		Month 4		VCt n=10		CCt n=10		VLr n=9		CLr n=11		Pvalues of each 2by2 comparison					
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	81.2%	100%	63.6%	100%	62.2%	100%	48.2%	100%	0.863	0.052	0.780	0.223	1.000	0.156
Actinobacteria	Actinomycetaceae		F.Actinomycetaceae	0.0%	0%	0.0%	50%	0.1%	56%	0.1%	55%	0.404	0.033	0.641	0.012	0.481	0.011
Actinobacteria	Actinomycetaceae	Actinomyces	G.Actinomyces	0.0%	0%	0.0%	40%	0.1%	56%	0.1%	55%	0.276	0.087	0.455	0.012	0.481	0.011
Actinobacteria	Actinomycetaceae	Varibaculum	G.Varibaculum	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	Bifidobacteriaceae		F.Bifidobacteriaceae	57.3%	100%	54.9%	90%	53.4%	100%	40.8%	100%	0.918	0.579	0.780	0.387	1.000	0.315
Actinobacteria	Bifidobacteriaceae	Aeriscardovia	G.Aeriscardovia	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	Bifidobacteriaceae	Bifidobacterium	G.Bifidobacterium	56.6%	100%	54.4%	90%	53.1%	100%	40.1%	100%	0.973	0.579	0.780	0.426	1.000	0.400
Actinobacteria	Bifidobacteriaceae	f_Bifidobacteriaceae_g_Unc	G.f_Bifidobacteriaceae_g_Unc	0.8%	90%	0.5%	90%	0.6%	100%	0.6%	82%	0.704	0.363	0.968	0.314	0.694	0.968
Actinobacteria	c_Actinobacteria_f_Unc		F.c_Actinobacteria_f_Unc	0.2%	70%	0.0%	50%	0.2%	89%	0.0%	45%	0.911	0.135	0.013	0.099	0.022	0.507
Actinobacteria	c_Actinobacteria_f_Unc	c_Actinobacteria_g_Unc	G.c_Actinobacteria_g_Unc	0.2%	70%	0.0%	50%	0.2%	89%	0.0%	45%	0.911	0.135	0.013	0.099	0.022	0.507
Actinobacteria	Coriobacteriaceae		F.Coriobacteriaceae	21.1%	80%	0.2%	70%	10.2%	89%	0.4%	64%	0.647	0.042	0.138	0.078	0.366	0.210
Actinobacteria	Coriobacteriaceae	Asaccharobacter	G.Asaccharobacter	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	Coriobacteriaceae	Atopobium	G.Atopobium	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	Collinsella	G.Collinsella	20.9%	80%	0.0%	40%	9.7%	67%	0.0%	45%	0.546	0.018	0.157	0.052	0.696	0.152
Actinobacteria	Coriobacteriaceae	Eggerthella	G.Eggerthella	0.2%	70%	0.0%	40%	0.1%	67%	0.0%	36%	0.966	0.124	0.287	0.201	0.206	0.671
Actinobacteria	Coriobacteriaceae	f_Coriobacteriaceae_g_Unc	G.f_Coriobacteriaceae_g_Unc	0.0%	10%	0.0%	40%	0.0%	33%	0.0%	27%	0.571	0.141	0.826	0.449	0.727	0.141
Actinobacteria	Coriobacteriaceae	Olsenella	G.Olsenella	0.0%	10%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	0.476	0.450	0.737
Actinobacteria	Coriobacteriaceae	Slackia	G.Slackia	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	Corynebacteriaceae		F.Corynebacteriaceae	0.0%	0%	0.0%	10%	0.0%	22%	0.0%	0%	0.476	1.000	0.458	1.000	0.189	0.211
Actinobacteria	Corynebacteriaceae	Corynebacterium	G.Corynebacterium	0.0%	0%	0.0%	10%	0.0%	22%	0.0%	0%	0.476	1.000	0.458	1.000	0.189	0.211
Actinobacteria	Dermabacteraceae		F.Dermabacteraceae	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	Dermabacteraceae	Dermabacter	G.Dermabacter	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	Micrococcaceae		F.Micrococcaceae	0.0%	20%	0.0%	40%	0.0%	0%	0.0%	9%	0.103	0.271	0.087	0.724	1.000	0.474
Actinobacteria	Micrococcaceae	Rothia	G.Rothia	0.0%	20%	0.0%	40%	0.0%	0%	0.0%	9%	0.103	0.271	0.087	0.724	1.000	0.474
