**Supplementary Figure S2** Phylogenetic analysis of Arabidopsis and kiwifruit XTH protein sequences. Reference Arabidopsis proteins are in blue and predicted XTH proteins from *Actinidia chinensis* var. *chinensis* gene models are in black. Amino acid sequences were aligned using the Geneious software package (https://www.geneious.com).The phylogenetic tree was produced using PHYML. Bootstraps values from 1000 iterations are given.

