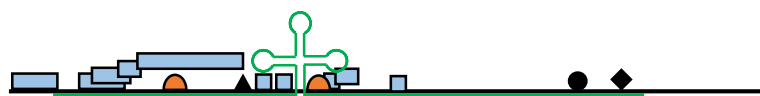


eRTBVL and eRTBVL2



ePVCVL and ePVCVL2



S5 Fig. Predicted elements of noncoding regulatory sequences in IGRs. The putative short ORF (blue box), stem-loop structure (green line), promoter (orange semi-circle), polyadenylation signal (black triangle), primer binding site (black circle), and polypurine tract (black diamond) in the IGR (black line) are shown for eRTBVL/eRTBVL2 and ePVCVL/ePVCVL2. Consensus sequences of eRTBVL-X, eRTBVL2, ePVCVL, and ePVCVL2 constructed from the *O. sativa* genome were used for predictions.