



Supplemental Fig. S8. Comparison of SNP calling between SAMtools and GATK tool. About 8.11 M and ~7.77 M SNPs for each individual were detected using SAMtools³ and GATK tool⁴, respectively. About 7.41 M SNPs were concurrently identified by two algorithms, which accounted for 91.24% and 95.34% SNPs identified by SAMtools and GATK, respectively.