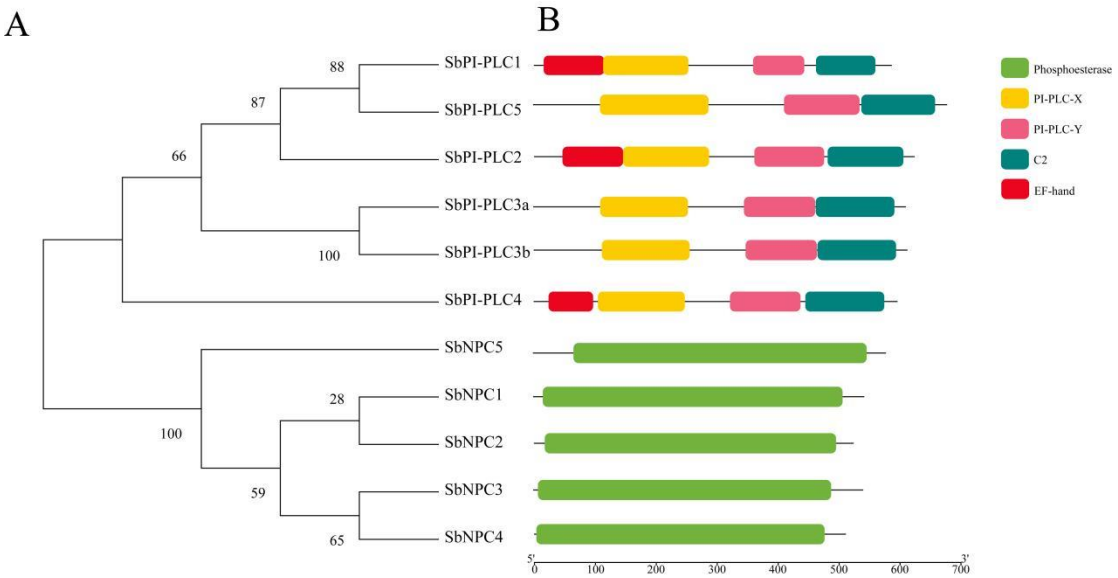


Fig. S1 Conserved Domain of SbPLC Proteins. (A) Phylogenetic tree of SbPLCs. The tree was generated using MEGA7 through the Maximum Likelihood method based on the protein sequences of SbPLCs. (B) Conserved Domain of SbPLC Proteins.

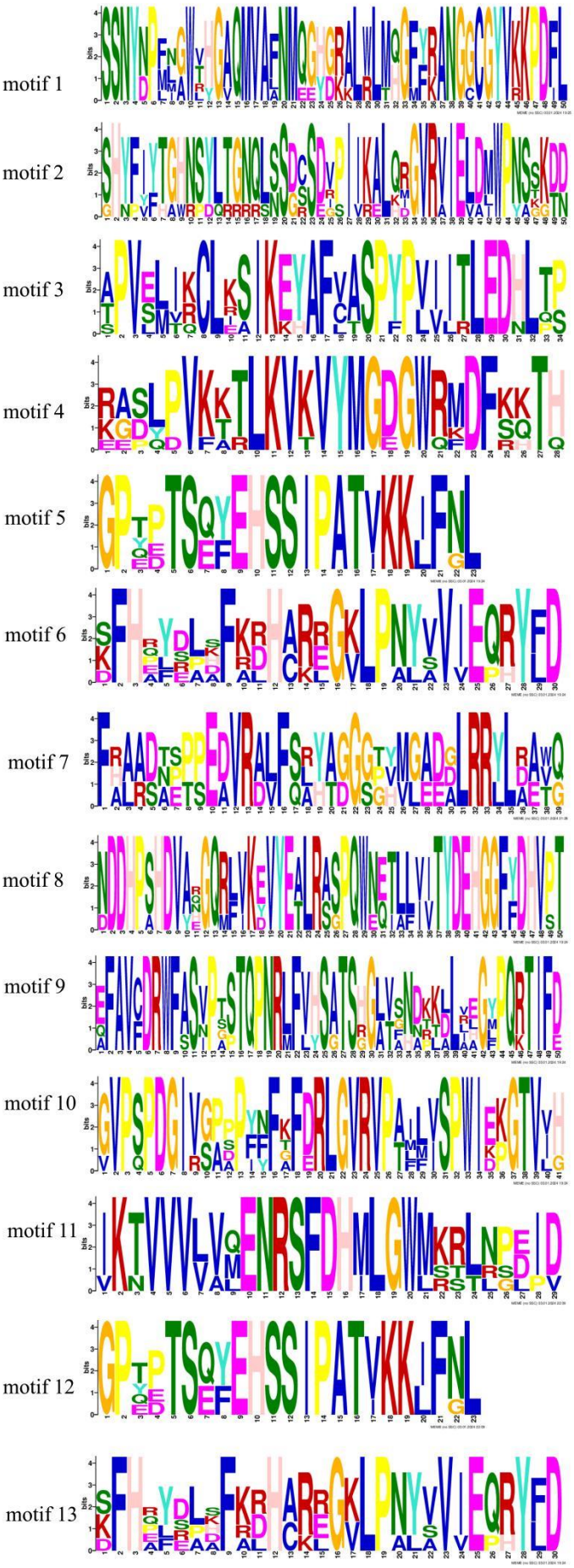


Fig. S2 The *SbPLCs* motifs. The overall height of each stack indicates theconservation of the sequence at that position, whereas the height of letters within eachstack represents the relative frequency of the corresponding amino acid.