%	9%	1.000	1.000	0.474	0.738	0.739	1.000
Actinobacteria	Coriobacteriaceae	f_Coriobacteriaceae_g_Unc	G.f_Coriobacteriaceae_g_Unc	0.0%	10%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	0.476	1.000	1.000
Actinobacteria	Coriobacteriaceae	Olsenella	G.Olsenella	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Actinobacteria	Coriobacteriaceae	Slackia	G.Slackia	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Actinobacteria	Corynebacteriaceae		F.Corynebacteriaceae	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Actinobacteria	Corynebacteriaceae	Corynebacterium	G.Corynebacterium	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Actinobacteria	Dermabacteraceae		F.Dermabacteraceae	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Actinobacteria	Dermabacteraceae	Dermabacter	G.Dermabacter	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Actinobacteria	Micrococcaceae		F.Micrococcaceae	0.1%	60%	0.1%	50%	0.0%	33%	0.0%	36%	0.353	0.613	0.193	0.737	0.558	0.200
Actinobacteria	Micrococcaceae	Rothia	G.Rothia	0.1%	60%	0.1%	50%	0.0%	33%	0.0%	36%	0.353	0.613	0.193	0.737	0.558	0.200
Actinobacteria	Nocardiaceae		F.Nocardiaceae	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Actinobacteria	Nocardiaceae	Rhodococcus	G.Rhodococcus	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Actinobacteria	o_Actinomycetales_f_Unc		F.o_Actinomycetales_f_Unc	0.0%	20%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	0.474	1.000	0.214	1.000	0.474
Actinobacteria	o_Actinomycetales_f_Unc	o_Actinomycetales_g_Unc	G.o_Actinomycetales_g_Unc	0.0%	20%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	0.474	1.000	0.214	1.000	0.474
Actinobacteria	Propionibacteriaceae		F.Propionibacteriaceae	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Actinobacteria	Propionibacteriaceae	Propionibacterium	G.Propionibacterium	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Bacteroidetes			P.Bacteroidetes	0.1%	50%	0.0%	20%	0.0%	33%	0.0%	9%	0.338	0.224	0.536	0.021	0.142	0.628
Bacteroidetes	Bacteroidaceae		F.Bacteroidaceae	0.1%	50%	0.0%	20%	0.0%	22%	0.0%	0%	0.214	0.224	0.930	0.012	0.189	0.350
Bacteroidetes	Bacteroidaceae	Bacteroides	G.Bacteroides	0.1%	50%	0.0%	20%	0.0%	22%	0.0%	0%	0.214	0.224	0.930	0.012	0.189	0.350
Bacteroidetes	o_Bacteroidales_f_Unc		F.o_Bacteroidales_f_Unc	0.0%	10%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	0.476	1.000	1.000
Bacteroidetes	o_Bacteroidales_f_Unc	o_Bacteroidales_g_Unc	G.o_Bacteroidales_g_Unc	0.0%	10%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	0.476	1.000	1.000
Bacteroidetes	Porphyromonadaceae		F.Porphyromonadaceae	0.0%	30%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	0.211	1.000	0.090	1.000	0.211
Bacteroidetes	Porphyromonadaceae	Barnesiella	G.Barnesiella	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Bacteroidetes	Porphyromonadaceae	f_Porphyromonadaceae_g_Unc	G.f_Porphyromonadaceae_g_Unc	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Bacteroidetes	Porphyromonadaceae	Parabacteroides	G.Parabacteroides	0.0%	30%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	0.211	1.000	0.090	1.000	0.211
