MiRNA-seq-based profiles of miRNAs in mulberry phloem sap provide insight into the pathogenic mechanisms of mulberry yellow dwarf disease

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Supplementary table 1. Primers used RT-PCR and RT-qPCR for mRNA abundance analysis.

Gene	Forward primer (5′-3′)	Reverse primer (5′-3′)
Actin	**CAGTGCTTCTCACTGAGGCTC**	**GGAAGAGGACTTCTGGGCATC**
MmPP16	**GGACCCGATTTCCTGTTCTG**	**ATGTCAGAGATTACCTCAGGC**
RuBisCo	**AATGATGGTGTGACTGTGGCG**	**CAGTGAGAATAGCAATATCGTC**
Regulator of chromosome condensation family protein gene	**CTTAGCGGGATTTTGGATCTG**	**AAGATCAACGGCACTGAGCCT**
Trehalose 6-phosphate synthase gene	**GGACAGCAAATCGTCGAAGTT**	**AGCAATCCGCAGTATGCTTTC**
Inositol 1,3,4-trisphosphate 5/6-kinase family protein gene	**ACCTCAAGTTGCTTGTGGTGT**	**TGGTGTAGACTTCTTAACCGC**
Pri-mul-miR482a	**GGAAAGGGAGATTTGAGCTAC**	**AGAGAATACGGAAGGGAAAGG**