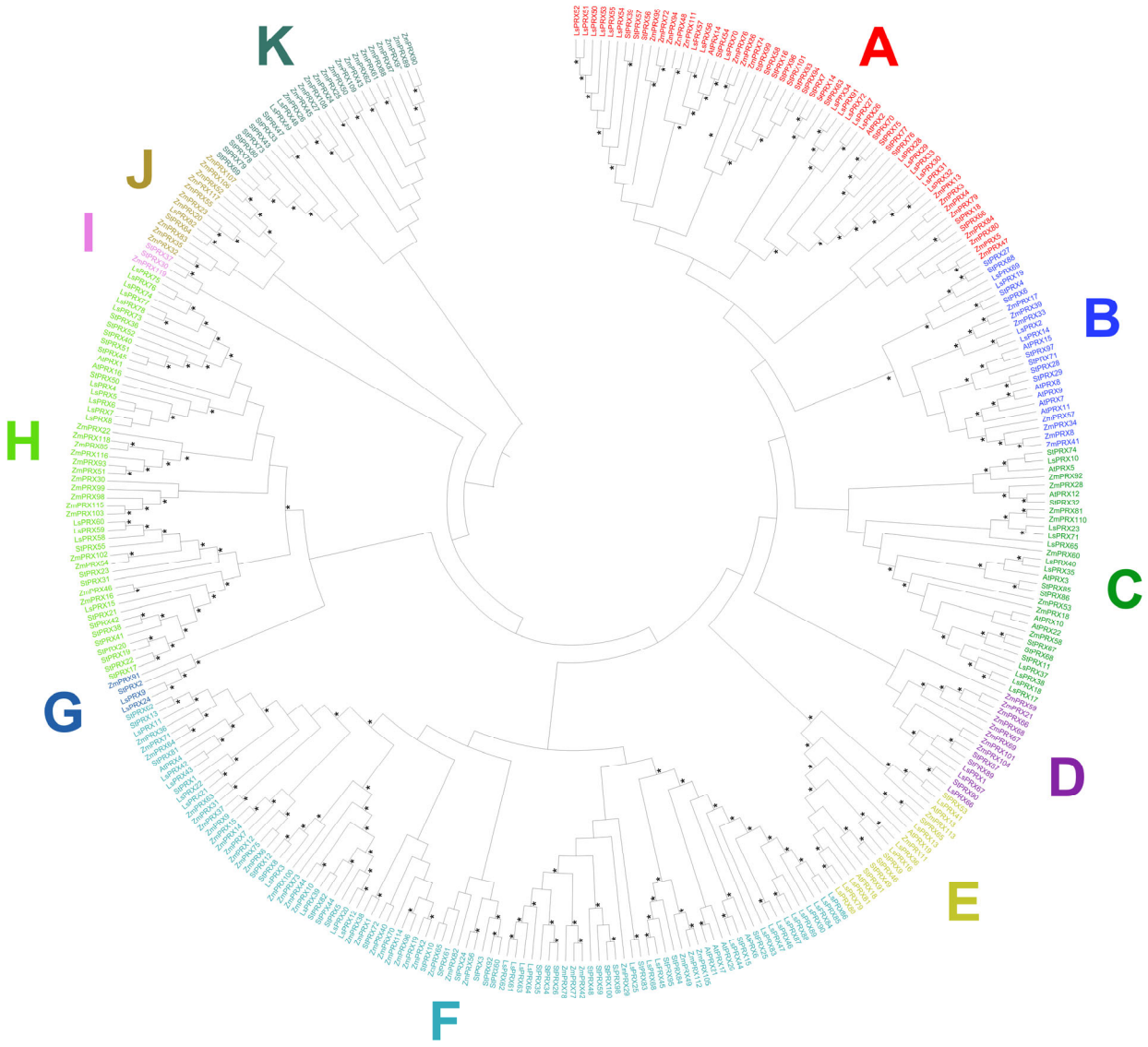


Supplementary Figure 3. Phylogenetic tree of PRX protein sequences from *L. sativa*, *A. thaliana*, *Z. mays*, and *S. tuberosum*, constructed using the maximum likelihood method. * indicates Bootstrap values above 60%.