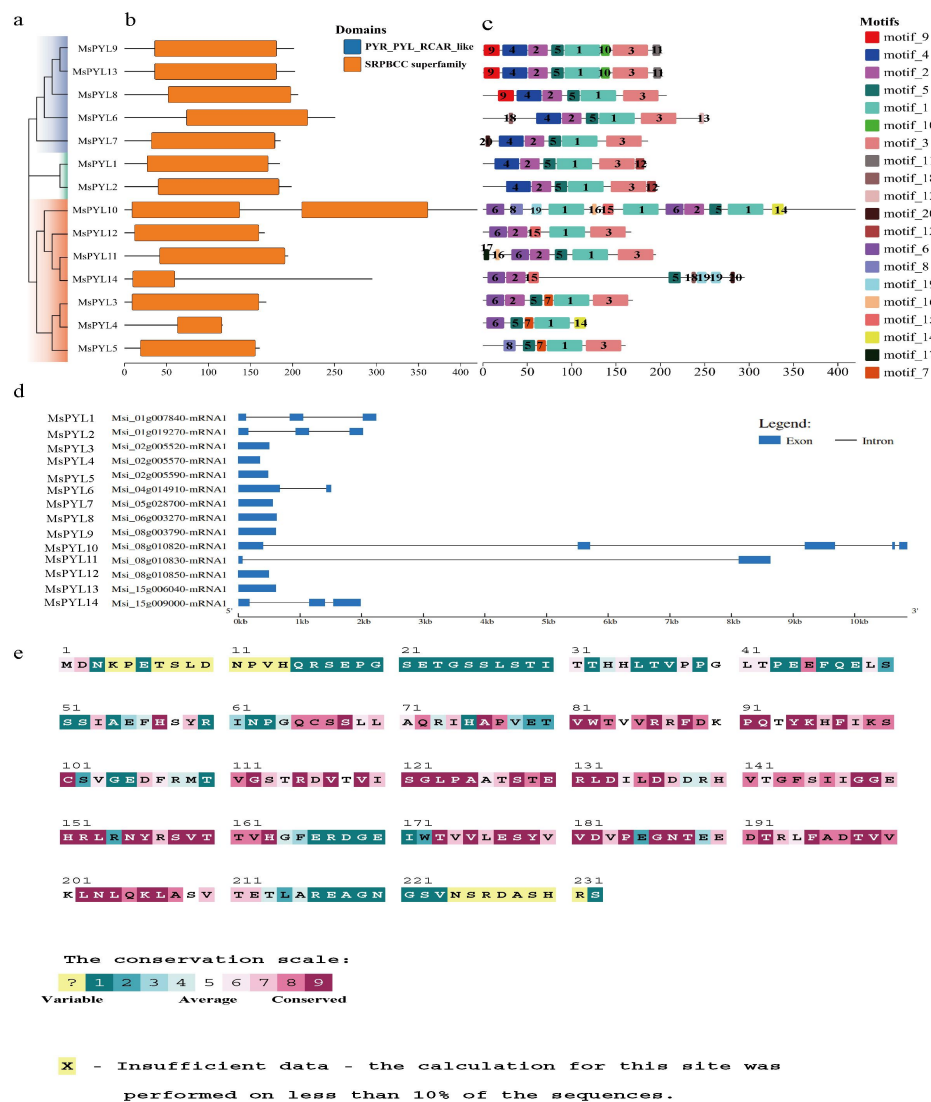




Supplemental Fig.S2 Conserved motifs, gene structure and amino acid sequence conservation of PYL in *Malus sieversii*. (a) Phylogenetic tree. (b) Distribution of conserved domains. (c) Conserved motif distribution. (d) Gene structure: exon-intron distribution map. Blue boxes indicate exons, black lines indicate introns. (e) Identification of amino acid conservation of homologous PYL proteins.