Actinobacteria	Nocardiaceae		F.Nocardiaceae	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	Nocardiaceae	Rhodococcus	G.Rhodococcus	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	o_Actinomycetales_f_Unc		F.o_Actinomycetales_f_Unc	0.0%	20%	0.0%	10%	0.0%	11%	0.0%	0%	0.476	0.737	0.737	0.214	0.450	1.000
Actinobacteria	o_Actinomycetales_f_Unc	o_Actinomycetales_g_Unc	G.o_Actinomycetales_g_Unc	0.0%	20%	0.0%	10%	0.0%	11%	0.0%	0%	0.476	0.737	0.737	0.214	0.450	1.000
Actinobacteria	Propionibacteriaceae		F.Propionibacteriaceae	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	Propionibacteriaceae	Propionibacterium	G.Propionibacterium	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Bacteroidetes			P.Bacteroidetes	0.3%	70%	0.0%	10%	0.0%	33%	0.0%	36%	0.136	0.003	0.141	0.134	0.709	0.062
Bacteroidetes	Bacteroidaceae		F.Bacteroidaceae	0.3%	70%	0.0%	10%	0.0%	33%	0.0%	27%	0.449	0.003	0.195	0.051	1.000	0.032
Bacteroidetes	Bacteroidaceae	Bacteroides	G.Bacteroides	0.3%	70%	0.0%	10%	0.0%	33%	0.0%	27%	0.449	0.003	0.195	0.051	1.000	0.032
Bacteroidetes	o_Bacteroidales_f_Unc		F.o_Bacteroidales_f_Unc	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	o_Bacteroidales_f_Unc	o_Bacteroidales_g_Unc	G.o_Bacteroidales_g_Unc	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	Porphyromonadaceae		F.Porphyromonadaceae	0.0%	20%	0.0%	0%	0.0%	11%	0.0%	18%	0.476	0.474	0.474	0.724	1.000	0.861
Bacteroidetes	Porphyromonadaceae	Barnesiella	G.Barnesiella	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	Porphyromonadaceae	f_Porphyromonadaceae_g_Unc	G.f_Porphyromonadaceae_g_Unc	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	Porphyromonadaceae	Parabacteroides	G.Parabacteroides	0.0%	20%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	0.474	0.474	0.214	0.450	0.861
Bacteroidetes	Prevotellaceae		F.Prevotellaceae	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	Prevotellaceae	f_Prevotellaceae_g_Unc	G.f_Prevotellaceae_g_Unc	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	Prevotellaceae	Prevotella	G.Prevotella	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	Rikenellaceae		F.Rikenellaceae	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	0%	1.000	1.000	1.000	0.476	1.000	1.000
Bacteroidetes	Rikenellaceae	Alistipes	G.Alistipes	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			P.Firmicutes	15.4%	100%	33.0%	100%	36.3%	100%	51.2%	100%	0.809	0.019	0.842	0.085	0.824	0.043
Firmicutes	c_Bacilli_f_Unc		F.c_Bacilli_f_Unc	0.0%	30%	0.0%	10%	0.0%	33%	0.0%	27%	0.586	0.334	0.443	0.672	0.801	0.851
Firmicutes	c_Bacilli_f_Unc	c_Bacilli_g_Unc	G.c_Bacilli_g_Unc	0.0%	30%	0.0%	10%	0.0%	33%	0.0%	27%	0.586	0.334	0.443	0.672	0.801	0.851
Firmicutes	Carnobacteriaceae		F.Carnobacteriaceae	0.0%	0%	0.0%	40%	0.0%	11%	0.0%	27%	0.571	0.087	0.238	0.214	0.438	0.474
Firmicutes	Carnobacteriaceae	f_Carnobacteriaceae_g_Unc	G.f_Carnobacteriaceae_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	Carnobacteriaceae	Granulicatella	G.Granulicatella	0.0%	0%	0.0%	40%	0.0%	11%	0.0%	27%	0.571	0.087	0.238	0.214	0.438	0.474
Firmicutes	Clostridiaceae		F.Clostridiaceae	0.2%	90%	0.7%	80%	0.7%	67%	0.1%	64%	0.372	0.314	0.717	0.755	0.761	0.410