Bacteroidetes	Prevotellaceae		F.Prevotellaceae	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Bacteroidetes	Prevotellaceae	f_Prevotellaceae_g_Unc	G.f_Prevotellaceae_g_Unc	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Bacteroidetes	Prevotellaceae	Prevotella	G.Prevotella	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Bacteroidetes	Rikenellaceae		F.Rikenellaceae	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Bacteroidetes	Rikenellaceae	Alistipes	G.Alistipes	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes			P.Firmicutes	49.0%	100%	59.5%	100%	60.3%	100%	58.5%	100%	0.809	0.315	0.549	0.314	0.503	0.905
Firmicutes	c_Bacilli_f_Unc		F.c_Bacilli_f_Unc	0.1%	70%	0.1%	70%	0.0%	22%	0.0%	45%	0.914	0.640	0.039	0.801	0.172	0.061
Firmicutes	c_Bacilli_f_Unc	c_Bacilli_g_Unc	G.c_Bacilli_g_Unc	0.1%	70%	0.1%	70%	0.0%	22%	0.0%	45%	0.914	0.640	0.039	0.801	0.172	0.061
Firmicutes	Carnobacteriaceae		F.Carnobacteriaceae	0.0%	0%	0.0%	30%	0.0%	11%	0.0%	0%	0.090	0.211	0.319	1.000	0.450	0.474
Firmicutes	Carnobacteriaceae	f_Carnobacteriaceae_g_Unc	G.f_Carnobacteriaceae_g_Unc	0.0%	0%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	1.000	1.000	1.000
Firmicutes	Carnobacteriaceae	Granulicatella	G.Granulicatella	0.0%	0%	0.0%	20%	0.0%	11%	0.0%	0%	0.214	0.474	0.598	1.000	0.450	0.474
Firmicutes	Clostridiaceae		F.Clostridiaceae	0.2%	60%	6.3%	100%	0.5%	78%	6.9%	100%	0.605	0.001	0.021	0.002	0.055	0.263
Firmicutes	Clostridiaceae	Anaerobacter	G.Anaerobacter	0.0%	30%	0.4%	90%	0.0%	44%	0.3%	91%	0.571	0.003	0.028	0.009	0.031	0.561
Firmicutes	Clostridiaceae	Clostridium	G.Clostridium	0.1%	60%	3.1%	100%	0.5%	78%	3.9%	100%	0.809	0.001	0.034	0.003	0.055	0.229
Firmicutes	Clostridiaceae	f_Clostridiaceae_g_Unc	G.f_Clostridiaceae_g_Unc	0.0%	50%	3.0%	100%	0.1%	67%	1.6%	100%	0.426	0.000	0.035	0.001	0.046	0.351
Firmicutes	Clostridiaceae	Sarcina	G.Sarcina	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	27%	0.214	1.000	0.474	0.214	0.591	0.474
Firmicutes	Enterococcaceae		F.Enterococcaceae	1.8%	100%	9.4%	100%	3.1%	89%	14.9%	100%	0.132	0.015	0.400	0.010	0.038	0.447
Firmicutes	Enterococcaceae	Catellicoccus	G.Catellicoccus	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	27%	0.214	1.000	1.000	0.214	0.218	1.000
Firmicutes	Enterococcaceae	Enterococcus	G.Enterococcus	1.7%	90%	8.9%	100%	2.9%	89%	14.4%	100%	0.152	0.011	0.447	0.008	0.046	0.326
Firmicutes	Enterococcaceae	f_Enterococcaceae_g_Unc	G.f_Enterococcaceae_g_Unc	0.0%	50%	0.3%	100%	0.2%	67%	0.6%	91%	0.349	0.214	0.133	0.072	0.026	0.782
Firmicutes	Enterococcaceae	Melissococcus	G.Melissococcus	0.0%	10%	0.0%	10%	0.0%	0%	0.0%	9%	0.738	1.000	1.000	0.738	1.000	1.000
Firmicutes	Enterococcaceae	Pilibacter	G.Pilibacter	0.0%	20%	0.0%	10%	0.0%	33%	0.0%	0%	0.476	0.737	0.195	0.214	0.074	0.536
Firmicutes	Erysipelotrichaceae		F.Erysipelotrichaceae	0.0%	40%	0.0%	20%	0.1%	56%	0.0%	0%	0.214	0.563	0.114	0.035	0.008	0.506
