## **Description of Additional Supplementary Files**

File Name: Supplementary Data 1

Description: Read mapping statistics and grouping information of each sRNA-seq experiment.

File Name: Supplementary Data 2

Description: Information of MIRNA loci identified by ShortStack.

File Name: Supplementary Data 3

Description: The number of DEMs in each comparison.

File Name: Supplementary Data 4

Description: Mean expression levels of identified MIRNA loci and DME analysis for each MIRNA loci.

File Name: Supplementary Data 5

Description: Mean, median, SD for ratios of miRNAs with extreme values (ratio >6 or < 1/6)

excluded.

File Name: Supplementary Data 6

Description: Predicted targets of miRNAs identified by psRNATarget.

File Name: Supplementary Data 7

Description: Significant correlations between expression levels of miRNAs and their mRNA targets

predicted by psRNATarget.

File Name: Supplementary Data 8

Description: Significant correlations between expression levels of miRNAs and their mRNA targets

found in degradome sequencing data.