

Supplementary Fig. 1s, Ploting result of a genome-wide BSA-seq analysis. The plot was created using the QTL-seq package of R (Core Team, 2020) with a 2.0-Mb window size and 10-kb step size. Red line,  $\Delta$ SNP index; Orange line,  $\rho$  < 0.01; Green line,  $\rho$  < 0.05.