Supplemental Fig. S1. Standard errors associated with the genetic effects of 308 single-cross hybrids and 4 common checks, obtained via best linear unbiased predictor (BLUP) through the multi-environment trial analysis (model [2]) performed separately for well-watered (WW) and water-stressed (WS) groups of trials. GY: grain yield (t/ha); EPP: number of ears per plot; ASI: anthesis-silking interval (days); FFT: female flowering time (days); and MFT: male flowering time (days). The checks correspond to the last four hybrids in the curves.
Supplemental Fig. S2. Genetic effects of 308 single-cross hybrids and 4 common checks, obtained via best linear unbiased predictor (BLUP) through the multi-environment trial analysis (model [2]) performed separately for well-watered (WW) and water-stressed (WS) groups of trials. GY: grain yield (t/ha); EPP: number of ears per plot; ASI: anthesis-silking interval (days); FFT: female flowering time (days); and MFT: male flowering time (days).