

Table S2. Primers used for genome sequencing of rose ilarvirus 2.

Primer name	Sequence (5' → 3')	Position
RNA1		
RIV2-R1-27F	CGACCCGGATTAATCAGTTTTCT	27-49
RIV2-R1-3255R	ACATCCCAGAGGACCCCTTACT	3234-3255
RIV2-R1-2688F	CACATTGAAACACCCCTCTATCA	2688-2710
RIV2-R1-3052F	GCGGGATCTTCACACAACCTTAG	3052-3073
RIV2-R1-577R	AACGCAACTCTCTCTCTGGTTGT	555-577
RIV2-R1-343R	CGTGTGAAGAAGTAGCTGAGTTG	321-343
RNA2		
RIV2-R2-234F	CGTTAGTGGAGAAGCTATCATGTG	234-257
RIV2-R2-2911R	CAAAGGTGGGCATATGGACTTAG	2889-2911
RIV2-R2-2412F	CGTTTCATGGACGTCACCTAGTA	2412-2434
RIV2-R2-2626F	GTCGTGGAGTATTTCCCTCCTATGG	2626-2649
RIV2-R2-616R	CAAAACGGCTAGACCAGACAGC	595-616
RIV2-R2-393R	GTATGCCTCTTCAGGCACATCTC	371-393
RNA3		
RIV2-R3-1F	CGCTGAAGCTGTTATCCTACCTGA	339-362
RIV2-R3-1R	TGGAGAAAGGGACGCAAGATAGAG	1395-1418
RIV2-R3-2F	GGAAGTGTCAACCGCTTCGAGGA	1257-1279
RIV2-R3-2R	AGCGCAAAAGCAATGACAGTACGG	2104-2127
RIV2-R3-1837F	GCTCCTAATAGAGTCCGTGTGAAG	1837-1860
RIV2-R3-2036F	CTCCTCCAGACTTTCTTGTCTGTG	2036-2059
RIV2-R3-590R	GAGCGGGTATTTTCATCACACA	569-590
RIV2-R3-441R	GGTCAAGACGACAGGCTTAGTT	420-441