Supplemental Figure 1. Coalescent simulation model. $N_a$, the ancestral population size; $\mu$, the mutation rate; $\rho$, the population-recombination parameter; $t$, the time of the split between maize and teosinte; $d$, the duration of the maize population bottleneck; $k$, the severity of the bottleneck; $N_b$, the population size of maize during bottleneck.