

Supplementary Table S2. List of primers used in this study for genome amplification and detection of babacovirus 1 (BabIV-1) and babacov rhabdovirus 1 (BabRV-1).

Virus	Primer name	Sequence 5' to 3'	Target genome* position (amplicon size nt)
BabIV-1	1Race-5'end-1	CGTTGCTAAAGCATCATGATCC	1-56
	1Race-5'end-2	CGAGTGACTTGAAGTGTCACTC	1-281
	RNA1-F1	CCAATCTCTCTCGTAATGGATC	18-866 (849)
	RNA1-R1	CCTCTCAATTCGGTAAGCCTT	
	RNA1-F2	GGTATGAGCACAAGTTTCTGT	778-1697 (920)
	RNA1-R2	GGTTAGTTTCTCGTCTTTCAGC	
	RNA1-F3	GTGGTTCTGGAACTACCTGG	1554-2968 (1415)
	RNA1-R3	GGGTCCTTACCACTGTATATGT	
	1Race-3'end-1	CGAATGTTGCACTGGTCCGTA	2908-3299 (392)
	1Race-3'end-2	CGTACCTGCACACTGATCATAAG	3115-3299 (185)
	2Race-5'end-1	CGAACAAGACCTTAGTGGCCG	1-59
	2Race-5'end-2	GGTCTAAGGAAAACCTACTAACCT	1-186
	RNA2-F1	CCATCACGATTTGAGAGAATTCC	64-939 (876)
	RNA2-R1	CGTTTCGACTAACTCCTGGAA	
	RNA2-F2	CCAACAACATCCGACATACTCA	808-1612 (805)
	RNA2-R2	GATGTAACCTCTGTTGAGACTTG	
	RNA2-F3	GAGAGAGCATTACCAGCTACTA	1378-2605 (1228)
	RNA2-R3	CTCTATATCCACACGGTGGAG	
	2Race-3'end-1	CGAGTGATTTCGTCGTATGGAGTGG	2342-2675 (334)
	2Race-3'end-2	CGAAAAGCAGAATGTGTTGAAC	2400-2675 (276)
	3Race-5'end-1	GATCTGGACCAGGAGAAGTAGT	1-159
	3Race-5'end-2	CGACCTACATCTCTATAGATCT	1-358
	RNA3-F1	GGAAGAGCTTCAGCAAGTCAC	152-1118 (967)
	RNA3-R1	CTTGAGCTCCAGTCGGATGAC	
	RNA3-F2	GGTTAGAAGCAGGATTGCGTC	722-1688 (967)
	RNA3-R2	CCTAGACCTCCAACAAGTCTTC	
	3Race-3'end-1	CGATTTGCACGAGAGTGCAGA	1526-1802 (277)
	3Race-3'end-2	CGTAGTAACGACTGAGTTACCACG	1609-1802 (194)
	Detection_F	GGTTAGAAGCAGGATTGCGTC	722-1118 (397)
	Detection_R	CTTGAGCTCCAGTCGGATGAC	

BabRV-1	Race-5'	CCCATCAGCTGAAGGAATTGC	1-251
	F1	GACTCTATGCCTACTGTGGTAC	80-1152
	R1	CGACACTGAGATCAACAAAGTAG	(1073)
	F2	GCAACTGTTCATGACAATCCCA	1056-2062
	R2	GGTGCCTATCTCTCAGTTTCC	(1007)
	F3	CTATCAGTGCCTGAGCTGAACT	1962-2991
	R3	GCAATTGCAATAGGAAGGACT	(1030)
	F4	CCTCAAAGATGTCTAAACCTCT	2907-3850
	R4	GTAGGTAGGATTAAGGCTGTAG	(944)
	F5	GAGCATCGAGGAACACCCTCA	3793-4670
	R5	GATCGGATCCTTCTGATTTTCTC	(878)
	F6	CCAGACTGAACATCATGGTCT	4563-5511
	R6	GACCAAACATCTGTGTCCATGG	(949)
	F7	GACATGCTTGACCTATGTCTCA	5305-6233
	R7	CTGAGTGATCACAACCTCATCTG	(929)
	F8	GCAGATTGGGTTTCATCAGATGT	6153-7059
	R8	CCATATCTCCAACAGTCTGTCT	(907)
	F9	CCACGATGGAAGACTGGAAG	6862-7930
	R9	CCATCTATATCTATGATCGGATG	(1069)
	F10	GAGATGATATATCAGAGACAGGC	7795-8790
	R10	GGAGGTCGTGAGTGATGTCAAA	(996)
	F11	GAAGTGGAATCAGCAAATGAGG	8687-9719
	R11	GAATGACATTCCCATCAGTTGTC	(1033)
	F12	GAGAGCTACATATGATCAATCCC	9655-10570
	R12	CGATGATCATAGCTACCTCTGA	(916)
	F13	CACAAACATTGATCCCTCACAAAC	10505-11437
	R13	CCAAGACAATCTAAACAAGAGAG	(933)
	F14	GCAGATTGTCCTTCATGTTATAG	11375-12359
	R14	CTCAATCTTCTCCAGTATTCTCT	(985)
	F15	CCAGATACAAGTTCTTGAATAGG	12292-12765
	R15	GGTGGTGATAGGGATTAAGTCC	(474)
	Race 3'	CCTAGACTGGTTCGCTTCTGAT	12712-12802 (91)
	Detection_F	GGAGTCGGCATTGGTATAGCA	1347-2269
	Detection_R	CCGTTGTCCTTTAACACCCTAG	(923)

* For babaco rhabdovirus-1 the primer positions correspond to the complementary (positive-sense) genome.