

Article

# Deciphering microRNAs and Their Associated Hairpin Precursors in a Non-Model Plant, *Abelmoschus esculentus*

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Supplementary Fig. S1. Sliding pattern of Small RNA Reads within a precursor.

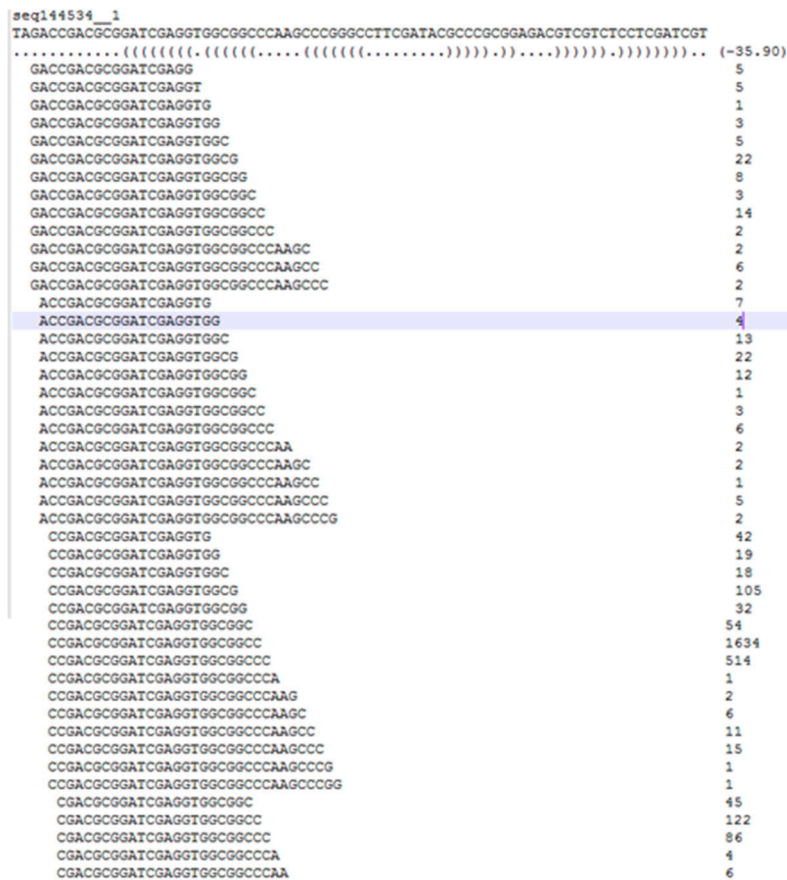


Fig. S1. Sliding pattern of Small RNA Reads within a precursor. Short reads from small RNA sequencing were mapped to the long reads of precursor sequencing by using perl scripts from miRGrep along with in-house built shell scripts. A sliding pattern of small reads was observed for many precursors as illustrated in figure S1.

