

表 2 测序数据与参考基因组比对情况

Table 2 Alignment statistics of sequencing data to the reference genome

编号 ID	测序深度 Ave-depth	1 个 read 覆盖率 (%) Cov-ratio-1x (%)	5 个 read 覆盖率(%) Cov-ratio-5x (%)	10 个 read 覆盖率 Cov-ratio-10x (%)
301	21	92.43	88.37	83.13
294	20	92.62	88.29	82.41
长柄混池 SP-pool	21	95.74	91.28	84.52
短柄混池 LP-pool	20	96.20	91.11	83.69