

Supplemental Table S6. Summary of HIFI genome sequencing data.

Library id	Total bases(nt)	Total reads	Mean length(nt)	Max length(nt)	N50 length(nt)	>10kb rate(%)	>20kb rate(%)	>40kb rate(%)
m64062_221024_06 4757	31,028,652,285	1,723,553	18,003	49,230	17,797	99.94	23.25	0.01
m64062_221031_18 1436	34,304,552,556	1,893,868	18,113	50,019	17,936	99.93	24.5	0.01