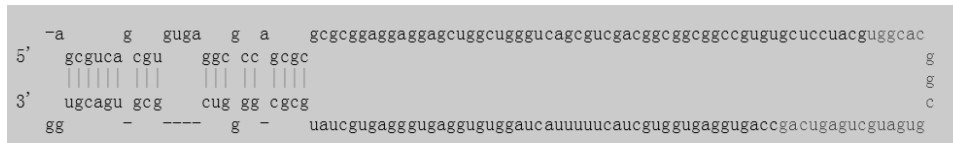


Appendix A PCR primers used in this study

Purpose	Accession number	Forward primer (5'-)	Reverse primer (5'-)
NtMIR1119 expression	MI0006181	AGCGUCAGCGUGUGAGGC	CCACGTCACGCGACCCCGC
<i>Tatubulin</i> expression	U76558	CATGCTATCCCTCGTCTCGACCT	CGCACTTCATGATGGAGTTGTAT
<i>Nttubulin</i> expression	U91563	TACACAGGGGAAGGAATGG	CTCGAAACCAACGGTATC
<i>TabHLH49</i>	XM_020294420	GGCTTGACCTCAACATAGAAG	GTACTCGTGCAGGTGGAGATT
<i>TaLZ</i>	XM_020312111	TGTCCAACAGGGAGTCGGCC	TCCTATGGGTGACCAAGGCC
<i>TaPCF</i>	XM_010236026	GTGGGGGAATGGTTAAGAGCA	TGGGTCGTTCTGGCACTGGT
<i>TaCS</i>	XM_003569426	ACACAGATATGAGGTGAACCC	ATCCATCATCATGTTCTCAAGG
<i>TaMNT</i>	XM_020325679	CATCGGCGTCTCCATGATCCT	TATAACTTGGTTAACCTACACG
<i>TaGT</i>	XM_020302607	CTAGCAGTCAAGTCAAGCACC	TTATGCCGTGCCGAACAGAG
NtMIR1119 overexpression cassette	MI0006181	AAACCATGGAGCGUCAGCGUGUGAGG	AAAGGTAACCACGTCACGCGACCCCGC
<i>NtMnSOD1</i> expression	X14482	TTGGGCTATCGACACTAACTTT	TCAGCCAGCGACTACATGCA
<i>NtMnSOD2</i> expression	AB093097	GACGGACCTTAGCAACAGGG	ACCAATGGGTCCTGATTAGCAG
<i>NtSOD1</i> expression	KJ874395	GTGGACATGTCGTGTCAAGG	TTCTCACCAACTCCTGCACTT
<i>NtSOD2</i> expression	EU123521	ATGTCACGGGACCACATTAC	AACCCTTCCACCAGCATTTC
<i>NtFeSOD</i> expression	KF724056	CATCACAGAGCTTATGTCGACA	CTAGAAGTACTGCTTCCCA
<i>NtCAT</i> expression	EF532799	CAAGGATCTCTACGACTCGATT	CTTGAGGGCAAATAATCCACCT
<i>NtCAT1</i> expression	NTU07627	GTCTCAGGCTGACAAGTCTT	ACGGAAGACAGAGTAGCAGC
<i>NtCAT3</i> expression	HF564633	GTCTTGGGCCAAACTATCTGCA	TCAGCTTCACATTGTGGGCC
<i>NtCAT1;1</i> expression	NTU93244	TCCTGCTAATGCTCCAAAGTGT	AATGCATATGTATTAGGAATGCTC
<i>NtCAT1;2</i> expression	HF564632	GGTATCGACTTGGACCAAATA	GGTCTCACATTAAGCCTAGAAG
<i>NtCAT1;3</i> expression	HF564631	TTGCAGCCGGTGGGAAGATT	GGTCTCACATTAAGCCTAGAAG
<i>NtPOD1;1</i> expression	L02124	GGAATTTGTCCTCAAGGTGGAA	CTTATTGGAATTGCCATTTTCAGC
<i>NtPOD1;2</i> expression	AB044154	CTGACATGGTCTGTGCCTAC	TCAGTTGATAGCAGAGCAAACCT
<i>NtPOD1;3</i> expression	AB044153	AAGATCTTGTGCTCTTACTGG	AATTGGATTTTCCAGCTTGCG
<i>NtPOD1;4</i> expression	D11396	TGCTGGTAGTCAAAGTCAGTTTT	CCCATGTTGAACACGTTCTTACC
<i>NtPOD1;5</i> expression	AB178953	AACAGCAACAACGTTAACCCAGC	TTAATTTTGACCACATTCAGGA
<i>NtPOD1;6</i> expression	AB027753	TCAACTCCACTGGTGGCCCT	AATTCGATTTTGCAGCTTGCGC
<i>NtPOD1;7</i> expression	AB027752	GCCCAAGAAGTTCAGGCTCA	ATACAAATACAGTCCTTACTCG

