

Supplementary Figure Captions

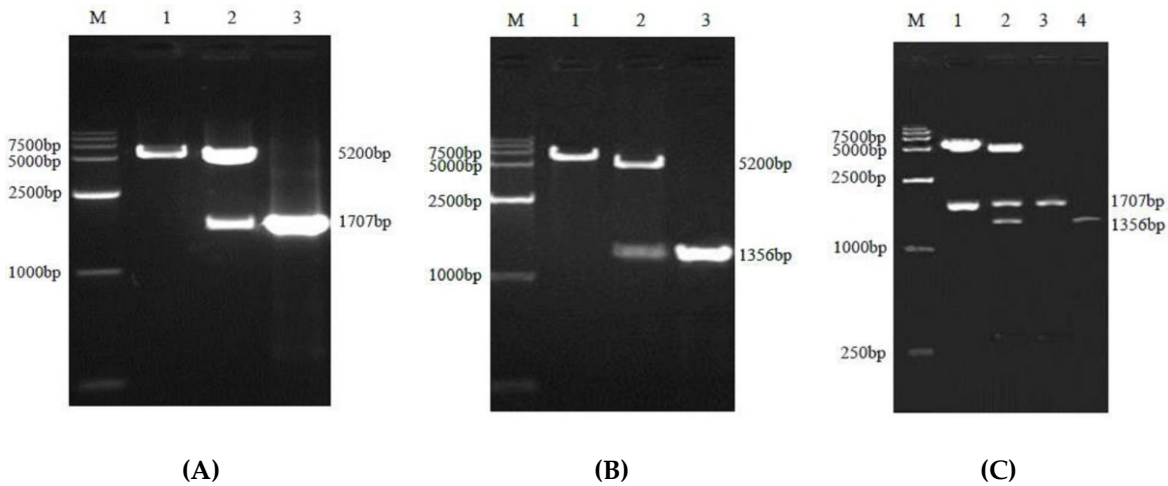


Figure S1. Identification of recombinant transposon vectors pFast-HBM-S1, pFast-HBM-N and pFast-HBM-S1-N by PCR and restriction endonuclease digestion. **(A)** Identification of recombinant transposon vectors pFast-HBM-S1. Lane M. DL15000 DNA Marker; Lane 1. pFast-HBM-S1 was digested with *Pst* I, Lane 2. pFast-HBM-S1 digested with *Bam*H I and *Pst* I. Lane 3. PCR identification of pFast-HBM-S1 with HBM-S1 specific primers. **(B)** Identification of recombinant transposon vector pFast-HBM-N. Lane M. DL15000 DNA Marker; Lane 1. pFast-HBM-N was digested with *Kpn* I. Lane 2. pFast-HBM-N was digested with *Xho* I and *Kpn* I. Lane 3. PCR identification of pFast-HBM-N with HBM-N specific primers. **(C)** Identification of recombinant transposon vector pFast-HBM-S1-N. Lane M. DL15000 DNA Marker; Lane 1. pFast-HBM-S1-N was digested with *Bam*H I and *Pst* I, Lane 2. pFast-HBM-S1-N was digested with *Bam*H I, *Pst* I, *Xho* I and *Kpn* I. Lane 3. PCR identification of pFast-HBM-S1-N with HBM-S1 specific primers. Lane 4. PCR identification of pFast-HBM-S1-N with HBM-N specific primers.

