

Table 4 Summary of sequencing data

Sample	Clean_Reads	Q20 (%)	Q30 (%)	Sequencing depth	GC content (%)
ZK	51457328	94.43	90.09	33	37.03
DDX	55249477	93.23	87.9	36	35.86
L_pool	93914419	93.41	88.21	64	36.03
H_pool	78432324	93.83	88.97	51	36.33