Firmicutes	Clostridiaceae	Anaerobacter	G.Anaerobacter	0.0%	20%	0.0%	20%	0.0%	22%	0.0%	45%	0.237	0.861	1.000	0.106	0.153	0.930
Firmicutes	Clostridiaceae	Clostridium	G.Clostridium	0.2%	80%	0.5%	70%	0.7%	67%	0.0%	45%	0.328	0.389	0.859	0.399	0.481	0.395
Firmicutes	Clostridiaceae	f_Clostridiaceae_g_Unc	G.f_Clostridiaceae_g_Unc	0.0%	30%	0.1%	60%	0.0%	56%	0.0%	45%	0.673	0.152	0.574	0.459	0.859	0.398
Firmicutes	Clostridiaceae	Sarcina	G.Sarcina	0.0%	10%	0.0%	40%	0.0%	33%	0.0%	18%	0.205	0.087	0.591	0.600	0.557	0.141
Firmicutes	Enterococcaceae		F.Enterococcaceae	0.7%	80%	7.9%	90%	1.3%	78%	1.4%	64%	0.070	0.035	0.156	1.000	0.864	0.646
Firmicutes	Enterococcaceae	Catellicoccus	G.Catellicoccus	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	Enterococcaceae	Enterococcus	G.Enterococcus	0.7%	80%	7.7%	90%	1.2%	78%	1.2%	64%	0.096	0.035	0.156	0.956	0.805	0.646
Firmicutes	Enterococcaceae	f_Enterococcaceae_g_Unc	G.f_Enterococcaceae_g_Unc	0.0%	30%	0.1%	60%	0.1%	56%	0.1%	55%	0.639	0.348	0.824	0.633	0.754	0.505
Firmicutes	Enterococcaceae	Melissococcus	G.Melissococcus	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	Enterococcaceae	Pilibacter	G.Pilibacter	0.0%	0%	0.0%	20%	0.0%	11%	0.0%	9%	0.338	0.474	0.598	1.000	1.000	0.474
Firmicutes	Erysipelotrichaceae		F.Erysipelotrichaceae	0.1%	50%	0.1%	60%	2.2%	67%	1.9%	73%	0.092	0.908	0.145	0.092	0.779	0.173
Firmicutes	Erysipelotrichaceae	Allobaculum	G.Allobaculum	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	Catenibacterium	G.Catenibacterium	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	18%	0.476	1.000	0.474	0.476	0.855	0.474
Firmicutes	Erysipelotrichaceae	Coprobacillus	G.Coprobacillus	0.0%	40%	0.1%	60%	0.4%	56%	1.1%	73%	0.107	0.701	0.463	0.085	0.445	0.403
Firmicutes	Erysipelotrichaceae	f_Erysipelotrichaceae_g_Unc	G.f_Erysipelotrichaceae_g_Unc	0.0%	30%	0.0%	40%	0.0%	33%	0.1%	55%	0.214	0.926	0.826	0.156	0.416	0.647
Firmicutes	Erysipelotrichaceae	Solobacterium	G.Solobacterium	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	Erysipelotrichaceae	Turicibacter	G.Turicibacter	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	Eubacteriaceae		F.Eubacteriaceae	0.0%	40%	0.0%	30%	0.1%	56%	0.0%	27%	0.781	0.721	0.346	0.465	0.205	0.619
Firmicutes	Eubacteriaceae	Anaerofustis	G.Anaerofustis	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	Eubacterium	G.Eubacterium	0.0%	40%	0.0%	20%	0.1%	56%	0.0%	27%	0.822	0.336	0.139	0.465	0.237	0.679
Firmicutes	Eubacteriaceae	f_Eubacteriaceae_g_Unc	G.f_Eubacteriaceae_g_Unc	0.0%	0%	0.0%	20%	0.0%	22%	0.0%	9%	0.724	0.474	0.930	1.000	0.566	0.211
Firmicutes	Incertae_Sedis_XI		F.Incertae_Sedis_XI	0.0%	40%	0.0%	10%	0.0%	22%	0.0%	9%	1.000	0.141	0.458	0.080	0.450	0.368
Firmicutes	Incertae_Sedis_XI	Anaerococcus	G.Anaerococcus	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	Incertae_Sedis_XI	f_Incertae_Sedis_XI_g_Unc	G.f_Incertae_Sedis_XI_g_Unc	0.0%	10%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	0.476	0.450	1.000
Firmicutes	Incertae_Sedis_XI	Finegoldia	G.Finegoldia	0.0%	10%	0.0%	10%	0.0%	11%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Incertae_Sedis_XI	Peptoniphilus	G.Peptoniphilus	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	Incertae_Sedis_XIV		F.Incertae_Sedis_XIV	0.0%	20%	0.0%	40%	0.1%	56%	1.7%	73%	0.023	0.206	0.403	0.004	0.104	0.063
