

Table S1. Summary of results of genome finishing for <i>E. rectale</i> strain ATCC 33656 and <i>E. eligens</i> strain ATCC 27750.		
	<i>E. rectale</i>	<i>E. eligens</i>
Genome size (bp)	3,449,685	2,831,389
Plasmids	0	2 (626 kb, 60 kb)
Predicted proteins	3,606	2,773
tRNAs	57	47
rRNAs	15	15
other RNAs	30	17
ABI 3730xl Plasmid reads	120,005 reads (9.6x)	37,846 reads (4.0x)
454 GS20 Pyrosequencer	120 Mb (40x)	114 Mb (40x)
Finishing reactions	204	104