

An integrated view of protein homologue groups and functional genomic development of baculoviruses: towards understanding baculoviral infection mechanism and improving baculoviral expression vector

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Summary. – This study aimed to gain a comprehensive view of protein homologue groups of baculoviruses and elaborate the correlation between protein conversation and functional roles in the viral biphasic infection cycle. A comprehensive identification of homologue groups among 76 baculoviral genomes was performed, supplying a global view of the distribution of 992 homologue groups among baculoviruses. Fifty-six missed proteins from previous annotated genomes were added, updating AC76 as a new number found in all baculoviruses except *Deltabaculovirus* and the LEF 10 (AC53a) as a new number present in all the *Alphabaculovirus* and *Betabaculovirus*. Accompanied by an updated encyclopedia of functional genomic development, an integrated view of the correlation between protein conversation and functions was summarized for *Autographa californica* nucleopolyhedrovirus (AcMNPV). The AC106/107 and ODV-E18 were screened out as underestimated proteins and would need investigations on their detailed roles in viral infection cycle. A baculovirus-host network was constructed based on archive exploration, indicating that the ODV-E18 would be the core of viral interaction network. Further, potential functional domains and sites of ODV-E18 were further analyzed, supplying future research targets for exploring their detailed role in the regulation of the BV/ODV formation mechanism. This would facilitate further baculoviral genome annotation, functional genomic investigation and construction of a baculoviral minigenome that could be exploited as high-capacity expression vector.

Keywords: baculovirus; biphasic infection; baculoviral expression vector; phylogenomic; virus-host network

Introduction

The family *Baculoviridae* is a group of insect-specific large DNA viruses isolated from members of the orders of Lepidoptera, Hymenoptera, and Diptera, and divided into

four genera, including *Alphabaculovirus*, *Betabaculovirus*, *Gammabaculovirus*, and *Deltabaculovirus* (Harrison *et al.*, 2018a). The *Alphabaculovirus* can be subdivided into two groups, I and II based on phylogenetic analysis (Harrison *et al.*, 2018b; Herniou and Jehle, 2007). Members of *Alphabaculovirus* have been characterized by a typical biphasic infection cycle, in which two kinds of progeny virions are produced: occlusion-derived viruses (ODVs), which are embedded in occlusion bodies (OBs) and are responsible for *per os* infection in the insect midgut, and the budded viruses (BVs) responsible for systemic infection inside the insect host (Slack and Arif, 2007).

The BV and ODV differ in their envelope proteins that are responsible for their distinct functions (Hou

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Abbreviations: AcMNPV = *Autographa californica* nucleopolyhedrovirus; BV = budded viruses; IAP = inhibitors of apoptosis; ML = maximum likelihood; NLS = nuclear localization signal; NES = nuclear export signal; OB = occlusion bodies; ODV = occlusion-derived viruses; PIF = *per os* infectivity factor; TM = transmembrane domain

et al., 2013). Their nucleocapsids contain the covalently closed, double-stranded DNA (dsDNA) with a genome size ranging from 80 to 180 kb with 80–190 open reading frames (ORFs) (Hayakawa *et al.*, 1999; Lauzon *et al.*, 2004). So far, a total of 38 conserved core genes have been found to be present in all sequenced baculoviral genomes (2012; Javed *et al.*, 2017; van Oers and Vlak, 2007; Zhu *et al.*, 2018). Among the 38 conserved core genes, 37 genes were identified by Garavaglia *et al.* (2012). Recently, the AC110 missed in some previous annotated baculoviral genomes (Javed *et al.*, 2017), was identified as the 38th gene shared by all baculoviruses. In addition, 26 genes were found to be present at least in both, the *Alphabaculovirus* and *Betabaculovirus* (Garavaglia *et al.*, 2012).

Analyses of protein conservation have promoted investigations on roles of viral proteins in infection processes, such as ODV envelopment, virion entry and transport, DNA replication and processing (Braunagel and Summers, 2007; Li *et al.*, 2018; Liu *et al.*, 2020; Vanarsdall *et al.*, 2007; Volkman, 2007; Wang *et al.*, 2019a; Zhang *et al.*, 2018b). It also has promoted the improvement of baculoviral expression vector. The baculoviral expression vector of a large size genome of about 133 kbp provides multiple insertion sites and high capacities for expression of multiple protein complexes. It has been modified by the deletion of BV non-essential proteins of both viral chitinase (V-ChiA) and viral cathepsin (V-CATH) to improve secretion and stability of recombinant proteins (Hitchman *et al.*, 2011). Furthermore, the baculoviral expression vector has been improved as different powerful systems for recombinant expression and post-translational modifications of multiple proteins, such as MultiBac (Berger *et al.*, 2004), SweetBac (Palmerberger *et al.*, 2012) and OmniBac systems (Vijayachandran *et al.*, 2013). Based on a combination of baculoviral protein conservation and functional genomic studies, the minimal and functional expression vector, which would contain less non-essential gene content but still be infectious and able to propagate and produce heterologous protein in cell culture has been already proposed by Vijayachandran *et al.* (2013).

With the development of functional genomics of baculoviruses and increasing sequenced baculoviral genomes, in this study, we firstly performed phylogenomic analysis of 76 baculoviral genomes based on 38 core proteins, and further supply a global and integrated view of baculoviral protein conservation and functional genomic investigations. The AC106/107 and potential functional domains and sites of ODV-E18 were supplied as future research targets for exploring BV/ODV formation mechanism. These results would facilitate further baculoviral genome annotation, understanding of baculoviral infection

mechanism and future construction of a more powerful baculoviral expression vector with a minigenome.

Materials and Methods

Genome and protein databases. A total of 76 individual genomes of baculoviruses with known reports were collected from the GenBank database (www.ncbi.nlm.nih.gov) (Table S1). The genomes of 53 alphabaculoviruses, 19 betabaculoviruses, 3 gammabaculoviruses, and 1 deltabaculovirus were included. The non-redundant annotated protein dataset of each baculovirus genome was acquired from NCBI. Recently, the AC110 was identified as a conservative *per os* factor (PIF) in all baculoviruses, which rectified previous missing annotations of their presence in the gammabaculoviruses and deltabaculoviruses (Javed *et al.*, 2017). To avoid such issue, the proteins were re-predicted for all the viral genome sequences using the getorf (<http://emboss.bioinformatics.nl/cgi-bin/emboss/getorf>) and proteins ≥ 50 aa were kept. Thus, each baculovirus had three different datasets: a) the Individual Genome DataSet (IGD) containing the corresponding genome sequence, b) the Individual Proteome DataSet (IPD) containing the corresponding annotated proteins, and c) the Individual Repredicted Proteome DataSet (IRPD) containing the corresponding re-predicted proteins.

Identification of homologue groups among baculoviruses. An overview of the identification of baculoviral homologue groups was demonstrated in a schematic diagram (Fig. S1). In the first round, the OrthoFinder (v2.3.14) (Emms and Kelly, 2019) was used for all-versus-all BLAST search among all the IPDs to identify homologue groups present in at least two baculoviral genomes, with the settings `-S blast -M msa`. The disability of orthology prediction tools for identification of all orthologs has been described (Garavaglia *et al.*, 2012), and also found by the OrthoFinder in this study. Thus, the OrthoFinder identified homologue groups were used as the Primary Homologous Group Database (PHGD), and the PHGD was further manually annotated and merged according to references (Table S1) of the genome sequences and previous identification of the remote orthology (Garavaglia *et al.*, 2012), and resulted in a Revised Homologous Group Database (RHGD). In the second round, the RHGD was subjected to all-versus-all search against the IRPDs with the settings `"-S blast -M msa -os"` in the second round to screen candidate homologues that were not annotated in the IPDs. The candidate homologues overlapping ≥ 25 aa with the ORFs present in the IPD were excluded and the remaining candidate homologues were then manually checked using BLASTP (McGinnis and Madden, 2004) in NCBI to search homologues in baculoviruses with an inclusion threshold of $1e-5$. Further, local TBLASTN was performed by searching against the corresponding viral genome sequences using BioEdit 7.5.2 (Alzohairy, 2011) with an expected threshold of $1e-5$ to

identify candidate homologues, and all the selected candidate homologues were manually checked using BLASTP (McGinnis and Madden, 2004) as described above. The Final Homologous Group Database (FHGD) containing all the sequences of individual homologue groups and a statistical matrix of presence of homologous proteins in individual baculoviruses were used for further analysis below.

Phylogenomic analysis of 76 baculoviruses. All the homologue groups present in all of the 76 baculoviral genomes were subjected to MAFFT (v7.310) alignment using the iterative refinement options G-INS-i with the settings --globalpair --maxiterate 1000 (Nakamura *et al.*, 2018). The alignment proteins were filtered using the Gblocks (v0.91b) (Talavera and Castresana, 2007) with the settings -t = p -b1 = 39 -b2 = 40 -b3 = 10 -b4 = 2 -b5 = h -e = -gb, and then concatenated in a stationary order (AcMNPV ORF order) using TBtools (v1.046) (Chen *et al.*, 2020) and PhyloSuite (v1.2.1) (Zhang *et al.*, 2020). A full tree search for protein models was then performed using the IQ-TREE (v1.6.12) (Minh *et al.*, 2020) with the settings -m MF -mtree -nt 4 (Kalyaanamoorthy *et al.*, 2017). Then, phylogenetic inference using maximum likelihood (ML) was performed by IQ-TREE using the best protein model with the settings iqtree -m LG+F+R7 -bb 10000 -alrt 10000 -nt 4 -bnni, supplying branch support with SH-aLRT value (Shimodaira-Hasegawa approximate likelihood ratio test) (Anisimova *et al.*, 2011) and ultrafast bootstrap (UFboot) value. The resulted file was used for further construction of an ultrametric tree (also known as time tree) with the RelTime method (Tamura *et al.*, 2012) implemented in MEGA X (Kumar *et al.*, 2018) following the well-illustrated protocol (Mello, 2018). The temporal information of the calibration points for NeleNPV (*Neodiprion lecontei* NPV) and CpGV (*Cydia pomonella* granulosis virus) and that for CpGV and AcMNPV (Theze *et al.*, 2011) was used for calibrating the relative node ages. The FigTree (<http://tree.bio.ed.ac.uk/software/figtree/>) was used for viewing the acquired tree file and further construction of phylogenetic tree with manual modifications.

Gain and loss of proteins among the baculoviruses. The analysis of expansion and contraction of protein homologue groups was performed using the CAFÉ 4 (De Bie *et al.*, 2006) following the software tutorial. The major input files used were the newick-formatted ultrametric tree above and a tab-delimited txt file containing gene family data (De Bie *et al.*, 2006). The parameter *p*-value was set as 0.01. The command 'lambda -s' was used for estimation of the λ -value (defined as the probability of both gene gain and loss per gene per unit time in the phylogeny).

Construction of baculovirus-host interaction network. For comprehensive understanding of viral infection mechanism, a literature survey was performed for collecting viral DNA/protein-protein interactions of the baculoviruses and virus-host protein interactions (Table S2). The interaction network was constructed using the Cytoscape (v3.8.0) (Shannon *et al.*, 2003) following the software tutorial.

Bioinformatics analysis of ODV-E18. The conserved motifs in proteins were searched using the MEME (Bailey and Elkan, 1994). The putative signal peptide (SP), transmembrane (TM) domain, nuclear localization signal (NLS) and nuclear export signal (NES) were predicted using SignaIP 5.0 (Almagro Armenteros *et al.*, 2019), TMHMM 2.0 (Krogh *et al.*, 2001), SeqNLS (Lin and Hu, 2013) and NetNES 1.1 (la Cour *et al.*, 2004), respectively. In addition, N-glycosylation sites and phosphorylation sites were predicted using NetNGlyc 1.0 (<http://www.cbs.dtu.dk/services/NetNGlyc/>) and NetPhos 3.1 (Blom *et al.*, 1999), respectively.

Results

Phylogenomic tree of baculoviruses based on 38 core genes

The homologue groups of 38 core genes were identified among the 76 genome-sequenced baculoviruses, with 11 newly annotated proteins (in baculoviral IRPDs) in 3 homologue groups of AC78, P6.9 (AC100) and PIF7 (AC110) (Table S3). Phylogenomic inference with concatenated 38 core genes was generated (Fig. 1) and then molecular clock inference was computed according to the previous age estimation of baculoviruses (Theze *et al.*, 2011). The topology of the resulted ML phylogenetic tree with timeline classified all the 76 baculoviruses into four clades corresponding to the four genus of the family *Baculoviridae*, consistent with previous tree inference (Harrison *et al.*, 2017; Santos *et al.*, 2018).

Comprehensive identification of baculoviral homologues

In combination of application of the orthology prediction tool OrthoFinder, reannotation of baculoviral genomes, manual BLASTP analysis and literature exploration (Garavaglia *et al.*, 2012; Javed *et al.*, 2017), a total of 992 homologue groups were identified, among which, 429 homologue groups were with the taxon occupancy of ≥ 2 species. Among the 429 homologue groups, 56 proteins in 25 homologue groups were newly annotated (Table S3) in baculoviral IRPDs, including homologues of AC78, P6.9 (AC100), PIF7 (AC110), PIF9 (AC108) present in all baculoviruses, and three proteins present in all the alphabaculoviruses and betabaculoviruses (Garavaglia *et al.*, 2012), including LEF3 (AC67), TLP (AC82) and P24 (AC129). In addition, the new annotation of the AC76 homologue in the *Peridroma* sp. NPV (PespNPV) IRPD facilitated identification of AC76 as a new number found in all baculoviruses without the CuniNPV (Table S3), while the LEF 10 (AC53a) was identified as a new number that

was present in all the members of genera *Alphabaculovirus* and *Betabaculovirus*.

Additionally, the literature-based manual checking of fused proteins was performed to refine the statistical matrix of number of homologues present in the individual baculoviruses. Each of both proteins in the fusion protein

form was considered to be present in the corresponding baculovirus. Taking the GP37/V-Ubi (viral ubiquitin) in *Spodoptera littoralis* MNPV (SpliMNPV) as an example, both of numbers of GP37 and V-Ubi in statistical matrix were manually revised as “1”, but not as “1” for only one protein. Thus, this supplied a comprehensive identifica-

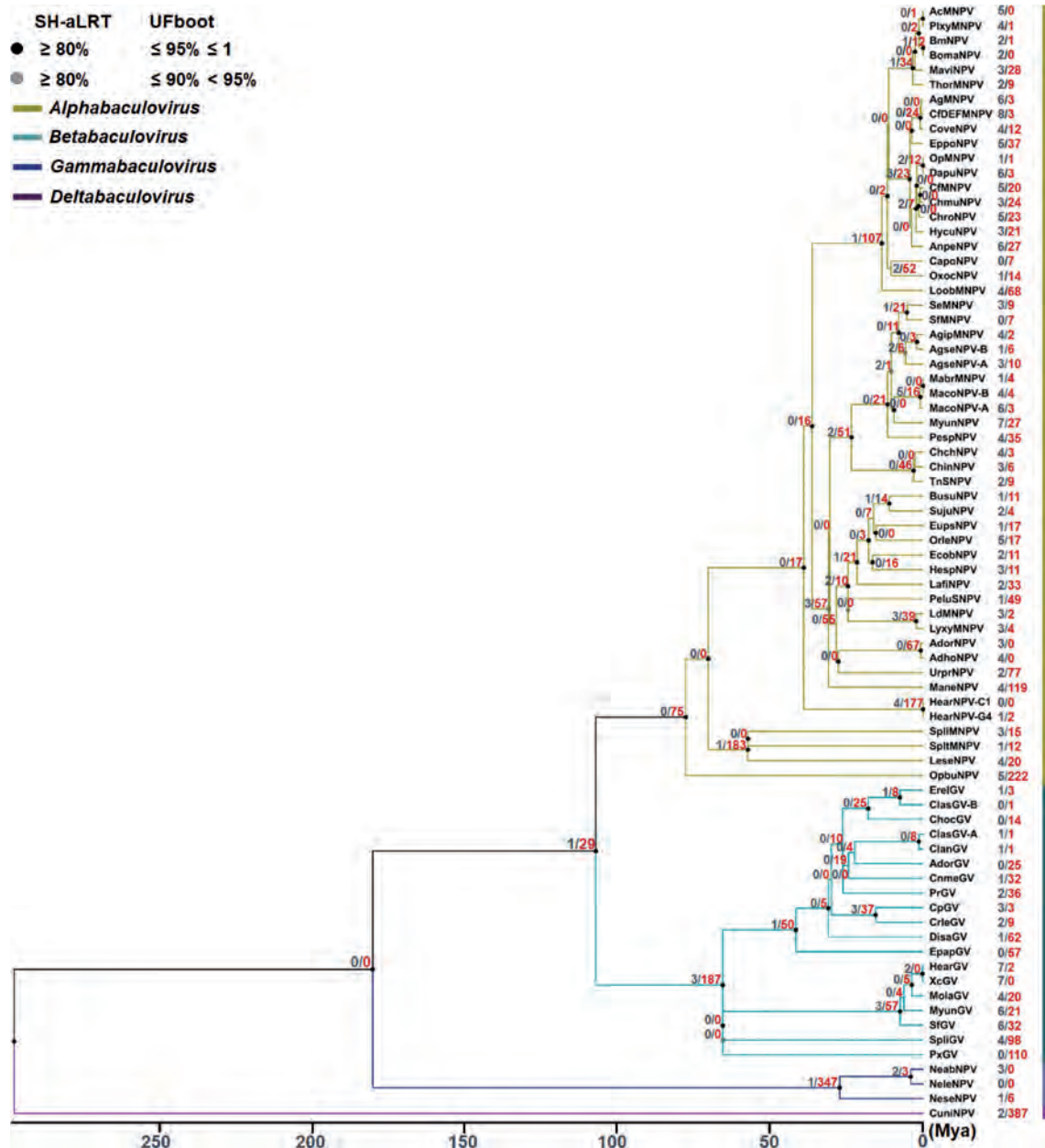


Fig. 1

TimeTree with mapped gain and loss of homologue groups (taxon occupancy of ≥ 2 species) based on the maximum likelihood (ML) inference using the concatenated sequences of 38 core proteins from 76 baculoviruses (Table S1)

The branch support with SH-aLRT value (measurement of confidence in given branches) and ultrafast bootstrap (UFboot) value was determined using a SH-like method implemented in the IQ-TREE software. Mya: millions of years ago. -/-: amount of gain/loss of homologue groups (indicated at the far right after the virus species acronyms).

tion of baculoviral protein homologues and a global and updated view of their distribution in genomes of the baculovirus species. Further, all the 429 homologue groups with the taxon occupancy of ≥ 2 species (Fig. 1) were mapped on the phylogenetic tree (Fig. 1), demonstrating a global view of gene gain and loss which are indicated at the far right after the virus species acronyms (Fig. 1).

Integrated view of protein conservation and functions of baculoviral homologue groups

All the distributions of 429 homologue groups with a taxon occupancy of ≥ 2 species (Table S3) were further summarized (Table S4): 38 core proteins present in all baculoviruses (Garavaglia *et al.*, 2012; Javed *et al.*, 2017), and 5, 7 and 25 conserved in alphabaculoviruses, betabaculoviruses, and gammabaculoviruses, respectively (Table S4). For understanding of the correlation between conservation and functional importance of proteins, a comprehensive literature view of functional genomic studies of baculoviruses was summarized (Table 1 and Table S5) and focused on studies on AcMNPV proteins and their homologues in BmNPV and HearNPV (Table S5).

For AcMNPV, most of the 65 (38+11+1+15, see Table 1) homologue groups (covering 69 AcMNPV ORFs) found at least in all the alphabaculoviruses and betabaculoviruses are essential/important for BV, ODV and OB production and infectivity, with the exception of the 2 dispensable protein homologue groups of P24 (AC129) and PEP (AC131) (Table 1 and Table S5). Apart from the 65 homologue groups above, 9 (4+5, see Table 1) homologue groups (covering 9 AcMNPV ORFs) were found at least in all the alphabaculoviruses, among which, VP80 (AC104), AC34, EXON0 (AC141) and P10 (AC137) are essential/important for BV, ODV and OB, while EGT (AC15), AC18, AC19, AC29 and AC55 are dispensable proteins for virus infection. Furthermore, 73 (47+25+1, see Table 1) homologue groups (covering the 78 AcMNPV ORFs) are not conserved in the alphabaculoviruses. Among the 73 homologue groups, 6 homologue groups including PKIP (AC24), BJDP (AC51), AC69, AC73, GP64 (AC128), AC132 are essential for BV and ODV production (Table 1 and Table S5). The 9 homologue groups including PTP (AC1), BRO (AC2), AC11, ARIF-1 (AC20/AC21), AC52, AC79, CG30 (AC88)/PE38 (AC153), P35 (AC135), IE2 (AC151) and AC152, are important for BV and ODV production (Table 1 and Table S5). The AC5 and AC43 are important for OB formation, while V-Chi (AC126) and V-CATH (AC127) are important for host liquefaction to facilitate OB release. Nevertheless, the four homologue groups and the remaining 54 of the 73 homologue groups are not dispensable for BV and ODV production. It was noted that for the proteins essential or important for virion infection and production, nearly all the proteins

had demonstrated their functional roles in infection process. Nevertheless, for the ODV-E18 (McCarthy and Theilmann, 2008) and the AC106/107 (Chen *et al.*, 2021), no investigation had been performed to reveal their detailed functional roles in the processes of baculoviral biphasic infection cycle.

Integrated baculovirus-host interaction network

To gain a global perspective on how a baculovirus assembles its progeny virions and interfaces with its host, several researches focused on the exploration of interactions among virion structural proteins and host proteins (Braunagel and Summers, 2007; Peng *et al.*, 2010; Yue *et al.*, 2018; Zhang *et al.*, 2018a). For a comprehensive view of baculovirus-host interaction network, a literature exploration based on functional genomic studies of baculoviruses was performed, supplying a list of 156 viral protein interactions (Table S2A), 12 viral protein-DNA interactions (Table S2B) and 72 virus-host protein interactions (Table S2C). With the exception of 24 self-interactions of viral proteins, a baculovirus-host interaction network was illustrated in Fig. 2, comprising of 98 nodes (including 26 host proteins nodes) and 214 edges (interactions).

In the interaction network, proteins including AC76 (Hu *et al.*, 2010), AC93 (Yuan *et al.*, 2011) and P48 (AC103) (Wang *et al.*, 2019b) are required for virus-induced intranuclear microvesicle formation, and interact with multiple host proteins involved in the ESCRT (endosomal complexes required for transport) pathway. These viral proteins interact with each other and other viral proteins including AC11 (Tao *et al.*, 2015), AC78 (Li *et al.*, 2014), GP41 (AC80) (Li *et al.*, 2018), ODV-EC43 (AC109) (Lin *et al.*, 2009), 49K (AC142) (McCarthy *et al.*, 2008) and AC146 (Dickison *et al.*, 2012), (Fig. 2). All of the viral proteins above are essential for BV production and ODV envelopment.

