

**Table S2.** Relative quantification by RT-qPCR of viruses other than GFLV in grapevines or nematodes.

The vines, virus source plants, are VA6, VA7 and VA8

$\Delta$ Ct : Ct Gene of Interest - Ct GAPDH grapevine

$\Delta$ Ct : Ct Gene of Interest - Ct Actin Nematode

rel. Quant: relative quantification compared to GFLV RNA1 accumulation within the same sample

av.: average of positive samples, with  $\pm$  corresponding to standard error (SE)

N/A: Not applicable

\*: not applicable since sample was negative for GFLV

		GFLV	ArMV	GVB	GRVFV
Grapevine	Vines, virus source plants	3/3	1/3	2/3	1/3
	av. $\Delta$ Ct	-0.0 $\pm$ 0.6	3.8	2.4 $\pm$ 0.5	2.2
	av. rel. Quant.	1	0.135	0.133 $\pm$ 0.009	0.399
Nematode	Pool	5/5	2/5	0/5	5/5
	av. $\Delta$ Ct	4.8 $\pm$ 0.6	13.7 $\pm$ 3.4	N/A	10.4 $\pm$ 1.3
	av. rel. Quant.	1	0.005 $\pm$ 0.005	N/A	0.190 $\pm$ 0.167
	Single	9/11	2/11	0/11	1/11
	av. $\Delta$ Ct	6.7 $\pm$ 0.6	11.7 $\pm$ 1.0	N/A	0.9
	av. rel. Quant.	1	0.113 $\pm$ 0.111	N/A	N/A*