**Supplementary Fig. S2. Reverse Transcription PCR and restriction digestion of miRNAs**

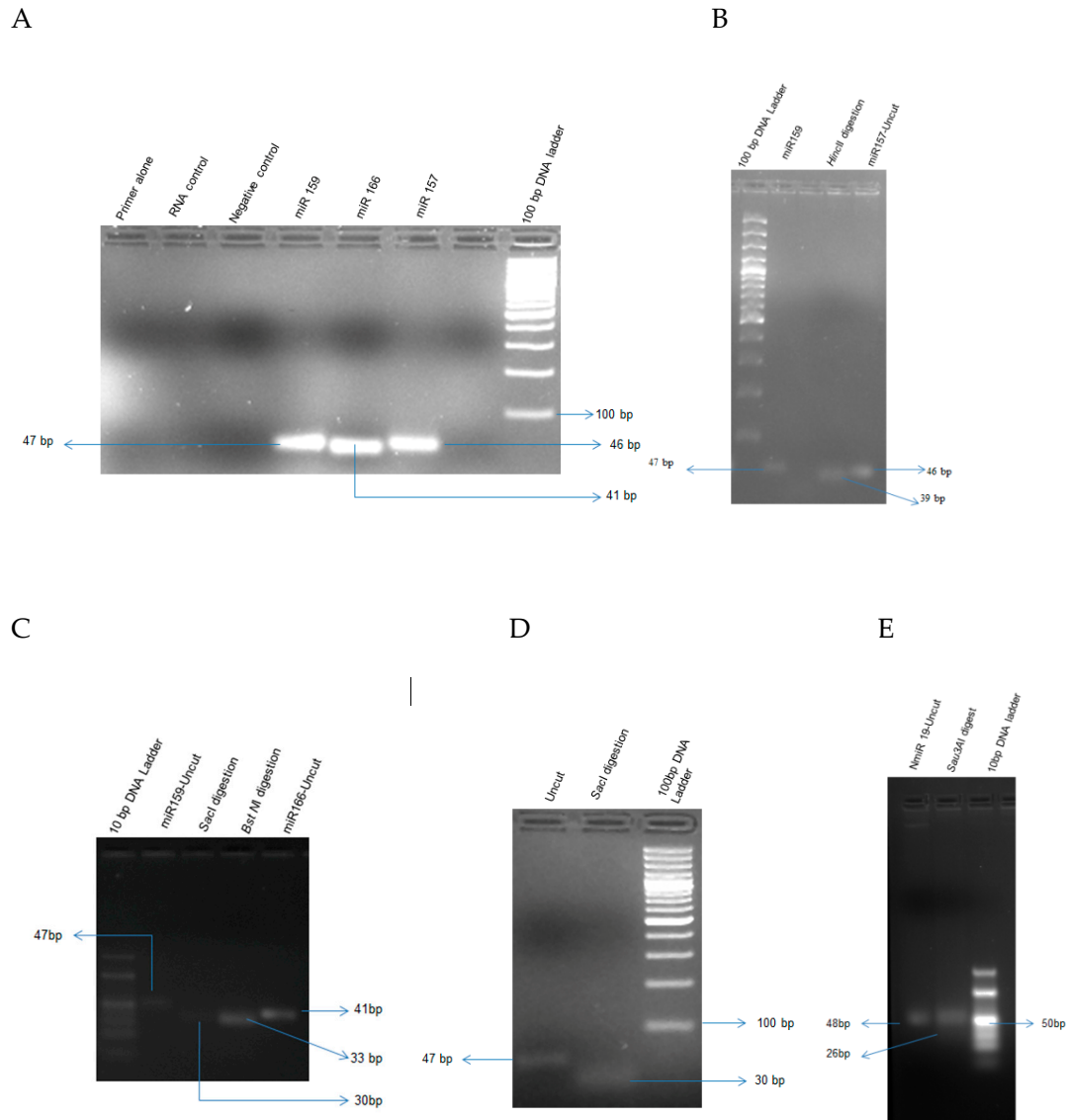


Fig. S2. Reverse transcription PCR and restriction digestion of miRNAs. PCR amplification was carried out for the randomly selected miRNAs from the reverse transcribed RNA (A). Amplified products were eluted and digested with specific restriction enzymes: *HincII* digestion for miRNA157 (B); *BstNI* digestion for miRNA166 (C); *SacI* digestion for miRNA159 (D); and *Sau3AI*

digestion for NmiRNA19 (E), to ensure genuine amplification prior to qRT-PCR. All gels are 2.5%.

**Table S1.** List of Novel miRNAs and their precursors

miRNA ID	miRNA Sequence	Precursor Sequence	MFEI
NmiRNA1	TCTTTC CTACTC CTCCCA T	TCTTTCCTACTCCTCCCATTCACGGGACGGAGGAGGCTAGGT	-0.750038
NmiRNA2	GCAAG GCGCA AGGAA GCTG	TTCCACAGCTTCTTGAACCTGCGAGTCAACGGGTTAGCAAACCCGCAAGGC GCAAGGAAGCTGATTGGCGGGATCCCTCGCGGGTGCACCGCCGA	-0.750814
NmiRNA3	CGCGG ATCGA GGTGG CGGC	ACGCGGATCGAGGTGGCGGCCCAAGCCCGGGCCTTCGATACGCCCCGCGGAG ACGTCGTCTCCTCGATCTGCGTTGATGAAGCAATTTTATTGTGGACTAGAGTTT CTGATCTGG	-0.756122
NmiRNA4	TTTGGA TGAA GGGAG CTCTA	TTTGGATTGAAGGGAGCTCTATGCCATGGTGGTCTGCAGAGATTATCGAACC TGGTATATCAGGGTTCCATATTGATCCA	-0.767551
NmiRNA5	TATTTTC CCCAG ATGATC GA	TCCATTGTCGTCCAGTGGTTAGGATATCTGGGGAAGTGATGATTATTTCCCCA GATGATCGAAGCTGAACAAACTGTGCGATTTT	-0.905393
NmiRNA6	TGTAGT ATAGT GGTAA GTATTC	GTCGTTGTAGTATAGTGGAAGTATTCCC CGGCAAGGGATGAATATTTATCT TATCCTACACT	-0.715473
NmiRNA7	TTGTAG TATAGT GGTGA GTAT	GTCGTTGTAGTATAGTGGTGAGTATTCTGGCAAGAGATGAATACTTATTCTA TCTTACACT	-0.770827
NmiRNA8	TGTGTT TGGAG AGGTG GAT	TTGGCATTCTGTCCACCTCCAGGTA CTCAAGATATTGTGTTTGGAGAGGTGGA TAGATAGGACT	-0.813838
NmiRNA9	GTGGCT GTAGTT TAGTGG	GGTGGCTGTAGTTTAGTGGTGAGAATTCCACGTTGTGGAATTCTCGGGTGCC AAGGAACTCCAGTCACGTCCGC	-0.827552
NmiRNA10	ATGTA GCCAA GTGGAT CAAGG C	TGTTTTCTCTGCCTTGATCTATATGCGCATGGCGGATGTAGCCAAGTGGATCA AGGCAGTGG	-0.867742
NmiRNA11	TCGATA AACCT CTGCAT CCAG	TCGATAAACCTCTGCATCCAGGAGCAATGAGGATAATCTGCTCTTGTGATGAT AGGGTTATC	-0.881449
NmiRNA12	CTTTTC TTTTTC TTATCC	CTTTTCTTTTTCTTATCCTACATTTGAGGATGATGAAAAAGATCAGACGGAT	-0.941243
NmiRNA13	GACCA AAGAG TCGGA GGGGG	ACCAGCATTCTTAAGACCAAAGAGTCCGAGGGGGGAAAGCTCTCCGTTCCCTG GTTCTCCTGTAGCTGGA	-0.700042

