

Supplementary Table 1: Total number of reads, after trimming, associated with each dataset analyzed in this study, as well as the genome length of each virus, the average coverage of the assembled contigs and the GenBank accession associated with each genome. GVN – grapevine virus N, GVO – grapevine virus O, nt – nucleotides.

Novel vitivirus	Cultivar of <i>Vitis vinifera</i>	Total number of reads after trim	Genome length (nt)	Average coverage (x)	GenBank accession number
GVN	Azal tinto	6,785,861	7486	7.8	MZ682351
	Bourboulenc	4,690,821	7486	12.3	MZ682352
	Cinsault noir	4,614,103	7486	40.5	MZ682353
	Crouchen blanc	4,582,363	7486	6.7	MZ682354
	Palamino	6,044,233	7486	7.1	MZ682355
GVO	Roter Zierfandler	3,505,570	7560	6	MZ682356
	Royal Molenberg	3,130,186	7560	9.1	MZ682357