

Supplemental figure 1-3

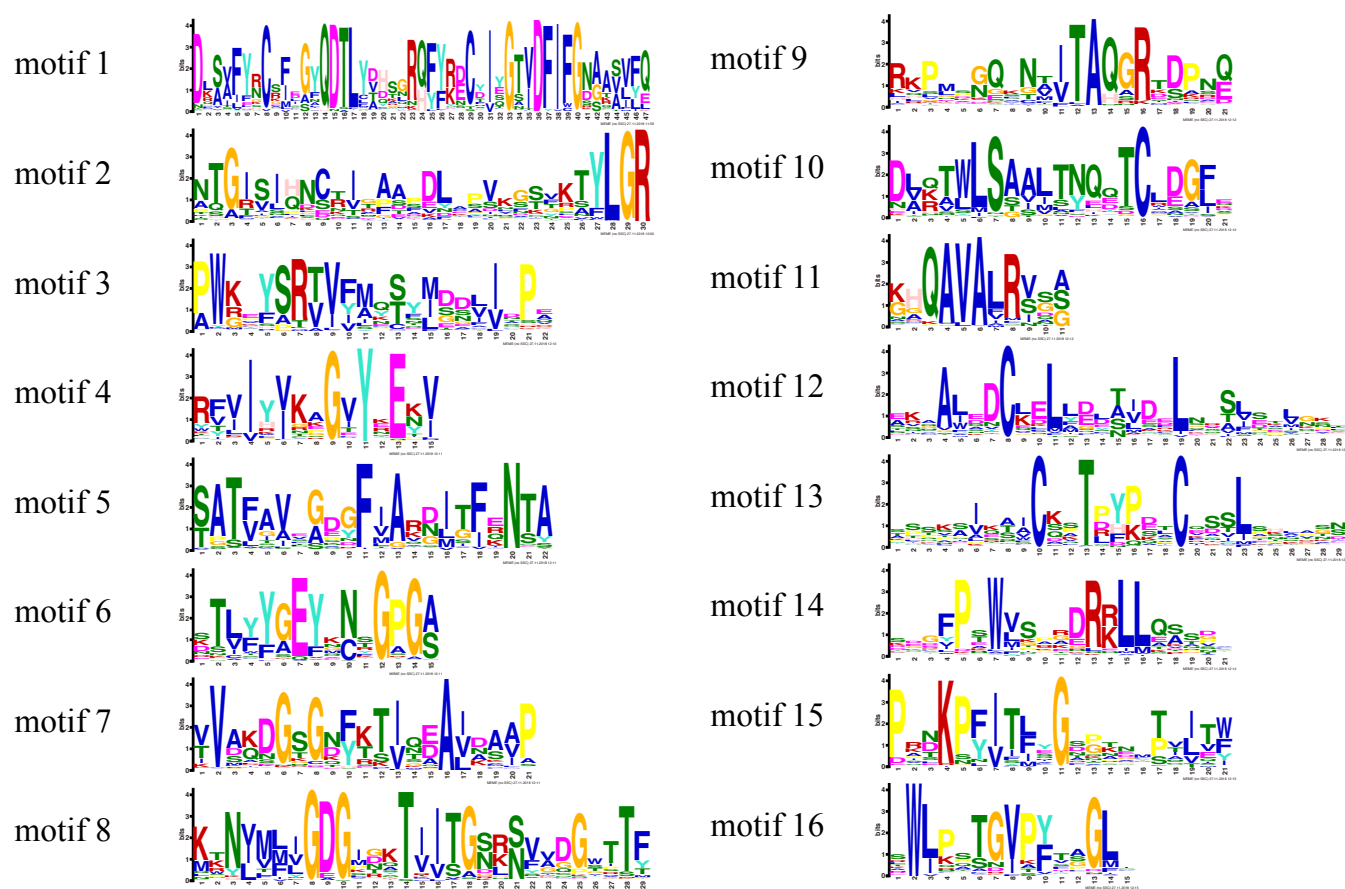


Figure S1 Logo of conserved motifs in Figure 3.