FP25K interacts with BV major envelope protein GP64 (Zhang *et al.*, 2018a) while ODV-E25 interacts with BV protein ME53 (Zhang *et al.*, 2018a) (Fig. 2). Both of the two proteins interact with nucleocapsid proteins (Zhang *et al.*, 2018a) and ODV major envelope protein ODV-E66 (Braunagel *et al.*, 2013). FP25K (Braunagel *et al.*, 1999; Li *et al.*, 2015) and ODV-E25 (Chen *et al.*, 2012) are important for BV infectivity and ODV formation, implicating their capability of regulation of the BV/ODV formation mechanism. Nevertheless, the two proteins are not essential for BV and ODV production (Braunagel *et al.*, 1999; Chen *et al.*, 2012; Li *et al.*, 2015). The ODV-E18 interacts not only with the FP25K and ODV-E25, but also with BV proteins GP64 and ME53, multiple PIFs and ODV-E66, nucleocapsid proteins and P48 (AC103) (Fig. 2). It seems that the ODV-E18

Table 1. Overview of the conservation and functions of AcMNPV ORFs

No. [#]	Alpha-	Beta-	Gamma-	Delta-	ORFs	Function annotation ^b
38	++	++	++	++	PIF0 (AC138), PIF1 (AC119), PIF2 (AC22), PIF3 (AC115), PIF4 (AC96), PIF5 (AC148), PIF6 (AC68), PIF7 (AC110)	Essential for <i>per os</i> infection
					PIF8 (AC83), LEF1 (AC14), LEF2 (AC6), LEF4 (AC90), LEF5 (AC99), LEF8 (AC50), LEF9 (AC62), P47 (AC40), AC53, VP1054 (AC54), DNA POL (AC65), Desmoplakin (AC66), VLF-1 (AC77), GP41 (AC80), VP39 (AC89), P33 (AC92), P18 (AC93), Helicase (AC95), 38K (AC98), P6.9 (AC100), ODV/BV-C42 (AC101), ODV-EC43 (AC109), ALK-EXO (AC133), 49K (AC142), ODV-E18 (AC143), ODV-EC27 (AC144)	Essential for BV and ODV
					AC78, AC81, ODV-E25 (AC94), P48 (AC103)	Important for BV and ODV
11	++	++	++	-	DBP (AC25), LEF11 (AC37), AC75, AC76, AC106/107	Essential for BV and ODV
					FP25K (AC61), IAPs (AC27/AC71) ^a	Important for BV and ODV
					PIF9 (AC108)	Essential for <i>per os</i> infection
					AC145/AC150	Important for <i>per os</i> infection
					POLH (AC8)	Essential for OB formation
					PEP (AC131)	Dispensable
1	++	++		++	F protein (AC23)	Essential for BV ^c
15	++	++	-	-	P78/83 (AC9), PK1 (AC10), LEF10 (AC53a), LEF3 (AC67), TLP (AC82), P12 (AC102), AC146	Essential for BV and ODV
					AC13, LEF6 (AC28), V-Ubi (AC35), PP31 (AC36), AC38, ME53 (AC139), IE0 (AC147-0)/IE1 (AC147)	Important for BV and ODV
					P24 (AC129)	Dispensable
4	++	+	-	-	EXON0 (AC141)	Important for BV and ODV
					P10 (AC137)	Important for OB release
					EGT (AC15), AC29	Dispensable
5	++	-	-	-	VP80 (AC104)	Essential for BV and ODV
					AC34	Important for BV and ODV
					AC18, AC19, AC55	Dispensable
47	+	-	-	-	PKIP (AC24), BJDp (AC51), AC69, AC73, AC132	Essential for BV and ODV
					AC11, ARIF-1 (AC20/AC21), AC52, IE2 (AC151), AC152	Important for BV and ODV
					AC5, AC43	Important for OB formation
					AC4, AC7, BV/ODV-E26 (AC16), AC17, AC26, AC30, LEF12 (AC41), GTA (AC42), AC44, AC45, AC47, AC48, PCNA (AC49), AC56, AC57, HCF-1 (AC70), AC72, AC74, AC85, AC91, AC97, AC114, AC116, AC117, AC118, AC120, AC121, AC122, PK2 (AC123), AC124, GP16 (AC130), P26 (AC136), AC140, AC149, AC154	Dispensable
25					GP64 (AC128)	Essential for BV
					PTP (AC1), BRO (AC2), AC79, CG30 (AC88)/PE38 (AC153), P35 (AC135) ^a	Important for BV and ODV
					V-Chi (AC126), V-CATH (AC127)	Important for host liquefaction
					CTX (AC3), AC12, SOD (AC31), FGF (AC32), HisP (AC33), P43 (AC39), AC46 (ODV-E66), ChaBs (AC58/AC59/AC60), AC63, GP37 (AC64), AC84, PNK/PNL (AC86), HE65 (AC105), AC111, AC112/AC113, LEF7 (AC125), 94K (AC134)	Dispensable
1	+	-	-	++	P15 (AC87)	Dispensable

Note: #: No. of homologue groups. ++: found in all the baculoviruses among the genus. +: found in the baculoviruses but not found in all the baculoviruses among the genus. -: not found in any of the baculoviruses among the genus. ^ainvolvement in antiapoptosis and may be important for BV and ODV. ^bDispensable: dispensable for BV and ODV production. ^cnot essential for AcMNPV due to the presence of GP64.

groups that include the 56 newly identified proteins from reannotated genomes. This would be used as a checklist for future annotation of increasing sequenced genomes and baculoviral comparative genomics. This study also supplied a comprehensive literature view of protein functions, and proposed research targets for future functional investigations, including the two proteins AC106/107, ODV-E18, and potential functional domains and post-translational sites of ODV-E18.

Most of the 992 protein homologue groups are distributed in each baculoviral genome as single copy. Nevertheless, a few proteins are found to be multigene family proteins, such as BROs (baculoviral repeat ORFs), ChaBs, IAPs (inhibitors of apoptosis), AC145/AC150 homologue

groups. Positive synergy effects of genes, such as ac145 and ac150 (important for *per os* infection) (Lapointe et al., 2004), and *ie1* and *ie-0* genes (Lu et al., 2005; Stewart et al., 2005), have been noted. It has been found that proteins in the same homologue group or with functional similarity may function with positive synergy effects and thus be beneficial for virus infection, such as AC145 and AC150 (Lapointe et al., 2004), both of which were noted as “important for *per os* infection” here. It also has been noted that the *iap* genes, which belong to a large gene family with eukaryote members (Kuzio et al., 1999), would function in mutual complementation, and result in a misconception that the genes may be not essential for BV and ODV production during deletion of each of the

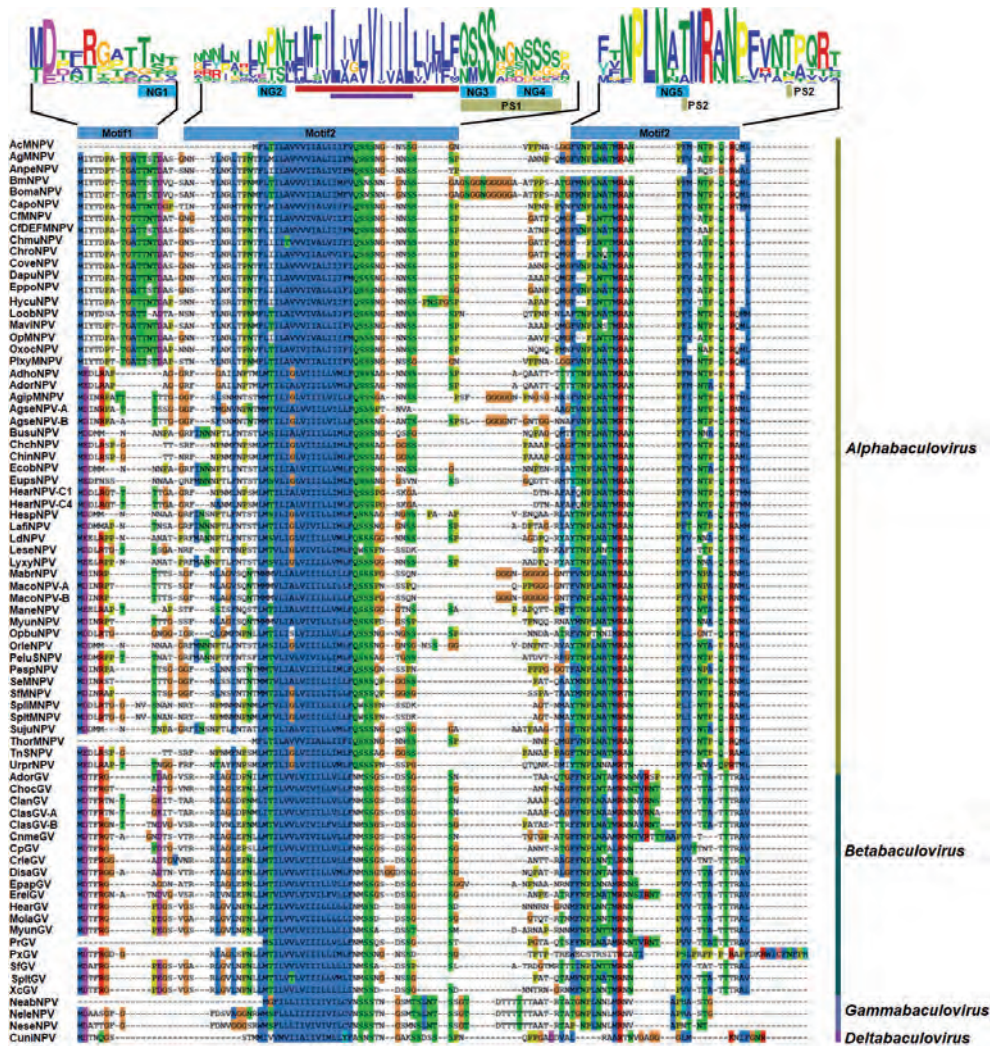


Fig 3

Analysis of functional domains and posttranslational modification sites of ODV-E18

The transmembrane (TM) domain is indicated as red horizontal bar, while the nuclear localization signal (NLS) domain is indicated as red horizontal bar. PS: indicates phosphorylation sites, NG: indicates N-glycosylation sites.

individuals. The IAP homologues (AC27 and AC71) and P35 are involved in antiapoptosis, assuring viral normal replication since deletions of individual IAP, two IAPs or combination of one IAP and P35 did not result in deficient AcMNPV replication in permissive cells (Griffiths *et al.*, 1999). Here we noted all the IAPs and P35 as “important for BV and ODV” for an awareness during construction minimum AcMNPV genome.

The virus-host network components (Fig. 2) would work together to facilitate accomplishment of viral infection cycle. The interactions among *per os* infectivity factors (PIFs) including P74 (PIF0), PIF1-5, PIF8 function in formation and/or assembly of PIF complex (Table S5) and thus would facilitate the interaction of PIF complex with host receptor(s) for mediating ODV entry into host midgut cells. The interactions among nucleocapsid proteins, ODV envelop proteins and DNA interaction proteins (Fig. 2) would facilitate virion assembly and stability. Multiple viral proteins essential for viral-induced intranuclear microvesicle formation and the egress of BV are involved in interactions with the host proteins in ESCRT pathway (Yue *et al.*, 2018). The ESCRT pathway has been found to be essential for membrane deformation (Hurley, 2015), entry and release of many enveloped viruses including AcMNPV (Votteler and Sundquist, 2013; Yue *et al.*, 2018). Taking protein conservation in to consideration, the summarized virus-host network would be shared by most of the baculoviruses to accomplish viral infection cycle.

Due to the correlation between protein conservation and functional importance, highly conserved proteins have attracted much attention on their functional roles in the baculoviral infection cycle (Garavaglia *et al.*, 2012; van Oers and Vlak, 2007). Here, a comprehensive collection of literature on the investigation of baculoviral genes using systematical (Chen *et al.*, 2021; Ono *et al.*, 2012) and individual gene knockout supplied an updated encyclopedia of genes of the reference baculovirus, AcMNPV (David *et al.*, 2009) (Table S5). All the 156 AcMNPV ORFs (and their homologues) were investigated using knockout analysis. Functions of 145 ORFs have been supported by at least two referred papers using knock-out analysis of each gene. This supplies a rapid and global view for further construction of BV minigenome by deleting non-essential/dispensable genes for BV, in spite of the highly conserved genes specifically essential or important for ODV infection (Chen *et al.*, 2021), such as multiple PIFs and Polyhedrin (POLH). We also anticipate future construction of the baculovirus containing a minigenome that carries minimum genetic information capable of supporting the baculoviral biphasic cycle. This would facilitate simulation of the natural evolution of baculoviral genome in laboratory through serial passage of viruses, and thus

gain valuable information on gain-and-loss of genes and virus-host interactions.

Furthermore, ODV-E18, as a conserved and essential protein for BV and ODV (McCarthy and Theilmann, 2008), interacts with multiple viral proteins in the virus-host interaction network. Nevertheless, its detailed role in viral infection cycle has not been revealed, likewise the AC106/107 which is also essential protein for BV and ODV (Chen *et al.*, 2021). Further investigations of posttranslational modifications (Fig. 3) of ODV-E18 also supply potential research targets for further functional studies of ODV-E18. It had been demonstrated (Hou *et al.*, 2013) that the ODV-E18 on HearNPV BV virions were *N*-glycosylated at the N64 site in the N-A-T sequon corresponding to the N5 region (Fig. 3), while the ODV-E18 on HearNPV ODV virions were phosphorylated at the T75 and T79 sites in the NTPQRTMM sequon corresponding to the PS3 (Fig. 3). It is anticipated that the further investigations of AC106/107 and both of *N*-glycosylation and phosphorylation of ODV-E18 would supply more insights into their functional roles in the processes of viral biphasic infection cycle.

Towards construction of a minimal and functional baculoviral expression vector supporting maximum expression of functional proteins and protein complexes (Vijayachandran *et al.*, 2013), serial knockout of large regions distributed between AcMNPV BV-essential genes can be conducted. However, further laboratory validation to investigate their effects on BV production, such as the regions *ac15-ac23*, *ac26-ac33*, *ac41-ac49*, and even the larger regions *ac15-ac33*, *ac55-ac74* and *ac110-ac138* while keeping essential/important genes (located in the regions) for BV are required. It has been suggested that all the *hr* (homologous repeat) regions, which are beneficial for enhanced transcription/expression of genes, should be kept, in spite of their non-essential role in DNA replication and BV production (Carstens and Wu, 2007; Leisy and Rohrmann, 1993; Rodems and Friesen, 1993; Venkaiah *et al.*, 2004).

In summary, this study provided a global and integrated view of protein conservation and functional genomic investigations of baculoviruses. It will facilitate further baculoviral genome annotation, understanding baculoviral infection mechanism and future construction of a powerful baculoviral expression vector with a minigenome.

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SUPPLEMENTARY INFORMATION

An integrated view of protein homologue groups and functional genomic development of baculoviruses: towards understanding baculoviral infection mechanism and improving baculoviral expression vector

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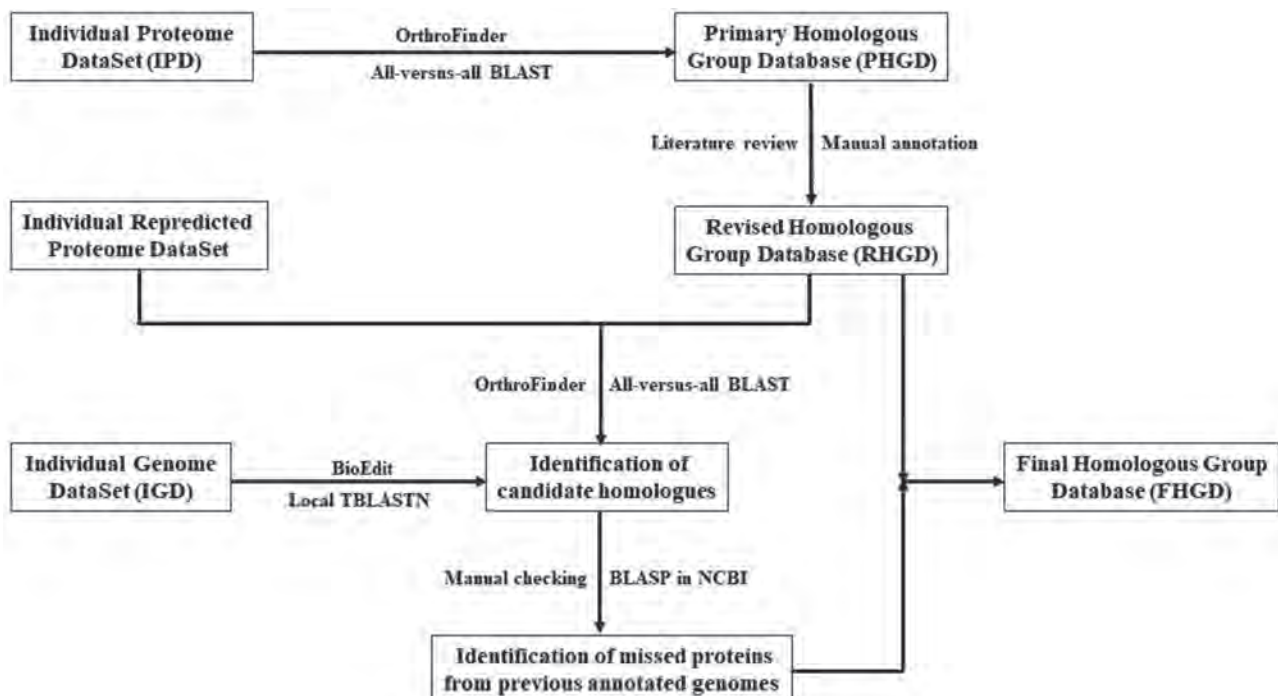


Fig. S1

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Table S1. Overview of 76 baculoviral genomes used in this study

Virus name	Abbreviation	Genbank accession	Length (bp)	G+C (%)	No. of ORFs	Reference^b
<i>Alphabaculovirus 43</i>						
<i>Antheraea pernyi</i> NPV	AnpeNPV	NC_008035.3/DQ486030.3	126,629	53.4	147	(Nie <i>et al.</i> , 2007)
<i>Anticarsia gemmatalis</i> MNPV	AgMNPV	NC_031761.1/KR815466.1	131,855	44.5	156	(Brito <i>et al.</i> , 2015)
<i>Autographa californica</i> MNPV	AcMNPV	NC_001623.1/L22858.1	133,894	40.7	156	(Ayres <i>et al.</i> , 1994)
<i>Bombyx mori</i> NPV	BmNPV	NC_001962.1/L33180.1	128,413	40.4	143	(Gomi <i>et al.</i> , 1999)
<i>Bombyx mandarina</i> NPV	BomaNPV	FJ882854.1	126,770	40.2	141	(Xu <i>et al.</i> , 2010)
<i>Catopsilia Pomona</i> NPV	CapoNPV	NC_030240.1/ KU565883.1	128,058	39.7	130	(Wang <i>et al.</i> , 2016)
<i>Choristoneura fumiferana</i> MNPV	CfMNPV	NC_004778.3/ AF512031.3	129,593	50.1	146	(de Jong <i>et al.</i> , 2005)
<i>Choristoneura fumiferana</i> DEFMNPV	CfDEFMNPV	NC_005137.2/ AY327402.2	131,160	45.8	149	(Lauzon <i>et al.</i> , 2005)
<i>Choristoneura murinana</i> NPV	ChmuNPV	NC_023177.1/ KF894742.1	124689	50.0	147	(Rohrmann <i>et al.</i> , 2014)

<i>Choristoneura rosaceana</i> NPV	ChroNPV	NC_021924.1/KC961304.1	129,052	48.6	149	(Thumbi <i>et al.</i> , 2013)
<i>Condylorrhiza vestigialis</i> MNPV	CoveNPV	NC_026430.1/ KJ631623.1	125,767	42.9	138	(Castro <i>et al.</i> , 2017)
<i>Dasychira pudibunda</i> NPV	DapuNPV	KP747440.1	136,761	54.4	161	(Krejmer <i>et al.</i> , 2015)
<i>Epiphyas postvittana</i> NPV	EppoNPV	NC_003083.1/ AY043265.1	118,584	40.7	136	(Hyink <i>et al.</i> , 2002)
<i>Hyphantria cunea</i> NPV	HycuNPV	NC_007767.1/ AP009046.1	132,959	45.1	148	(Ikeda <i>et al.</i> , 2006)
<i>Lonomia oblique</i> MNPV	LoobMNPV	NC_043520.1/KP763670.1	120,022	35.7	134	(Aragao-Silva <i>et al.</i> , 2016)
<i>Maruca vitrata</i> NPV	MaviNPV	NC_008725.1/ EF125867.1	111,953	38.6	126	(Chen <i>et al.</i> , 2008)
<i>Orgyia pseudotsugata</i> MNPV	OpMNPV	NC_001875.2/ U75930.2	131,990	55.1	152	(Ahrens <i>et al.</i> , 1997)
<i>Oxyplax ochracea</i> NPV	OxocNPV	NC_043529.1/MF143631.1	113.97	31.2	124	(Wang <i>et al.</i> , 2018)
<i>Plutella xylostella</i> MNPV	PlxyMNPV	DQ457003.1	134,417	40.7	153	(Harrison and Lynn, 2007)
<i>Thysanoplusia orichalcea</i> MNPV	ThorMNPV	NC_019945.1/ JX467702.1	132,978	37.9	145	(Wang <i>et al.</i> , 2012)
<i>Adoxophyes honmai</i> NPV	AdhoNPV	NC_004690.1/AP006270	113,220	35.6	125	(Nakai <i>et al.</i> , 2003)
<i>Adoxophyes orana</i> NPV	AdorNPV	NC_011423.1/ EU591746.1	111,724	35	121	(Hilton and Winstanley, 2008)

<i>Agrotis ipsilon</i> MNPV	AgipMNPV	NC_011345.1/EU839994.1	155,122	48.6	163	(Harrison, 2009)
<i>Agrotis segetum</i> NPV-A	AgseNPV-A	NC_007921.1/DQ123841.1	147,544	45.7	153	(Jakubowska <i>et al.</i> , 2006)
<i>Agrotis segetum</i> NPV-B	AgseNPV-B	NC_025960.1/KM102981.1	148,981	45.7	150	(Wennmann <i>et al.</i> , 2015)
<i>Buzura suppressaria</i> NPV	BusuNPV	NC_023442.1/ KF611977.1	120,420	36.8	127	(Zhu <i>et al.</i> , 2014)
<i>Chrysodeixis chalcites</i> SNPV	ChchNPV	NC_007151.1/ AY864330.1	149,622	39.1	151	(van Oers <i>et al.</i> , 2005)
<i>Chrysodeixis includens</i> NPV	ChinNPV	NC_026268.1/KJ631622.1	139,132	39.3	141	(Craveiro <i>et al.</i> , 2015)
<i>Ectropis obliqua</i> NPV	EcobNPV	NC_008586.1/ DQ837165.1	131,204	37.6	126	(Ma <i>et al.</i> , 2007)
<i>Euproctis pseudoconspersa</i> NPV	EupsNPV	NC_012639.1/ FJ227128.1	141,291	40.4	139	(Tang <i>et al.</i> , 2009)
<i>Helicoverpa armigera</i> SNPV-G4	HearNPV-G4	NC_002654.2/AF271059.2	131,405	39.1	135	(Chen <i>et al.</i> , 2001)
<i>Helicoverpa armigera</i> SNPV-C1	HearNPV-C1	NC_003094.2/AF303045.2	130,759	38.9	137	(Zhang <i>et al.</i> , 2005)
<i>Hemileuca</i> sp. NPV	HespNPV	NC_021923.1/ KF158713.1	140,633	38.1	137	(Rohrmann <i>et al.</i> , 2013)
<i>Lambdina fiscellaria</i> NPV	LafiNPV	NC_026922.1/ KP752043.1	157,989	43.7	137	(Rohrmann <i>et al.</i> , 2015b)
<i>Leucania separata</i> NPV	LeseNPV	NC_008348.1/ AY394490.1	168,041	48.6	169	(Xiao and Qi, 2007)

<i>Lymantria dispar</i> MNPV	LdMNPV	NC_001973.1/ AF081810.1	161,046	57.5	164	(Kuzio <i>et al.</i> , 1999)
<i>Lymantria xyli</i> MNPV	LyxyMNPV	NC_013953.1/GQ202541.1	156,344	53.4	157	(Nai <i>et al.</i> , 2010)
<i>Malacosoma neustria</i> NPV	ManeNPV	NC_040606.1/KY968317.1	130.2	38.2	131	(Gencer <i>et al.</i> , 2018)
<i>Mamestra brassicae</i> MNPV	MabrMNPV	NC_023681.1/ JQ798165.1	152,710	39.8	158	(Choi <i>et al.</i> , 2013)
<i>Mamestra configurata</i> NPV-A	MacoNPV-A	NC_003529.1/U59461.2	155,060	41.7	169	(Li <i>et al.</i> , 2002b)
<i>Mamestra configurata</i> NPV-B	MacoNPV-B	NC_004117.1/ AY126275.1	158,482	40.4	168	(Li <i>et al.</i> , 2002a)
<i>Mythimna unipuncta</i> NPV	MyunNPV	NC_043530.1/MF375894.1	148.48	48.6	158	(Harrison <i>et al.</i> , 2018)
<i>Operophtera brumata</i> NPV	OpbuNPV	NC_040621.1/MF614691.1	119.05	39.8	130	(Harrison <i>et al.</i> , 2017b)
<i>Orgyia leucostigma</i> NPV	OrleNPV	NC_010276.1/ EU309041.1	156,179	39.9	135	(Thumbi <i>et al.</i> , 2011)
<i>Spodoptera exigua</i> MNPV	SeMNPV	NC_002169.1/AF169823.1	135,611	43.8	139	(WF <i>et al.</i> , 1999)
<i>Spodoptera frugiperda</i> MNPV	SfMNPV	NC_009011.2/ EF035042.2	131,331	40.2	143	(Harrison <i>et al.</i> , 2008)
<i>Spodoptera littoralis</i> MNPV	SpliMNPV	NC_038369.1/JX454574.1	138	44.7	132	(Breitenbach <i>et al.</i> , 2013)
<i>Spodoptera litura</i> MNPV	SpltMNPV	NC_003102.1/ AF325155.1	139,342	42.7	141	(Pang <i>et al.</i> , 2001)