NmiRNA14	TTCGTC CCTCTA ATATGT	TCTTTTATGGTTCATATTCTGGATTGGGTTTCGTCCTCTAATATGTAATATAAG ATGTAATAATGGGATGAACTAAGTTATAGGCATGAAAGTGAAGAACTCAAC GG	-0.702688
NmiRNA15	ACTGG AGGTCT GTAGTT CG	TAAACTGGAGGTCTGTAGTTCGATCCTGCATGGGGGCACCTGGAATTCTCAGG TGCCAAGGAACTCCAGTCACGTCCGC	-0.704497
NmiRNA16	TTCGAT ACGCC CGCGG AGA	ACGTTCCCGTGGGCGGATCGAGGTGGCGGCCCAAGCCCGGGCCTTCGATACG CCCGCGGAGACGTCGTCTCCTCGATCGT	-0.744643
NmiRNA17	AGACG AAGCG GAAGC GAG	CCCTTTGTCGCTTCGATTTCGTCGTTCGCTTAGGGGAAGCACCCGCGTGAG CGGGAGTAGACGAAGCGGAAGCGAGAATGT	-0.746012
NmiRNA18	CCAGA TATAG AGATG GCGAC	AAACCAGATATAGAGATGGCGACTAAGGTTGCTGTTCCATTATTATATAATT TCAAGATCACAAATGGATCTATGA	-0.748108

Novel miRNA sequences along with their corresponding precursor sequences are shown. NmiRNA denotes- Novel miRNA.

**Table S2.** List of Precursors of miRNAs from PMRD which are not present in miRBase

miRNA ID	miRNA Sequence	Precursor Sequence
MIR35-npr	GGAATATGAGTG TGTGACTT	GGTAGTTCGACCCGCAAATTTATTTGTTTCTGTATTTCCGGAATA TGAGTGTGTGACTTGTTAGAATTGATC
MIRf10239-npr	CGCTTGGTGCAG GTCGGG	TCGCTTGGTGCAGGTCCGGAAATTACGATAGGTGTCAAGTGG AGTGCA
MIRf10238-npr	TGCCTGGCTCCCT GTATGCCA	TGCCTGGCTCCCTGTATGCCACAATGTAGGCAAGGGAAGTCCG CAAAATGG
MIRf11025-akr	TGCATTTGCACCT GCACCT	TGCATTTGCACCTGCACCTTCTCATTACGATAGGTGTCAAGTGG AAGTGCA
MIRf10271-akr	TGGATTGAAGGG AGCTCT	TTTGGATTGAAGGGAGCTCTATGCCATGGTGGTCTGCAGAGAT TATCGAACCTGGTATATCAGGGTTCCATATTGATCCA
MIRf10082-akr	TGCATTTGCACCT GCACCT	TGCATTTGCACCTGCACCTTCTCATTACGATAGGTGTCAAGTGG AAGTGCA
MIRf11010-npr	TTCCACAGCTTTC TTGAAC	TTCCACAGCTTTCCTGAACCTTATTACGATAGGTGTCAAGTGGAA GTGCA