

**Supp. Table 4.** Classification of all the metagenomic reads using MGTAXA

<b>Kingdom</b>	<b>Phylum</b>	<b>Class</b>	<b>Percentage of reads</b>
Bacteria	<i>Actinobacteria</i>	<i>Actinobacteria</i>	1
	<i>Bacteroidetes</i>	<i>Bacteroidia</i>	1
	<i>Cyanobacteria</i>		1
	<i>Firmicutes</i>	<i>Bacili</i>	13
		<i>Clostridia</i>	3
		Other	1
	<i>Proteobacteria</i>	<i>Alphaproteobacteria</i>	3
		<i>Betaproteobacteria</i>	2
		<i>Delta/epsilon</i>	3
		<i>Gammaproteobacteria</i>	39
	Other		5
Archaea			1
Eukaryote			2
Viruses			14