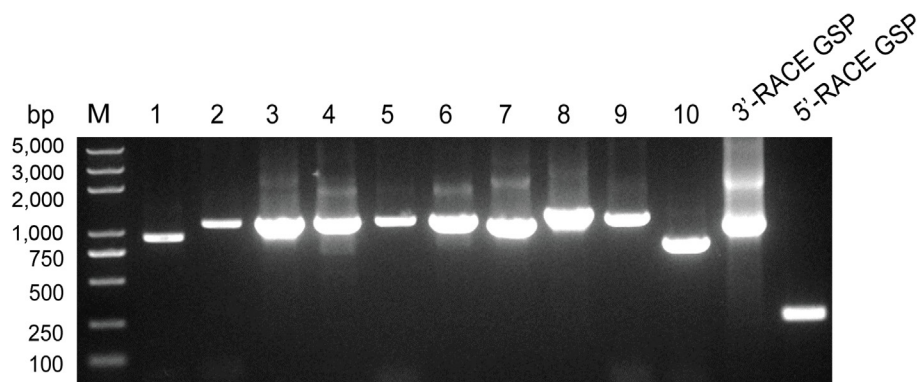


Supplementary Table 1. Primers used in this study

Primer	Sequence (5' to 3')	Position
Long primer	CTAATACGACTCACTATAGGGCAAGCAGTGGTATCAACGCAGAGT	/
Short primer	CTAATACGACTCACTATAGGGC	/
5'-RACE GSP	CCTACCGTAATGCTCTCCATAATTTGC	260-286
3'-RACE GSP	CAGGGAAGGGCCAGGAAACAGCAGCCG	9,038-9,064
PotyF	ATGGTHTGGTGYATHGARAAYGG	/
PotyR	TGCTGCKGCYTTTCATYTG	/
BCMV-1-F	GCAATTCAAGCTCTACGTACAC	73-94
BCMV-1-R	CTTGAAGCTCCTACGCCAGTAAG	997-1,019
BCMV-2-F	TTGGTACGTAAGAGGCAAGAAG	870-891
BCMV-2-R	GCCTTTGACGATCTCTTGTTTC	2,005-2,026
BCMV-3-F	CATCAGTGACTCAGGAAGAA	1,894-1,913
BCMV-3-R	TGAGCTGTCCCAACAGTACA	3,047-3,066
BCMV-4-F	TGTGAAACAGCCCTCATTAC	2,859-2,878
BCMV-4-R	TCATCTAAGCTTTGGATTTGA	4,022-4,042
BCMV-5-F	TGGAACACCTTGAGAAGATACG	3,793-3,814
BCMV-5-R	CACAGTCAACATCTAGTGTGAC	4,965-4,986
BCMV-6-F	GAGTTACAACGAAGTTGATCA	4,799-4,819
BCMV-6-R	ATGGTGTCGTTTCATGAATCTC	5,987-6,007
BCMV-7-F	CTGTTAGAATTCTCAAGCACA	5,847-5,867
BCMV-7-R	TCCTCTTTGATGGGTGGTCTG	6,962-6,982
BCMV-8-F	GCAATAATGGTGTGCTCAACA	6,811-6,831
BCMV-8-R	TGAGAACCGTCTGCGTGACAG	8,120-8,140
BCMV-9-F	CAACTTGGAGTGCCATGGAC	8,030-8,050
BCMV-9-R	CATCTTTGTTGCTCTTGTGTTG	9,239-9,260
BCMV-10-F	AGAGACAGGGATGTGAATGCAGGT	9,087-9,110
BCMV-10-R	TGTTGGTGGTCCAACCACTCTATC	9,919-9,942

RACE, rapid amplification of cDNA ends; BCMV, bean common mosaic virus.



Supplementary Fig. 1. Electrophoresis of the amplicon of reverse transcription polymerase chain reaction amplification and rapid amplification of cDNA ends (RACE).