

Table S2. Primers used for dsRNA synthesis template amplification, qPCR assays of expression or segment abundance, and protein expression.

Target	Accession numbers	Primer name	Sequence 5'-3'
dsRNA synthesis primers			
<i>lef-9</i>	JO947537.1- JO947542.1	Mdlef9RN AiF	TAATACGACTCACTATAGGGGTCGTCCGTATAACATTGGT
		Mdlef9RN AiR	TAATACGACTCACTATAGGGTGGATTTATCAAATCGAAGC
<i>lef-4</i>	JO932223.1- JO932224.1	Mdlef4RN AiF	TAATACGACTCACTATAGGGATGTTTCATACCCCTCACCAG
		Mdlef4RN AiR	TAATACGACTCACTATAGGGACGGCAGGGTATGTTTAGTA
<i>vp39</i>	JR139428.1- JR139430.1	vp39_RNA iF	TAATACGACTCACTATAGGGCGGCATCATGCAGTTGTATC
		vp39_RNA iR	TAATACGACTCACTATAGGGATTTGCTGCTTGAGGGTTG
<i>vlf-1</i>	JO963955.1	vlf1_RNAi F	TAATACGACTCACTATAGGGTTCGGTGTTCGTCCGTATT
		vlf1_RNAi R	TAATACGACTCACTATAGGGACGGAATTTACGCCACCATA
<i>vlf-1 #2</i>	JO963955.1	vlf- IRNAi2F	TAATACGACTCACTATAGGGAACATGACGGACGCATTAAG
		vlf- IRNAi2R	TAATACGACTCACTATAGGGGAAATCATTGGATACTGATTTTGAA
<i>int-1</i>	JO956053.1	Int_RNAiF	TAATACGACTCACTATAGGGTCTTCCAACACACGATTCCA
		Int_RNAi R	TAATACGACTCACTATAGGGTTTTGGTTTCATCAGCAATAAAGC
<i>p74</i>	JO979915.1	Mdp74RN AiF	TAATACGACTCACTATAGGGCCCCACATAATCTCACATCT
		Mdp74RN AiR	TAATACGACTCACTATAGGGTCTTTTGATCACCCTTCTT
<i>pif-1</i>	JO979916.1	pif- IRNAiF	TAATACGACTCACTATAGGGATTTATCTGCGACGGTGAGG
		pif- IRNAiR	TAATACGACTCACTATAGGGTAATTGCCTTCTTGGGTGG
<i>eGFP</i>		T7pGEM- Forward	TAATACGACTCACTATAGGGCACGACGTTGTAAAACGAC
		T7pGEM- Reverse	TAATACGACTCACTATAGGGGATAACAATTCACACAGG
qPCR primers			
<i>lef-9</i>	JO947537.1- JO947542.1	lef9qPCR F*	CATCTTGATCAGCGTGCAAT
		lef9qPCR R*	ACGTCAGTATTCCCCAGCAC
<i>lef-4</i>	JO932223.1- JO932224.1	lef4qPCR F*	ACCCTTACCAGGACAACCTG
		lef4qPCR R*	AAATAGTACGCGCCACCTTG
<i>vp39</i>	JR139428.1- JR139430.1	vp39qPCR F	CGCTACACGCTATCGATTTG
		vp39qPCR R	CACTGACTGTGCACAAAATTCA
<i>vlf-1</i>	JO963955.1	vlf1_qPCR F	TTTAAAGCCCAAAGCCACAG
		vlf1_qPCR R	AATGGCTTAAAGTCGGGATG
<i>int-1</i>	JO956053.1	IntqPCR F	GCGCAGCAGCTAAGTCATT
		IntqPCR R	AAAATATCGACGCACGGGTA
<i>p74</i>	JO979915.1	p74F*	TCCGTAATTGATTTGGGGAGA
		p74R*	TGCAGCACCAACAACAAT
<i>pif-1</i>	JO979916.1	pif- lqPCR F	CTGCAAGTCGTGTCCCTACA
		pif- lqPCR R	ATAACCAATCGCCGTGATA

		IqPCR	
Segment B	AY848690.1	SegBLL*	CGTGGATTGACAACGCGTTT
		SegBRR*	TTCTTAGCAGATGATGTCATCGC
		SegBRL*	AGCTTATGTCGACAAGCGCT
		SegBLR2*	TGATTAATTTGTGATACTTCCATGTT
<i>lef-9</i>	JO947537.1- JO947542.1	LEF-9F2	GACGACGACAAGATGTCAATTTTCAACCACTTTATTG
		LEF-9R	GAGGAGAAGCCCGTTTCAATATTCGGACGAATCTAAATCC

*Primer sequences have been previously published in:
 Burke GR, Strand MR (2012) Deep sequencing identifies viral and wasp genes with potential roles in replication of *Microplitis demolitor* Bracovirus. J Virol 86: 3293-3306.