

Description of Additional Supplementary Files

File name: Supplementary Data 1

Description: Twenty-seven annotated genes were differentially expressed between HKW9^{Mc} and HKW9^{V671} within 73-Mb mapping region.

File name: Supplementary Data 2

Description: The maize inbred lines list in this study.

File name: Supplementary Data 3

Description: Primers and probes used in this study.