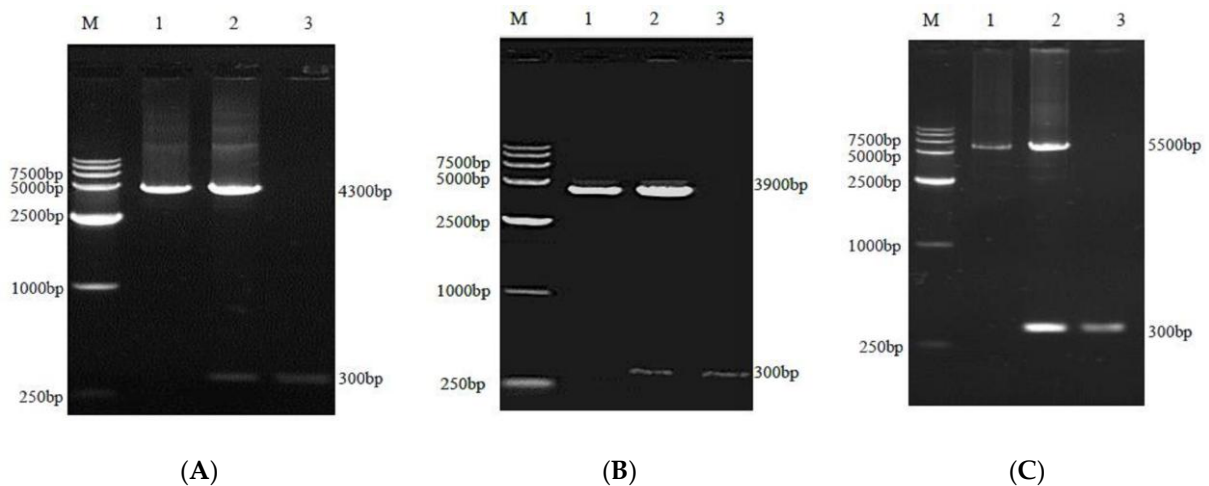


Figure S2. Identification of recombinant bacmids rHBM-S1, rHBM-N and rHBM-S1-N by PCR with M13 primers. **(A)** Identification of recombinant bacmids rHBM-S1. Lane M. DL15000 DNA Marker; Lane 1. PCR product of the purified bacmids rHBM-S1; Lane 2. PCR product of the unpurified bacmids rHBM-S1; Lane 3. PCR product of the wild bacmid. **(B)** Identification of recombinant bacmids rHBM-N. Lane M. DL15000 DNA Marker; Lane 1. PCR product of the purified bacmids rHBM-N; Lane 2. PCR product of the unpurified bacmids rHBM-N; Lane 3. PCR product of the wild bacmid. **(C)** Identification of recombinant bacmids rHBM-S1-N. Lane M. DL15000 DNA Marker; Lane 1. PCR product of the purified bacmids rHBM-S1-N; Lane 2. PCR product of the unpurified bacmids rHBM-S1-N; Lane 3. PCR product of the wild bacmid.

