



Figure S2 Phylogenetic analysis of histone deacetylase homolog protein in plants.

The Bayesian inference tree was constructed with MrBayes v3.2.7a. At: *Arabidopsis thaliana*; Cs: *Camellia sinensis*; Gm: *Glycine max*; Os: *Oryza sativa*; Sl: *Solanum lycopersicum*; Sb: *Sorghum bicolor*.