Supplemental Figure S2

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5' UAUUUUUGGAGAGUUCGAGCA 3'
nta-miRX27
           QPT2 2,652 bp 3' GUAGAAGUCUCUCACGUUCAA 5' 2,632 bp
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Figure S2. Validating nta-miRX27-mediated cleavage of the predicted target site (the first one at exon #2 in Fig. S1) using degradome data from roots of topping-treated tobacco plants. Base pairing between nta-miRX27 and its target OPT2 is shown, in which a vertical line means a Watson-Crick pair, two dots represent a G-U pair and 0 represents mismatch. The numbers below the alignment indicate the numbers of degradomic reads at the positions indicated. The expected cleavage site is indicated by a red arrow.