

Table S1. Description of the primers specifically designed to fulfil gaps into the genome sequences generated after NGS sequencing.

Virus	Primer	Sequence	Size (nt)	Position
BBOV_9716RCA	9716_GSP1_smart	5' GGATCGGCATTTTGTGAAGAGGTAGAGG 3'	384	1 nt – 384 nt (N gene)
	9716_NGSP1_smart	5' CATTCCTTGAACCGATCCTCCGACC 3'		
OUAV_9718RCA	9718_GSP1_smart	5' CAGCCCGCAACAGAACACATACATCC 3'	461	1 nt – 461 nt (N gene)
	9718_NGSP1_smart	5' GGATTTATGTCCTCACCTTGCCTTCC 3'		
	9718-F1	5' GATCACATTTTCATTGGCAATGC 3'	347	4287 nt – 4633 nt (G gene)
	9718-R1	5' GAGTCTTGTTGGTTTCTCTGAAGATG 3'		
	9718-F2	5' GGATTGATTACCACAAGTTAG 3'		
	9718-R2	5' CAAGTTCAGATATTCTGTATTCC 3'		
MTYV_09027EGY	MTYV_GSP1_smart	5' CAATCCGATGCAATAAACAAACATCCAC 3'	459	1 nt – 459 nt (N gene)
	MTYV_NGSP1_smart	5' CCTCTATTTACAAAAGATTCCAAGGGGC 3'		