<i>Perigonia lusca</i> SNPV	PeluSNPV	NC_027923.1/KM596836.1	132,831	39.6	145	(Ardisson-Araujo <i>et al.</i> , 2016)
<i>Sucra jujuba</i> NPV	SujuNPV	NC_028636.1/ KJ676450.1	135.95	38.7	131	(Liu <i>et al.</i> , 2014)
<i>Peridroma sp.</i> NPV	PespNPV	NC_024625.1/KM009991.1	151,110	53.3	139	(Rohrmann <i>et al.</i> , 2015a)
<i>Trichoplusia ni</i> SNPV	TnSNPV	NC_007383.1/DQ017380	134,394	39	145	(Willis <i>et al.</i> , 2005)
<i>Urbanus proteus</i> NPV	UrprNPV	NC_029997.2/KR011717.2	105,555	34.7	119	(Santos <i>et al.</i> , 2018)

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<i>Adoxophyes orona</i> GV	AdorGV	NC_005038.1/ AF547984.1	99,657	34.5	119	(Wormleaton <i>et al.</i> , 2003)
<i>Choristoneura occidentalis</i> GV	ChocGV	NC_008168.1/DQ333351.1	104,710	32.7	116	(Escasa <i>et al.</i> , 2006)
<i>Clostera anachoreta</i> GV	ClanGV	NC_015398.1/ HQ116624.1	101,487	44.4	123	(Liang <i>et al.</i> , 2011)
<i>Clostera anastomosis (L.)</i> GV	ClasGV-A	NC_022646.1/ KC179784.1	101,818	46.7	123	(Liang <i>et al.</i> , 2013)
<i>Clostera anastomosis (L.)</i> GV	ClasGV-B	NC_038371.1/KR091910.1	107,439	37.8	123	(Yin <i>et al.</i> , 2015)
<i>Cnaphalocrocis medinalis</i> GV	CnmeGV	NC_029304.1/KP658210.1	112,060	35.2	133	(Han <i>et al.</i> , 2016)
<i>Cryptophlebia leucotreta</i> GV	CrleGV	NC_005068.1/AY229987.1	110,907	32.4	128	(Lange and Jehle, 2003)

<i>Cydia pomonella</i> GV	CpGV	NC_002816.1/ U53466.2	123,500	45.3	143	(Luque <i>et al.</i> , 2001)
<i>Diatraea saccharalis</i> GV	DisaGV	NC_028491.1/ KP296186.1	98,392	34.9	125	(Ardisson-Araujo <i>et al.</i> , 2015)
<i>Epinotia aporema</i> GV	EpapGV	NC_018875.1/ JN408834.1	119,082	41.5	133	(Ferrelli <i>et al.</i> , 2012)
<i>Erinnyis ello</i> GV	ErelGV	NC_025257.1/ KJ406702.1	102,76	38.7	130	(Ardisson-Araujo <i>et al.</i> , 2014)
<i>Helicoverpa armigera</i> GV	HearGV	NC_010240.1/ EU255577.1	169,794	40.8	179	(Harrison and Popham, 2008)
<i>Mocis latipes</i> GV	MolaGV	NC_029996.1/ KR011718.1	134,27	38.3	145	(Ardisson-Araujo <i>et al.</i> , 2018)
<i>Mythimna unipuncta</i> GV	MyunGV	NC_033780.2/ KX855660.2	144,673	49.9	153	(Harrison <i>et al.</i> , 2017a)
<i>Pieris brassicae</i> GV	PrGV	NC_013797.1/GQ884143.1	108,592	33.2	120	(Zhang <i>et al.</i> , 2012)
<i>Plutella xylostella</i> GV	PxGV	NC_002593.1/ AF270937.1	100,999	40.7	120	(Hashimoto <i>et al.</i> , 2000)
<i>Spodoptera frugiperda</i> GV	SfGV	NC_026511.1/KM371112.1	140,913	46.2	146	(Cuartas <i>et al.</i> , 2015)
<i>Spodoptera litura</i> GV	SpliGV	NC_009503.1/ DQ288858.2	124,121	38.8	136	(Wang <i>et al.</i> , 2011)
<i>Xestia c-nigrum</i> GV	XcGV	NC_002331.1/ AF162221.1	178,733	40.7	181	(Hayakawa <i>et al.</i> , 1999)

***Gammabaculovirus*, 3**

<i>Neodiprion abietis</i> NPV	NeabNPV	NC_008252.1/ DQ317692.1	84,264	33.4	93	(Duffy <i>et al.</i> , 2006)
<i>Neodiprion lecontei</i> NPV	NeleNPV	NC_005906.1/ AY349019.1	81,755	33.4	89	(Lauzon <i>et al.</i> , 2004)
<i>Neodiprion sertifer</i> NPV	NeseNPV	NC_005905.1/ AY430810.1	86,462	33.8	90	(Garcia-Maruniak <i>et al.</i> , 2004)
<i>Deltabaculovirus, 1</i>						
<i>Culex nigripalpus</i> NPV	CuniNPV	NC_003084.1/ AF403738.1	108,252	50.9	109	(Afonso <i>et al.</i> , 2001)

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Table S2. Summary of protein-protein and DNA-protein interactions of baculoviruses

A. Baculoviral protein-protein interactions

Protein	Protein	Method^a	Reference
LEF2	LEF1	Y2H, Co-IP	(Evans <i>et al.</i> , 1997)
POLH	POLH	Y2H	(Braunagel and Summers, 2007)
POLH	P78/83	Y2H	(Braunagel and Summers, 2007)
POLH	BV/ODV-E26	Y2H	(Braunagel and Summers, 2007)
POLH	ODV-E66	Y2H	(Braunagel and Summers, 2007)
POLH	FP25K	Y2H	(Braunagel and Summers, 2007)
POLH	VP39	Y2H	(Braunagel and Summers, 2007)
POLH	AC91	Y2H	(Braunagel and Summers, 2007)
POLH	ODV-E25	Y2H	(Braunagel and Summers, 2007)
POLH	BV/ODV-C42	Y2H	(Braunagel and Summers, 2007)
POLH	PIF3	Y2H	(Braunagel and Summers, 2007)
POLH	ODV-EC27	Y2H	(Braunagel and Summers, 2007)
POLH	PIF5	Y2H	(Braunagel and Summers, 2007)
P78/83	VP39	Y2H	(Braunagel and Summers, 2007)
P78/83	BV/ODV-C42	Y2H, Native PAGE	(Braunagel <i>et al.</i> , 2001)
PK1	PKIP	Y2H, Pull down	(Fan <i>et al.</i> , 1998)
AC11	AC11	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC11	GP41	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC11	AC93	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
BV/ODV-E26	ODV-E66	Y2H, Co-IP	(Braunagel <i>et al.</i> , 1999)
			(Beniya <i>et al.</i> , 1998; Braunagel <i>et al.</i> , 1999; Braunagel and Summers 2007)
BV/ODV-E26	FP25K	Y2H, Co-IP	
BV/ODV-E26	PEP	Y2H	(Zhang <i>et al.</i> , 2018)

BV/ODV-E26	IE1	Y2H, Co-IP, TAP	(Braunagel and Summers, 2007; Nie <i>et al.</i> , 2009)
BV/ODV-E26	IE0	Y2H, Co-IP, TAP	(Nie <i>et al.</i> , 2009)
PKIP	PKIP	Y2H, Co-IP	(Zhang <i>et al.</i> , 2018)
PKIP	FP25K	Y2H, Co-IP	(Zhang <i>et al.</i> , 2018)
PKIP	38K	Y2H	(Zhang <i>et al.</i> , 2018)
PKIP	P6.9	Co-IP	(Lai <i>et al.</i> , 2020)
PKIP	P48	Y2H	(Zhang <i>et al.</i> , 2018)
PKIP	GP64	Y2H, Co-IP	(Zhang <i>et al.</i> , 2018)
PKIP	49K	Y2H	(Zhang <i>et al.</i> , 2018)
PKIP	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
PKIP	ODV-EC27	Y2H, Co-IP	(Zhang <i>et al.</i> , 2018)
V-Ubi	FP25K	Y2H	(Zhang <i>et al.</i> , 2018)
V-Ubi	VP39	Y2H	(Zhang <i>et al.</i> , 2018)
V-Ubi	V-CATH	Y2H	(Zhang <i>et al.</i> , 2018)
V-Ubi	PEP	Y2H	(Zhang <i>et al.</i> , 2018)
V-Ubi	EXON0	Co-IP	(Biswas <i>et al.</i> , 2018)
V-Ubi	49K	Y2H	(Zhang <i>et al.</i> , 2018)
LEF11	LEF11	Co-IP, BiFC, Non-reducing SDS-PAGE	(Dong <i>et al.</i> , 2015; Zhang <i>et al.</i> , 2014)
LEF11	LEF3	BiFC	(Zhang <i>et al.</i> , 2014)
ODV-E66	FP25K	Y2H, Co-IP	(Braunagel <i>et al.</i> , 1999)
ODV-E66	VP39	Y2H, Co-IP	(Braunagel <i>et al.</i> , 1999; Braunagel and Summers, 2007)
ODV-E66	ODV-E25	Y2H, Co-IP	(Braunagel <i>et al.</i> , 1999)
ODV-E66	ODV-EC27	Y2H	(Braunagel and Summers, 2007)
LEF10	LEF10	Y2H, Non-reducing	(Nan <i>et al.</i> , 2019)

SDS-PAGE			
VP1054	38K	Y2H, Co-IP	(Wu <i>et al.</i> , 2008)
VP1054	BV/ODV-C42	Co-IP	(Guan <i>et al.</i> , 2016)
VP1054	VP80	Co-IP	(Guan <i>et al.</i> , 2016)
FGF	V-CATH	Y2H	(Zhang <i>et al.</i> , 2018)
FGF	FP25K	Y2H	(Zhang <i>et al.</i> , 2018)
FP25K	FP25K	Y2H	(Zhang <i>et al.</i> , 2018)
FP25K	GP37	Y2H	(Zhang <i>et al.</i> , 2018)
FP25K	38 K	Y2H	(Zhang <i>et al.</i> , 2018)
FP25K	P6.9	Y2H	(Zhang <i>et al.</i> , 2018)
FP25K	BV/ODV-C42	Y2H	(Zhang <i>et al.</i> , 2018)
FP25K	GP64	Y2H, Co-IP	(Braunagel <i>et al.</i> , 1999; Zhang <i>et al.</i> , 2018)
FP25K	EXON0	Y2H, Co-IP	(Fang <i>et al.</i> , 2007; Fang <i>et al.</i> , 2008)
FP25K	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
FP25K	IE1	Y2H	(Braunagel and Summers, 2007)
GP37	VP39	Y2H	(Zhang <i>et al.</i> , 2018)
GP37	ODV-E25	Y2H	(Zhang <i>et al.</i> , 2018)
GP37	38 K	Y2H	(Zhang <i>et al.</i> , 2018)
GP37	BV/ODV-C42	Y2H	(Zhang <i>et al.</i> , 2018)
AC66	AC66	Y2H	(Peng <i>et al.</i> , 2010)
AC66	EXON0	Co-IP	(Biswas <i>et al.</i> 2018)
LEF3	LEF3	Y2H, Pull down, Cross linking	(Evans and Rohrmann, 1997; Peng <i>et al.</i> , 2010)
LEF3	Helicase	Y2H	(Evans <i>et al.</i> , 1999; Peng <i>et al.</i> , 2010)
LEF3	ALK-EXO	AUC	(Mikhailov <i>et al.</i> , 2003)

AC76	AC76	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC76	AC78	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC76	AC93	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC76	P48	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC78	AC78	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC78	AC93	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
GP41	GP41	Y2H, BiFC, Non-reducing SDS-PAGE	(Li <i>et al.</i> , 2018; Peng <i>et al.</i> , 2010; Yue <i>et al.</i> , 2018)
GP41	38K	Y2H	(Peng <i>et al.</i> , 2010)
GP41	P12	Y2H	(Peng <i>et al.</i> , 2010)
CG30	CG30	Y2H	(Peng <i>et al.</i> , 2009)
VP39	ODV-EC27	Y2H	(Braunagel and Summers, 2007; Zhang <i>et al.</i> , 2018)
VP39	VP39	Y2H	(Braunagel and Summers, 2007)
VP39	38K	Y2H, Co-IP	(Wu <i>et al.</i> , 2008)
VP39	AC91	Y2H	(Braunagel and Summers, 2007)
VP39	P6.9	Y2H	(Zhang <i>et al.</i> , 2018)
VP39	BV/ODV-C42	Y2H	(Braunagel and Summers, 2007)
VP39	PIF3	Y2H	(Braunagel and Summers, 2007)
VP39	GP64	Y2H, Co-IP	(Zhang <i>et al.</i> , 2018)
VP39	ME53	Y2H	(Zhang <i>et al.</i> , 2018)
VP39	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
VP39	PIF5	Y2H	(Braunagel and Summers, 2007)
AC91	IE1	Y2H	(Braunagel and Summers, 2007)
P33	P33	GF, AUC	(Hakim <i>et al.</i> , 2011; Kuang <i>et al.</i> , 2017)
P33	PIF5	Pull down	(Zhang <i>et al.</i> , 2020)

AC93	AC93	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
ODV-E25	38 K	Y2H	(Zhang <i>et al.</i> , 2018)
ODV-E25	ME53	Y2H	(Zhang <i>et al.</i> , 2018)
ODV-E25	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
38K	PIF2	Y2H	(Peng <i>et al.</i> , 2010)
38K	PP31	Y2H	(Zhang <i>et al.</i> , 2018)
38K	38K	Y2H, Co-IP	(Peng <i>et al.</i> , 2010; Wu <i>et al.</i> , 2008; Zhang <i>et al.</i> , 2018)
38K	BV/ODV-C42	Y2H	(Zhang <i>et al.</i> , 2018)
38K	P48	Y2H	(Zhang <i>et al.</i> , 2018)
38K	VP80	Y2H, Co-IP	(Wu <i>et al.</i> , 2008)
38K	PIF3	Y2H	(Peng <i>et al.</i> , 2010)
38K	V-CATH	Y2H	(Zhang <i>et al.</i> , 2018)
38K	GP64	Y2H, Co-IP	(Zhang <i>et al.</i> , 2018)
38K	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
38K	ODV-EC27	Y2H, Co-IP	(Zhang <i>et al.</i> , 2018)
38K	PIF5	Y2H	(Peng <i>et al.</i> , 2010)
P6.9	V-CATH	Y2H	(Zhang <i>et al.</i> , 2018)
P6.9	AC132	Co-IP	(Fang <i>et al.</i> , 2016)
BV/ODV-C42	EXON0	Y2H, Co-IP	(Fang <i>et al.</i> , 2007; Fang <i>et al.</i> , 2008)
BV/ODV-C42	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
BV/ODV-C42	ODV-EC27	Y2H, Native PAGE	(Braunagel <i>et al.</i> , 2001)
BV/ODV-C42	PIF5	Y2H	(Braunagel and Summers, 2007)
P48	P48	Y2H	(Zhang <i>et al.</i> , 2018)
P48	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
P48	AC146	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
VP80	HA100	Y2H, Co-IP	(Peng <i>et al.</i> , 2010)

ODV-EC43	49K	Y2H	(Peng <i>et al.</i> , 2010)
V-ChiA	V-CATH	Pull down, BiFC	(Hodgson <i>et al.</i> , 2011)
V-CATH	ME53	Y2H	(Zhang <i>et al.</i> , 2018)
V-CATH	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
V-CATH	ODV-EC27	Y2H	(Zhang <i>et al.</i> , 2018)
GP64	GP64	Y2H	(Zhang <i>et al.</i> , 2018)
GP64	PEP	Y2H	(Zhang <i>et al.</i> , 2018)
GP64	ODV-E18	Co-IP	(Zhang <i>et al.</i> , 2018)
P24	P24	Y2H	(Peng <i>et al.</i> , 2010)
AC132	ODV-E18	Co-IP	(Fang <i>et al.</i> , 2016)
ALK-EXO	ALK-EXO	GF	(Mikhailov <i>et al.</i> , 2003)
P26	P26	Y2H	(Goenka and Weaver, 2008)
ME53	ODV-E18	Y2H	(Zhang <i>et al.</i> , 2018)
ME53	ODV-EC27	Y2H	(Zhang <i>et al.</i> , 2018)
49K	ODV-EC27	Y2H	(Zhang <i>et al.</i> , 2018)
EC27	ODV-EC27	Y2H, Co-IP	(Braunagel and Summers, 2007; Zhang <i>et al.</i> , 2018)
AC146	AC146	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
ODV-E66	PIF2	Y2H	(Peng <i>et al.</i> , 2010)
ODV-E66	PIF3	Y2H	(Peng <i>et al.</i> , 2010)
ODV-E66	PIF4	Co-IP, BiFC	(Dong <i>et al.</i> , 2014)
ODV-E66	PIF5	Y2H	(Peng <i>et al.</i> , 2010)
P74	PIF5	Y2H	(Peng <i>et al.</i> , 2010)
PIF1	PIF2	Y2H, BiFC	(Zheng <i>et al.</i> , 2017)
PIF1	PIF3	Y2H, BiFC	(Zheng <i>et al.</i> , 2017)
PIF1	PIF4	Y2H, Co-IP, BiFC	(Dong <i>et al.</i> , 2014; Zheng <i>et al.</i> , 2017)
PIF1	PIF5	Y2H	(Peng <i>et al.</i> , 2010)

PIF1	VP91	Y2H, BiFC	(Zheng <i>et al.</i> , 2017)
PIF2	PIF3	Y2H, BiFC	(Peng <i>et al.</i> , 2010; Zheng <i>et al.</i> , 2017)
PIF2	PIF4	Co-IP, BiFC	(Dong <i>et al.</i> , 2014)
PIF2	PIF5	Y2H	(Peng <i>et al.</i> , 2010)
PIF3	PIF3	Y2H	(Peng <i>et al.</i> , 2010)
PIF3	PIF4	Y2H, Co-IP, BiFC	(Dong <i>et al.</i> , 2014; Zheng <i>et al.</i> , 2017)
PIF3	PIF5	Y2H	(Peng <i>et al.</i> , 2010)
PIF5	PIF5	Y2H	(Peng <i>et al.</i> , 2010)
PIF5	HA107	Y2H	(Peng <i>et al.</i> , 2010)
49K	HA122	Y2H	(Peng <i>et al.</i> , 2010)
HA44	HA44	Y2H, Co-IP	(Peng <i>et al.</i> , 2010)

B. Baculoviral DNA-protein interactions

Protein	Method ^a	Reference
BRO	DNA-C	(Zemskov <i>et al.</i> , 2000)
DBP	EMSA	(Mikhailov <i>et al.</i> , 1998; Mikhailov <i>et al.</i> , 2008)
PP31	Southwestern blot	(Guarino <i>et al.</i> , 1992)
AC59 (HA51)	DNA-C	(Li <i>et al.</i> , 2006a; Li <i>et al.</i> , 2006b)
AC60 (HA52)	DNA-C	(Li <i>et al.</i> , 2006a; Li <i>et al.</i> , 2006b)
DNA POL	DNA-C	(Hang and Guarino, 1999)
LEF3	DNA-C/ChIP	(Hang <i>et al.</i> , 1995; Ito <i>et al.</i> , 2004)
VLF-1	EMSA	(Mikhailov and Rohrmann, 2002)
Helicase	EMSA/ChIP	(Ito <i>et al.</i> , 2004; McDougal and Guarino, 2000)
P6.9	EM	(Tweeten <i>et al.</i> , 1980)
ALK-EXO	DNA digestion	(Mikhailov <i>et al.</i> , 2003)
IE1	ChIP	(Ito <i>et al.</i> , 2004)

C. Baculovirus-host protein-protein interactions

Viral Protein	Host Protein	Method ^a	Reference
BRO	Laminin	Y2H, Pull down	(Kang <i>et al.</i> , 2003)
LEF2	TER94	BiFC	(Li <i>et al.</i> , 2020)
P78/83	Actin	Actin binding	(Lanier and Volkman, 1998)
P78/83	Arp2/3	Actin polymerization	(Goley <i>et al.</i> , 2006)
AC11	VPS2B	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC11	VPS20	BiFC	(Yue <i>et al.</i> , 2018)
AC11	VPS24	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC11	VPS46	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC11	VPS60	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
LEF1	TER94	BiFC	(Li <i>et al.</i> , 2020)
AC34	P40	Co-IP	(Mu <i>et al.</i> , 2016)
AC34	P34	Co-IP	(Mu <i>et al.</i> , 2016)
AC34	P20	Co-IP	(Mu <i>et al.</i> , 2016)
LEF11	ATAD3A	Co-IP	(Dong <i>et al.</i> , 2017)
LEF11	HSP60	Co-IP	(Dong <i>et al.</i> , 2017)
LEF11	Importin α -3	Co-IP	(Zhang <i>et al.</i> , 2014)
DNA POL	TER94	BiFC	(Li <i>et al.</i> , 2020)
LEF3	TER94	Co-IP, Cross-linking, BiFC	(Li <i>et al.</i> , 2020)
AC76	NSF	Co-IP, BiFC	(Guo <i>et al.</i> , 2017)
AC76	VPS2B	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC76	VPS20	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC76	VPS24	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC76	SNF7	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC76	VPS46	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC76	VPS60	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)

AC78	NSF	Co-IP, BiFC	(Guo <i>et al.</i> , 2017)
AC78	VPS2B	BiFC	(Yue <i>et al.</i> , 2018)
AC78	VPS20	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC78	VPS24	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC78	SNF7	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC78	VPS46	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC78	VPS60	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
GP41	NSF	Co-IP, BiFC	(Guo <i>et al.</i> , 2017)
GP41	VPS2B	Co-IP	(Yue <i>et al.</i> , 2018)
GP41	VPS20	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
GP41	VPS24	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
GP41	SNF7	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
GP41	VPS46	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
GP41	VPS60	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
GP41	VPS4	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
VP39	Actin	Actin overlay	(Lanier and Volkman, 1998)
P33	P53	Co-IP	(Prikhod'ko <i>et al.</i> , 1999; Wu <i>et al.</i> , 2013)
AC93	NSF	Co-IP, BiFC	(Guo <i>et al.</i> , 2017)
AC93	SNF7	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC93	VPS4	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC93	VPS2B	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC93	VPS20	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC93	VPS24	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC93	VPS46	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC93	VPS60	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
Helicase	TER94	Co-IP, Cross-linking, BiFC	(Li <i>et al.</i> , 2020)
P48	NSF	Co-IP, BiFC	(Guo <i>et al.</i> , 2017)

P48	VPS24	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
P48	VPS4	BiFC	(Yue <i>et al.</i> , 2018)
VP80	Actin	Co-IP	(Marek <i>et al.</i> , 2011)
PK2	eIF2 α kinase	Y2H	(Li <i>et al.</i> , 2015)
LEF7	SKP1	Co-IP	(Mitchell <i>et al.</i> , 2013)
GP64	NPC1	Co-IP	(Li <i>et al.</i> , 2019)
GP64	SINAL10	Y2H, Co-IP	(Feng <i>et al.</i> , 2018)
P10	Tubulin	Y2H, Pull down	(Patmanidi <i>et al.</i> , 2003)
EXON0	Tubulin	Co-IP, TAP	(Fang <i>et al.</i> , 2009)
49K	VPS20	BiFC	(Yue <i>et al.</i> , 2018)
49K	VPS24	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
49K	VPS46	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
ODV-EC27	Cdc2	Co-IP	(Belyavskiy <i>et al.</i> , 1998)
ODV-EC27	Cdk6	Co-IP	(Belyavskiy <i>et al.</i> , 1998)
AC146	SNF7	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC146	VPS20	BiFC	(Yue <i>et al.</i> , 2018)
AC146	VPS24	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC146	VPS46	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
AC146	VPS60	Co-IP, BiFC	(Yue <i>et al.</i> , 2018)
IE1	TER94	BiFC	(Li <i>et al.</i> , 2020)

Note:

a: AUC: analytical ultracentrifugation; BiFC: bimolecular fluorescent complimentary; ChIP: chromatin immunoprecipitation; Co-IP: co-immunoprecipitation; DNA-C: DNA-cellulose chromatography; EM: Electron microscopy; EMSA: Electrophoretic mobility shift assay; GF: Gel filtration chromatography; TAP: tandem affinity purification; Y2H: yeast two-hybrid; Native PAGE: native polyacrylamide gel electrophoresis; Non-reducing SDS-PAGE: non-reducing sodium dodecyl sulfate polyacrylamide gel electrophoresis.

