

Figure S1. Phylogenetic tree based on multiple sequence alignment of GLK proteins from *Arabidopsis thaliana*, *Gossypium hirsutum*, *Nicotiana tabacum*, and *Zea mays*. See the main text of the paper for corresponding references of the known genes from *Arabidopsis* (At-), cotton (Gh-), tobacco (Nt-), and maize (Zm-).