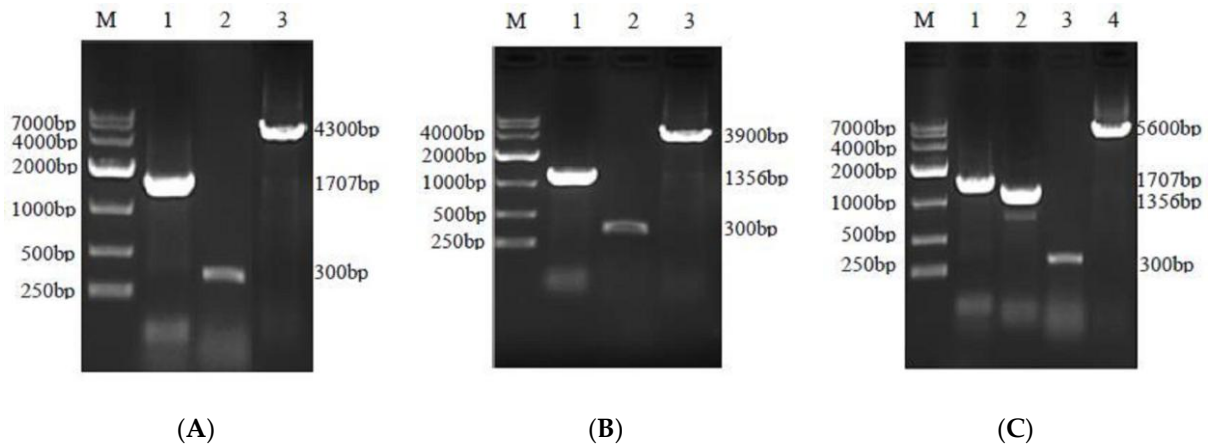


Figure S3. Identification of recombinant baculovirus by PCR with IBV specific primers and M13 primers. **(A)** Identification of recombinant baculovirus rHBM-S1. Lane M. DL10000 DNA Maker; Lane 1, PCR product of baculovirus rHBM-S1 with S1 gene specific primers; Lane 2, PCR product of wild baculovirus with M13 primers; Lane 3, PCR product of baculovirus rHBM-S1 with M13 primers. **(B)** Identification of recombinant baculovirus rHBM-N. Lane M. DL10000 DNA Maker; Lane 1, PCR product of baculovirus rHBM-N with N gene specific primers; Lane 2, PCR product of wild baculovirus with M13 primers; Lane 3, PCR product of baculovirus rHBM-N with M13 primers. **(C)** Identification of recombinant baculovirus rHBM-S1-N. Lane M. DL10000 DNA Maker; Lane 1, PCR product of baculovirus rHBM-S1-N with S1 gene specific primers; Lane 2. PCR product of baculovirus rHBM-S1-N with N gene specific primers; Lane 3, PCR product of wild baculovirus with M13 primers; Lane 4, PCR product of baculovirus rHBM-S1-N with M13 primers.

Table S1. All the primers used in this study

Primer ID	Sequence(5'→3')	Gene size
HBM-F1	CGCGGATCCATGAAATTCCTAGTCAACGTTGCCCTTGTTTTATGGTC (<i>Bam</i> H I)	78 bp
HBM-R1	TCGATCCGCATAGATGTAAGAAATGTATACGACCATAAAAAACAAGGGC	
S1-F	CGCGGATCCATGTTGGGGAAGTCACTGTTTTAGTG (<i>Bam</i> H I)	1692 bp
S1-R	CCCAAGCTTTCAGTGATGGTGATGGTGATGGCCTTGAAAGTACAAGTTTTACGTCTGTGACGACGACTTTCAT (<i>Hind</i> III)	
HBM-S1-F	CATTTCTTACATCTATGCGGATCGAAATTTGTTTCATTCTGGT	1707 bp
HBM-S1-R	CAAACTGCAGTCAGTGATGGTGATGGTGATGGCCTTGAAAGTACAAGTTTTACGTCTGTGACGACGACTTTC (<i>Pst</i> I)	
HBM-F2	CCGCTCGAGATGAAATTCCTAGTCAACGTTGCCCTTGTTTTATGGTC (<i>Xho</i> I)	78 bp
HBM-R2	TCGATCCGCATAGATGTAAGAAATGTATACGACCATAAAAAACAAGGGC	
N-F	CATTTCTTACATCTATGCGGATCGAATGGCAAGCGGTAAGCAGCT	1303 bp
HBM-N-R	CGGGGTACCTCAGTGATGGTGATGGTGATGGCCTTGAAAGTACAAGTTTTCAAGTTCATTCTCTCCAAGTGCTGAATCG (<i>Kpn</i> I)	
HBM-N-F	CCGCTCGAGATGAAATTCCTAGTCAACGTTGCCCTTGTTTTATGGTCGTATACATTTCTTACATCTATGCGGATCGAATGGCAA GCGGTAAGCAGCT (<i>Xho</i> I)	1356 bp
HBM-N-R	CGGGGTACCTCAGTGATGGTGATGGTGATGGCCTTGAAAGTACAAGTTTTCAAGTTCATTCTCTCCAAGTGCTGAATCG (<i>Kpn</i> I)	
M13-F	CCCAGTCACGACGTTGTAAAACG	≈2560 bp + size of insert
M13-R	AGCGGATAACAATTCACACAGG	
M-F	GGTAGAAAACCTTAACAATCC	740 bp
M-R	AAGACTACTTCCTCCTGTTG	

Note: The italicized is the restriction enzyme site, the underlined are the 6 × His tag and the TEV protease cleavage site added downstream of the target gene