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No.	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	
ID of protein homologue	AC74	AC114	AC120	AC124	AC130 GP16	AC132	AC4	AC5	AC11	AC26	AC30	AC42 GTA	AC43	AC44	AC47	AC48	
Distribution [#]	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	
<i>Alphabaculovirus</i>	AcMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	
	AgMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1
	BmNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0
	BomaNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0
	CapoNPV	1	1	1	1	1	1	1	1	1	1	0	0	1	1	1	1
	CoveNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1
	CfDEFMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1
	CfMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1
	ChmuNPV	1	1	1	1	1	1	1	0	1	0	1	1	0	1	0	1
	ChroNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1
	DapuNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
	EppoNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1
	HycuNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	0	1	1
	LoobMNPV	1	1	1	1	1	1	0	1	0	1	1	0	1	1	0	0
	MaviNPV	1	1	1	1	1	1	1	1	1	1	1	0	1	1	1	0
	OxocNPV	1	1	1	1	1	1	0	1	1	1	1	1	1	1	1	0
	PlyxMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
	ThorMNPV	1	1	1	1	1	1	1	1	1	1	1	1	0	1	1	0
	AdhoNPV	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	0	0	1	0	1	0	2	1	0	1	0	0	1	0	0	0
	AgseNPV-A	0	0	1	0	1	0	1	1	0	1	0	0	1	0	0	0
	AgseNPV-B	0	0	1	0	1	0	1	1	0	1	0	0	1	0	0	0
	AnpeNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
	BusuNPV	0	0	1	0	1	0	0	1	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	0	1	0	0	0	0	1	0	0	1	0	0	0
	ChinNPV	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0
	EcobNPV	0	0	1	0	1	0	0	1	0	0	0	0	1	0	0	0
	EupsNPV	0	0	1	0	1	0	0	1	1	0	0	0	1	0	0	0
	HearNPV-C1	0	0	1	0	1	0	0	1	0	1	0	0	1	0	0	0
	HearNPV-G4	0	0	1	0	1	0	0	0	0	1	0	0	1	0	0	0
	HespNPV	0	0	1	0	1	0	0	1	0	0	0	0	0	0	0	0
	LafiNPV	0	0	0	0	1	0	0	1	1	0	0	0	0	0	0	0
	LdMNPV	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0
	LeseNPV	0	0	1	0	0	0	0	1	0	1	0	0	0	0	0	0
	LyxyMNPV	0	0	0	0	0	0	0	1	1	1	0	0	0	0	0	0
	MabrMNPV	0	0	1	0	1	0	2	1	0	1	0	0	0	0	0	0
	MacoNPV-A	0	0	1	0	1	0	2	1	0	1	0	0	1	0	0	0
	MacoNPV-B	0	0	1	0	1	0	2	1	0	1	0	0	1	0	0	0
	ManeNPV	0	0	1	0	1	0	0	1	0	0	0	0	1	0	0	0
	MyunNPV	0	0	1	0	1	0	1	1	0	1	0	0	1	0	0	0
	OpMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
	OpbuNPV	0	0	0	0	0	0	0	1	0	1	0	0	1	0	0	0
	OrleNPV	0	0	0	0	1	0	0	1	1	0	0	0	1	0	0	0
	PelusNPV	0	0	1	0	1	0	0	1	1	0	0	0	1	0	0	0
PespNPV	0	0	1	0	0	0	1	0	0	1	0	0	0	0	0	0	
SeMNPV	0	0	1	0	1	0	1	1	0	1	0	0	1	1	0	0	
SfMNPV	0	0	1	0	1	0	1	1	0	1	0	0	1	0	0	0	
SpliMNPV	0	0	1	0	0	0	0	1	0	1	0	0	0	0	0	0	
SpltMNPV	0	0	1	0	0	0	0	1	0	1	0	0	0	0	0	0	
SujuNPV	0	0	1	0	0	0	1	1	0	0	0	0	0	0	0	0	
TnSNPV	0	0	0	0	1	0	0	0	0	1	0	0	1	0	0	0	
UrprNPV	0	0	1	0	1	0	0	1	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeseNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
No. of homologues	20	20	45	20	45	20	33	44	25	40	19	18	36	20	12	14	
Strain occupancy	20	20	45	20	45	20	29	44	25	40	19	18	36	20	12	14	

No.	97	98	99	100	101	102	103	104	105	106	107	108	109	110	111	112	
ID of protein homologue	AC49 PCNA	AC52	AC56	AC57	AC69	AC72	AC73	AC91	AC117	AC122	AC136 P26	AC151 IE2	AC152	AC7 ORF603	AC45	AC70 HCF-1	
Distribution [#]	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	
<i>Alphabaculovirus</i>	AcMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	
	AgMNPV	0	0	1	1	1	1	1	1	2	1	1	0	0	0	0	
	BmNPV	0	1	1	1	1	1	1	1	1	1	1	0	0	1	0	
	BomaNPV	0	1	1	1	1	1	1	1	1	1	1	0	0	1	0	
	CapoNPV	0	1	1	1	0	1	1	1	1	1	0	1	1	0	1	0
	CoveNPV	0	0	1	1	1	1	1	1	1	1	1	1	0	0	0	0
	CfDEFMNPV	0	0	1	1	1	1	1	1	1	1	1	1	0	0	0	0
	CfMNPV	1	0	1	1	1	1	1	1	1	1	2	1	0	0	0	0
	ChmuNPV	1	0	0	1	1	1	1	1	1	1	1	1	0	0	0	0
	ChroNPV	1	0	1	1	1	1	1	1	1	1	2	1	0	0	0	0
	DapuNPV	1	0	1	1	0	1	1	1	1	1	1	1	0	0	0	0
	EppoNPV	0	0	1	1	1	1	1	1	2	0	1	1	0	0	0	0
	HycuNPV	1	0	1	1	1	1	1	1	1	1	1	1	0	0	0	0
	LoobMNPV	1	1	1	1	0	0	0	0	0	1	1	0	1	0	0	0
	MaviNPV	0	1	1	0	1	1	1	0	1	0	1	1	1	0	1	0
	OxocNPV	0	1	0	1	1	1	1	0	1	0	1	0	0	0	1	0
	PlyxMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
	ThorMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	0	0	1
	AdhoNPV	0	1	1	0	1	0	0	0	0	0	1	0	0	0	0	0
	AdorNPV	0	1	1	0	1	0	0	0	0	0	1	0	0	0	0	0
	AgipMNPV	0	1	1	1	1	0	0	0	1	0	2	0	2	0	0	0
	AgseNPV-A	0	1	1	1	1	0	0	0	1	0	2	0	0	0	0	0
	AgseNPV-B	0	1	1	1	1	0	0	0	1	0	2	0	1	0	0	0
	AnpeNPV	1	0	0	1	1	1	1	1	1	1	1	1	0	0	0	0
	BusuNPV	0	1	0	1	0	0	0	0	1	0	2	0	0	0	0	0
	ChchNPV	1	1	1	1	0	0	0	0	0	0	2	0	0	0	0	0
	ChinNPV	1	1	1	1	0	0	0	0	0	0	2	0	1	0	0	0
	EcobNPV	0	1	1	1	0	0	0	0	0	0	2	0	0	0	0	0
	EupsNPV	0	1	0	1	0	0	0	0	1	0	2	0	0	0	0	0
	HearNPV-C1	0	1	1	1	1	0	0	0	1	0	1	0	1	0	0	0
	HearNPV-G4	0	1	1	1	1	0	0	0	1	0	1	0	0	0	0	0
	HespNPV	0	1	0	1	0	0	0	0	1	0	2	0	0	0	0	0
	LafiNPV	0	1	1	1	0	0	0	0	1	0	3	0	0	0	0	0
	LdMNPV	0	1	0	1	0	0	0	0	0	0	1	0	0	0	0	0
	LeseNPV	0	1	1	1	0	0	0	0	0	0	1	0	0	0	0	0
	LyxyMNPV	0	2	0	1	0	0	0	0	0	0	1	0	0	0	0	0
	MabrMNPV	0	1	1	1	1	0	0	0	1	0	2	0	0	0	0	0
	MacoNPV-A	0	1	1	1	1	0	0	0	1	0	2	0	1	0	0	0
	MacoNPV-B	0	1	1	1	1	0	0	0	1	0	2	0	0	0	0	0
	ManeNPV	0	1	0	1	0	0	0	0	0	0	2	0	0	0	0	0
	MyunNPV	0	1	1	1	1	0	0	0	1	0	2	0	0	0	0	0
	OpMNPV	1	0	1	1	0	1	1	1	1	1	1	1	0	0	0	0
	OpbuNPV	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0
	OrleNPV	0	1	1	1	0	0	0	0	1	0	2	0	0	0	0	0
	PelusNPV	0	0	1	1	1	0	0	0	1	0	2	0	0	0	0	0
PespNPV	0	1	1	1	1	0	0	0	1	0	2	0	0	0	0	0	
SeMNPV	0	1	1	1	1	0	0	0	1	0	2	0	0	0	0	0	
SfMNPV	0	1	1	1	1	0	0	0	1	0	2	0	0	0	0	0	
SpliMNPV	0	1	0	1	1	0	0	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	1	0	1	1	0	0	0	0	0	0	0	1	0	0	0	
SujuNPV	0	1	1	1	0	0	0	0	1	0	2	0	0	0	0	0	
TnSNPV	1	1	1	1	0	0	0	0	0	0	2	0	0	0	0	0	
UrprNPV	0	1	0	1	0	0	0	0	0	0	2	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
NeleNPV		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
NeseNPV		0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
No. of homologues		14	42	40	50	34	19	19	17	39	18	75	18	13	3	7	3
Strain occupancy		14	41	40	50	34	19	19	17	38	17	49	18	12	3	7	3

No.	113	114	115	116	117	118	119	120	121	122	123	124	125	126	127	128	
ID of protein homologue	AC85	AC116	AC118	AC121	AC123 PK2	AC149	AC154	AC97	AC140	AC32 FGF	AC58/AC59/AC60 ChaBs	AC88 CG30/AC153 PE38	AC1 PTP	AC128 GP64	AC2 BRO	AC12	
Distribution [#]	(α -)	(α -)	(α -)	(α -)	(α -)	(α -)	(α -)	Species-unique	Species-unique	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	
<i>Alphabaculovirus</i>	AcMNPV	1	1	1	1	1	1	1	1	1	3	2	1	1	1	1	
	AgMNPV	0	0	0	0	0	0	0	0	1	2	2	2	1	7	0	
	BmNPV	0	1	0	1	1	1	1	0	0	1	2	1	1	5	0	
	BomaNPV	0	1	0	1	1	1	1	0	0	1	2	1	1	3	0	
	CapoNPV	0	0	0	0	0	1	0	0	0	0	2	2	1	1	1	
	CoveNPV	0	0	0	0	0	0	0	0	0	1	2	2	2	1	9	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	1	2	2	2	1	4	0
	CfMNPV	0	0	0	0	0	0	0	0	0	1	2	2	2	1	3	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	1	2	2	2	1	1	0
	ChroNPV	0	0	0	0	0	0	0	0	0	1	2	2	2	1	2	0
	DapuNPV	0	0	0	0	0	0	0	0	0	1	2	2	2	1	3	1
	EppoNPV	0	0	0	0	0	0	0	0	0	1	2	2	1	1	1	0
	HycuNPV	0	0	0	0	0	0	0	0	0	1	2	2	1	1	5	0
	LoobMNPV	0	0	0	0	0	0	0	0	0	1	2	3	1	1	1	1
	MaviNPV	0	0	0	0	0	1	1	0	0	0	2	1	1	1	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	1	2	1	1	1	0	0
	PlyxMNPV	1	1	1	1	1	1	1	0	0	1	2	2	1	1	2	1
	ThorMNPV	1	0	1	0	1	1	1	0	0	1	2	2	1	1	3	1
	AdhoNPV	0	0	0	0	0	0	0	0	0	1	2	0	0	0	4	0
	AdorNPV	0	0	0	0	0	0	0	0	0	1	2	0	0	0	3	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	1	2	1	1	0	5	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	1	2	1	1	0	4	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	1	2	1	1	0	2	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	1	2	2	2	1	2	0
	BusuNPV	0	0	0	0	0	0	0	0	0	1	3	0	0	0	3	0
	ChchNPV	0	0	0	0	0	0	0	0	0	1	2	0	1	0	4	0
	ChinNPV	0	0	0	0	0	0	0	0	0	1	2	0	1	0	2	0
	EcobNPV	0	0	0	0	0	0	0	0	0	2	2	1	0	0	2	0
	EupsNPV	0	0	0	0	0	0	0	0	0	1	2	1	0	0	2	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	1	2	1	0	0	3	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	1	2	1	0	0	3	0
	HespNPV	0	0	0	0	0	0	0	0	0	1	3	1	0	0	2	0
	LafiNPV	0	0	0	0	0	0	0	0	0	1	2	0	0	0	0	0
	LdMNPV	0	0	0	0	0	0	0	0	0	1	2	0	0	0	16	2
	LeseNPV	0	0	0	0	0	0	0	0	0	1	1	1	0	0	10	0
LyxyMNPV	0	0	0	0	0	0	0	0	0	1	2	0	0	0	14	2	
MabrMNPV	0	0	0	0	0	0	0	0	0	1	2	1	1	0	6	0	
MacoNPV-A	0	0	0	0	0	0	0	0	0	1	2	1	1	0	8	0	
MacoNPV-B	0	0	0	0	0	0	0	0	0	1	2	1	1	0	8	0	
ManeNPV	0	0	0	0	0	0	0	0	0	1	2	1	0	0	2	1	
MyunNPV	0	0	0	0	0	0	0	0	0	1	2	1	1	0	7	0	
OpMNPV	0	0	0	0	0	0	0	0	0	1	2	2	2	1	2	1	
OpbuNPV	0	0	0	0	0	0	0	0	0	1	2	0	0	0	2	0	
OrleNPV	0	0	0	0	0	0	0	0	0	1	3	1	0	0	5	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	1	2	1	0	0	1	0	
PespNPV	0	0	0	0	0	0	0	0	0	1	1	1	0	0	6	0	
SeMNPV	0	0	0	0	0	0	0	0	0	1	2	1	1	0	0	0	
SfMNPV	0	0	0	0	0	0	0	0	0	1	2	1	1	0	1	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	1	2	1	0	0	1	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	1	2	1	0	0	2	0	
SujuNPV	0	0	0	0	0	0	0	0	0	1	2	2	0	0	4	1	
TnSNPV	0	0	0	0	0	0	0	0	0	1	2	0	1	0	1	0	
UrprNPV	0	0	0	0	0	0	0	0	0	1	0	1	0	0	7	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	3	0	0	0	0	0	0	
	ChocGV	0	0	0	0	0	0	0	0	3	0	0	0	0	0	0	
	ClanGV	0	0	0	0	0	0	0	0	3	0	0	0	0	0	1	
	ClasGV-A	0	0	0	0	0	0	0	0	3	0	0	0	0	0	1	
	ClasGV-B	0	0	0	0	0	0	0	0	3	0	0	0	0	0	1	
	CnmeGV	0	0	0	0	0	0	0	0	3	0	0	0	0	3	0	
	CpGV	0	0	0	0	0	0	0	0	3	0	0	1	2	0	0	0
	CrleGV	0	0	0	0	0	0	0	0	3	0	0	2	1	0	0	0
	DisaGV	0	0	0	0	0	0	0	0	3	0	0	0	0	1	0	1
	EpapGV	0	0	0	0	0	0	0	0	3	1	0	0	0	0	0	0
	ErelGV	0	0	0	0	0	0	0	0	3	0	0	0	0	0	0	1
	PrGV	0	0	0	0	0	0	0	0	3	0	0	1	0	0	0	0
	HearGV	0	0	0	0	0	0	0	0	3	1	0	0	0	0	11	0
	MolaGV	0	0	0	0	0	0	0	0	3	1	0	0	0	0	2	0
	MyunGV	0	0	0	0	0	0	0	0	3	1	0	0	0	0	5	0
	PxGV	0	0	0	0	0	0	0	0	3	0	1	0	0	0	0	0
	SfGV	0	0	0	0	0	0	0	0	3	1	0	0	0	0	6	0
SpliGV	0	0	0	0	0	0	0	0	3	1	2	0	0	0	6	1	
XcGV	0	0	0	0	0	0	0	0	3	1	0	0	0	0	9	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0	
No. of homologues	3	4	3	4	5	7	6	1	1	106	113	70	44	21	243	19	
Strain occupancy	3	4	3	4	5	7	6	1	1	69	59	48	34	21	57	17	

No.	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143	144	
ID of protein homologue	AC31 SOD	AC33 HisP	AC46 ODV-E66	AC64 GP37	AC79	AC105 HE65	AC111	AC125 LEF7	AC126 V-Chi	AC127 V-CATH	AC39 P43	AC63	AC135 P35	AC3 CTX	AC84	AC86 PNK/PNL	
Distribution [#]	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	
<i>Alphabaculovirus</i>	AcMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	
	AgMNPV	1	1	1	0	1	1	1	1	0	0	0	0	1	0	1	
	BmNPV	1	0	1	1	1	1	1	1	1	1	1	1	0	0	0	
	BomaNPV	1	0	1	1	1	1	1	1	1	1	1	1	1	0	0	1
	CapoNPV	1	1	0	1	1	0	1	0	1	1	0	0	0	0	0	0
	CoveNPV	1	0	0	0	1	0	1	1	0	0	0	0	0	1	0	0
	CfDEFMNPV	1	1	1	1	1	2	1	1	1	1	0	0	0	0	0	0
	CfMNPV	1	0	1	1	1	0	1	3	1	1	0	0	0	1	0	0
	ChmuNPV	1	0	1	1	1	0	1	3	1	1	0	0	0	2	0	0
	ChroNPV	1	0	1	1	1	0	1	7	1	1	0	0	0	2	0	0
	DapuNPV	1	1	1	1	1	1	0	4	1	1	0	0	0	2	0	0
	EppoNPV	0	0	1	1	1	0	1	1	1	1	0	0	0	0	0	0
	HycuNPV	1	0	1	1	1	0	1	1	1	1	0	0	0	2	0	0
	LoobMNPV	1	0	1	0	0	1	1	1	0	0	0	1	0	2	0	0
	MaviNPV	1	0	0	1	1	1	1	1	1	1	1	1	1	0	0	0
	OxocNPV	1	0	1	1	1	0	1	0	1	1	0	0	0	1	0	0
	PlyxMNPV	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0
	ThorMNPV	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
	AdhoNPV	1	0	1	0	1	0	0	0	0	1	1	0	0	1	0	0
	AdorNPV	1	0	1	0	1	0	0	0	0	1	1	0	0	1	0	0
	AgipMNPV	1	1	2	1	1	1	0	0	1	1	0	0	0	0	0	0
	AgseNPV-A	1	1	1	1	0	1	0	1	1	1	0	0	0	0	1	0
	AgseNPV-B	1	1	2	1	1	1	0	0	1	1	0	0	0	0	0	0
	AnpeNPV	1	0	1	1	1	1	1	1	1	1	0	0	0	2	0	1
	BusuNPV	1	1	0	1	1	0	1	0	1	1	1	0	0	1	0	0
	ChchNPV	1	1	1	1	0	1	1	0	1	1	0	1	0	1	1	0
	ChinNPV	1	1	1	1	0	1	1	0	1	1	0	1	0	1	2	0
	EcobNPV	1	1	2	1	1	0	0	0	1	1	1	1	0	1	0	0
	EupsNPV	1	1	1	1	0	0	1	0	1	1	1	0	0	0	0	0
	HearNPV-CI	1	0	1	1	0	1	1	0	1	1	0	1	0	0	0	0
	HearNPV-G4	1	0	1	1	0	1	1	0	1	1	0	1	0	0	0	0
	HespNPV	1	1	1	1	0	0	0	0	1	1	0	0	0	1	0	0
	LafiNPV	1	1	1	1	0	0	0	0	1	1	0	1	0	0	0	0
	LdMNPV	1	1	1	1	0	0	1	0	1	1	0	1	0	2	0	0
	LeseNPV	1	1	1	1*	1	2	1	0	1	1	0	0	1	0	0	0
	LyxyMNPV	1	1	1	1	0	0	1	0	1	1	0	1	0	1	0	0
	MabrMNPV	1	1	2	1	1	1	0	0	1	1	0	0	0	1	0	0
	MacoNPV-A	1	1	2	1	1	1	0	1	1	1	0	0	0	1	0	0
	MacoNPV-B	1	1	2	1	1	1	0	0	1	1	0	0	0	1	0	0
	ManeNPV	1	2	1	1	0	0	0	0	1	1	0	0	0	0	0	0
	MyunNPV	1	1	2	1	0	0	0	2	2	1	0	0	0	0	0	0
	OpMNPV	1	1	1	1	1	0	1	1	1	1	0	0	0	2	0	0
	OpbuNPV	1	2	1	0	0	0	0	0	0	0	0	0	0	1	0	0
	OrleNPV	1	1	2	1	1	0	0	0	1	1	0	0	0	2	0	0
PelusNPV	1	1	1	1	0	1	1	1	1	1	0	0	0	2	0	0	
PespNPV	1	1	1	1	1	0	0	0	1	1	0	0	0	0	1	0	
SeMNPV	1	1	2	1	0	0	0	2	1	1	0	0	0	0	0	0	
SfMNPV	1	1	2	1	0	0	0	1	1	1	0	0	0	0	0	0	
SpliMNPV	0	1	1	1*	0	0	0	0	1	1	0	0	1	0	0	0	
SpltMNPV	0	1	1	1*	0	0	0	0	1	1	0	0	1	0	0	0	
SujuNPV	1	1	0	1	1	0	1	0	1	1	1	0	0	0	0	0	
TnSNPV	1	1	1	1	0	0	1	1	1	1	0	1	0	0	1	0	
UrprNPV	1	0	1	0	0	0	1	0	0	0	0	0	0	1	0	0	
<i>Betabaculovirus</i>	AdorGV	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	
	ChocGV	1	0	1	0	0	0	0	0	0	0	0	0	1	0	0	
	ClanGV	1	0	0	1	0	0	1	0	1	1	0	0	1	0	0	
	ClasGV-A	1	0	0	1	0	0	1	0	1	1	0	0	1	0	0	
	ClasGV-B	1	1	0	0	0	1	0	1	0	0	1	0	0	0	0	
	CnmeGV	1	0	1	0	1	0	0	0	0	0	0	0	0	0	0	
	CpGV	1	1	1	1	1	0	0	0	1	1	0	0	0	0	0	
	CrleGV	1	1	1	0	0	0	0	0	0	1	0	0	0	0	0	
	DisaGV	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	
	EpapGV	1	1	1	1	0	0	0	0	1	1	0	1	0	0	0	
	ErelGV	1	1	0	0	0	1	0	0	0	0	1	0	0	0	0	
	PrGV	1	0	2	1	0	0	0	0	1	1	0	0	0	0	0	
	HearGV	1	0	1	1	1	1	0	1	1	0	0	0	0	1	1	0
	MolaGV	1	0	1	0	1	0	1	0	0	0	0	0	0	0	0	0
	MyunGV	1	1	1	0	1	0	1	0	0	0	0	0	0	0	0	0
	PxGV	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	SfGV	1	0	1	0	1	0	0	2	0	0	0	0	0	0	0	0
	SpliGV	0	0	1	0	0	0	1	0	0	0	0	0	0	0	0	1
XcGV	1	0	1	1	1	1	1	1	1	1	0	0	0	1	0	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
No. of homologues	66	45	74	53	39	29	38	45	54	55	14	17	12	42	10	6	
Strain occupancy	66	43	63	53	39	27	38	29	53	55	14	17	12	32	9	6	

No.	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160	
ID of protein homologue	AC112/AC113	AC134 94K	AC87 P15	CpGV ORF4	CpGV ORF33	CpGV ORF39	CpGV ORF46 MP-nase	CpGV ORF99	CpGV ORF135	CpGV ORF82	Neab5	Neab14	Neab17	Neab23	Neab24	Neab25	
Distribution [#]	(α , β -)	(α , β -)	(α , δ)	(β)	(β)	(β)	(β)	(β)	(β)	(β)	(γ)	(γ)	(γ)	(γ)	(γ)	(γ)	
<i>Alphabaculovirus</i>	AcMNPV	2	1	1	0	0	0	0	0	0	0	0	0	0	0	0	
	AgMNPV	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	
	BmNPV	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	
	BomaNPV	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PixyMNPV	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-B	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AnpeNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChinNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EcobNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EupsNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-CI	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HespNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LafiNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LdMNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
LeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
LyxyMNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	
MabrMNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-A	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-B	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ManeNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MyunNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpMNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OrleNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PelusNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SeMNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SujuNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
TnSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	ChocGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	ClanGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	ClasGV-A	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	ClasGV-B	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	CnmeGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	CpGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	CrleGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	DisaGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	EpapGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	ErelGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	PrGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	HearGV	1	1	0	1	1	1	1	1	1	0	0	0	0	0	0	
	MolaGV	1	1	0	1	1	1	1	1	1	0	0	0	0	0	0	
	MyunGV	1	0	0	2	1	1	1	1	1	0	0	0	0	0	0	
	PxGV	0	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
	SfGV	1	0	0	1	1	1	1	1	1	0	0	0	0	0	0	
SpliGV	1	0	0	1	1	1	1	1	1	0	0	0	0	0	0		
XcGV	1	1	0	1	1	1	1	1	1	0	0	0	0	0	0		
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	1	1	1	1	1	1	
	NeleNPV	0	0	0	0	0	0	0	0	0	1	1	1	1	1	1	
	NeseNPV	0	0	0	0	0	0	0	0	0	1	1	1	1	5	1	
<i>Deltabaculovirus</i>	CuniNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	
No. of homologues		23	16	15	20	19	19	19	19	19	3	3	3	3	9	3	
Strain occupancy		22	16	15	19	19	19	19	19	19	3	3	3	3	5	3	

