

-1172

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TAACATAATTTGAATTAAACAGAAAACGAAAGCTATAGTTATAAGAATAATTAAGTAAATAA TAGTACATTTATCTTAAA
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-783
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ACATATAAATATATTGTTGCTCCTATTTTATATAAATTTT CCGTCTATTTTACATGTTTCATCGTATATTAAAAATATA
-576
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-316
GAAATGAGTTAGAGAAATCATGTTTTATAGTTCAAATTTTTTACTAAGTCTGTTGGAATTACTAATA ACTTTTGCTTGT
GGA AATTATAGTAAAATTTTCATTTTAAAGAACGTA TATTAAGTTAATTATAA ACAAGTAAAAATGA ACCGACAAAATA
TCATTTATTTTTTGGACACCCTTTTAATTTCTT CACCAATAT TTATTAATTTCTCTTTTAAAGTGATTATTCCTTTT
+1
ATTTTTCTTCTCACACCCCTATTTATACCCCTAGCCCTCTTTGTTGAGAAAGTC ATATAAAGTTAGGTCTCTTCTTCT
TSS
TCTTCTTCAACAATATTAGGGTTTTGTCATATTGATAGCGTAGACTAC

	L1-Box		Heat shock elements
	GT-Box elements		W-Box
	CAAT-Box elements		GA regulated MYB
	CRT/DRE elements		Plant I-Box
	Evening elements		NAC2 elements

Supplementary figure 5. Promoter sequence of *MIR167a* gene showing putative transcription factor binding sites. Boundary of different deletion fragments of the promoter are marked with blue bigger font and the length is shown below the letter. TSS, transcription start site.