Firmicutes	Incertae_Sedis_XIV	Blautia	G.Blautia	0.0%	20%	0.0%	40%	0.1%	56%	1.7%	73%	0.023	0.206	0.403	0.004	0.104	0.063
Firmicutes	Lachnospiraceae		F.Lachnospiraceae	0.2%	70%	20.3%	90%	15.5%	78%	12.4%	73%	0.521	0.066	0.842	0.173	0.548	0.063
Firmicutes	Lachnospiraceae	Anaerostipes	G.Anaerostipes	0.0%	0%	0.0%	30%	0.0%	11%	0.0%	9%	0.201	0.211	0.582	1.000	0.739	0.474
Firmicutes	Lachnospiraceae	Coprococcus	G.Coprococcus	0.0%	30%	3.1%	70%	4.5%	78%	1.1%	64%	0.860	0.026	0.904	0.028	0.533	0.005
Firmicutes	Lachnospiraceae	Dorea	G.Dorea	0.0%	0%	0.0%	0%	0.0%	22%	0.0%	9%	1.000	1.000	0.211	1.000	0.334	0.211
Firmicutes	Lachnospiraceae	f_Lachnospiraceae_g_Unc	G.f_Lachnospiraceae_g_Unc	0.1%	60%	6.7%	90%	7.5%	78%	5.6%	73%	0.615	0.073	0.660	0.165	0.548	0.086
Firmicutes	Lachnospiraceae	Robinsoniella	G.Robinsoniella	0.0%	10%	0.0%	50%	0.1%	67%	0.2%	55%	0.648	0.054	0.911	0.021	1.000	0.013
Firmicutes	Lachnospiraceae	Roseburia	G.Roseburia	0.0%	20%	4.0%	70%	1.5%	67%	0.3%	64%	0.304	0.052	0.555	0.065	0.761	0.063
Firmicutes	Lachnospiraceae	Syntrophococcus	G.Syntrophococcus	0.0%	0%	0.0%	10%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Lactobacillaceae		F.Lactobacillaceae	0.0%	30%	0.0%	30%	0.6%	78%	0.3%	73%	0.051	0.889	0.032	0.051	0.499	0.012
Firmicutes	Lactobacillaceae	f_Lactobacillaceae_g_Unc	G.f_Lactobacillaceae_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	Lactobacillaceae	Lactobacillus	G.Lactobacillus	0.0%	30%	0.0%	30%	0.6%	78%	0.3%	73%	0.051	0.889	0.032	0.051	0.499	0.012
Firmicutes	Leuconostocaceae		F.Leuconostocaceae	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	Leuconostocaceae	Weissella	G.Weissella	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	o_Clostridiales_f_Unc		F.o_Clostridiales_f_Unc	0.0%	50%	0.2%	70%	0.2%	78%	0.4%	64%	0.377	0.631	0.492	0.203	0.584	0.170
Firmicutes	o_Clostridiales_f_Unc	o_Clostridiales_g_Unc	G.o_Clostridiales_g_Unc	0.0%	50%	0.2%	70%	0.2%	78%	0.4%	64%	0.377	0.631	0.492	0.203	0.584	0.170
Firmicutes	o_Lactobacillales_f_Unc		F.o_Lactobacillales_f_Unc	0.2%	90%	0.1%	70%	0.2%	78%	0.2%	73%	0.989	0.695	0.545	0.289	0.548	0.968
Firmicutes	o_Lactobacillales_f_Unc	o_Lactobacillales_g_Unc	G.o_Lactobacillales_g_Unc	0.2%	90%	0.1%	70%	0.2%	78%	0.2%	73%	0.989	0.695	0.545	0.289	0.548	0.968
Firmicutes	p_Firmicutes_f_Unc		F.p_Firmicutes_f_Unc	0.0%	40%	0.1%	60%	0.2%	78%	0.1%	55%	0.746	0.210	0.639	0.311	0.401	0.035
Firmicutes	p_Firmicutes_f_Unc	p_Firmicutes_g_Unc	G.p_Firmicutes_g_Unc	0.0%	40%	0.1%	60%	0.2%	78%	0.1%	55%	0.746	0.210	0.639	0.311	0.401	0.035
Firmicutes	Peptostreptococcaceae		F.Peptostreptococcaceae	0.4%	80%	0.7%	90%	1.2%	100%	0.8%	82%	0.918	0.630	0.604	0.692	0.444	0.268
Firmicutes	Peptostreptococcaceae	f_Peptostreptococcaceae_g_Unc	G.f_Peptostreptococcaceae_g_Unc	0.0%	40%	0.3%	80%	0.2%	100%	0.1%	55%	0.144	0.106	0.533	0.851	0.093	0.038
Firmicutes	Peptostreptococcaceae	Sporacetigenium	G.Sporacetigenium	0.4%	70%	0.4%	80%	0.6%	100%	0.5%	82%	0.641	0.970	0.643	0.755	0.982	0.719
