

Table S2. Statistical data of the analysis of amplicons.

	Total number of reads	Median number of reads	Average number of reads	Number of ASVs	Number of singletons	Average sequence length
Bacteria	449694	20816	18737	1628	177	417
Archaea	17342	993	1445	87	2	339
Protists	171722	3432	7155	562	58	382