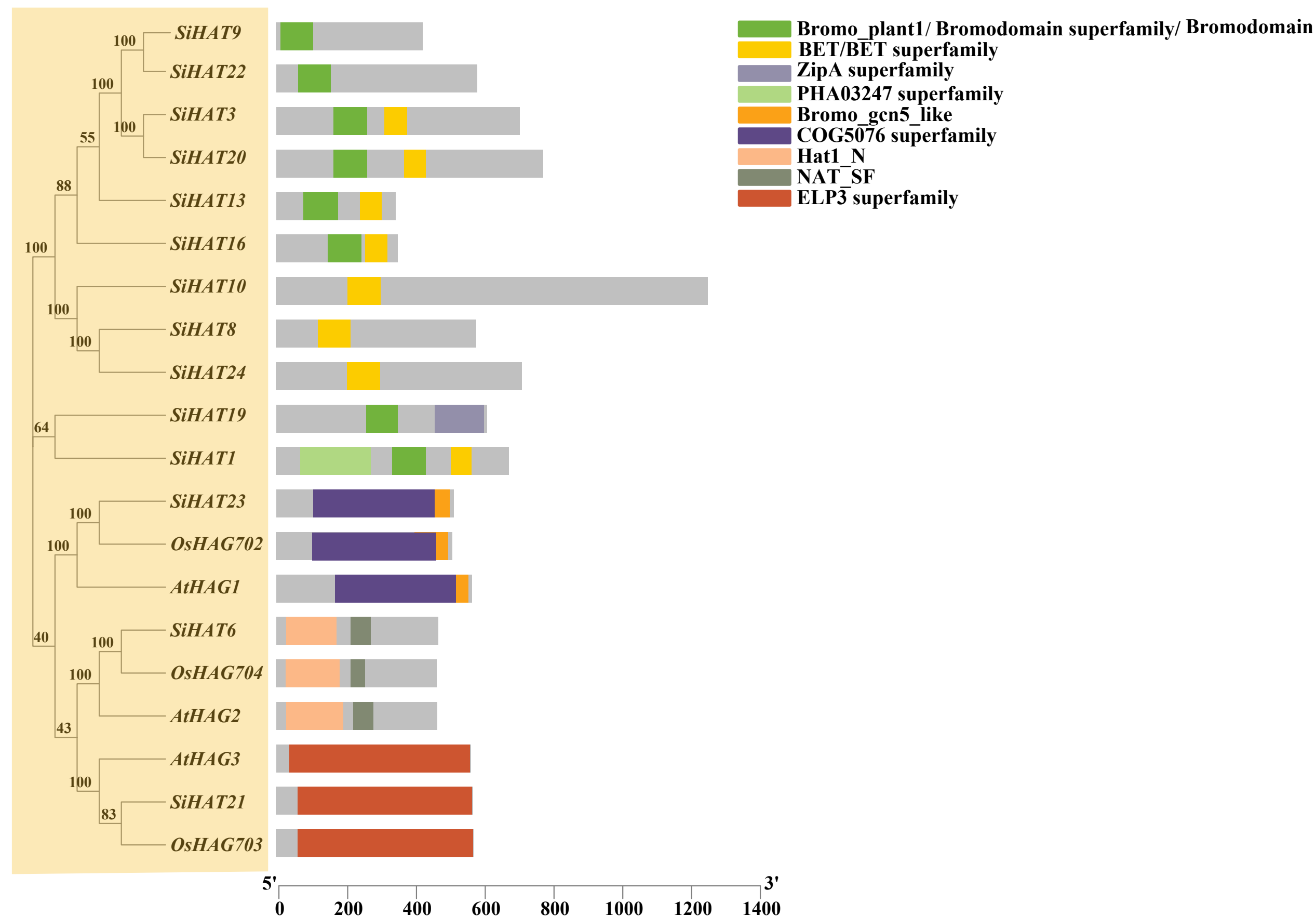


GNAT



Additional file 3. Phylogenetic trees and domain composition of GNAT subfamily. Phylogenetic tree and domain composition of GNAT subfamily predicted proteins from *Arabidopsis thaliana* (At), *Oryza sativa* (Os) and *Setaria italica* (Si). Conserved domains include Bromo_plant1/Bromodomain superfamily/Bromodomain/Bromo_gcn5_like, BET/BET superfamily, ZipA superfamily, PHA03247 superfamily, COG5076 superfamily, Hat1_N, NAT_SF and ELP3 superfamily