Firmicutes	Erysipelotrichaceae	Allobaculum	G.Allobaculum	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Firmicutes	Erysipelotrichaceae	Catenibacterium	G.Catenibacterium	0.0%	0%	0.0%	0%	0.0%	22%	0.0%	0%	1.000	1.000	0.211	1.000	0.189	0.211
Firmicutes	Erysipelotrichaceae	Coprobacillus	G.Coprobacillus	0.0%	40%	0.0%	0%	0.0%	33%	0.0%	0%	1.000	0.087	0.087	0.035	0.074	0.893
Firmicutes	Erysipelotrichaceae	f_Erysipelotrichaceae_g_Unc	G.f_Erysipelotrichaceae_g_Unc	0.0%	10%	0.0%	20%	0.0%	33%	0.0%	0%	0.214	0.474	0.851	0.476	0.074	0.319
Firmicutes	Erysipelotrichaceae	Solobacterium	G.Solobacterium	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Erysipelotrichaceae	Turicibacter	G.Turicibacter	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Eubacteriaceae		F.Eubacteriaceae	0.0%	20%	0.0%	20%	0.0%	33%	0.0%	18%	1.000	0.721	0.336	0.655	0.310	0.638
Firmicutes	Eubacteriaceae	Anaerofustis	G.Anaerofustis	0.0%	0%	0.0%	10%	0.0%	0%	0.0%	18%	0.600	1.000	1.000	0.476	0.479	1.000
Firmicutes	Eubacteriaceae	Eubacterium	G.Eubacterium	0.0%	20%	0.0%	10%	0.0%	22%	0.0%	0%	0.476	0.474	0.334	0.214	0.189	1.000
Firmicutes	Eubacteriaceae	f_Eubacteriaceae_g_Unc	G.f_Eubacteriaceae_g_Unc	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Firmicutes	Incertae_Sedis_XI		F.Incertae_Sedis_XI	0.0%	0%	0.0%	10%	0.0%	11%	0.0%	0%	0.476	1.000	1.000	1.000	0.450	0.474
Firmicutes	Incertae_Sedis_XI	Anaerococcus	G.Anaerococcus	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Incertae_Sedis_XI	f_Incertae_Sedis_XI_g_Unc	G.f_Incertae_Sedis_XI_g_Unc	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Incertae_Sedis_XI	Finegoldia	G.Finegoldia	0.0%	0%	0.0%	10%	0.0%	11%	0.0%	0%	0.476	1.000	1.000	1.000	0.450	0.474
Firmicutes	Incertae_Sedis_XI	Peptoniphilus	G.Peptoniphilus	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Incertae_Sedis_XIV		F.Incertae_Sedis_XIV	0.0%	10%	0.0%	10%	0.0%	22%	0.0%	18%	0.600	1.000	0.458	0.600	0.864	0.334
Firmicutes	Incertae_Sedis_XIV	Blautia	G.Blautia	0.0%	10%	0.0%	10%	0.0%	22%	0.0%	18%	0.600	1.000	0.458	0.600	0.864	0.334
Firmicutes	Lachnospiraceae		F.Lachnospiraceae	0.0%	30%	0.0%	40%	0.7%	89%	0.0%	45%	0.779	0.721	0.086	0.507	0.205	0.037
Firmicutes	Lachnospiraceae	Anaerostipes	G.Anaerostipes	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Firmicutes	Lachnospiraceae	Coprococcus	G.Coprococcus	0.0%	10%	0.0%	20%	0.0%	44%	0.0%	18%	1.000	0.737	0.303	0.862	0.344	0.152
Firmicutes	Lachnospiraceae	Dorea	G.Dorea	0.0%	0%	0.0%	10%	0.0%	22%	0.0%	9%	1.000	1.000	0.458	1.000	0.450	0.211
Firmicutes	Lachnospiraceae	f_Lachnospiraceae_g_Unc	G.f_Lachnospiraceae_g_Unc	0.0%	30%	0.0%	40%	0.3%	89%	0.0%	45%	0.842	0.721	0.104	0.507	0.178	0.046
Firmicutes	Lachnospiraceae	Robinsoniella	G.Robinsoniella	0.0%	0%	0.0%	20%	0.0%	22%	0.0%	18%	0.724	0.474	0.930	0.476	0.864	0.211
Firmicutes	Lachnospiraceae	Roseburia	G.Roseburia	0.0%	10%	0.0%	20%	0.0%	33%	0.0%	18%	0.931	1.000	0.433	1.000	0.310	0.443