No.	177	178	179	180	181	182	183	184	185	186	187	188	189	190	191	192
ID of protein homologue	Neab90	Neab93	Nele70	Neab2/Neab29	Neab3/Neab4/Neab20/Neab80/Neab81/Neab84	Neab15/Neab18/Neab19	Neab27	Neab38	Neab91	AdhoNPV ORF9	AdhoNPV ORF44	AgipMNPV ORF69	AnpeNPV ORF2	AnpeNPV ORF3	AnpeNPV ORF11	AnpeNPV ORF30 P118
Distribution [#]	(γ)	(γ)	(γ)	(γ-)	(γ-)	(γ-)	(γ-)	(γ-)	(γ-)	(α-)	(α-)	(α-)	(α-)	(α-)	(α-)	(α-)
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	1	1
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	2	1
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	1	1
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	1	1
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
	EppoNPV	0	0	0	0	0	0	0	0	0	1	0	0	1	0	1
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	1	1
	BusuNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	ChchNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	ChinNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	EcobNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	EupsNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LafiNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	LdMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LeseNPV	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0
LyxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MabrMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
MacoNPV-A	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
MacoNPV-B	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
MyunNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OrleNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	
SeMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
SujuNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
TnSNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Gammabaculovirus</i>	NeabNPV	1	1	1	2	6	3	1	1	1	0	0	0	0	0	0
	NeleNPV	1	1	1	1	2	2	1	1	1	0	0	0	0	0	0
	NeseNPV	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	3	3	3	3	8	5	2	2	2	3	27	6	11	2	7	10
Strain occupancy	3	3	3	2	2	2	2	2	2	3	27	6	11	2	6	10

No.		193	194	195	196	197	198	199	200	201	202	203	204	205	206	207	208
ID of protein homologue		AnpeNPV ORF32	AnpeNPV ORF130	AnpeNPV ORF143	BusuNPV ORF99	CfDEFMNPV ORF79	CfDEFNPV ORF107	CfDEFNPV ORF119	CfMNPV ORF121	CfDEFNPV ORF144	CfDEFNPV ORF145	ChchNPV ORF118 E3 ubiquitin-protein ligase	ChmuNPV ORF148	DapuNPV ORF8 P25	DapuNPV ORF129	LdNPV ORF67 HRF-1	HycuNPV ORF8 EP32
	Distribution [#]	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	1	0	0	1	0	1	1	0	1	1	1	0	0	0	0	0
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	1	1	0	1	1	1	0	0	0	0	0
	CfDEFMNPV	1	0	1	1	1	1	1	0	1	1	1	0	0	0	0	0
	CfMNPV	1	0	0	1	0	1	1	1	0	0	0	0	0	0	0	0
	ChmuNPV	1	1	0	1	0	0	1	0	0	0	0	0	1	0	0	0
	ChroNPV	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	1	1	0	0	1	0	0	0	1	0	0	0	1	1	1	1
	EppoNPV	1	0	0	1	0	1	0	0	0	0	0	0	1	0	0	0
	HycuNPV	1	1	0	1	0	1	0	0	0	0	0	0	0	0	0	1
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	AnpeNPV	1	1	1	0	0	1	1	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0
	ChinNPV	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0
	EcobNPV	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0
	EupsNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	LafiNPV	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
	LdMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1
	LeseNPV	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
LyxyMNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	
MabrMNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
MacoNPV-A	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
MacoNPV-B	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
ManeNPV	0	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
MyunNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpMNPV	1	1	0	1	1	0	0	0	1	0	0	0	1	1	1	1	
OpbuNPV	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	1	
OrleNPV	0	0	0	1	0	0	0	0	0	0	0	2	0	0	0	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	
SeMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SujuNPV	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	
TnSNPV	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChocGV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
	ClanGV	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
	ClasGV-A	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues		11	7	2	17	3	8	6	3	5	8	18	4	2	2	4	3
Strain occupancy		11	7	2	17	3	8	6	3	5	8	17	4	2	2	4	3

No.	209	210	211	212	213	214	215	216	217	218	219	220	221	222	223	224
ID of protein homologue	HycuNPV ORF87	BmNPV ORF7a	BmNPV ORF111	CapoNPV ORF76	CfDEFMNPV ORF116	CfMNPV ORF89/ORF90	ChmuNPV ORF120	ChmuNPV ORF133	MaviNPV ORF108	AdhoNPV ORF6 HOAR	AdhoNPV ORF7	AdhoNPV ORF15	AdhoNPV ORF44	AdhoNPV ORF45	AdhoNPV ORF96	AdhoNPV ORF97
Distribution [#]	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
	BmNPV	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	0	0	0	2	1	1	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	1	1	1	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	1	0	0	0	0	0	0	1	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	1	1	1	1	1	1	1
	AdorNPV	0	0	0	0	0	0	0	0	1	1	1	1	1	1	1
	AgipMNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0
	AgseNPV-A	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0
	AgseNPV-B	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0
	ChchNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0
	ChinNPV	0	0	0	0	0	0	0	0	2	0	1	1	1	1	0
	EcobNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0
	EupsNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0
	HearNPV-C1	0	0	0	0	0	0	0	0	1	0	0	1	1	1	0
	HearNPV-G4	0	0	0	0	0	0	0	0	1	0	0	1	1	1	0
	HespNPV	0	0	0	0	0	0	0	0	1	0	0	0	1	1	0
	LafiNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0
LdMNPV	0	0	0	0	0	0	0	0	1	0	0	0	1	1	0	
LeseNPV	0	0	0	0	0	0	0	0	1	0	0	1	1	1	0	
LyxyMNPV	0	0	0	0	0	0	0	0	1	0	0	0	1	1	0	
MabrMNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
MacoNPV-A	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
MacoNPV-B	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
ManeNPV	0	0	0	0	0	0	0	0	1	0	0	1	1	1	0	
MyunNPV	0	0	0	0	0	0	0	0	1	0	0	1	1	1	0	
OpMNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	
OrleNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
PeluSNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
PespNPV	0	0	0	0	0	0	0	0	1	0	1	0	1	1	0	
SeMNPV	0	0	0	0	0	0	0	0	1	0	1	0	1	1	0	
SfMNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
SpliMNPV	0	0	0	0	0	0	0	0	1	0	0	1	1	1	0	
SpltMNPV	0	0	0	0	0	0	0	0	1	0	0	1	1	1	0	
SujuNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
TnSNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
UrprNPV	0	0	0	0	0	0	0	0	1	0	1	1	1	1	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	2	3	2	2	2	5	2	2	2	33	2	22	27	33	3	3
Strain occupancy	2	3	2	2	2	3	2	2	2	32	2	22	27	33	3	3

No.	225	226	227	228	229	230	231	232	233	234	235	236	237	238	239	240	
ID of protein homologue	AdhoNPV	AdhoNPV	AdhoNPV	AdhoNPV	AdhoNPV	AdhoNPV	AdhoNPV	AdhoNPV	AdhoNPV	AdhoNPV	AgipMNPV	AgipMNPV	AgipMNPV	AgipMNPV	AgipMNPV	AgipMNPV	
	ORF105	ORF107	ORF109	ORF111	ORF114	ORF119	ORF121	ORF123	ORF116	ORF8	ORF13	ORF24	ORF28	ORF51	ORF52 RR2B	ORF58	
Distribution [#]	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AdhoNPV	1	1	1	1	1	1	1	1	1	0	0	0	0	0	0	
	AdorNPV	1	1	1	1	1	1	1	1	1	0	0	0	0	0	0	
	AgipMNPV	1	0	1	1	0	1	1	1	0	1	1	1	1	1	1	1
	AgseNPV-A	1	0	1	1	0	1	1	1	0	0	1	1	1	1	1	1
	AgseNPV-B	1	0	1	1	0	1	1	0	0	1	1	0	0	1	1	1
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	1	1	0	1	1	1	0	0	0	0	0	0	0	1
	ChchNPV	0	0	1	1	0	1	1	1	0	0	0	0	0	1	1	1
	ChinNPV	0	0	1	1	0	1	1	1	0	0	0	0	0	1	1	1
	EcobNPV	0	0	1	1	0	1	1	1	0	0	0	0	0	0	1	1
	EupsNPV	0	0	1	1	0	1	1	1	0	0	0	0	0	0	1	1
	HearNPV-C1	1	0	1	0	0	1	1	0	0	0	0	1	0	0	0	0
	HearNPV-G4	1	0	1	0	0	1	1	0	0	0	0	1	0	0	0	0
	HespNPV	1	0	1	1	0	1	1	1	0	0	0	0	0	0	1	1
	LafiNPV	0	0	1	1	0	0	1	1	0	0	0	0	0	0	1	1
	LdMNPV	0	0	1	0	0	1	1	1	0	0	0	0	0	0	1	0
	LeseNPV	1	0	1	0	0	1	0	1	0	0	0	0	0	0	0	0
LyxyMNPV	1	0	1	0	0	1	1	1	0	0	0	0	0	0	1	0	
MabrMNPV	1	0	1	1	0	1	1	1	0	0	1	0	1	1	1	1	
MacoNPV-A	1	0	1	2	0	1	1	1	0	0	1	0	1	1	1	1	
MacoNPV-B	1	0	1	1	0	1	1	1	0	0	1	0	1	1	1	1	
ManeNPV	1	0	1	0	0	1	1	0	0	0	0	0	0	1	0	0	
MyunNPV	1	0	1	1	0	1	1	1	0	0	1	0	1	1	1	1	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	1	0	1	0	0	0	0	0	0	0	0	1	0	
OrleNPV	0	0	1	0	0	1	1	1	0	0	0	0	0	0	1	1	
PeluSNPV	1	0	1	1	0	1	1	1	0	0	0	0	0	0	1	0	
PespNPV	1	0	1	1	0	1	1	1	0	0	1	0	0	1	1	1	
SeMNPV	0	0	1	1	0	1	1	1	0	0	0	2	0	1	1	1	
SfMNPV	0	0	1	1	0	1	1	1	0	0	0	0	0	1	0	0	
SpliMNPV	1	0	1	0	0	1	0	1	0	0	0	1	0	0	1	0	
SpltMNPV	1	0	1	0	0	1	0	1	0	0	0	1	0	0	1	0	
SujuNPV	0	0	1	1	0	1	1	1	0	0	0	0	0	0	0	1	
TnSNPV	0	0	1	1	0	1	1	1	0	0	0	0	0	1	1	1	
UrprNPV	0	1	1	0	0	1	1	1	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
No. of homologues	19	3	32	24	2	32	29	28	2	2	8	8	6	14	23	19	
Strain occupancy	19	3	32	23	2	32	29	28	2	2	8	7	6	14	23	19	

No.	241	242	243	244	245	246	247	248	249	250	251	252	253	254	255	256	
ID of protein homologue	AgipMNPV ORF60	AgipMNPV ORF84	AgipMNPV ORF131	AgipMNPV ORF135	AgipMNPV ORF136	AgipMNPV ORF152	AgseNPV-B ORF17	AgseNPV-A ORF61	AgseNPV-B ORF46	AgseNPV-B ORF50	BusuNPV ORF5	ChchNPV ORF5	ChchNPV ORF6	ChchNPV ORF90	ChchNPV ORF34	ChchNPV ORF36	
Distribution [#]	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	1	1	1	1	1	1	0	0	1	1	0	0	0	0	0	0
	AgseNPV-A	1	1	1	1	1	1	0	1	1	1	0	0	0	0	0	0
	AgseNPV-B	1	1	1	1	1	1	1	0	1	1	0	0	0	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0
	ChchNPV	0	0	0	0	1	0	1	0	0	0	0	1	1	1	1	1
	ChinNPV	0	0	0	0	0	0	1	0	0	0	0	0	1	1	1	1
	EcobNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EupsNPV	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
	HearNPV-C1	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LafiNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
LdMNPV	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	
LeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	
LyxyMNPV	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	
MabrMNPV	1	1	1	2	1	1	0	0	0	0	0	0	0	0	0	0	
MacoNPV-A	1	1	1	2	1	1	0	0	0	0	0	0	0	0	1	0	
MacoNPV-B	1	1	1	2	1	1	0	0	0	0	0	0	0	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MyunNPV	1	1	0	1	0	1	0	0	0	0	0	0	0	0	1	0	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	
OrleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PeluSNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	1	1	0	1	1	1	0	1	0	0	0	0	0	0	0	0	
SeMNPV	1	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	1	1	0	0	2	1	0	0	0	0	0	0	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	
SujuNPV	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	
TnSNPV	0	0	0	0	1	0	0	0	0	0	0	1	1	1	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
No. of homologues	14	9	6	11	13	14	3	2	3	3	2	2	3	3	12	2	
Strain occupancy	14	9	6	8	11	14	3	2	3	3	2	2	3	3	11	2	

No.		257	258	259	260	261	262	263	264	265	266	267	268	269	270	271	272	
ID of protein homologue		ChchNPV ORF103	ChchNPV ORF104	ChchNPV ORF105	ChchNPV ORF116	ChchNPV ORF117	ChchNPV ORF132	ChchNPV ORF135	ChchNPV ORF144	ChchNPV ORF149	EcobNPV ORF23	EcobNPV ORF100	EupsNPV ORF5	EupsNPV ORF116	HearNPV-C1 ORF30 EL25	HearNPV-C1 ORF34	LdMNPV ORF42	
	Distribution [#]	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	(a-)	
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0
	ChchNPV	1	1	1	1	1	1	1	1	1	0	0	0	0	0	0	0	0
	ChinNPV	1	1	1	1	1	1	1	1	0	0	0	0	0	0	0	0	0
	EcobNPV	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0
	EupsNPV	0	0	0	0	0	0	0	0	0	0	1	1	1	0	0	0	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	1	1	0	1	1	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	1	1	0	1	0	0	0	0
	LafiNPV	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0
LdMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	
LeseNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	1	1	0	0	
LyxyMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	
MabrMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1	
MyunNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	
OrleNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	0	
SeMNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	
SfMNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	
SujuNPV	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	
TnSNPV	1	1	0	1	1	1	1	1	1	1	0	0	0	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
No. of homologues		3	3	2	3	3	3	3	2	2	15	15	2	3	3	3	3	
Strain occupancy		3	3	2	3	3	3	3	2	2	15	15	2	3	3	3	3	

No.	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336
ID of protein homologue	CDEFNPV ORF110	AgipMNPV ORF65	XcGV ORF66	AgipMNPV ORF64	AdorGV ORF102	AdhoNPV ORF87 P13	AdorGV ORF108 Helicase-2	CpGV ORF100	XcGV ORF114	AdorGV ORF40	AnpeNPV ORF125 P22.2	CpGV ORF128 RR2A	AgipMNPV ORF82	ChchNPV ORF113	ChmuNPV ORF10	ChmuNPV ORF120
Distribution [#]	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0
	CfDEFMNPV	1	0	0	0	0	0	0	0	0	1	0	0	0	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	1	0	1	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1
	ChroNPV	0	0	0	0	0	0	0	0	0	1	0	1	0	1	1
	DapuNPV	0	0	0	0	0	0	0	1	1	0	1	0	0	0	0
	EppoNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1
	HycuNPV	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0
	LoobMNPV	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
	PlxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	1	0	1	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	1	0	1	0	0	0	0	0	0	0
	AgipMNPV	0	1	0	1	1	1	0	0	0	0	0	0	1	0	0
	AgseNPV-A	0	0	0	1	1	1	0	0	0	0	0	0	3	0	0
	AgseNPV-B	0	1	0	1	1	1	0	0	0	0	0	0	2	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	1	1	1	0	0	0	0	0	0	1	0	0
	ChinNPV	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0
	EcobNPV	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
	EupsNPV	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0
	HearNPV-C1	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0
	HespNPV	0	0	1	1	2	0	0	1	1	0	0	0	0	0	0
	LafiNPV	0	0	0	1	0	0	1	0	1	0	0	0	0	0	0
	LdMNPV	0	0	0	1	0	0	1	0	0	0	1	2	0	0	0
	LeseNPV	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0
LyxyMNPV	0	0	0	1	0	0	1	0	0	0	0	2	0	0	0	
MabrMNPV	0	0	0	1	1	1	1	0	0	0	0	1	0	0	0	
MacoNPV-A	0	0	0	1	1	1	0	0	0	0	0	1	0	0	0	
MacoNPV-B	0	0	0	1	1	1	1	0	0	0	0	1	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	2	0	0	0	0	0	0	
MyunNPV	0	1	0	1	1	1	0	0	0	0	0	1	0	0	0	
OpMNPV	0	0	0	0	0	0	0	0	1	0	1	0	0	0	0	
OpbuNPV	0	0	0	1	0	1	0	0	0	0	0	1	0	0	0	
OrleNPV	0	0	0	1	0	0	1	0	0	1	0	0	1	0	0	
PeluSNPV	0	0	0	0	1	1	0	0	1	0	0	0	0	0	0	
PespNPV	0	2	0	1	1	1	1	0	0	0	0	1	0	0	0	
SeMNPV	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	
SfMNPV	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	
SpliMNPV	0	1	0	1	0	1	1	0	0	0	0	0	0	0	0	
SpltMNPV	0	1	0	1	0	1	1	0	0	0	0	0	0	0	0	
SujuNPV	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	
TnSNPV	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	1	1	1	0	1	0	0	0	0	0	0
	ChocGV	0	0	0	0	1	1	1	1	1	0	0	0	0	1	0
	ClanGV	0	0	0	0	1	1	0	1	1	0	0	0	0	0	0
	ClasGV-A	0	0	0	0	1	1	0	0	1	0	0	0	0	0	0
	ClasGV-B	0	0	0	0	0	1	1	1	0	1	0	0	0	0	0
	CnmeGV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0
	CpGV	1	2	0	0	1	1	1	1	0	0	1	0	0	0	0
	CrleGV	1	0	1	0	1	1	1	1	0	1	0	0	0	0	0
	DisaGV	0	0	0	1	0	1	1	0	0	0	1	0	0	0	0
	EpapGV	0	0	0	1	1	1	1	0	0	0	1	0	0	0	1
	ErelGV	0	0	0	0	0	1	1	1	0	2	0	0	0	0	0
	PrGV	1	0	0	0	1	1	1	1	0	0	1	0	0	0	0
	HearGV	1	0	1	0	1	1	1	1	1	0	0	0	4	0	0
	MolaGV	0	0	0	0	0	1	1	1	1	0	0	0	3	0	0
	MyunGV	1	1	0	1	0	1	1	1	1	0	0	3	0	0	0
	PxGV	0	0	0	0	1	1	1	1	1	0	0	0	0	0	0
SfGV	1	0	0	1	0	1	1	1	1	0	0	2	0	0	0	
SpliGV	0	0	0	1	0	1	1	1	1	0	0	0	0	0	0	
XcGV	2	0	1	0	1	1	1	1	1	0	0	4	2	0	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	10	10	6	27	26	41	26	19	12	18	10	6	36	5	2	4
Strain occupancy	9	8	6	27	25	41	26	19	12	16	10	6	20	4	2	4

No.	337	338	339	340	341	342	343	344	345	346	347	348	349	350	351	352
ID of protein homologue	AgipMNPV ORF40	AgipMNPV ORF163 RR1	CpGV ORF40	ClanGV ORF42	ClasGV-B ORF50	ChorGV ORF11	ChorGV ORF36	DisaGV ORF54	AdhoNPV ORF3	AdhoNPV ORF115	AgipMNPV ORF26	AgipMNPV ORF20	AnpeNPV ORF145	BusuNPV ORF12	ChchNPV ORF68 Phr-1/ChchNPV ORF72 Phr-2	ChchNPV ORF109
Distribution [#]	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	DapuNPV	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0
	AgipMNPV	1	1	0	0	0	0	0	0	0	0	1	1	0	0	0
	AgseNPV-A	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-B	1	1	0	0	0	0	0	0	0	0	1	0	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	BusuNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0
	ChchNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	2
	ChinNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1
	EcobNPV	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	EupsNPV	1	1	0	0	0	0	0	0	0	0	0	0	0	1	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HespNPV	1	1	0	0	1	0	0	0	0	0	0	0	0	0	0
	LafiNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	LdMNPV	1	1	0	0	0	0	0	1	0	0	0	0	0	1	0
LeseNPV	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	
LyxyMNPV	1	0	0	0	0	0	0	1	0	0	0	0	0	1	0	
MabrMNPV	1	1	0	0	0	0	0	0	0	0	1	0	0	0	0	
MacoNPV-A	1	1	0	0	0	0	0	0	0	2	1	0	0	0	0	
MacoNPV-B	1	1	0	0	0	0	0	0	0	1	1	0	0	0	0	
ManeNPV	1	0	0	2	0	0	1	0	0	0	0	0	0	0	0	
MyunNPV	0	1	0	0	0	0	0	0	0	0	1	1	0	0	0	
OpMNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	
OrleNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	
PeluSNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	
SeMNPV	1	1	0	0	0	0	0	0	0	0	1	0	0	0	0	
SfMNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliMNPV	0	1	0	0	0	1	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	1	0	0	0	1	0	0	0	0	0	0	0	0	0	
SujuNPV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	
TnSNPV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	
UrprNPV	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChocGV	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0
	ClanGV	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-A	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-B	1	0	1	0	1	0	1	0	0	0	0	0	0	0	0
	CnmeGV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	CpGV	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
	CrleGV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
	DisaGV	0	1	0	0	0	0	0	1	0	0	0	0	0	0	0
	EpapGV	0	1	0	0	0	0	0	0	1	0	0	0	0	0	0
	ErelGV	1	0	1	0	1	0	1	0	0	0	0	0	0	0	0
	PrGV	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
	HearGV	0	0	0	0	0	0	0	0	1	1	1	1	0	0	0
	MolaGV	0	0	0	0	0	0	0	0	1	1	0	0	0	0	0
	MyunGV	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	SfGV	0	0	0	0	0	0	0	0	2	0	1	0	0	0	1
SpliGV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	
XcGV	0	0	0	0	0	0	0	0	1	1	1	1	0	0	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	21	29	5	4	5	4	4	3	9	9	10	4	9	4	8	4
Strain occupancy	21	29	5	3	5	4	4	3	8	8	10	4	9	4	7	4

No.	353	354	355	356	357	358	359	360	361	362	363	364	365	366	367	368
ID of protein homologue	MabrMNPV ORF14 XE-1/ORF68 XE-2	MabrMNPV ORF23	MabrMNPV ORF24	SeMNPV ORF20	HearGV ORF53/ORF157	MyunGV B ORF49	XcGV ORF42	XcGV ORF57	XcGV ORF61	XcGV ORF65	CpGV ORF35	CpGV ORF42	CpGV ORF5	CpGV ORF30	CpGV ORF45	CpGV ORF50
Distribution [#]	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(α -, β -)	(β -)	(β -)	(β -)	(β -)	(β -)	(β -)
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChinNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EcobNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EupsNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LafiNPV	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
LdMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
LeseNPV	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	
LyxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MabrMNPV	2	1	1	0	0	0	0	1	0	0	0	0	0	0	0	
MacoNPV-A	2	1	1	0	0	0	0	1	0	0	0	0	0	0	0	
MacoNPV-B	2	1	1	0	1	0	0	1	1	1	0	0	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MyunNPV	1	1	1	0	0	0	0	1	0	1	0	0	0	0	0	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OrleNPV	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	3	0	0	0	0	0	
SeMNPV	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	
SpliMNPV	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	0	1	0	0	0	0	0	0	1	0	0	0	0	0	
SujuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
TnSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	1	1	1	1	1	1
	ChocGV	0	0	0	0	0	0	0	0	0	0	1	1	0	1	0
	ClanGV	0	0	0	0	0	0	0	0	0	0	1	1	0	1	1
	ClasGV-A	0	0	0	0	0	0	0	0	0	1	1	1	0	1	1
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	1	1	0	1	1
	CnmeGV	0	0	0	0	0	0	0	0	0	1	1	1	0	1	1
	CpGV	0	0	0	0	0	0	0	0	0	1	1	1	1	1	1
	CrleGV	0	0	0	0	0	0	0	0	0	1	1	1	1	1	1
	DisaGV	0	0	0	0	0	0	0	0	0	1	1	0	0	0	1
	EpapGV	0	0	0	0	0	0	0	0	0	0	1	1	0	0	1
	ErelGV	0	0	0	0	0	0	0	0	0	0	1	1	0	1	1
	PrGV	0	0	0	0	0	0	0	0	0	1	1	1	1	1	1
	HearGV	1	0	0	0	2	0	1	1	1	1	1	1	1	1	1
	MolaGV	0	0	0	0	0	0	1	1	1	0	1	1	1	1	1
	MyunGV	0	0	0	0	0	1	1	1	0	0	1	1	1	1	1
	PxGV	0	0	0	0	0	0	0	0	0	0	0	1	1	1	1
SfGV	0	0	1	1	0	0	1	2	0	0	1	1	1	1	1	
SpliGV	0	0	0	0	0	0	0	0	0	1	1	1	1	1	1	
XcGV	0	1	0	0	0	0	1	1	1	1	1	1	1	1	2	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	8	5	8	2	4	2	6	11	4	9	12	18	18	11	17	19
Strain occupancy	5	5	8	2	3	2	6	10	4	7	12	18	18	11	17	18

