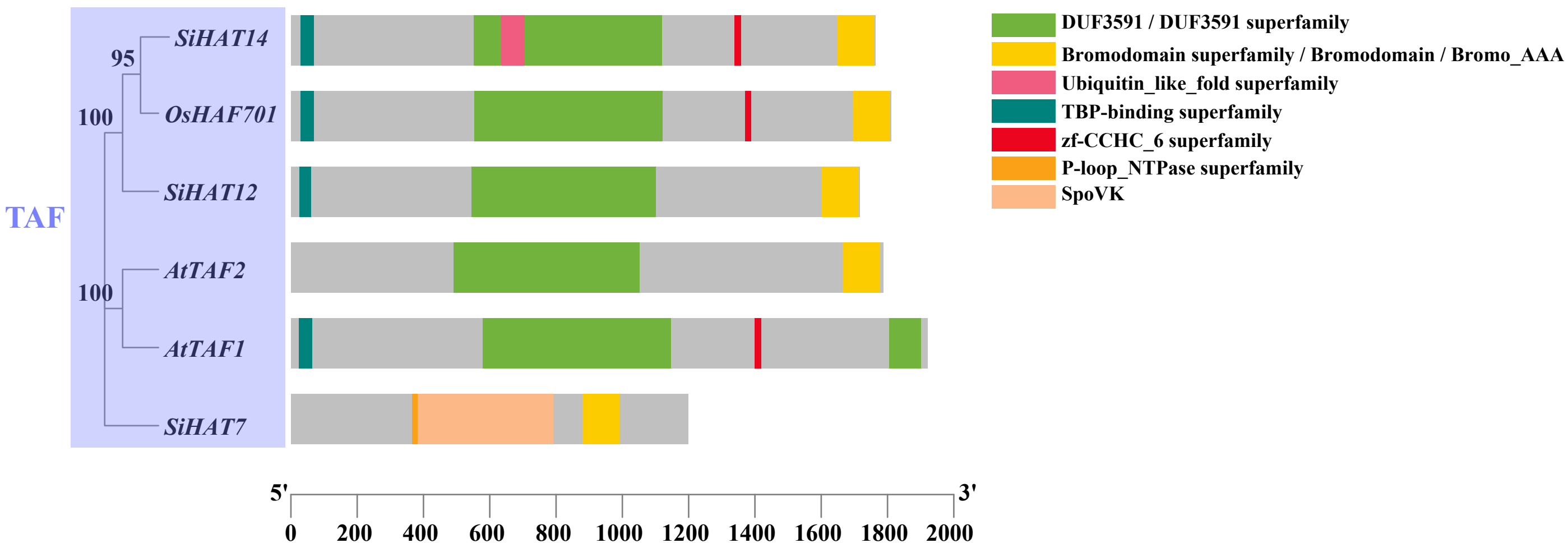




Additional file 6. Phylogenetic trees and domain composition of TAF subfamily. Phylogenetic tree and domain composition of TAF subfamily predicted proteins from *Arabidopsis thaliana* (*At*), *Oryza sativa* (*Os*) and *Setaria italica* (*Si*). Conserved domains include DUF3591/DUF3591 superfamily, Bromodomain superfamily/Bromodomain/Bromo_AAA, Ubiquitin_like_fold superfamily, TBP-binding superfamily, zf-CCHC_6 superfamily, P-loop_NTPase superfamily and SpoVK.