Firmicutes	Lachnospiraceae	Syntrophococcus	G.Syntrophococcus	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Lactobacillaceae		F.Lactobacillaceae	0.1%	70%	0.0%	40%	2.8%	89%	4.4%	91%	0.027	0.669	0.086	0.005	0.469	0.045
Firmicutes	Lactobacillaceae	f_Lactobacillaceae_g_Unc	G.f_Lactobacillaceae_g_Unc	0.0%	0%	0.0%	20%	0.0%	11%	0.0%	0%	0.214	0.474	0.598	1.000	0.450	0.474
Firmicutes	Lactobacillaceae	Lactobacillus	G.Lactobacillus	0.1%	70%	0.0%	40%	2.8%	89%	4.4%	91%	0.027	0.669	0.086	0.005	0.469	0.045
Firmicutes	Leuconostocaceae		F.Leuconostocaceae	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Firmicutes	Leuconostocaceae	Weissella	G.Weissella	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Firmicutes	o_Clostridiales_f_Unc		F.o_Clostridiales_f_Unc	0.0%	40%	0.2%	80%	0.1%	56%	0.4%	82%	0.744	0.053	0.477	0.024	0.637	0.272
Firmicutes	o_Clostridiales_f_Unc	o_Clostridiales_g_Unc	G.o_Clostridiales_g_Unc	0.0%	40%	0.2%	80%	0.1%	56%	0.4%	82%	0.744	0.053	0.477	0.024	0.637	0.272
Firmicutes	o_Lactobacillales_f_Unc		F.o_Lactobacillales_f_Unc	0.5%	90%	0.6%	100%	0.5%	89%	0.5%	91%	1.000	0.631	0.968	0.571	0.959	0.620
Firmicutes	o_Lactobacillales_f_Unc	o_Lactobacillales_g_Unc	G.o_Lactobacillales_g_Unc	0.5%	90%	0.6%	100%	0.5%	89%	0.5%	91%	1.000	0.631	0.968	0.571	0.959	0.620
Firmicutes	p_Firmicutes_f_Unc		F.p_Firmicutes_f_Unc	0.0%	40%	0.3%	80%	0.1%	56%	0.0%	36%	0.029	0.285	0.058	0.459	0.710	0.806
Firmicutes	p_Firmicutes_f_Unc	p_Firmicutes_g_Unc	G.p_Firmicutes_g_Unc	0.0%	40%	0.3%	80%	0.1%	56%	0.0%	36%	0.029	0.285	0.058	0.459	0.710	0.806
Firmicutes	Peptostreptococcaceae		F.Peptostreptococcaceae	0.1%	50%	0.0%	40%	0.0%	22%	0.2%	64%	0.122	0.486	0.610	0.231	0.055	0.393
Firmicutes	Peptostreptococcaceae	f_Peptostreptococcaceae_g_Unc	G.f_Peptostreptococcaceae_g_Unc	0.0%	40%	0.0%	30%	0.0%	22%	0.0%	45%	0.279	0.981	0.819	0.360	0.172	0.610
Firmicutes	Peptostreptococcaceae	Sporacetigenium	G.Sporacetigenium	0.0%	50%	0.0%	20%	0.0%	22%	0.2%	55%	0.084	0.224	0.930	0.816	0.187	0.271
Firmicutes	Ruminococcaceae		F.Ruminococcaceae	0.0%	10%	0.0%	0%	0.0%	22%	0.0%	0%	1.000	1.000	0.211	0.476	0.189	0.334
Firmicutes	Ruminococcaceae	AnaerotrUncus	G.AnaerotrUncus	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Firmicutes	Ruminococcaceae	Butyricicoccus	G.Butyricicoccus	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Ruminococcaceae	f_Ruminococcaceae_g_Unc	G.f_Ruminococcaceae_g_Unc	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Ruminococcaceae	Oscillibacter	G.Oscillibacter	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Firmicutes	Ruminococcaceae	Subdoligranulum	G.Subdoligranulum	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Firmicutes	Staphylococcaceae		F.Staphylococcaceae	0.7%	70%	0.1%	80%	0.2%	56%	0.1%	64%	0.587	0.389	0.583	0.339	0.864	0.123
Firmicutes	Staphylococcaceae	f_Staphylococcaceae_g_Unc	G.f_Staphylococcaceae_g_Unc	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Firmicutes	Staphylococcaceae	Gemella	G.Gemella	0.0%	10%	0.0%	20%	0.0%	0%	0.0%	0%	0.214	1.000	0.474	0.476	1.000	1.000