<i>NtPOD2;1</i> expression	AB178954	AGGGGAAAAGACCTCACCAC	AGTTTCCCATCTTGATCATAGCA
<i>NtPOD2;2</i> expression	KF701483	AGACAGTGAGTATGCAGCTAA	AAAAAAGCTGCCCTTGGCACC
<i>NtPOD4</i> expression	AY032675	AGACTCAAAGATAGCAAACCTCA	CTTCCTGATGTCACCCTTGA
<i>NtPOD9</i> expression	AY032674	CACCACCTTCATTCAACGCTA	ACATCTCAGACAAAACACTTGTC

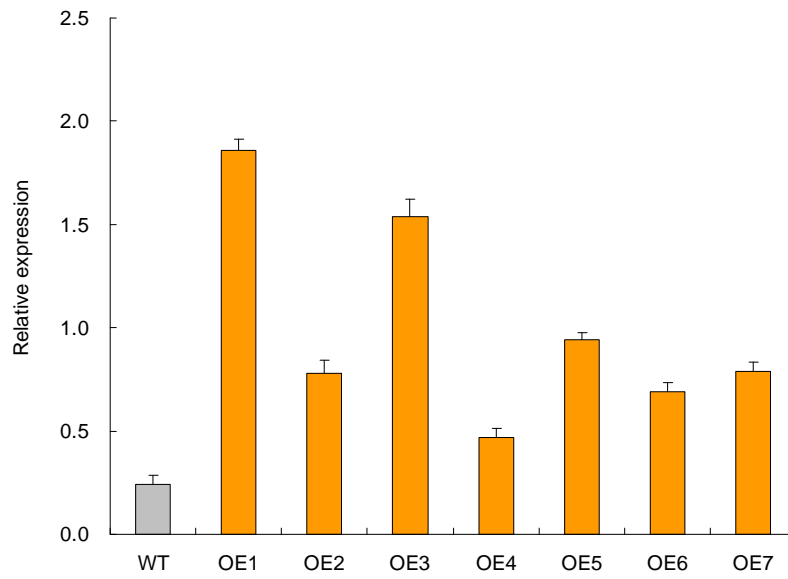
A



B

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AGCGT CAGCG TGTG AGGC GCGC CAGC GCGC GCGG AGGAG GAGC TGGC TGGGT
      10      20      30      40      50
1  A G C G T C A G C G T G T G A G G C C G C C A G C G C G C G C G G A G G A G G A G C T G G C T G G G T
1  A G C G T C A G C G T G T G A G G C C G C C A G C G C G C G C G G A G G A G G A G C T G G C T G G G T
      C A G C G T C G A C G G C G G C G G C C G T G T G C T C C T A C G T G G C A C G G C G G A T G C T
      60      70      80      90      100
51 C A G C G T C G A C G G C G G C G G C C G T G T G C T C C T A C G T G G C A C G G C G G A T G C T
51 C A G C G T C G A C G G C G G C G G C C G T G T G C T C C T A C G T G G C A C G G C G G A T G C T
      G A G T C A G C C A G T G G A G T G G T G C T A C T I T T T A C T A G G T G T G G A G T G G G A G T
      110      120      130      140      150
101 S A G T C A G C C A G T G G A G T G G T G C T A C T I T T T A C T A G G T G T G G A G T G G G A G T
101 S A G T C A G C C A G T G G A G T G G T G C T A C T I T T T A C T A G G T G T G G A G T G G G A G T
      G C T A T G C G C G G G G T C G C G T G A C G T G G
      160      170
151 G C T A T G C G C G G G G T C G C G T G A C G T G G
151 G C T A T G C G C G G G G T C G C G T G A C G T G G
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Appendix B Characterization of TaMIR1119. A, a secondary stem-loop structure initiated by the TaMIR1119 precursor. B, alignment result between TaMIR1119 and its ortholog NtMIR1119 in tobacco.



Appendix C Expression levels of the target gene in tobacco lines with TaMIR1119 overexpression OE1 to OE7, seven lines with TaMIR1119 overexpression; WT, wild type. Data are normalized by internal standard and shown by average plus standard error.