Firmicutes	Ruminococcaceae		F.Ruminococcaceae	0.0%	30%	0.0%	10%	0.0%	22%	0.0%	27%	0.325	0.334	0.334	0.854	0.770	0.819
Firmicutes	Ruminococcaceae	Anaerotruncus	G.Anaerotruncus	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	Ruminococcaceae	Butyricicoccus	G.Butyricicoccus	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	Ruminococcaceae	f_Ruminococcaceae_g_Unc	G.f_Ruminococcaceae_g_Unc	0.0%	20%	0.0%	10%	0.0%	11%	0.0%	18%	0.862	0.737	0.737	0.724	0.855	0.861
Firmicutes	Ruminococcaceae	Oscillibacter	G.Oscillibacter	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	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	0.0%	30%	0.0%	30%	0.0%	11%	0.0%	27%	1.000	1.000	0.512	0.854	0.361	0.512
Firmicutes	Staphylococcaceae	f_Staphylococcaceae_g_Unc	G.f_Staphylococcaceae_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	Staphylococcaceae	Gemella	G.Gemella	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Staphylococcaceae	Staphylococcus	G.Staphylococcus	0.0%	20%	0.0%	30%	0.0%	11%	0.0%	27%	1.000	0.861	0.512	0.578	0.361	0.861
Firmicutes	Streptococcaceae		F.Streptococcaceae	3.0%	100%	4.1%	100%	0.8%	78%	1.8%	100%	0.605	0.971	0.235	0.426	0.195	0.109
Firmicutes	Streptococcaceae	f_Streptococcaceae_g_Unc	G.f_Streptococcaceae_g_Unc	0.0%	10%	0.0%	40%	0.0%	11%	0.0%	18%	0.177	0.195	0.130	1.000	0.711	1.000
Firmicutes	Streptococcaceae	Lactococcus	G.Lactococcus	0.0%	10%	0.0%	20%	0.0%	11%	0.0%	0%	0.214	0.474	0.598	0.476	0.450	1.000
Firmicutes	Streptococcaceae	Lactovum	G.Lactovum	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	Streptococcaceae	Streptococcus	G.Streptococcus	3.0%	100%	4.0%	100%	0.7%	78%	1.8%	100%	0.705	1.000	0.268	0.426	0.195	0.092
Firmicutes	Veillonellaceae		F.Veillonellaceae	0.4%	80%	0.1%	60%	0.1%	56%	0.1%	55%	0.746	0.155	1.000	0.339	0.815	0.071
Firmicutes	Veillonellaceae	Allisonella	G.Allisonella	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	Veillonellaceae	Dialister	G.Dialister	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	f_Veillonellaceae_g_Unc	G.f_Veillonellaceae_g_Unc	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Firmicutes	Veillonellaceae	Megasphaera	G.Megasphaera	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	Veillonellaceae	Veillonella	G.Veillonella	0.4%	80%	0.1%	60%	0.0%	44%	0.1%	55%	0.746	0.155	0.899	0.339	0.647	0.058
Unc			P.Unc	0.1%	50%	0.3%	80%	0.6%	78%	0.2%	64%	0.636	0.047	0.766	0.328	0.533	0.104
Unc	p_Unc_f_Unc		F.p_Unc_f_Unc	0.1%	50%	0.3%	80%	0.6%	78%	0.2%	64%	0.636	0.047	0.766	0.328	0.533	0.104
Unc	p_Unc_f_Unc	p_Unc_g_Unc	G.p_Unc_g_Unc	0.1%	50%	0.3%	80%	0.6%	78%	0.2%	64%	0.636	0.047	0.766	0.328	0.533	0.104
Proteobacteria			P.Proteobacteria	0.7%	100%	0.6%	80%	1.0%	100%	1.2%	82%	0.456	0.424	0.345	0.793	0.923	0.905
Proteobacteria	Alcaligenaceae		F.Alcaligenaceae	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	Alcaligenaceae	Achromobacter	G.Achromobacter	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	Alcaligenaceae	f_Alcaligenaceae_g_Unc	G.f_Alcaligenaceae_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
Proteobacteria	c_Gammaproteobacteria_f_Unc		F.c_Gammaproteobacteria_f_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
