

Table S1 The result of random sequencing

singlet number	92
Average sequence length	656 bp
Results of sequences blast in NCBI	1 singlet had $\geq 97\%$ similarity with <i>Bacteroides thetaiotaomicron</i> (1.08% of all sequenced singlets). 2 singlets had $\geq 97\%$ similarity with uncultured <i>Bacteria</i> (2.16% of all sequenced singlets). 89 singlets didn't have high similarity with known sequences (96.76% of all sequenced singlets).