No.	369	370	371	372	373	374	375	376	377	378	379	380	381	382	383	384
ID of protein homologue	CpGV ORF62	CpGV ORF115	CpGV ORF122	CpGV ORF132	CpGV ORF121	CpGV ORF136	CpGV ORF43	CpGV ORF129/ORF130	CpGV ORF133	PxGV ORF3	XcGV ORF173	CpGV ORF28/29	HearGV ORF24	XcGV ORF124	XcGV ORF176	AdorGV ORF56
Distribution [#]	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChinNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EcobNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EupsNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LafiNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LdMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
LyxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MabrMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MyunNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OrleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SeMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SujuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
TnSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	1	1	0	0	1	1	0	1	0	1	0	0	0	1
	ChocGV	1	1	1	0	1	1	1	1	1	0	0	1	0	0	1
	ClanGV	1	0	1	1	1	1	1	1	1	1	0	1	0	0	0
	ClasGV-A	1	0	1	1	1	1	1	1	1	1	0	1	0	0	0
	ClasGV-B	1	1	1	0	1	1	1	1	1	1	0	1	0	0	1
	CnmeGV	1	1	0	1	0	1	0	1	1	0	0	1	0	0	1
	CpGV	1	1	1	1	1	1	1	2	1	0	0	2	0	0	0
	CrleGV	1	1	1	1	1	1	1	2	0	0	0	2	0	0	0
	DisaGV	1	1	0	0	0	1	0	0	0	0	0	1	0	0	1
	EpapGV	1	1	1	1	0	0	0	1	1	0	1	1	0	0	0
	ErelGV	1	1	1	0	1	1	1	1	0	1	0	1	0	0	1
	PrGV	1	1	1	1	1	1	1	1	1	1	0	1	0	0	1
	HearGV	1	1	1	1	1	0	0	0	1	0	1	0	1	1	0
	MolaGV	1	1	1	1	1	0	0	0	1	0	1	0	1	1	0
	MyunGV	1	1	1	1	1	0	0	0	1	0	1	0	1	1	0
	PxGV	1	1	1	1	1	1	0	1	0	1	0	0	1	1	0
SfGV	1	1	1	1	1	0	0	0	0	0	1	0	1	1	0	
SpliGV	1	1	1	1	0	0	1	1	0	0	1	0	1	1	0	
XcGV	1	1	1	1	1	0	0	0	1	0	1	0	1	1	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	18	17	17	15	14	12	10	15	12	7	7	14	7	7	7	7
Strain occupancy	18	17	17	15	14	12	10	13	12	7	7	12	7	7	7	7

No.	385	386	387	388	389	390	391	392	393	394	395	396	397	398	399	400
ID of protein homologue	ClanGV	ClanGV	ClanGV	ClanGV	ClanGV	ClanGV	ClanGV	ClasGV-B	ClasGV-B	ClasGV-B	CpGV	CpGV	CpGV	CpGV	CpGV	CpGV
	ORF67	ORF120	ORF10	ORF40	ORF53	ORF79	ORF93	ORF24	ORF63	ORF93	ORF12	ORF27	ORF32	ORF44	ORF67	ORF77
Distribution [#]	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChinNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EcobNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EupsNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LafiNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LdMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
LyxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MabrMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MyunNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OrleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SeMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SujuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
TnSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
	ChocGV	1	0	0	0	0	0	0	0	0	1	0	1	0	0	1
	ClanGV	1	1	1	1	1	1	1	0	0	0	0	1	0	0	1
	ClasGV-A	1	1	1	1	1	1	1	0	0	0	0	1	0	0	1
	ClasGV-B	1	0	0	0	0	0	0	1	1	1	0	1	0	0	1
	CnmeGV	1	0	0	0	0	0	0	0	0	0	0	1	0	0	1
	CpGV	0	0	0	0	0	0	0	0	0	0	1	1	1	1	1
	CrleGV	0	0	0	0	0	0	0	0	0	0	1	0	1	1	1
	DisaGV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EpapGV	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1
	ErelGV	1	1	0	0	0	0	0	1	1	1	0	1	0	0	1
	PrGV	1	1	0	0	0	0	0	0	0	0	1	0	1	0	1
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
XcGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	9	4	2	2	2	2	2	2	2	3	3	4	7	2	3	11
Strain occupancy	9	4	2	2	2	2	2	2	2	3	3	4	7	2	3	11

No.	401	402	403	404	405	406	407	408	409	410	411	412	413	414	415	416
ID of protein homologue	CpGV ORF110	CpGV ORF124	CpGV ORF139	HearGV ORF104	MyunGV ORF17	PxGV ORF77	XcGV ORF4	XcGV ORF23	XcGV ORF24	XcGV ORF28	XcGV ORF33	XcGV ORF37	XcGV ORF38	XcGV ORF44	XcGV ORF70	XcGV ORF71
Distribution [#]	401	402	403	404	405	406	407	408	409	410	411	412	413	414	415	416
<i>Alphabaculovirus</i>	AcMNPV	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChinNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EcobNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EupsNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LafiNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LdMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
LyxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MabrMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MyunNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OrleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SeMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SujuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
TnSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChocGV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClanGV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-A	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-B	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	CnmeGV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	CpGV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	CrleGV	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
	DisaGV	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
	EpapGV	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PrGV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearGV	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	MolaGV	0	0	0	1	0	0	1	1	1	1	1	1	1	1	1
	MyunGV	0	0	0	1	0	0	1	1	0	1	1	0	0	1	1
	PxGV	0	0	0	0	1	0	1	1	1	1	1	0	1	0	1
SfGV	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	
SpliGV	0	0	0	0	0	0	1	1	1	1	0	0	1	0	1	
XcGV	0	0	0	0	1	1	0	0	1	1	0	0	1	0	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	1	1	1	1	1	1	1	1	1	1
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	3	11	2	2	3	2	5	5	5	6	4	2	5	3	3	5
Strain occupancy	3	11	2	2	3	2	5	5	5	6	4	2	5	3	3	5

No.	417	418	419	420	421	422	423	424	425	426	427	428	429	430	431	432	
ID of protein homologue	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	XcGV	DisaGV	
	ORF72	ORF73/ORF155/ORF161	ORF74	ORF86	ORF90	ORF106	ORF115	ORF117	ORF157	ORF162	ORF164	ORF167	ORF168	ORF177	ORF181	ORF106/ORF107/ORF108	
Distribution [#]	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	(β-)	Species-unique	
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PixyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AgipMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AgseNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	BusuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChchNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChinNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EcobNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	EupsNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	LafiNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
LdMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
LeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
LyxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
MabrMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
MacoNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
MacoNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
ManeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
MyunNPV	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0		
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
OrleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
PeluSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
PespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
SeMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
SfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
SpliMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
SpltMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
SujuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
TnSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
UrprNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	HearGV	1	3	1	1	1	1	1	1	1	1	1	1	1	1	1	0
	MolaGV	1	3	0	1	1	1	1	1	0	1	0	0	0	1	1	0
	MyunGV	0	2	2	1	1	0	1	1	0	1	1	1	0	1	1	0
PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SfGV	0	2	0	0	1	0	1	0	0	1	0	1	0	1	1	0	
SpliGV	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	
XcGV	1	4	1	1	1	1	1	1	1	1	1	1	1	1	1	0	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
No. of homologues	3	14	4	5	6	3	5	4	2	6	3	4	2	5	5	3	
Strain occupancy	3	5	3	5	6	3	5	4	2	6	3	4	2	5	5	1	

No.	977	978	979	980	981	982	983	984	985	986	987	988	989	990	991	992
ID of protein homologue	UrprNPV ORF5	UrprNPV ORF6	UrprNPV ORF18	UrprNPV ORF23	UrprNPV ORF34	UrprNPV ORF40	UrprNPV ORF63	UrprNPV ORF118	XcGV ORF6	XcGV ORF41	XcGV ORF49	XcGV ORF69	XcGV ORF104	XcGV ORF153	XcGV ORF156	XcGV ORF163
Distribution [#]	Species-unique															
<i>Alphabaculovirus</i>	AcMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BmNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BomaNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CapoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CoveNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfDEFMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChmuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChroNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DapuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EppoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HycuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LoobMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MaviNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	OxocNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PlyxMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ThorMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdhoNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AdorNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgipMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AgseNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	AnpeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	BusuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChchNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChinNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EcobNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EupsNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-C1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearNPV-G4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	LafiNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
LdMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
LeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
LyxyMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MabrMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MacoNPV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ManeNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MyunNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OpbuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
OrleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PeluSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
PespNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SeMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SfMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpliMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SpltMNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
SujuNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
TnSNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
UrprNPV	1	1	1	1	1	1	1	1	0	0	0	0	0	0	0	
<i>Betabaculovirus</i>	AdorGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ChocGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClanGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ClasGV-B	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CnmeGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CpGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	CrleGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	DisaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	EpapGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	ErelGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PrGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	HearGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MolaGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	MyunGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	PxGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	SfGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
SpliGV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
XcGV	0	0	0	0	0	0	0	0	1	1	1	1	1	1	1	
<i>Gammabaculovirus</i>	NeabNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeleNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	NeseNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Deltabaculovirus</i>	CuniNPV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
No. of homologues	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
Strain occupancy	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1

#: "-" indicates that the protein homologue is not present in all the corresponding group of baculoviruses.
The newly identified homologues in in baculoviral IRPDs were indicated as blue box.

Table S4. Distribution of homologue groups with a species occupancy ≥ 2 among baculoviruses

No.	<i>Alphabaculovirus</i>	<i>Betabaculovirus</i>	<i>Gammabaculovirus</i>	<i>Deltabaculovirus</i>	No. of homologue groups
1	++	++	++	++	38
2	++	++	++		11
3	++	++	-	++	1
4	++	++	-	-	15
5	++	+	-	-	4
6	+	++	-	-	1
7	+	+	-	-	67
8	++	-	-	-	5
9	+	-	-	++	1
10	+	-	-	-	179
11	-	++	-	-	7
12	-	+	-	-	69
13	-	-	++	-	25
14	-	-	+	-	6

Note:

++: found in all the baculoviruses among the genus.

+: found in the baculoviruses but not found in all the baculoviruses among the genus.

-: not found in any of the baculoviruses among the genus.

Table S5. Overview of functions of AcMNPV ORFs and their homologues

Identities	AcMNPV ORF	Mut & Del ^a			Function
		AcMNPV	BmNPV	HearNPV	
PTP	1	Mut (Gross and Shuman, 1998a; Li and Miller, 1995b; Takagi <i>et al.</i> , 1998), Del (Li and Guarino, 2008)	Mut (Katsuma, 2015), Del (Gomi <i>et al.</i> , 1999; Kamita <i>et al.</i> , 2005; Katsuma, 2015; Katsuma <i>et al.</i> , 2008b; Ono <i>et al.</i> , 2012)		Important but not essential for baculovirus infection by promoting viral replication (Kamita <i>et al.</i> , 2005; Katsuma <i>et al.</i> , 2008b; Wang <i>et al.</i> , 2016) and capable of inducing enhanced locomotory activity in a lepidopteran host (Kamita <i>et al.</i> , 2005) and affecting ODV production in SF-21 but not TN-368 cell lines (Li and Miller, 1995b); exhibiting the RNA 5'-triphosphatase activity involved in RNA cap formation (Gross and Shuman, 1998a; Li and Guarino, 2008; Martins and Shuman, 2002; Takagi <i>et al.</i> , 1998) and the protein tyrosine phosphatases activity (Li and Miller, 1995b; Sheng and Charbonneau, 1993) which is not required for ELA induction in BmNPV-infected <i>B. mori</i> larvae (Katsuma, 2015)
BRO	2		Mut (Kang <i>et al.</i> , 2006), Del (Kang <i>et al.</i> , 1999; Ono <i>et al.</i> , 2012)		Important for BV production but not essential (Gomi <i>et al.</i> , 1999; Ono <i>et al.</i> , 2012) as a DNA binding protein (Zemskov <i>et al.</i> , 2000) and nucleocytoplasmic shuttling proteins (Kang <i>et al.</i> , 2006) and involved in interacting with host protein laminin (Kang <i>et al.</i> , 2003)
CTX	3	Del (Eldridge <i>et al.</i> , 1992)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Eldridge <i>et al.</i> , 1992)
AC4	4	Del (Gandhi <i>et al.</i> , 2012)	Del (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2018a)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Gandhi <i>et al.</i> , 2012; Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2018a)
AC5	5	Del (Wang <i>et al.</i> , 2018)	Del (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2018a)		Dispensable for BV and ODV production but indispensable to the embedding of ODVs into polyhedra (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2018a; Wang <i>et al.</i> , 2018) as a ODV- and OB-associated protein (Braunagel <i>et al.</i> , 2003; Wang <i>et al.</i> , 2018)
LEF2	6	Mut (Merrington <i>et al.</i> , 1996), Del (Wu <i>et al.</i> , 2010)	Del (Gomi <i>et al.</i> , 1997; Ono <i>et al.</i> , 2012)		Essential for DNA replication (Kool <i>et al.</i> , 1994; Lu and Miller, 1995; Wu <i>et al.</i> , 2010) but not its initiation (Wu <i>et al.</i> , 2010) and involved in interacting with LEF1 (Evans <i>et al.</i> , 1997)
AC7	7	Del (Gearing and Possee, 1990)			Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Gearing and Possee, 1990),
POLH	8	Mut (Lopez <i>et al.</i> , 2011; Smith <i>et al.</i> , 1983), Del (Luckow <i>et al.</i> , 1993)	Del (Maeda <i>et al.</i> , 1985; Ono <i>et al.</i> , 2012)	Del (Wang <i>et al.</i> , 2003)	Dispensable for DNA and BV production (Rohrmann, 1986) but essential for ODV specific occlusion and polyhydra formation (Ji <i>et al.</i> , 2010; Lopez <i>et al.</i> , 2011) as the major polyhedron protein (Ji <i>et al.</i> , 2010; Rohrmann, 1986)
P78/83	9	Mut (Goley <i>et al.</i> , 2006; Ohkawa <i>et al.</i> , 2010; Wang <i>et al.</i> , 2015), Del (Wang <i>et al.</i> , 2015)	Del (Ono <i>et al.</i> , 2012)	Del (Wang <i>et al.</i> , 2008c)	Essential (Wang <i>et al.</i> , 2008c) and involved mediating actin-driven mobility of nucleocapsid and actin nucleation through interaction with Arp2/3 complex and BV/ODV-C42 (Goley <i>et al.</i> , 2006; Lanier and Volkman, 1998; Ohkawa <i>et al.</i> , 2010; Ohkawa and Welch, 2018; Wang <i>et al.</i> , 2015); also a component of the virus-induced RNA polymerase (Iorio <i>et al.</i> , 1998)
PK1	10	Mut (Fan <i>et al.</i> , 1996; Liang <i>et al.</i> , 2017), Del	Del (Katsuma <i>et al.</i> , 2008b);		Essential (Katsuma <i>et al.</i> , 2008b), associated with a very late transcription complex, and involved in LEF8 phosphorylation (Mishra <i>et al.</i> , 2008b) and regulating the hyperexpression of very late

		(Liang <i>et al.</i> , 2017)	Ono <i>et al.</i> , 2012)	gene expression (Liang <i>et al.</i> , 2017); may be a very late gene transcription factor (Mishra <i>et al.</i> , 2008a; Mishra <i>et al.</i> , 2007)
AC11	11	Del (Tao <i>et al.</i> , 2015)	Del (Ono <i>et al.</i> , 2012; Wang <i>et al.</i> , 2019a)	Important but not essential for OB production, ODV embedding and BV infectivity <i>in vivo</i> for BmNPV (Ono <i>et al.</i> , 2012; Wang <i>et al.</i> , 2019a), but essential for BV production and ODV envelopment in AcMNPV as an early gene (Tao <i>et al.</i> , 2015).
AC12	12	Del (Costa Navarro <i>et al.</i> , 2019)		Dispensable for BV as a F box-like protein capable of interacting with cellular SKP1 (Costa Navarro <i>et al.</i> , 2019)
AC13	13	Del (Chen <i>et al.</i> , 2021; Tong <i>et al.</i> , 2021)	Del (Kokusho <i>et al.</i> , 2016; Ono <i>et al.</i> , 2012)	Important but not essential BV production and involved in efficient nuclear egress of nucleocapsids and regulation of viral gene expression (Chen <i>et al.</i> , 2021; Kokusho <i>et al.</i> , 2016; Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
LEF1	14	Mut (Evans <i>et al.</i> , 1997; Mikhailov and Rohrmann, 2002a), Del (Evans <i>et al.</i> , 1997)	Del (Ono <i>et al.</i> , 2012)	Essential for transient DNA replication (Kool <i>et al.</i> , 1994; Lu and Miller, 1995) and production of both BV and ODV (Evans <i>et al.</i> , 1997; Ono <i>et al.</i> , 2012) and capable of interacting with LEF2 (Evans <i>et al.</i> , 1997); has the DNA primase activity (Mikhailov and Rohrmann, 2002a)
EGT	15	Mut (Shikata <i>et al.</i> , 1998), Del (Flipsen <i>et al.</i> , 1995)	Del (Katsuma <i>et al.</i> , 2008b; Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV (Flipsen <i>et al.</i> , 1995; Katsuma <i>et al.</i> , 2008b) and capable of blocking insect molting and pupation (O'Reilly and Miller, 1989) by transferring glucose from UDP-glucose to ecdysteroids (O'Reilly, 1999; O'Reilly and Miller, 1989); deletions commonly occur in cell culture (O'Reilly and Miller, 1990); deletion results in early degeneration of the malpighian tubules and increased speed of killing host (Flipsen <i>et al.</i> , 1995) and affects the moulting and metamorphosis of larvae (Shikata <i>et al.</i> , 1998) and tree-top disease (Han <i>et al.</i> , 2015)
BV/ODV-E26	16	Mut (Nie <i>et al.</i> , 2009; O'Reilly <i>et al.</i> , 1990), Del (Nie and Theilmann, 2010)	Mut (Kang <i>et al.</i> , 2005), Del (Hikida <i>et al.</i> , 2018; Katsuma <i>et al.</i> , 2012; Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV (Nie and Theilmann, 2010); a protein in multiple forms, which is capable of interacting with IE0 (Nie <i>et al.</i> , 2009), IE1 (Kang <i>et al.</i> , 2005; Nie <i>et al.</i> , 2009) and FP25K (Beniya <i>et al.</i> , 1998; Braunagel <i>et al.</i> , 1999), and forming a complex with FP25K and cellular actin (Beniya <i>et al.</i> , 1998) and associated with sorting motif of ODV-E66 (Saksena <i>et al.</i> , 2006); deletion results in delayed viral gene expression in BV-infected cells but not transfected cells; determines tissue tropism and virulence in host lepidopteran insects (Hikida <i>et al.</i> , 2018; Katsuma <i>et al.</i> , 2012)
AC17	17	Del (Nie and Theilmann, 2010)	Del (Ono <i>et al.</i> , 2012; Yang <i>et al.</i> , 2009)	Dispensable for BV and ODV (Nie and Theilmann, 2010; Yang <i>et al.</i> , 2009) as a late protein which may interact with host proteins (An <i>et al.</i> , 2005); deletion results in delayed viral gene expression in BV-infected cells but not transfected cells
AC18	18	Del (Wang <i>et al.</i> , 2007)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV (Ono <i>et al.</i> , 2012; Wang <i>et al.</i> , 2007); deletion does not reduce AcMNPV infectivity in LD ₅₀ bioassay but results in delayed mortality in LT ₅₀ bioassay for <i>Trichoplusia ni</i> (Wang <i>et al.</i> , 2007)
AC19	19	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
ARIF-1	20/21	Mut (Dreschers <i>et al.</i> , 2001), Del (Dreschers <i>et al.</i> , 2001)	Del (Kokusho <i>et al.</i> , 2015; Ono <i>et al.</i> , 2012)	Important but not essential for systemic infection (Dreschers <i>et al.</i> , 2001; Kokusho <i>et al.</i> , 2015; Ono <i>et al.</i> , 2012) and capable of inducing early actin rearrangement (Dreschers <i>et al.</i> , 2001; Roncarati and Knebel-Morsdorf, 1997)

