



Figure S3: Neighbor-joining tree of the sequenced region in the 5' UTR (right; Region 1) and the 66,169 bp upstream region (left; Region 2) of *tb1*. Individuals with genotype data are colored: Homozygous for the teosinte (no *Hopsotch*) allele (red), homozygous for the maize (*Hopsotch*) allele (blue), heterozygotes (purple). TILs (teosinte inbred lines) are colored in green, with stars indicating the 3 TILs known to have the *Hopsotch* insertion. Black indicates individuals not genotyped for the *Hopsotch* insertion. EjuA refers to individuals from population Ejutla A, EjuB from Ejutla B, SLO from San Lorenzo, and MSA from La Mesa. Remaining individuals are lines of maize (*Zea mays* ssp. *mays*).