

Table S1. Summary of the sequencing statistics obtained from each sample.

Sample	Raw_Reads	Raw_Bases	Clean_Reads	Clean_Bases	Error_Rate	Q20	Q30	GC_PCT
CT1	48070372	7.21G	47231216	7.08G	0.01	99.26	97.63	47.05
CT2	42301748	6.35G	41509158	6.23G	0.01	99.16	97.34	46.62
CT3	52611534	7.89G	51776016	7.77G	0.01	98.54	96.05	46.63
MT1	47220058	7.08G	46251992	6.94G	0.01	99.15	97.39	46.84
MT2	48024112	7.2G	46880808	7.03G	0.01	99.16	97.42	46.93
MT3	43239912	6.49G	42366370	6.35G	0.01	99.27	97.62	47.28