



Figure S1. Logo of the conserved motifs of the HD-ZIP III proteins.

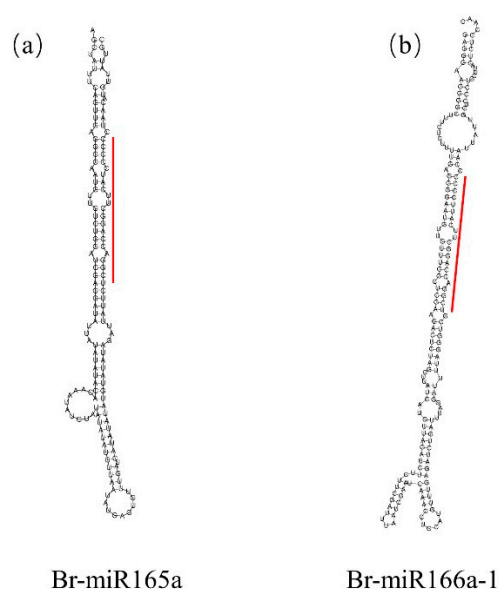


Figure S2. Prediction of miR165/166 stem-loop structures that produced miR165/166 in *B. rapa* using an RNAfold web server.

Predicted secondary stem-loop structures of miR165a (a) and miR166a-1(b) using the miRNA precursor sequences from *B. rapa*. Red lines indicate the mature miRNA sequences.