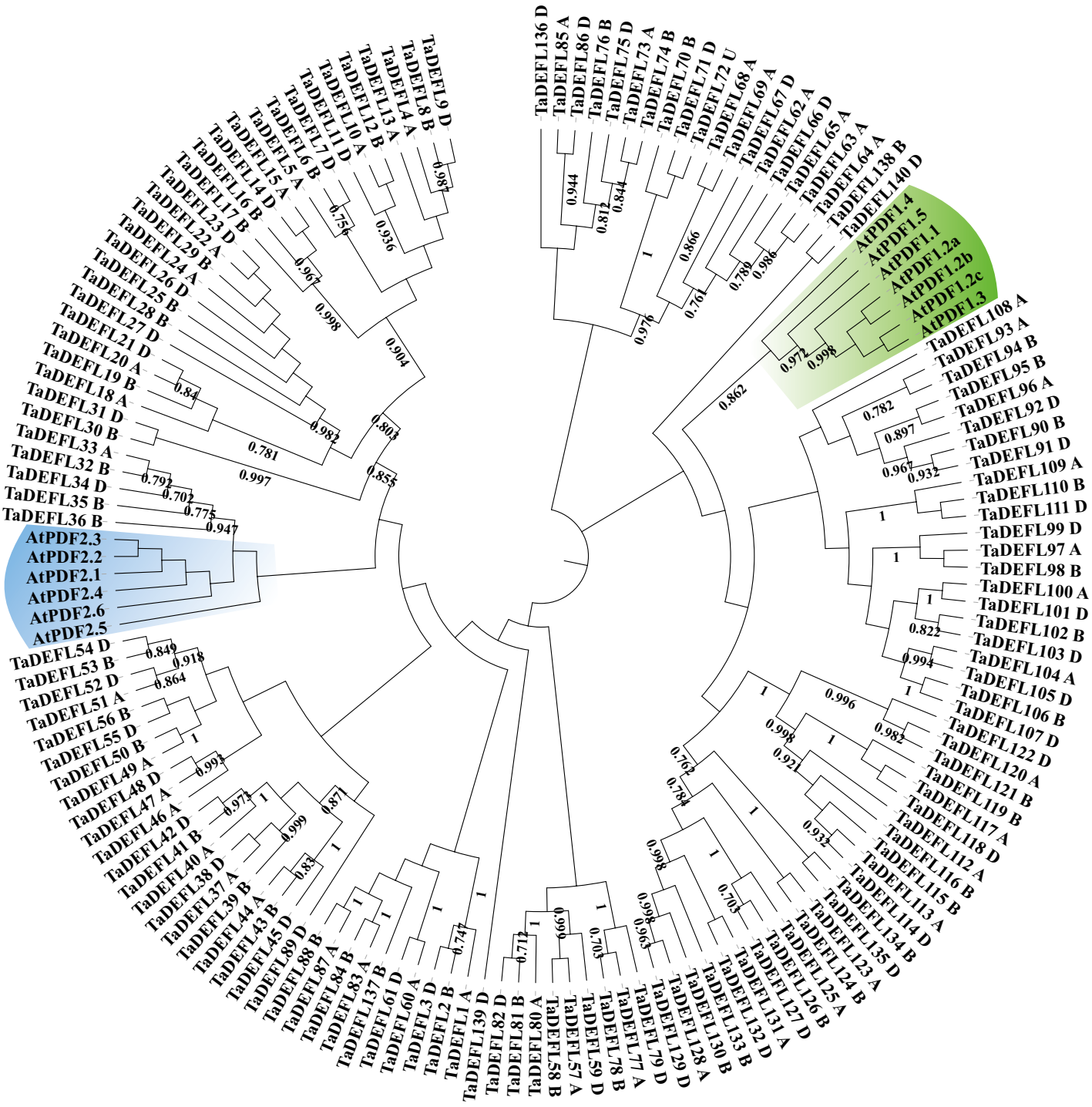

Figure S11. Plyhogenetic analysis of 140 TaDEFLs, 7 AtPDF1s and 6 AtPDF2s. The phylogenetic tree was constructed using MEGA by the NJ method with 1000 bootstrap replications, and the bootstrap values greater than 700 were displayed on the branches.