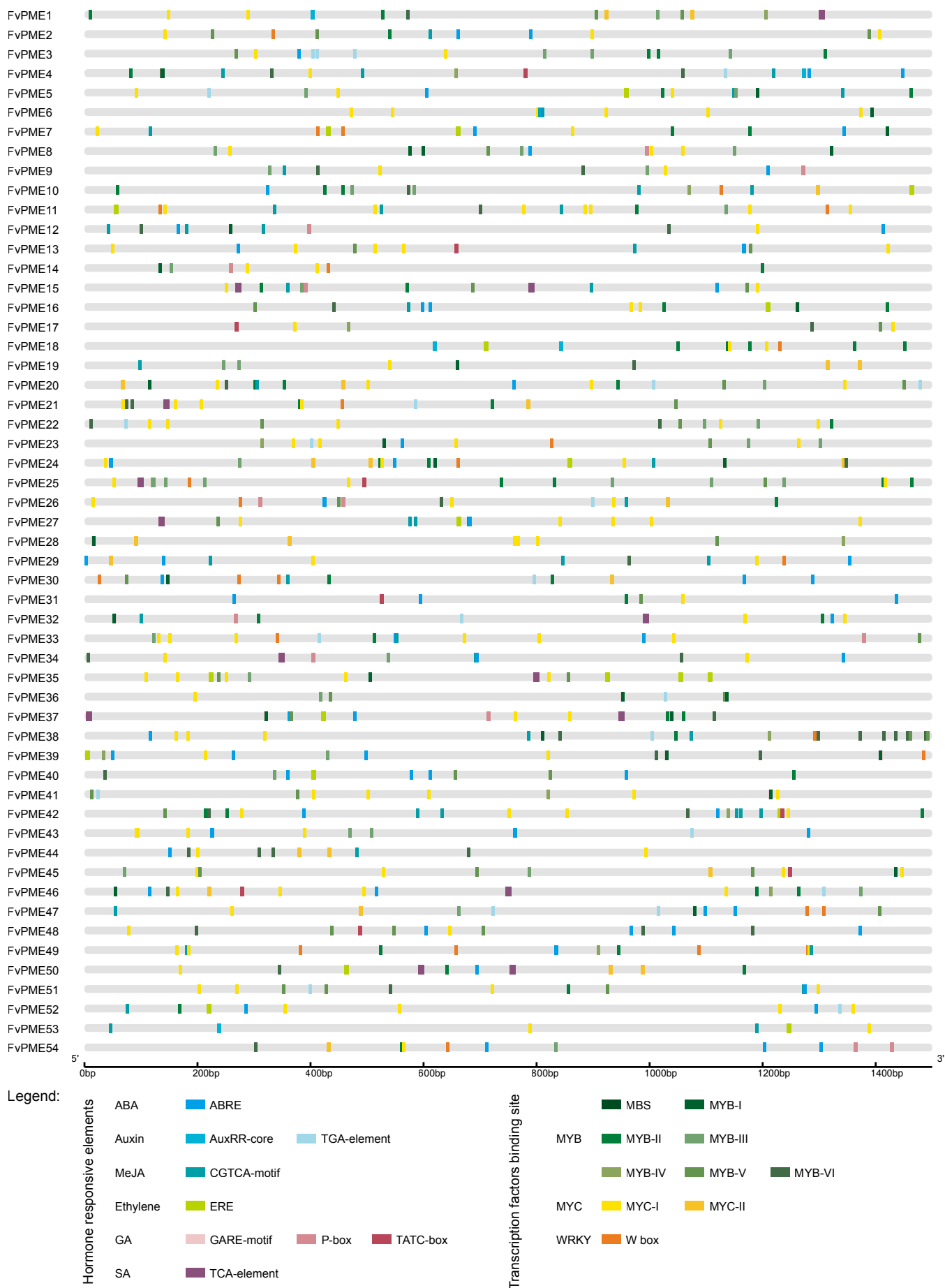


Figure S2 Putative cis-elements in the 1.5 kb promoter region of FvPMEs. Those cis-elements could be distinguished from each other with different color, and their relative position are displayed.

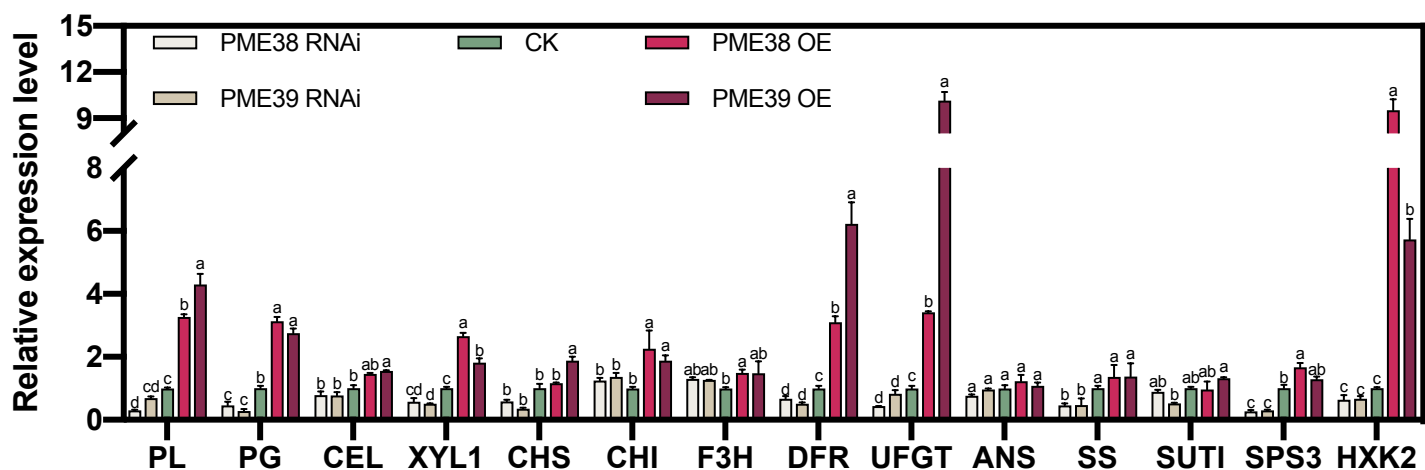


Figure S3 QRT-PCR analysis of transcript levels for ripening-related genes in fruits of transient overexpression or silencing of *FvPME*. Error bars represent SD of three independent replicates. Relative expression levels of each gene were normalized to internal control *FvGAPDH*.