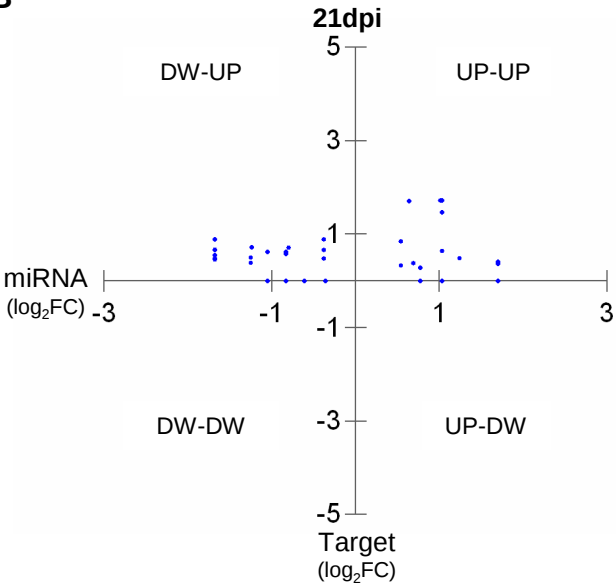


A

Number of miRNA-target pairs (PARE)			
miRNA	Target	14 dpi	21 dpi
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%)
TOTAL		3	43 (100%)

B



Additional file 9: Fig. S9. Expression levels of the DEmiRNAs 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 (log₂FC for the ratio TYLCV/mock) at 21 dpi of the DEmiRNAs (x axis) and their target genes (y axis). Only targets with a degradome category ≤ 3 and a p-value ≤ 0.05 are shown.