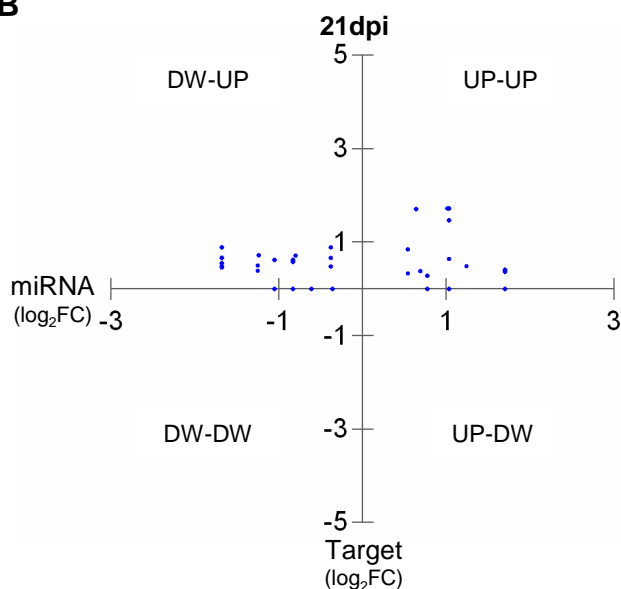


**A**

<i>Number of miRNA-target pairs (PARE)</i>			
<i>miRNA</i>	<i>Target</i>	<i>14 dpi</i>	<i>21 dpi</i>
DW	UP	1	17 (39%)
UP	DW	0	0
DW	DW	0	0
UP	UP	0	12 (28%)
DW	nc	1	9 (21%)
UP	nc	1	5 (12%)
<b>TOTAL</b>		<b>3</b>	<b>43 (100%)</b>

**B**

**Additional file 10: Fig. S10. Expression levels of the DE miRNAs and their predicted target genes in tomato according to the degradome analysis.** A) Classification based on their expression levels at 14 and 21 dpi of the miRNA-target pairs according to degradome's data [82]. UP: upregulation, DW: downregulation, nc (no changes): target gene not differentially expressed. B) Expression level ( $\log_2FC$  for the ratio TYLCV/mock) at 21 dpi of the DE miRNAs (x axis) and their target genes (y axis). Only targets with a degradome category  $\leq 3$  and a p-value  $\leq 0.05$  are shown.