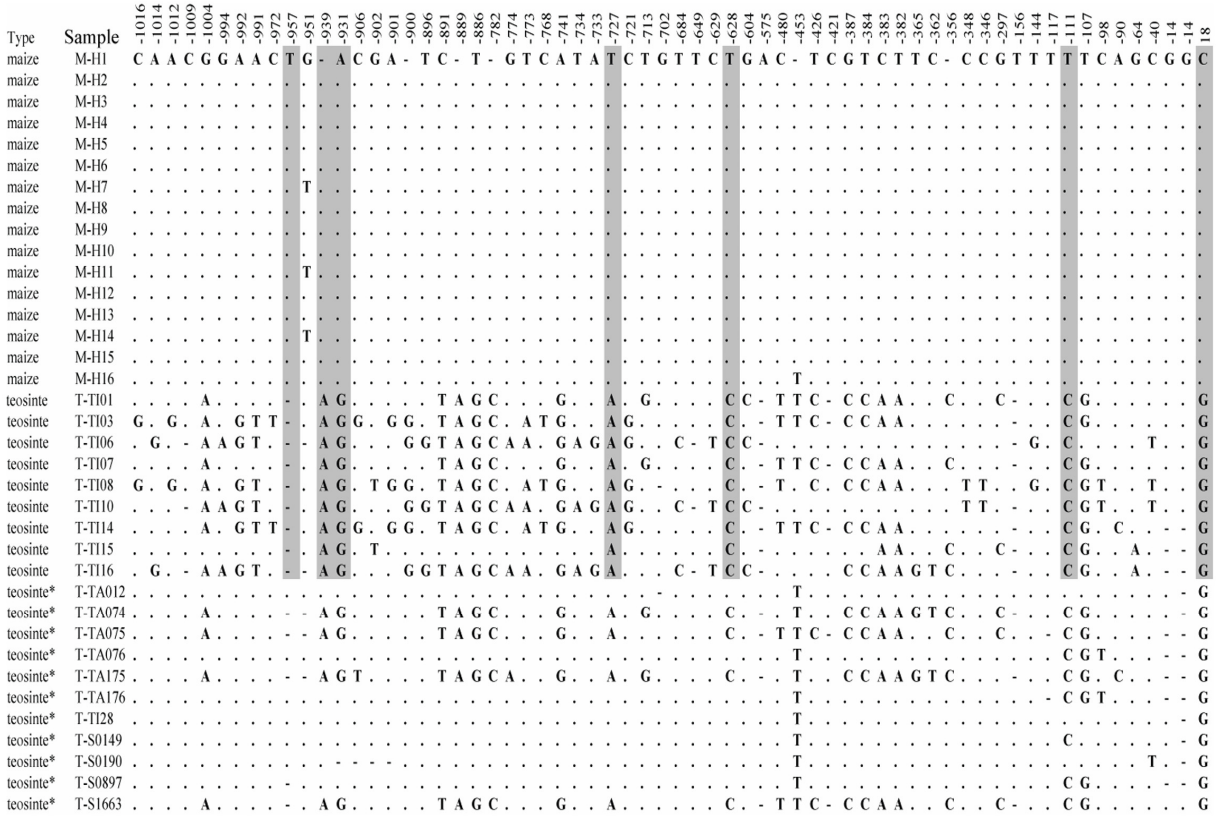

Figure S1 Nucleotide sequence alignment (polymorphic sites only) of the 1042-bp region of *tga1* identified as containing the causative site (WANG *et al.* 2005). A sample of maize and teosinte genotypes showing that the only fixed difference between maize and teosinte is in the 18th nucleotide of the open reading frame. Numbers in the top row indicate the nucleotide positions corresponding to the W22:*tga1* sequence. Indel polymorphisms were shown as single mutations. Dots indicate identity to the sequence in the top row. The seven "fixed" differences initially identified by Wang *et al.* (2005) are shown by shading. Eleven teosinte individuals not sampled by Wang *et al*. (2005) are indicated by asterisks.