

Table S11. Differentially expressed miRNAs targeting differentially expressed mRNAs during different developmental periods in pollen of Ogura CMS and maintained lines of cabbage

miRNA name	Target gene	NR_annotation
miR158	<i>Bra027656</i>	restorer-of-fertility
miR159	<i>Bra000531</i>	Predicted protein
	<i>Bra022962</i>	Hypothetical protein ARALYDRAFT_902483
	<i>Bra021791</i>	Putative transcription factor MYB101
	<i>Bra026958</i>	hypothetical protein
miR5654	<i>Bra031731</i>	hypothetical protein
	<i>Bra026836</i>	pentatricopeptide repeat-containing protein(PPR)
	<i>Bra016758</i>	DEAD-box ATP-dependent RNA helicase 47(mitochondrial)
	<i>Bra026884</i>	hypothetical protein
miR860	<i>Bra014184</i>	Cytochrome c biogenesis protein family (CCS1)
miR9569	<i>Bra030219</i>	IAA8
	<i>Bra008407</i>	Metacaspase 5
	<i>Bra012414</i>	Betv I allergen family protein
	<i>Bra013168</i>	H ⁺ -ATPase 6 (AHA6)
	<i>Bra012398</i>	Betv I allergen family protein
novel_mir_354	<i>Bra032694</i>	ATP binding kinase protein kinase
novel_mir_403	<i>Bra024504</i>	cyclin 2a protein [<i>Arabidopsis thaliana</i>]
	<i>Bra037265</i>	cyclin 2a protein [<i>Arabidopsis thaliana</i>]
novel_mir_448	<i>Bra007991</i>	Sucrose transporter(SUC1)
	<i>Bra016103</i>	Sucrose transporter(SUC1)
novel_mir_51	<i>Bra035074</i>	V-type proton ATPase catalytic subunit A-like
novel_mir_95	<i>Bra034358</i>	S-adenosyl-L-methionine-dependent methyltransferases superfamily protein