

Table 1 showing the difference in abundance of major phyla and families in diarrheal and non-diarrheal groups

Phyla	Actinobacteria	Proteobacteria	Firmicutes	Bacteroidetes	Verrucomicrobia	Fusobacteria	TM7	Tenericutes	Synergistetes
Diarrheal	17.122	21.7584	51.97764	7.72244	0.65012	0.20918	0.20682	0.00868	0.00126
Non-diarrheal	12.124	3.01725652	42.21951	41.7104	0.6184304	0.01335652 2	0.02928	0.22083	0.0110043
t-test	0.5894	6.74676.	1.17297	-4.53844	0.02848	3.741	2.11332	-0.93385	-0.69884
p-value	0.5606	< .00001	0.251443	0.000114	0.977495	0.000915	0.04433	0.358973	0.490857
Families	Lachnospiraceae	Lactobacillaceae	Erysipelotrichaceae	Prevotellaceae	Streptococcaceae	Enterobacteriaceae	Coriobacteriaceae	Veillonellaceae	Ruminococcaceae
Diarrheal	1.682	0.28276	1.370	0.361	30.1609	9.6501	1.409	1.22158	8.38546
Non-diarrheal	5.724 7478 26	0.335078 261	6.567 1391 3	27.17 99	1.72402 6087	2.0259	4.643 8347 83	7.26865 217 4	17.3614 087
t-test	-2.473	-0.096	-1.334	-3.065	4.710	4.248	-1.509	-2.1869	-1.72
p- value	.010 signifi- cant	.462303	.0968 signifi- cant	.00502 signifi- cant	.000036 significa- nt	.000122 significnt	.0716	.01896 significa- nt	.048693 significa- nt

Table 2 showing the correlation between different families in diarrheal and non-diarrheal groups

	Prevotellaceae Vs Streptococcaceae	Prevotellaceae Vs Enterobacteriaceae	Lachnospiraceae Vs Enterobacteriaceae	Lachnospiraceae Vs Streptococcaceae	Enterobacteriaceae vs Ruminococcaceae
Diarrheal					
R	-0.4897	-0.3172	-0.3548	0.7362	-0.1944
p-value	.4132	0.6	0.559	0.156	0.75
Non-diarrheal					
R	0.1198	0.2399	-0.4426	-0.288	-0.2575
p-value	0.586	.270	0.03 significant	0.183	0.23