Firmicutes	Staphylococcaceae	Staphylococcus	G.Staphylococcus	0.7%	70%	0.1%	70%	0.2%	56%	0.1%	64%	0.860	0.315	0.712	0.339	0.864	0.123
Firmicutes	Streptococcaceae		F.Streptococcaceae	30.1%	100%	13.1%	100%	10.4%	100%	6.5%	100%	0.152	0.684	0.400	0.036	0.795	0.278
Firmicutes	Streptococcaceae	f_Streptococcaceae_g_Unc	G.f_Streptococcaceae_g_Unc	0.0%	50%	0.0%	0%	0.0%	22%	0.0%	18%	0.476	0.033	0.211	0.149	0.932	0.156
Firmicutes	Streptococcaceae	Lactococcus	G.Lactococcus	0.0%	0%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	1.000	1.000	1.000
Firmicutes	Streptococcaceae	Lactovum	G.Lactovum	0.0%	0%	0.0%	10%	0.0%	11%	0.0%	0%	0.476	1.000	1.000	1.000	0.450	0.474
Firmicutes	Streptococcaceae	Streptococcus	G.Streptococcus	30.1%	100%	13.1%	100%	10.4%	100%	6.5%	100%	0.152	0.684	0.400	0.036	0.795	0.278
Firmicutes	Veillonellaceae		F.Veillonellaceae	1.4%	80%	0.4%	70%	0.2%	56%	0.1%	64%	0.416	0.627	0.651	0.107	0.844	0.300
Firmicutes	Veillonellaceae	Allisonella	G.Allisonella	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Veillonellaceae	Dialister	G.Dialister	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Firmicutes	Veillonellaceae	f_Veillonellaceae_g_Unc	G.f_Veillonellaceae_g_Unc	0.0%	20%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	0.474	1.000	0.214	1.000	0.474
Firmicutes	Veillonellaceae	Megasphaera	G.Megasphaera	0.0%	0%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	1.000	1.000	1.000
Firmicutes	Veillonellaceae	Veillonella	G.Veillonella	1.4%	80%	0.4%	70%	0.2%	56%	0.1%	64%	0.416	0.575	0.651	0.107	0.844	0.300
Unc			P.Unc	0.0%	50%	0.2%	70%	0.2%	89%	0.1%	82%	0.755	0.631	0.324	0.377	0.245	0.093
Unc	p_Unc_f_Unc		F.p_Unc_f_Unc	0.0%	50%	0.2%	70%	0.2%	89%	0.1%	82%	0.755	0.631	0.324	0.377	0.245	0.093
Unc	p_Unc_f_Unc	p_Unc_g_Unc	G.p_Unc_g_Unc	0.0%	50%	0.2%	70%	0.2%	89%	0.1%	82%	0.755	0.631	0.324	0.377	0.245	0.093
Proteobacteria			P.Proteobacteria	2.5%	100%	27.6%	100%	2.4%	100%	7.3%	100%	0.051	0.011	0.035	0.809	0.882	0.842
Proteobacteria	Alcaligenaceae		F.Alcaligenaceae	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Proteobacteria	Alcaligenaceae	Achromobacter	G.Achromobacter	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Alcaligenaceae	f_Alcaligenaceae_g_Unc	G.f_Alcaligenaceae_g_Unc	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Proteobacteria	c_Gammaproteobacteria_f_Unc		F.c_Gammaproteobacteria_f_Unc	0.0%	20%	0.0%	20%	0.0%	11%	0.0%	18%	0.931	0.861	0.861	0.931	1.000	1.000
Proteobacteria	c_Gammaproteobacteria_f_Unc	c_Gammaproteobacteria_g_Unc	G.c_Gammaproteobacteria_g_Unc	0.0%	20%	0.0%	20%	0.0%	11%	0.0%	18%	0.931	0.861	0.861	0.931	1.000	1.000
Proteobacteria	Caulobacteraceae		F.Caulobacteraceae	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Caulobacteraceae	Caulobacter	G.Caulobacter	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Comamonadaceae		F.Comamonadaceae	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Proteobacteria	Comamonadaceae	Delftia	G.Delftia	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Proteobacteria	Enterobacteriaceae		F.Enterobacteriaceae	2.5%	100%	27.6%	100%	2.4%	100%	7.3%	100%	0.051	0.009	0.035	0.705	0.882	0.842