Proteobacteria	c_Gammaproteobacteria_f_Unc	c_Gammaproteobacteria_g_Unc	G.c_Gammaproteobacteria_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
Proteobacteria	Caulobacteraceae		F.Caulobacteraceae	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	Caulobacteraceae	Caulobacter	G.Caulobacter	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	Comamonadaceae		F.Comamonadaceae	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	Delftia	G.Delftia	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		F.Enterobacteriaceae	0.7%	100%	0.6%	80%	1.0%	89%	1.2%	82%	0.456	0.424	0.496	0.793	0.882	0.905
Proteobacteria	Enterobacteriaceae	Citrobacter	G.Citrobacter	0.0%	20%	0.0%	10%	0.0%	22%	0.0%	18%	0.600	0.737	0.458	1.000	1.000	1.000
Proteobacteria	Enterobacteriaceae	Cronobacter	G.Cronobacter	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	Enterobacter	G.Enterobacter	0.0%	10%	0.0%	20%	0.0%	33%	0.0%	27%	0.716	1.000	0.368	0.586	0.946	0.443
Proteobacteria	Enterobacteriaceae	Escherichia/Shigella	G.Escherichia/Shigella	0.7%	100%	0.2%	70%	0.8%	78%	0.4%	73%	0.517	0.122	0.492	0.604	0.940	0.702
Proteobacteria	Enterobacteriaceae	f_Enterobacteriaceae_g_Unc	G.f_Enterobacteriaceae_g_Unc	0.0%	20%	0.0%	30%	0.0%	44%	0.0%	27%	0.945	0.861	0.393	0.822	0.609	0.368
Proteobacteria	Enterobacteriaceae	Klebsiella	G.Klebsiella	0.0%	30%	0.0%	20%	0.0%	33%	0.0%	36%	0.599	1.000	0.536	0.600	0.918	0.647
Proteobacteria	Enterobacteriaceae	Morganella	G.Morganella	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	Pantoea	G.Pantoea	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	Enterobacteriaceae	Pragia	G.Pragia	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	Proteus	G.Proteus	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	Raoultella	G.Raoultella	0.0%	0%	0.0%	10%	0.0%	44%	0.0%	18%	0.862	1.000	0.098	0.476	0.229	0.033
Proteobacteria	Enterobacteriaceae	Tatumella	G.Tatumella	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	Enterobacteriaceae	Trabulsiella	G.Trabulsiella	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	Phyllobacteriaceae		F.Phyllobacteriaceae	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	Defluvibacter	G.Defluvibacter	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	Rhizobiaceae		F.Rhizobiaceae	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	Rhizobiaceae	Rhizobium	G.Rhizobium	0.0%	0%	0.0%	0%	0.0%	11%	0.0%	0%	1.000	1.000	0.474	1.000	0.450	0.474
TM7			P.TM7	0.0%	10%	0.0%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	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%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	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%	0%	0.0%	0%	0.0%	9%	1.000	1.000	1.000	1.000	1.000	1.000
Verrucomicrobia			P.Verrucomicrobia	0.0%	10%	0.0%	20%	0.0%	0%	0.0%	0%	0.214	0.737	0.474	0.476	1.000	1.000
Verrucomicrobia	Verrucomicrobiaceae		F.Verrucomicrobiaceae	0.0%	10%	0.0%	20%	0.0%	0%	0.0%	0%	0.214	0.737	0.474	0.476	1.000	1.000
Verrucomicrobia	Verrucomicrobiaceae	Akkermansia	G.Akkermansia	0.0%	10%	0.0%	20%	0.0%	0%	0.0%	0%	0.214	0.737	0.474	0.476	1.000	1.000