

Table S1 Quality assessment of sample sequencing data

Sample name	Raw Reads	Clean Reads	Clean Bases(Gb)	Error(%)	Q20(%)	Q30(%)	GC(%)
SAC1	17719104	17425772	2.61G	0.03	97.45	92.72	35.3
SAC2	20159534	19641306	2.95G	0.03	97.33	92.45	35.85
SAC3	20159288	19752162	2.96G	0.03	97.47	92.77	35.21
SAQ_BITC1	17509930	17015102	2.55G	0.03	97.17	92.09	35.14
SAQ_BITC2	19672798	19245460	2.89G	0.03	97.24	92.26	35.37
SAQ_BITC3	13703234	13327608	2G	0.03	97.45	92.71	35.43