

Table S3 Quality statistics of sample sequencing data

Bulk	Total	Valid	Average Depth (×)	Q20 (20%)	Q30 (30%)	GC%	SNP Number
FL	102 431 528 (100.00%)	100 362 822 (97.98%)	28.55	98.04	95.16	37.88	2 556 275
RL	102 403 516 (100.00%)	100 320 998 (97.97%)	28.47	98.08	95.28	38.09	2 497 086