PIF2	22	Del (Clavijo <i>et al.</i> , 2009; Ohkawa <i>et al.</i> , 2005; Peng <i>et al.</i> , 2010; Wang <i>et al.</i> , 2019b)	Del (Ono <i>et al.</i> , 2012)	Del (Fang <i>et al.</i> , 2006; Song <i>et al.</i> , 2008)	<i>Per os</i> infectivity factor (Ohkawa <i>et al.</i> , 2005) as a component of PIF complex (Peng <i>et al.</i> , 2012; Peng <i>et al.</i> , 2010; Wang <i>et al.</i> , 2019b)
F protein	23	Del (Lung <i>et al.</i> , 2003; Wang <i>et al.</i> , 2008a; Yu <i>et al.</i> , 2009)	Del (Ono <i>et al.</i> , 2012; Xu <i>et al.</i> , 2019; Xu <i>et al.</i> , 2020a; Xu <i>et al.</i> , 2020b)	Mut (Long <i>et al.</i> , 2007; Tan <i>et al.</i> , 2008; Yin <i>et al.</i> , 2014), Del (Long <i>et al.</i> , 2007; Tan <i>et al.</i> , 2008; Wang <i>et al.</i> , 2008b; Wang <i>et al.</i> , 2010b; Yin <i>et al.</i> , 2014)	Essential for BV infectivity and production (Long <i>et al.</i> , 2006; Tan <i>et al.</i> , 2008; Wang <i>et al.</i> , 2010b) and capable of affecting ODV phenotype and OB morphogenesis in (Xu <i>et al.</i> , 2019; Yu <i>et al.</i> , 2009) and facilitating ODV attachment to the midgut epithelial cells (Xu <i>et al.</i> , 2020a) and GP64-Mediated BV efficient infection (Xu <i>et al.</i> , 2020b)
PKIP	24	Mut (McLachlin <i>et al.</i> , 1998), Del (Fan <i>et al.</i> , 1998; Lai <i>et al.</i> , 2020; Lai <i>et al.</i> , 2019)	Del (Ono <i>et al.</i> , 2012)		Essential (Fan <i>et al.</i> , 1998; Lai <i>et al.</i> , 2020; Lai <i>et al.</i> , 2019; McLachlin <i>et al.</i> , 1998) and involved in the hyperphosphorylation of the viral basic protein P6.9 (Lai <i>et al.</i> , 2020); interacts with and stimulate the activity viral PK and affect very late gene transcription (Fan <i>et al.</i> , 1998; McLachlin <i>et al.</i> , 1998);
DBP	25	Del (Guo <i>et al.</i> , 2019; Quadt <i>et al.</i> , 2007; Vanarsdall <i>et al.</i> , 2007a)	Del (Ono <i>et al.</i> , 2012; Zhao <i>et al.</i> , 2016)		Essential for nucleocapsid production (Quadt <i>et al.</i> , 2007; Vanarsdall <i>et al.</i> , 2007a) and BV production (Guo <i>et al.</i> , 2019) and involved in regulation of late gene expression (Guo <i>et al.</i> , 2019; Zhao <i>et al.</i> , 2016) as a ssDNA binding protein capable of both unwinding and annealing DNA (Mikhailov <i>et al.</i> , 1998; Mikhailov <i>et al.</i> , 2008)
AC26	26	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021) as a late protein (Shen <i>et al.</i> , 2012)
IAP1	27	Del (Griffiths <i>et al.</i> , 1999)	Del (Ono <i>et al.</i> , 2012)		May be important for BV and ODV (Griffiths <i>et al.</i> , 1999); involved in antiapoptosis (Clem and Miller, 1994)
LEF6	28	Del (Lin and Blissard, 2002a)	Del (Ono <i>et al.</i> , 2012)		Important but not essential for production of both BV and ODV (Lin and Blissard, 2002a), and involved in late gene expression (Lin and Blissard, 2002a; Passarelli and Miller, 1994a)
AC29	29	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC30	30		Del (Huang <i>et al.</i> , 2008; Ono <i>et al.</i> , 2012)		Dispensable for BV (Huang <i>et al.</i> , 2008); deletion results in delayed mortality of larvae (Huang <i>et al.</i> , 2008)
SOD	31	Del (Tomalski <i>et al.</i> , 1991)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tomalski <i>et al.</i> , 1991)
FGF	32	Del (Detvisitsakun <i>et al.</i> , 2007)	Mut (Katsuma <i>et al.</i> , 2006a), Del (Katsuma <i>et al.</i> , 2006b; Ono <i>et al.</i> , 2012)	Del (Yin <i>et al.</i> , 2016)	Dispensable for BV and ODV, but important for systematic infection (Detvisitsakun <i>et al.</i> , 2007; Katsuma <i>et al.</i> , 2006b; Means and Passarelli, 2010; Yin <i>et al.</i> , 2016) and involved in efficient BV

				production and stimulating host cell motility (Detvisitsakun <i>et al.</i> , 2007; Lehiy <i>et al.</i> , 2009)
AC33	33	Del (Tong <i>et al.</i> , 2021)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Tong <i>et al.</i> , 2021)
AC34	34	Del (Cai <i>et al.</i> , 2012; Mu <i>et al.</i> , 2016b; Qiu <i>et al.</i> , 2017)	Del (Ono <i>et al.</i> , 2012)	Important for BV production (Cai <i>et al.</i> , 2012; Ono <i>et al.</i> , 2012; Qiu <i>et al.</i> , 2017; Zhang <i>et al.</i> , 2019), capable of retaining and interacting with subunits of the actin-related protein 2/3 complex (Mu <i>et al.</i> , 2016a; Mu <i>et al.</i> , 2016b), and involved in triggering apoptosis via activation of the JNK pathway (Zhang <i>et al.</i> , 2019) and enhancing heterologous gene expression (Cai <i>et al.</i> , 2012)
V-Ubi	35	Mut (Reilly and Guarino, 1996)	Del (Ono <i>et al.</i> , 2012)	Not essential (Reilly and Guarino, 1996); May block destruction of viral proteins by the host degradative pathway (Haas <i>et al.</i> , 1996) and confers a slight growth advantage under certain conditions (Reilly and Guarino, 1996);
39K/PP31	36	Mut (Broussard <i>et al.</i> , 1996a; Broussard <i>et al.</i> , 1996b; Yan <i>et al.</i> , 2019), Del (Yamagishi <i>et al.</i> , 2007)	Del (Gomi <i>et al.</i> , 1997; Ono <i>et al.</i> , 2012; Yan <i>et al.</i> , 2019)	Important for late gene transcription (Gomi <i>et al.</i> , 1997; Yamagishi <i>et al.</i> , 2007; Yan <i>et al.</i> , 2019), BV production, and abnormal formation of the virogenic stroma (Gomi <i>et al.</i> , 1997) as a DNA binding protein associated with virogenic stroma (Guarino <i>et al.</i> , 1992), which is stimulated by IE1 (Guarino and Summers, 1986) and required for transient DNA replication (Kool <i>et al.</i> , 1994; Lu and Miller, 1995)
LEF11	37	Del (Dong <i>et al.</i> , 2017; Dong <i>et al.</i> , 2015; Lin and Blissard, 2002b)	Del (Ono <i>et al.</i> , 2012)	Essential for viral DNA replication and virion production (Dong <i>et al.</i> , 2015; Lin and Blissard, 2002b; Ono <i>et al.</i> , 2012) and capable of forming oligomers and interacting with host proteins (Dong <i>et al.</i> , 2017)
AC38	38	Del (Ge <i>et al.</i> , 2007)	Del (Ono <i>et al.</i> , 2012)	Important for BV production as a ADP-ribose pyrophosphatase (Chen <i>et al.</i> , 2013a; Ge <i>et al.</i> , 2007)
P43	39	Del (Yu and Carstens, 2011)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV production (Ono <i>et al.</i> , 2012; Yu and Carstens, 2011)
P47	40	Mut (Partington <i>et al.</i> , 1990)	Del (Ono <i>et al.</i> , 2012)	Essential (Ono <i>et al.</i> , 2012) and involved in very late gene expression (Partington <i>et al.</i> , 1990) as a potential component of the virus-induced RNA polymerase (Jin <i>et al.</i> , 1998)
LEF12	41	Del (Guarino <i>et al.</i> , 2002)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV but beneficial the late gene expression and BV production (Guarino <i>et al.</i> , 2002; Rapp <i>et al.</i> , 1998)
GTA	42	Del (Tong <i>et al.</i> , 2021)	Del (Katsuma <i>et al.</i> , 2008a; Ono <i>et al.</i> , 2012)	Dispensable for BV (Katsuma <i>et al.</i> , 2008a; Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021) and ODV but beneficial for killing infected larvae (Katsuma <i>et al.</i> , 2008a)
AC43	43	Del (Tao <i>et al.</i> , 2013b)	Del (Katsuma and Shimada, 2009; Ono <i>et al.</i> , 2012)	Dispensable for BV but for OB production (Katsuma and Shimada, 2009; Ono <i>et al.</i> , 2012; Tao <i>et al.</i> , 2013b), and possibly involved in efficient late and very late gene expression, the morphogenesis of occlusion body, and the assembly of virions occluded in occlusion bodies (Tao <i>et al.</i> , 2013b).
AC44	44	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC45	45	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> ,

					2012; Tong <i>et al.</i> , 2021); may be required for the expression of AC41 (Li <i>et al.</i> , 1999)
ODV-E66	46	Del (Xiang <i>et al.</i> , 2011b)	Del (Ono <i>et al.</i> , 2012)	Del (Hou <i>et al.</i> , 2019)	Dispensable for BV replication but involved in degrading larval peritrophic membrane to facilitate baculovirus oral infection (Hou <i>et al.</i> , 2019; Xiang <i>et al.</i> , 2011b) as the major ODV envelope protein (Hong <i>et al.</i> , 1994) exhibiting chondroitinase activities with distinct substrate specificity (Sugiura <i>et al.</i> , 2011)
AC47	47	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012; Zhang <i>et al.</i> , 2014)		Dispensable for BV (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021; Zhang <i>et al.</i> , 2014) and involved in regulating viral transcription (Zhang <i>et al.</i> , 2014)
AC48	48	Del (Tong <i>et al.</i> , 2021)			Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Tong <i>et al.</i> , 2021)
PCNA	49	Mut (Iwahori <i>et al.</i> , 2002), Del (Iwahori <i>et al.</i> , 2004)			Dispensable for BV (Iwahori <i>et al.</i> , 2002), involved in the expression of late genes and BV production (Crawford and Miller, 1988; Fu <i>et al.</i> , 2018; Iwahori <i>et al.</i> , 2004), and associated with OEV-EC27 and viral capsid (Belyavskiy <i>et al.</i> , 1998)
LEF8	50	Mut (Titterington <i>et al.</i> , 2003)	Del (Ioannidis <i>et al.</i> , 2016; Ono <i>et al.</i> , 2012)		Essential (Ioannidis <i>et al.</i> , 2016; Ono <i>et al.</i> , 2012) as a potential component of the virus-induced RNA polymerase (Iorio <i>et al.</i> , 1998; Jin <i>et al.</i> , 1998) and containing a conserved motif of (Passarelli <i>et al.</i> , 1994) which is required for the late gene expression (Titterington <i>et al.</i> , 2003)
AC51	51	Del (Qiu <i>et al.</i> , 2019; Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2018b)		Essential for nucleocapsid egress from the nucleus and BV/ODV production (Qiu <i>et al.</i> , 2019; Shen <i>et al.</i> , 2018b; Tong <i>et al.</i> , 2021); ODV nucleocapsid and envelope protein (Wang <i>et al.</i> , 2002) as a BV-associated (Wang <i>et al.</i> , 2002) proteins containing RNA recognition motif (Shen <i>et al.</i> , 2018b; Wang <i>et al.</i> , 2002)
AC52	52	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012; Tian <i>et al.</i> , 2009b)		Important for BV production, ODV envelopment and polyhedra formation in the nucleus (Ono <i>et al.</i> , 2012; Tian <i>et al.</i> , 2009b; Tong <i>et al.</i> , 2021)
AC53	53	Del (Liu <i>et al.</i> , 2008)	Del (Ono <i>et al.</i> , 2012)		Essential for nucleocapsid assembly and virus production but not DNA replication (Liu <i>et al.</i> , 2008)
LEF10	53a	Mut (Xu <i>et al.</i> , 2016), Del (Xu <i>et al.</i> , 2016)	Del (Ono <i>et al.</i> , 2012; Yu <i>et al.</i> , 2013)		Essential and involved in DNA replication, RNA transcription and expression of the viral early gene (Nan <i>et al.</i> , 2019; Ono <i>et al.</i> , 2012; Xu <i>et al.</i> , 2016; Yu <i>et al.</i> , 2013) as a prion (Nan <i>et al.</i> , 2019; Xu <i>et al.</i> , 2016)
VP1054	54	Del (Guan <i>et al.</i> , 2016; Marek <i>et al.</i> , 2013; Olszewski and Miller, 1997a)	Del (Ono <i>et al.</i> , 2012)		Essential for nucleocapsid formation and required for proper viral DNA encapsidation (Marek <i>et al.</i> , 2013; Olszewski and Miller, 1997a) and capable of interacting with BV/ODV-C42 and VP80 but not VP39 (Guan <i>et al.</i> , 2016)
AC55	55	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC56	56	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC57	57	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)	Del (Chen <i>et al.</i> , 2012b)	Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Chen <i>et al.</i> , 2012b; Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021) as an early gene (Chen <i>et al.</i> , 2012b)

AC58/59	58/59	Del (Tong <i>et al.</i> , 2021)	Mut (Guo <i>et al.</i> , 2010b), Del (Ono <i>et al.</i> , 2012; Zhang <i>et al.</i> , 2014)	Del (Zheng <i>et al.</i> , 2011)	Dispensable for BV and ODV but involved in viral transcription (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021; Zhang <i>et al.</i> , 2014; Zheng <i>et al.</i> , 2011) as a ChaB homologue, DNA-binding protein (Li <i>et al.</i> , 2006a; Li <i>et al.</i> , 2006b)
AC60	60	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021) as a ChaB homologue, DNA-binding protein (Li <i>et al.</i> , 2006a; Li <i>et al.</i> , 2006b);
FP25K	61	Mut (Braunagel <i>et al.</i> , 1999; Garretson <i>et al.</i> , 2016; Harrison and Summers, 1995; Kelly <i>et al.</i> , 2006), Del (Garretson <i>et al.</i> , 2016; Li <i>et al.</i> , 2015c; Rosas-Acosta <i>et al.</i> , 2001)	Mut (Katsuma <i>et al.</i> , 1999), Del (Katsuma <i>et al.</i> , 2009b; Nakanishi <i>et al.</i> , 2010; Ono <i>et al.</i> , 2012)	Mut (Lua and Reid, 2003), Del (Braunagel <i>et al.</i> , 1999)	Important for BV and ODV (Braunagel <i>et al.</i> , 2009; Katsuma <i>et al.</i> , 2009b; Kelly <i>et al.</i> , 2006; Rosas-Acosta <i>et al.</i> , 2001) and capable of regulating responsible for the regulation of BV/ODV (Braunagel <i>et al.</i> , 1999; Harrison and Summers, 1995; Li <i>et al.</i> , 2015c; Wu <i>et al.</i> , 2005) and involved in ODV membrane protein transport into nucleus and interacting with host importin α -16 (Saksena <i>et al.</i> , 2006), multifunctional
LEF9	62	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012; Zhang <i>et al.</i> , 2016),		Essential for BV and ODV production but not essential for viral genome replication (Ono <i>et al.</i> , 2012; Zhang <i>et al.</i> , 2016) (Tong <i>et al.</i> , 2021); significantly affects viral gene transcription and expression (Zhang <i>et al.</i> , 2016) and may be a components of the virus-induced RNA polymerase (Iorio <i>et al.</i> , 1998; Jin <i>et al.</i> , 1998)
AC63	63	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021) as an early protein associated with BV envelope (Tian <i>et al.</i> , 2009a)
GP37	64	Del (Cheng <i>et al.</i> , 2001)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV and ODV (Cheng <i>et al.</i> , 2001) as envelope protein of both BV and ODV (Li <i>et al.</i> , 2003) capable of binding host chitin to facilitate ODVs cross host peritrophic membranes and fuse with epithelia (Liu <i>et al.</i> , 2019)
DNA POL	65	Del (Chen <i>et al.</i> , 2018; Feng and Krell, 2014; Vanarsdall <i>et al.</i> , 2005), Mut (Chen <i>et al.</i> , 2018; Feng and Krell, 2014)	Del (Ono <i>et al.</i> , 2012)		Essential for BV and ODV DNA replication (Chen <i>et al.</i> , 2016; Chen <i>et al.</i> , 2019; Chen <i>et al.</i> , 2018; Feng and Krell, 2014; Vanarsdall <i>et al.</i> , 2005)
AC66	66	Del (Ke <i>et al.</i> , 2008)	Del (Katsuma <i>et al.</i> , 2008b; Ono <i>et al.</i> , 2012; Zhang <i>et al.</i> , 2012b)		Important for nuclear egress of nucleocapsid and BV & ODV production but not essential for AcMNPV (Ke <i>et al.</i> , 2008) as a viral desmoplakin; however, the deletion of its homologue in BmNPV does not affect assembly and release of nucleocapsids, but results in non-infectious BV and defective polyhedral (Zhang <i>et al.</i> , 2012b)
LEF3	67	Mut (Chen and Carstens, 2005), Del (Nie <i>et al.</i> , 2012)	Del (Ono <i>et al.</i> , 2012)		Essential for DNA replication (Hang <i>et al.</i> , 1995; Li <i>et al.</i> , 1993; Mikhailov <i>et al.</i> , 2005; Mikhailov <i>et al.</i> , 2006) and virion production (Ono <i>et al.</i> , 2012), as ssDNA binding protein involved in interacting with Alkaline Nuclease (Hang <i>et al.</i> , 1995; Li <i>et al.</i> , 1993; Mikhailov <i>et al.</i> , 2005; Mikhailov <i>et al.</i> , 2006), and required (Chen and Carstens, 2005; Wu and Carstens, 1998) but not essential (Nie <i>et al.</i> , 2012) for helicase transport into the nuclear
PIF6	68	Del (Li <i>et al.</i> , 2011; Li <i>et al.</i> , 2008; Nie <i>et al.</i> , 2012;	Del (Ono <i>et al.</i> , 2012; Xu <i>et al.</i> , 2008)		<i>Per os</i> infectivity factor (Nie <i>et al.</i> , 2012) as a component of PIF complex (Peng <i>et al.</i> , 2012; Wang <i>et al.</i> , 2019b)

		Wang <i>et al.</i> , 2019b)			
AC69	69	Del (Ke <i>et al.</i> , 2011; Wu and Guarino, 2003)	Del (Ono <i>et al.</i> , 2012)		Essential (Ke <i>et al.</i> , 2011; Wu and Guarino, 2003) with cap O-dependent methyltransferase activity (Wu and Guarino, 2003)
HCF-1	70	Mut (Wilson <i>et al.</i> , 2005), Del (Lu and Miller, 1996)			Dispensable for BV as a host cell-specific factor containing a RING finger domain required for its self-association and gene repression activity (Wilson <i>et al.</i> , 2005) and affecting virus replicate rate in tissue-specific manner (Lu and Miller, 1996; Tachibana <i>et al.</i> , 2017)
IAP2	71	Del (Griffiths <i>et al.</i> , 1999)	Del (Ono <i>et al.</i> , 2012)		May be important for BV and ODV (Griffiths <i>et al.</i> , 1999), involved in antiapoptosis (Clem and Miller, 1994)
AC72	72	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC73	73	Del (Shao <i>et al.</i> , 2019)	Del (Ono <i>et al.</i> , 2012)		Essential for BV production in BmNPV (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021) but not for BV/ODV production in AcMNPV (Shao <i>et al.</i> , 2019)
AC74	74	Del (Tong <i>et al.</i> , 2021)	Del (Guo <i>et al.</i> , 2010a; Ono <i>et al.</i> , 2012)		Dispensable for BV (Guo <i>et al.</i> , 2010a) as a late protein localizes to both the cytoplasm and nucleus of infected cells (Du <i>et al.</i> , 2006)
AC75	75	Del (Guo <i>et al.</i> , 2017; Shi <i>et al.</i> , 2018)	Del (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2009b)		Essential (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2009b; Shi <i>et al.</i> , 2018) as a BV and ODV protein localized on nuclear membrane and in intranuclear ring zone of infected cells (Shen <i>et al.</i> , 2009b) and required in nuclear egress of nucleocapsids and intranuclear microvesicle formation (Guo <i>et al.</i> , 2017; Shen <i>et al.</i> , 2009b; Shi <i>et al.</i> , 2018)
AC76	76	Del (Hu <i>et al.</i> , 2010; Zhang <i>et al.</i> , 2017)	Del (Ono <i>et al.</i> , 2012)		Essential for BV and ODV production as a dimeric type II integral membrane protein involved in intranuclear microvesicle formation (Hu <i>et al.</i> , 2010; Wei <i>et al.</i> , 2014)
VLF-1	77	Mut (Yang and Miller, 1998), Del (Li <i>et al.</i> , 2005; Todd <i>et al.</i> , 1996; Vanarsdall <i>et al.</i> , 2006)	Del (Ono <i>et al.</i> , 2012)		Essential for capsid assembly (Li <i>et al.</i> , 2005; Todd <i>et al.</i> , 1996; Vanarsdall <i>et al.</i> , 2006) and maturation of DNA (Todd <i>et al.</i> , 1996; Vanarsdall <i>et al.</i> , 2006) as a DNA binding protein (Mikhailov and Rohrmann, 2002b) involved in regulation of very late gene expression (Todd <i>et al.</i> , 1996; Yang and Miller, 1999); the integrase activity is not essential (Yang and Miller, 1998);
AC78	78	Del (Chen <i>et al.</i> , 2015)	Del (Chen <i>et al.</i> , 2015; Ono <i>et al.</i> , 2012)	Del (Huang <i>et al.</i> , 2014)	Important for BV (Chen <i>et al.</i> , 2015; Huang <i>et al.</i> , 2014; Ono <i>et al.</i> , 2012), ODV formation and <i>per os</i> infection (Chen <i>et al.</i> , 2015) but may be not essential (Li <i>et al.</i> , 2014a; Tao <i>et al.</i> , 2013a)
AC79	79	Del (Wu and Passarelli, 2012)	Del (Li <i>et al.</i> , 2019a; Ono <i>et al.</i> , 2012)		Important but not essential for BV production and involved in the repair of ultraviolet-induced DNA damage (Li <i>et al.</i> , 2019a; Ono <i>et al.</i> , 2012; Tang <i>et al.</i> , 2015; Tang <i>et al.</i> , 2017; Wu and Passarelli, 2012); mainly exists in mainly a tetrameric form and contains important NLS motif (Li <i>et al.</i> , 2019a).
GP41	80	Mut (Li <i>et al.</i> , 2018b; Olszewski and Miller, 1997b), Del (Li <i>et al.</i> , 2018b)	Del (Ono <i>et al.</i> , 2012)		Essential for BV and ODV and required for nucleocapsid egress from nucleus and important for BV production (Li <i>et al.</i> , 2018b; Olszewski and Miller, 1997b; Ono <i>et al.</i> , 2012)
AC81	81	Del (Dong <i>et al.</i> , 2016)	Del (Ge <i>et al.</i> , 2008; Ono <i>et al.</i> , 2012; Shi <i>et al.</i> , 2015)		Important for BV and ODV production (Ge <i>et al.</i> , 2008; Shi <i>et al.</i> , 2015) and ODV envelopment (Dong <i>et al.</i> , 2016)

Telokin	82	Del (Gauthier <i>et al.</i> , 2012)	Del (Iwanaga <i>et al.</i> , 2002; Ono <i>et al.</i> , 2012)		Essential for AcMNPV (Gauthier <i>et al.</i> , 2012) but not essential (Iwanaga <i>et al.</i> , 2002; Ono <i>et al.</i> , 2012) in BmNPV as a telokin homologue (Raynes <i>et al.</i> , 1994), BV-associated protein involved in entry and budding from cells (Iwanaga <i>et al.</i> , 2002); Need clarification.
VP91/PIF8	83	Mut (Huang <i>et al.</i> , 2017; Zhu <i>et al.</i> , 2013), Del (Huang <i>et al.</i> , 2017; Javed <i>et al.</i> , 2017; Wang <i>et al.</i> , 2019b; Zhu <i>et al.</i> , 2013)	Del (Ono <i>et al.</i> , 2012; Xiang <i>et al.</i> , 2013)	Mut (Zhou <i>et al.</i> , 2019), Del (Zhou <i>et al.</i> , 2019)	Essential for BV and ODV production and associated with both ODV envelope and nucleocapsid (Russell and Rohrmann, 1997) and involved in the interaction with IE1 for enhancing transcription (Lu <i>et al.</i> , 1998); a <i>per os</i> infectivity factor (Huang <i>et al.</i> , 2017; Javed <i>et al.</i> , 2017; Xiang <i>et al.</i> , 2013; Zhou <i>et al.</i> , 2019; Zhu <i>et al.</i> , 2013) as a component of PIF complex (Peng <i>et al.</i> , 2012; Wang <i>et al.</i> , 2019b);
AC84	84	Del (Tong <i>et al.</i> , 2021)			Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Tong <i>et al.</i> , 2021)
AC85	85	Del (Tong <i>et al.</i> , 2021)			Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Tong <i>et al.</i> , 2021)
PNK/PNL	86	Del (Durantel <i>et al.</i> , 1998)			Dispensable for viral replication as an immediate early gene (Durantel <i>et al.</i> , 1998)
AC87	87	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
CG30	88	Del (Passarelli and Miller, 1994b)	Mut (Ishihara <i>et al.</i> , 2013), Del (Ishihara <i>et al.</i> , 2013; Ono <i>et al.</i> , 2012; Zhang <i>et al.</i> , 2012a)		Important for BV (Ishihara <i>et al.</i> , 2013; Ono <i>et al.</i> , 2012; Passarelli and Miller, 1994b; Zhang <i>et al.</i> , 2012a) and containing a Zinc Finger and a leucine zipper domain (Thiem and Miller, 1989a)
VP39	89	Mut (Katsuma and Kokusho, 2017), Del (Katsuma and Kokusho, 2017)	Del (Ono <i>et al.</i> , 2012)		Essential for viral propagation (Katsuma and Kokusho, 2017) and involved in actin rearrangement and polymerization (Charlton and Volkman, 1993) as the major capsid protein (Pearson <i>et al.</i> , 1988; Thiem and Miller, 1989b) with a conserved glycine residue essential for correct nucleocapsid assembly, viral DNA packaging, and viral gene expression, especially of very late genes
LEF4	90	Mut (Martins and Shuman, 2001), Del (Knebel-Morsdorf <i>et al.</i> , 2006)	Del (Ono <i>et al.</i> , 2012)		Essential for viral replication (Knebel-Morsdorf <i>et al.</i> , 2006; Ono <i>et al.</i> , 2012) with RNA 5'-triphosphatase and ATPase and guanylyltransferase activities (Gross and Shuman, 1998b; Guarino <i>et al.</i> , 1998; Jin <i>et al.</i> , 1998; Martins and Shuman, 2001; Schrawat <i>et al.</i> , 2002)
AC91	91	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2009a)		Dispensable for BV (Ono <i>et al.</i> , 2012; Shen <i>et al.</i> , 2009a; Tong <i>et al.</i> , 2021)
P33	92	Del (Andersen <i>et al.</i> , 2011; Clem <i>et al.</i> , 2014; Kuang <i>et al.</i> , 2017; Wu and Passarelli, 2010)	Del (Ono <i>et al.</i> , 2012)	Del (Kuang <i>et al.</i> , 2016)	Essential for efficient BV production and multiply-ODV formation (Andersen <i>et al.</i> , 2011; Clem <i>et al.</i> , 2014; Kuang <i>et al.</i> , 2017; Kuang <i>et al.</i> , 2016; Wu and Passarelli, 2010) as a flavin adenine dinucleotide-linked sulfhydryl oxidase which enhance p53-mediated apoptosis (Prikhod'ko <i>et al.</i> , 1999b) through interacting with host P53 protein (Wu <i>et al.</i> , 2013);
AC93	93	Mut (Liu <i>et al.</i> , 2020), Del (Liu <i>et al.</i> , 2020; Yuan <i>et al.</i> , 2011)	Del (Ono <i>et al.</i> , 2012)	Del (Li <i>et al.</i> , 2014b)	Essential for ODV envelopment and nuclear egress of nucleocapsids but not DNA replication (Li <i>et al.</i> , 2014b; Liu <i>et al.</i> , 2020; Yuan <i>et al.</i> , 2011) and involved in interactions with the endosomal sorting complex required for transport (ESCRT) pathway complex (Liu <i>et al.</i> , 2020; Yue <i>et al.</i> , 2018)

