

Table S3 The whole genome sequencing reports of *mp* and CCMC pools.

Sample	Clean bases	Sequencing depth	Map reads rate (%)	Clean GC (%)	Clean Q20 (%)	Coverage rate (%)
CCMC-pool	111,165,212	82.26	58	37.13	96.44	93.43
<i>mp</i> -pool	177,657,038	131.47	51.73	39.7	92.99	93.35