



Supplemental Figure 3. Rooted phylogenetic tree of *Arabidopsis* group I bZIP proteins (Tsugama et al., 2014). Asterisks indicate null mutant lines of six VIP1 homologs that were tested for *Agrobacterium*-mediated transformation susceptibility. Bar: amino acid substitutions per site. Indicated expression levels are based on results of reverse transcription-PCR performed previously (Tsugama et al., 2014).