

Supplementary table 1: Number of reads (after trimming) for datasets that were found to be associated with macadamia ringspot virus . Length refers to complete genomic segment in nucleotides (nt),together with the average coverage (x). L - large segment, M – medium segment, S – small segment.

Sample	Total number of reads (after trim)	Length (nt) and average coverage (x)			GenBank accession RNA1/RNA2/RNA3
		L	M	S	
19-3002	2,003,304	8715 (24)	4779 (55)	2935 (616)	OP604037/8/9
19-3005	2,017,064	8714 (31)	4779 (38)	2934 (608)	OP604040/1/2
19-3006	1,882,062	8714 (58)	4779 (75)	2935 (1189)	OP604043/4/5
19-3007	1,168,372	-	4779 (13)	2936 (171)	OP604046/7
19-3008	3,149,606	8715 (19)	4779 (29)	2935 (308)	OP604048/9/50