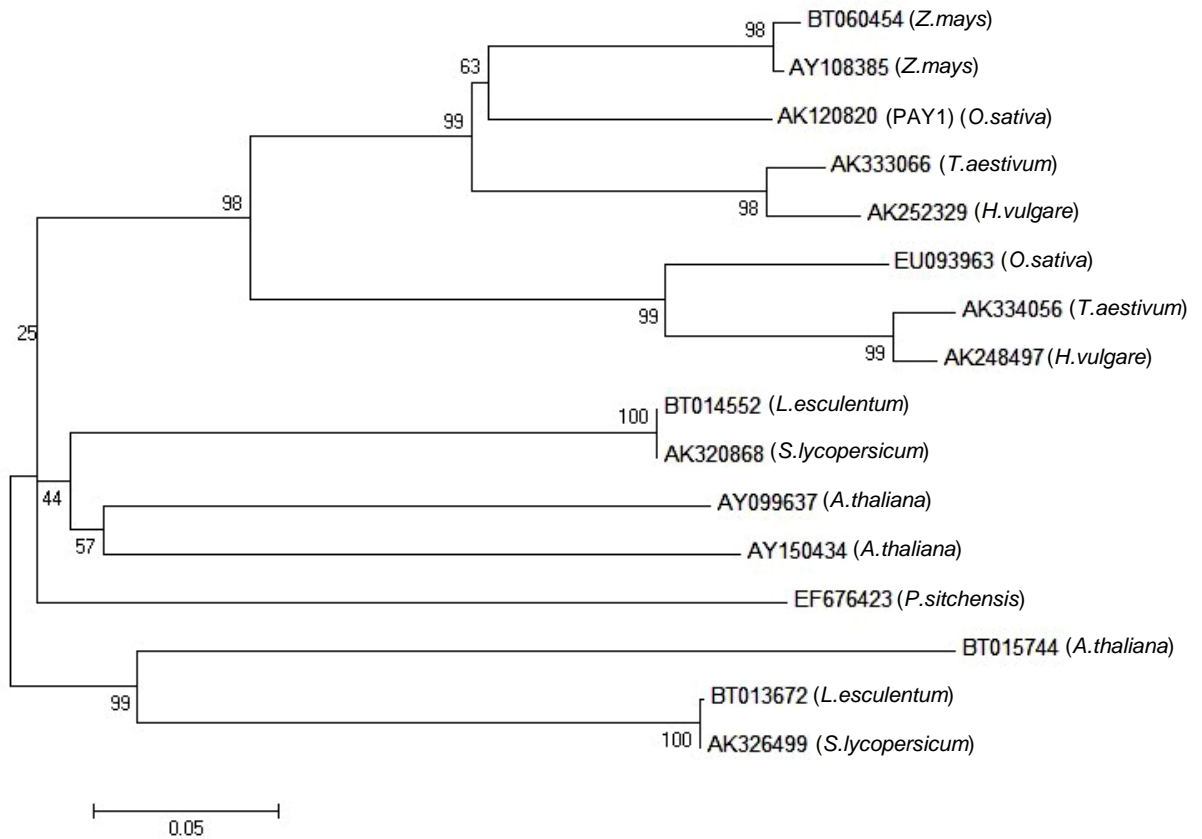


Figure S5



Supplementary Fig. 5 Phylogeny of the PAY1 protein family. A neighbor-joining tree was built by MEGA5 using a Poisson correction model with gaps to complete deletion. Topological robustness was assessed by bootstrap analysis with 500 replicates. The bar is an indicator of genetic distance based on branch length.