Proteobacteria	Enterobacteriaceae	Citrobacter	G.Citrobacter	0.0%	50%	0.4%	70%	0.0%	33%	0.3%	73%	0.351	0.115	0.093	0.066	0.206	0.755
Proteobacteria	Enterobacteriaceae	Cronobacter	G.Cronobacter	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Enterobacteriaceae	Enterobacter	G.Enterobacter	0.0%	50%	0.6%	90%	0.0%	44%	0.3%	82%	0.250	0.043	0.131	0.165	0.594	0.923
Proteobacteria	Enterobacteriaceae	Escherichia/Shigella	G.Escherichia/Shigella	1.3%	90%	9.3%	100%	2.3%	89%	1.0%	91%	0.152	0.353	0.356	0.772	0.823	0.367
Proteobacteria	Enterobacteriaceae	f_Enterobacteriaceae_g_Unc	G.f_Enterobacteriaceae_g_Unc	0.2%	80%	2.3%	100%	0.1%	89%	0.4%	100%	0.010	0.004	0.017	0.217	0.175	0.968
Proteobacteria	Enterobacteriaceae	Klebsiella	G.Klebsiella	0.0%	40%	0.5%	70%	0.0%	33%	0.1%	64%	0.416	0.145	0.076	0.353	0.288	0.740
Proteobacteria	Enterobacteriaceae	Morganella	G.Morganella	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Enterobacteriaceae	Pantoea	G.Pantoea	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Enterobacteriaceae	Pragia	G.Pragia	0.0%	10%	0.0%	10%	0.0%	11%	0.0%	0%	0.476	1.000	1.000	0.476	0.450	0.737
Proteobacteria	Enterobacteriaceae	Proteus	G.Proteus	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Enterobacteriaceae	Raoultella	G.Raoultella	0.0%	20%	0.0%	40%	0.0%	44%	0.0%	27%	0.465	0.563	0.960	0.927	0.487	0.503
Proteobacteria	Enterobacteriaceae	Tatumella	G.Tatumella	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Proteobacteria	Enterobacteriaceae	Trabulsiella	G.Trabulsiella	0.0%	0%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Phyllobacteriaceae		F.Phyllobacteriaceae	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Proteobacteria	Phyllobacteriaceae	Defluvibacter	G.Defluvibacter	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Proteobacteria	Rhizobiaceae		F.Rhizobiaceae	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
Proteobacteria	Rhizobiaceae	Rhizobium	G.Rhizobium	0.0%	0%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	1.000	1.000	1.000
TM7			P.TM7	0.0%	10%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	0.476	1.000	1.000
TM7	c_TM7_genera_incertae_sedis_f_Unc		F.c_TM7_genera_incertae_sedis_f_Unc	0.0%	10%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	0.476	1.000	1.000
TM7	c_TM7_genera_incertae_sedis_f_Unc	c_TM7_genera_incertae_sedis_g_Unc	G.c_TM7_genera_incertae_sedis_g_Unc	0.0%	10%	0.0%	10%	0.0%	0%	0.0%	0%	0.476	1.000	1.000	0.476	1.000	1.000
Verrucomicrobia			P.Verrucomicrobia	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Verrucomicrobia	Verrucomicrobiaceae		F.Verrucomicrobiaceae	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
Verrucomicrobia	Verrucomicrobiaceae	Akkermansia	G.Akkermansia	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474