

Table S1. Primers developed for the Sanger sequencing of little cherry virus 1 (LChV1) genome regions.

Primer name	Primer sequence, 5'-3'	Target ^a	PCR product, bp
lchv-F1 lchv-R1	ACCTCACTGTTGATCGCCGA AGTGTCTCGTCAGAGTGCCA	ORF1a (1312-1521)	209
lchv-F2 lchv-R2	TGAGTTGAGTTATGAGCGTGGT AGCCCTGAATTGGCAATCTTGA	ORF1a (4010-4190)	180
lchv-F3 lchv-R3	GATGGGATGGTTTCGAGTCTCT GGGACATCACTATGAATTAGAGA	ORF1a (5094-5887)	793
lchv-F4 lchv-R4	GAATCTGAAATAGAACGCTTAGT CGATAATGATTCTATTTCCCTGT	ORF3 - ORF4 (10237-10608)	371
lchv-F5 lchv-R5	AGCTATACGTGTGAACGAGAGA ATCATCGCCAATGTCTAAGGCA	ORF4 - ORF5 (11735-12253)	518
lchv-F6 lchv-R6	CTGTAGAAGATAGAAGGTCAGA GAGTCAATTGCACACCTGAAC	ORF5 (12372-12774)	402
lchv-F7 lchv-R7	GTGAAGTTTGGAGAATTCTGGT CTACTTCGACTTCATCATCACT	ORF6 (14135-14656)	521
lchv-F8 lchv-R8	GTAAATGACCAATTACTCGGTGGT TGCAACACAGGTGGTCTATCACA	ORF8 - 3'-UTR (16412-16907)	495

^a Target genes and positions (in brackets) in the genome of LChV1 isolate Stepnoe are indicated.

ORF - open reading frame; UTR - untranslated region.