ODV-E25	94	Del (Chen <i>et al.</i> , 2012a; Chen <i>et al.</i> , 2013b; Luo <i>et al.</i> , 2013a), Mut (Luo <i>et al.</i> , 2013b)	Del (Ono <i>et al.</i> , 2012)		Important for BV infectivity and ODV formation but not DNA replication (Chen <i>et al.</i> , 2012a; Chen <i>et al.</i> , 2013b; Luo <i>et al.</i> , 2013a) as a both BV and ODV associated (Wang <i>et al.</i> , 2010c) protein which may exist and function as dimmers or polymers (Luo <i>et al.</i> , 2013b) and may be regulated by host microRNA (Luo <i>et al.</i> , 2013a)
Helicase	95	Del (Argaud <i>et al.</i> , 1998; Liu and Carstens, 1999)	Del(Ono <i>et al.</i> , 2012)		Essential with DNA and ATP binding activities (McDougal and Guarino, 2001) and involved in DNA replication (Kool <i>et al.</i> , 1994; Liu and Carstens, 1999) and viral host range (Argaud <i>et al.</i> , 1998; Croizier <i>et al.</i> , 1994)
PIF4	96	Del (Fang <i>et al.</i> , 2009a; Wang <i>et al.</i> , 2019b)	Del (Dong <i>et al.</i> , 2014; Ono <i>et al.</i> , 2012)	Del (Huang <i>et al.</i> , 2012)	<i>Per os</i> infectivity factor (Dong <i>et al.</i> , 2014; Fang <i>et al.</i> , 2009a; Huang <i>et al.</i> , 2012) as a component of PIF complex (Dong <i>et al.</i> , 2014; Huang <i>et al.</i> , 2012; Peng <i>et al.</i> , 2012; Wang <i>et al.</i> , 2019b)
AC97	97	Del (Tong <i>et al.</i> , 2021)			Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Tong <i>et al.</i> , 2021)
38K	98	Mut (Lai <i>et al.</i> , 2018), Del (Lai <i>et al.</i> , 2018; Wu <i>et al.</i> , 2006)	Del (Ono <i>et al.</i> , 2012)		Essential for nucleocapsid assembly (Wu <i>et al.</i> , 2006) and involved in late gene expression (Hefferon, 2003), specific dephosphorylation of the core protein P6.9 (Lai <i>et al.</i> , 2018) and interactions with VP1054, VP39, VP80, and itself (Wu <i>et al.</i> , 2008)
LEF5	99	Mut (Chen <i>et al.</i> , 2020; Harwood <i>et al.</i> , 1998), Del (Chen <i>et al.</i> , 2020; Harwood <i>et al.</i> , 1998; Su <i>et al.</i> , 2011)	Del (Ono <i>et al.</i> , 2012)		Essential for BV production (Ono <i>et al.</i> , 2012; Su <i>et al.</i> , 2011), and containing a nucleolar localization signal (NoLS) (Chen <i>et al.</i> , 2020) and a zinc ribbon domain which is required for maximal late transcription activity (Harwood <i>et al.</i> , 1998)
P6.9	100	Mut (Wang <i>et al.</i> , 2010a), Del (Wang <i>et al.</i> , 2010a)	Del (Ono <i>et al.</i> , 2012)		Essential for of BV/ODV production but not for viral DNA replication (Wang <i>et al.</i> , 2010a) as a nucleocapsid core DNA-binding protein (Kelly <i>et al.</i> , 1983; Tweeten <i>et al.</i> , 1980; Wilson <i>et al.</i> , 1987; Wilson and Price, 1988) whose phosphorylation states are regulated by PKIP and 38K proteins (Lai <i>et al.</i> , 2018; Lai <i>et al.</i> , 2020; Li <i>et al.</i> , 2015a; Liu <i>et al.</i> , 2012)
BV/ODV-C42	101	Del (Vanarsdall <i>et al.</i> , 2007b)	Del (Ono <i>et al.</i> , 2012)		Essential (Vanarsdall <i>et al.</i> , 2007b), and involved in recruiting P78/83 to nucleus (Wang <i>et al.</i> , 2008d), nuclear actin polymerization (Li <i>et al.</i> , 2010) and nucleocapsid morphogenesis (Vanarsdall <i>et al.</i> , 2007b)
P12	102	Mut (Gandhi <i>et al.</i> , 2012), Del (Gandhi <i>et al.</i> , 2012)	Del (Ono <i>et al.</i> , 2012)		Essential for BV production (Gandhi <i>et al.</i> , 2012; Ono <i>et al.</i> , 2012) and mediating nuclear localization of G-Actin (Gandhi <i>et al.</i> , 2012; Hepp <i>et al.</i> , 2018) through modulating BV/ODV-C42 ubiquitination (Zhang <i>et al.</i> , 2018)
P48	103	Del (Wang <i>et al.</i> , 2019c; Yuan <i>et al.</i> , 2008)	Del (Ono <i>et al.</i> , 2012)		Important for BV production, intranuclear microvesicle and ODV envelopment (Ono <i>et al.</i> , 2012; Wang <i>et al.</i> , 2019c; Yuan <i>et al.</i> , 2008)
VP80	104	Del (Marek <i>et al.</i> , 2011)	Del (Ono <i>et al.</i> , 2012; Tang <i>et al.</i> , 2008)		Essential for BV production, nucleocapsid maturation but not for assembly of nucleocapsids (Marek <i>et al.</i> , 2011; Tang <i>et al.</i> , 2008)
HE65	105	Del (Gandhi <i>et al.</i> , 2012)	Del (Ono <i>et al.</i> , 2012)		Not essential and deletion may have no or mild effect on BV and ODV production (Gandhi <i>et al.</i> , 2012; Ono <i>et al.</i> , 2012) as an early gene (Becker and Knebel-Morsdorf, 1993) transcriptionally actiated by IE1 (Kremer and Knebel-Morsdorf, 1998)

AC106/107	106/107	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Essential for BV production (Ono <i>et al.</i> , 2012) (Tong <i>et al.</i> , 2021)
PIF9	108	Del (Boogaard <i>et al.</i> , 2019; Wang <i>et al.</i> , 2019b)	Del (Ono <i>et al.</i> , 2012; Tang <i>et al.</i> , 2013)		<i>Per os</i> infectivity factor (Boogaard <i>et al.</i> , 2019; Simon <i>et al.</i> , 2012; Tang <i>et al.</i> , 2013) as a component of PIF complex (Boogaard <i>et al.</i> , 2019; Wang <i>et al.</i> , 2019b)
ODV-EC43	109	Del (Alfonso <i>et al.</i> , 2017; Alfonso <i>et al.</i> , 2012; Fang <i>et al.</i> , 2009c; Lehiy <i>et al.</i> , 2013; Lin <i>et al.</i> , 2009)	Del (Ono <i>et al.</i> , 2012)		Essential for nucleocapsid formation but not viral DNA replication (Fang <i>et al.</i> , 2009c; Lin <i>et al.</i> , 2009), involved in nucleocapsid transport and envelopment in infected cell (Alfonso <i>et al.</i> , 2012; Lehiy <i>et al.</i> , 2013); may be tegument protein (Fang <i>et al.</i> , 2003)
PIF7	110	Del (Jiantao <i>et al.</i> , 2016; Wang <i>et al.</i> , 2019b)	Del (Ono <i>et al.</i> , 2012)		Essential for <i>per os</i> infectivity factor (Jiantao <i>et al.</i> , 2016) as a component of PIF complex (Wang <i>et al.</i> , 2019b)
AC111	111	Del (Li <i>et al.</i> , 2018a)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV and ODV; affects <i>per os</i> infectivity in a host-dependent manner (Li <i>et al.</i> , 2018a; Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC112/113	112/113	Del (Tong <i>et al.</i> , 2021)			Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Tong <i>et al.</i> , 2021)
AC114	114	Del (Wei <i>et al.</i> , 2012)	Del (Ono <i>et al.</i> , 2012)		Not essential for BV and ODV production (Ono <i>et al.</i> , 2012; Wei <i>et al.</i> , 2012) as a late gene, which distributed both in the cytoplasm and in the nucleus also a component of BmNPV ODV (Liang <i>et al.</i> , 2010);
PIF3	115	Del (Ohkawa <i>et al.</i> , 2005; Peng <i>et al.</i> , 2010; Wang <i>et al.</i> , 2019b)	Del (Ono <i>et al.</i> , 2012)	Del (Song <i>et al.</i> , 2008)	Essential for <i>per os</i> infectivity factor (Ohkawa <i>et al.</i> , 2005) as a component of PIF complex (Peng <i>et al.</i> , 2012; Peng <i>et al.</i> , 2010; Wang <i>et al.</i> , 2019b)
AC116	116	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC117	117	Del (Tong <i>et al.</i> , 2021)	Del (Gomi <i>et al.</i> , 1999; Ono <i>et al.</i> , 2012)		Dispensable for BV and ODV (Gomi <i>et al.</i> , 1999; Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021); deletion has a slight effect on the motility of infected larvae (Gomi <i>et al.</i> , 1999)
AC118	118	Del (Tong <i>et al.</i> , 2021)			Dispensable for BV; deletion may have mild effect on BV and ODV production (Tong <i>et al.</i> , 2021)
PIF1	119	Del (Clavijo <i>et al.</i> , 2009; Ohkawa <i>et al.</i> , 2005; Peng <i>et al.</i> , 2010; Wang <i>et al.</i> , 2019b)	Del (Ono <i>et al.</i> , 2012)	Del (Song <i>et al.</i> , 2008)	Essential for <i>per os</i> infectivity factor (Harrison <i>et al.</i> , 2010; Ohkawa <i>et al.</i> , 2005; Sparks <i>et al.</i> , 2011; Xiang <i>et al.</i> , 2011a) as a component of PIF complex (Peng <i>et al.</i> , 2012; Peng <i>et al.</i> , 2010; Wang <i>et al.</i> , 2019b)
AC120	120	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC121	121	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021); a transcriptional activator of some early genes, including IE1 and PP31, but not late genes (Li <i>et al.</i> , 1999) (Tong <i>et al.</i> , 2021)
AC122	122	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)		Dispensable for BV; deletion may have mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)

PK2	123	Mut (Li and Miller, 1995a), Del (Dever <i>et al.</i> , 1998)	Del (Katsuma <i>et al.</i> , 2008b; Ono <i>et al.</i> , 2012)	Dispensable for BV (Dever <i>et al.</i> , 1998; Katsuma <i>et al.</i> , 2008b; Ono <i>et al.</i> , 2012) but involved in interaction with and regulating eIF2 α family kinases for inactivating a host stress response to virus infection (Dever <i>et al.</i> , 1998; Li <i>et al.</i> , 2015b)
AC124	124	Del (Fang <i>et al.</i> , 2019)	Del (Chen <i>et al.</i> , 2014; Gomi <i>et al.</i> , 1999; Ono <i>et al.</i> , 2012)	Dispensable for BV (Fang <i>et al.</i> , 2019; Ono <i>et al.</i> , 2012) but in conflict with other study on Bm101 (Chen <i>et al.</i> , 2014) and involved in regulating the expression of chitinase gene (Fang <i>et al.</i> , 2019).
LEF7	125	Del (Chen and Thiem, 1997; Mitchell <i>et al.</i> , 2013)	Del (Gomi <i>et al.</i> , 1999; Gomi <i>et al.</i> , 1997; Ono <i>et al.</i> , 2012)	Dispensable for BV, involved in transient DNA replication and capable of suppressing DNA damage response to enhance virus multiplication (Chen and Thiem, 1997; Gomi <i>et al.</i> , 1999; Lu and Miller, 1995; Mitchell <i>et al.</i> , 2013; Morris <i>et al.</i> , 1994)
V-ChiA	126	Mut (Hodgson <i>et al.</i> , 2011; Saville <i>et al.</i> , 2004; Saville <i>et al.</i> , 2002), Del (Hawtin <i>et al.</i> , 1997)	Mut (Daimon <i>et al.</i> , 2007), Del (Gomi <i>et al.</i> , 1999; Katsuma <i>et al.</i> , 2008b; Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV production but crucial for V-CATH processing (Daimon <i>et al.</i> , 2007; Hom and Volkman, 2000) and the liquefaction of infected larvae (Daimon <i>et al.</i> , 2007; Hawtin <i>et al.</i> , 1997), with the interaction with V-CATH (Hodgson <i>et al.</i> , 2011)
V-CATH	127	Del (Hawtin <i>et al.</i> , 1997)	Mut (Katsuma <i>et al.</i> , 2009a), Del (Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV production but important for the liquefaction of infected larvae and viral horizontal transmission (D'Amico <i>et al.</i> , 2013; Hawtin <i>et al.</i> , 1997) with the interaction with V-ChiA (Hodgson <i>et al.</i> , 2011) and involvement in actin proteolysis (Lanier <i>et al.</i> , 1996);
GP64	128	Mut (Jarvis <i>et al.</i> , 1998; Li and Blissard, 2008; Li and Blissard, 2009a; Li and Blissard, 2009b; Li and Blissard, 2010; Monsma and Blissard, 1995; Zhang <i>et al.</i> , 2003; Zhou and Blissard, 2008), Del (Li and Blissard, 2010; Monsma <i>et al.</i> , 1996)	Del (Ono <i>et al.</i> , 2012)	Essential for cell-cell transmission as a BV major envelope fusion protein capable of interacting with host cholesterol transport protein (Kingsley <i>et al.</i> , 1999; Li <i>et al.</i> , 2019b; Monsma <i>et al.</i> , 1996; Whitford <i>et al.</i> , 1989)
P24	129	Mut (Schetter <i>et al.</i> , 1990)	Del (Gomi <i>et al.</i> , 1999; Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV (Gombart <i>et al.</i> , 1989; Gomi <i>et al.</i> , 1999; Schetter <i>et al.</i> , 1990); affects pathogenesis in larvae specifically (Gomi <i>et al.</i> , 1999)
GP16	130	Del (Yang <i>et al.</i> , 2014a)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV (Ono <i>et al.</i> , 2012; Yang <i>et al.</i> , 2014a) as a glycosylated protein localized near the nuclear membrane in the cytoplasm involved in membrane trafficking of BV (Gross <i>et al.</i> , 1993; Yang <i>et al.</i> , 2014a)
PEP	131	Del (Bischoff and Slavicek, 1999)	Del (Ono <i>et al.</i> , 2012)	Dispensable for polyhedral potency and BV production (Bischoff and Slavicek, 1999) as the major polyhedron membrane protein (Gross <i>et al.</i> , 1994; Whitt and Manning, 1988) which facilitates larval liquefaction and lysis of cultured cell (Bischoff and Slavicek, 1999)
AC132	132	Del (Fang <i>et al.</i> , 2016; Yang <i>et al.</i> , 2014b)	Del (Ono <i>et al.</i> , 2012)	Essential for BV production, nuclear entry of nucleocapsid and multiply enveloped ODV morphogenesis (Fang <i>et al.</i> , 2016; Ono <i>et al.</i> , 2012; Yang <i>et al.</i> , 2014b)
ALK-EXO	133	Mut (Li and Rohrmann, 2000), Del (Okano <i>et al.</i> ,	Del (Ono <i>et al.</i> , 2012)	Essential as baculovirus alkaline nuclease (Li and Rohrmann, 2000; Okano <i>et al.</i> , 2004) involved in the maturation of viral DNA or packaging of the DNA into virions (Okano <i>et al.</i> , 2004; Okano <i>et</i>

		2004; Okano <i>et al.</i> , 2007)		<i>al.</i> , 2007),	
94K	134	Del (Clem <i>et al.</i> , 1994)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Clem <i>et al.</i> , 1994; Ono <i>et al.</i> , 2012)	
35K/P35	135	Mut (Kelly <i>et al.</i> , 2006), Del (Clem and Miller, 1993; Clem <i>et al.</i> , 1994)	Del (Makino <i>et al.</i> , 2018; Ono <i>et al.</i> , 2012)	Essential for antiapoptosis in semi-permissive cells, beneficial for BV production (Bertin <i>et al.</i> , 1996; Clem and Miller, 1993; Clem and Miller, 1994; LaCount <i>et al.</i> , 2000; Makino <i>et al.</i> , 2018; Miao <i>et al.</i> , 2016; Zoog <i>et al.</i> , 1999) and capable of suppressing host RNA interference antiviral response (Mehrabadi <i>et al.</i> , 2015)	
P26	136	Del (Simon <i>et al.</i> , 2008; Wang <i>et al.</i> , 2009)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV and auxiliary for virus transmission (Simon <i>et al.</i> , 2008); may act in concert with P74 and P10 to regulate the virion occlusion process (Wang <i>et al.</i> , 2009)	
P10	137	Mut (Graves <i>et al.</i> , 2019; Raza <i>et al.</i> , 2017), Del (Graves <i>et al.</i> , 2019; Raza <i>et al.</i> , 2017; van Oers <i>et al.</i> , 1993)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV but important as phosphorylation-regulated protein involved in occlusion body maturation and their release (Carpentier <i>et al.</i> , 2008; Graves <i>et al.</i> , 2019; Gross <i>et al.</i> , 1994; Lee <i>et al.</i> , 1996; Raza <i>et al.</i> , 2017; van Oers <i>et al.</i> , 1993) and capable of interacting with host tubulin (Patmanidi <i>et al.</i> , 2003)	
P74	138	Del (Faulkner <i>et al.</i> , 1997; Haas-Stapleton <i>et al.</i> , 2004; Kuzio <i>et al.</i> , 1989; Wang <i>et al.</i> , 2019b)	Del (Ono <i>et al.</i> , 2012)	Del (Song <i>et al.</i> , 2008; Yao <i>et al.</i> , 2004)	Essential for oral infection (Haas-Stapleton <i>et al.</i> , 2004; Ohkawa <i>et al.</i> , 2005; Song <i>et al.</i> , 2008; Wang <i>et al.</i> , 2019b) as a component of PIF complex (Peng <i>et al.</i> , 2012; Peng <i>et al.</i> , 2010; Wang <i>et al.</i> , 2019b)
ME53	139	Mut (Liu <i>et al.</i> , 2016); Del (de Jong <i>et al.</i> , 2009; Xi <i>et al.</i> , 2007)	Del (Ono <i>et al.</i> , 2012)	Important for optimal BV and ODV production as a major early-transcribed gene containing NLS motif (de Jong <i>et al.</i> , 2009; Knebel-Morsdorf <i>et al.</i> , 1993; Liu <i>et al.</i> , 2016; Xi <i>et al.</i> , 2007), which interacts with P64 and may be providing a scaffold that bridges the viral envelope and nucleocapsid (de Jong <i>et al.</i> , 2011)	
AC140	140	Del (Tong <i>et al.</i> , 2021)		Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Tong <i>et al.</i> , 2021)	
EXON0	141	Del (Dai <i>et al.</i> , 2004a; Fang <i>et al.</i> , 2007; Fang <i>et al.</i> , 2008; Fang <i>et al.</i> , 2009b)	Del (Ono <i>et al.</i> , 2012)	Important for efficient nuclear egress of nucleocapsid and production of BV (Dai <i>et al.</i> , 2004a; Fang <i>et al.</i> , 2007; Fang <i>et al.</i> , 2008; Fang <i>et al.</i> , 2009b) through a potential E3 ubiquitin ligase activity via interactions with viral ubiquitin and AC66 (Biswas <i>et al.</i> , 2018), capable of interacting with β -tubulin (Fang <i>et al.</i> , 2009b), itself and associated with BV/ODV-C42 and FP25 (Fang <i>et al.</i> , 2008)	
49K	142	Del (McCarthy <i>et al.</i> , 2008; Vanarsdall <i>et al.</i> , 2007b)	Del (Gandhi <i>et al.</i> , 2012; Ono <i>et al.</i> , 2012)	Essential for BV production and nucleocapsid maturation (McCarthy <i>et al.</i> , 2008; Vanarsdall <i>et al.</i> , 2007b; Yang <i>et al.</i> , 2008) and ODV envelopment (McCarthy <i>et al.</i> , 2008),	
ODV-E18	143	Del (McCarthy and Theilmann, 2008)	Del (Ono <i>et al.</i> , 2012)	Essential for BV production but not for DNA replication (McCarthy and Theilmann, 2008)	
ODV-EC27	144	Del (Vanarsdall <i>et al.</i> , 2007b)	Del (Ono <i>et al.</i> , 2012)	Essential for BV production and nucleocapsid formation (Vanarsdall <i>et al.</i> , 2007b) as a multifunctional viral cyclin (Belyavskiy <i>et al.</i> , 1998) and capable of interacting with BV/ODV-C42 (Braunagel <i>et al.</i> , 2001)	

AC145	145	Del (Lapointe <i>et al.</i> , 2004)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV but involved in affecting oral infectivity in a host-dependent manner (Lapointe <i>et al.</i> , 2004; Ono <i>et al.</i> , 2012)
AC146	146	Del (Dickison <i>et al.</i> , 2012)	Del (Ono <i>et al.</i> , 2012)	Essential for BV production as a late virion associated protein (Dickison <i>et al.</i> , 2012; Ono <i>et al.</i> , 2012) which accumulates within the nuclear region of infected cells (Ge <i>et al.</i> , 2009)
IE1	147	Mut (Olson <i>et al.</i> , 2002), Del (Stewart <i>et al.</i> , 2005)	Del (Ono <i>et al.</i> , 2012)	Essential as ie0-ie1 gene complex form (Stewart <i>et al.</i> , 2005), essential for baculovirus early replication (Schultz <i>et al.</i> , 2009) and required for expression of delayed-early gene (Guarino and Summers, 1986) as a DNA binding protein (Ito <i>et al.</i> , 2004; Leisy and Rohrmann, 2000; Nagamine <i>et al.</i> , 2005; Olson <i>et al.</i> , 2003) involved in late and very late gene expression (Passarelli and Miller, 1993), initiation of viral DNA replication and assembly of viral DNA replication factories (Okano <i>et al.</i> , 1999)
IE0	147-0	Del (Lu <i>et al.</i> , 2005; Stewart <i>et al.</i> , 2005)	Del (Ono <i>et al.</i> , 2012)	Essential as ie0-ie1 gene complex form (Lu <i>et al.</i> , 2005; Stewart <i>et al.</i> , 2005) but functional for transcriptional activation (Dai <i>et al.</i> , 2004b; Huijskens <i>et al.</i> , 2004; Stewart <i>et al.</i> , 2005) and viral replication (Lu <i>et al.</i> , 2003; Lu <i>et al.</i> , 2005)
PIF5	148	Del (Harrison <i>et al.</i> , 2010; Sparks <i>et al.</i> , 2011; Wang <i>et al.</i> , 2019b)	Del (Ono <i>et al.</i> , 2012; Xiang <i>et al.</i> , 2011a)	Essential for <i>per os</i> infectivity factor (Harrison <i>et al.</i> , 2010; Sparks <i>et al.</i> , 2011; Xiang <i>et al.</i> , 2011a) as a component of PIF complex (Peng <i>et al.</i> , 2012; Wang <i>et al.</i> , 2019b) but not essential for binding and fusion of ODV to the host midgut (Sparks <i>et al.</i> , 2011), and also associated with BV (Peng <i>et al.</i> , 2012; Wang <i>et al.</i> , 2010c)
AC149	149	Del (Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV; deletion may have no or mild effect on BV and ODV production (Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021)
AC150	150	Del (Lapointe <i>et al.</i> , 2004; Zhang <i>et al.</i> , 2005)	Del (Hao <i>et al.</i> , 2009; Huang <i>et al.</i> , 2019; Ono <i>et al.</i> , 2012)	Dispensable for BV but involved in the regulation of viral gene expression (Huang <i>et al.</i> , 2019) and oral infectivity in a host-dependent manner (Lapointe <i>et al.</i> , 2004)
IE2	151	Mut (Prikhod'ko <i>et al.</i> , 1999a; Prikhod'ko and Miller, 1998; Tung <i>et al.</i> , 2016), Del (Tung <i>et al.</i> , 2016)	Del (Ono <i>et al.</i> , 2012)	Important for BV and ODV production (Kool <i>et al.</i> , 1994; Prikhod'ko <i>et al.</i> , 1999a), involved in DNA replication, cell cycle arrest and stimulating the expressions of host heat shock proteins (Prikhod'ko and Miller, 1998; Tung <i>et al.</i> , 2016)
AC152	152	Del (Gandhi <i>et al.</i> , 2012)		Dispensable for BV but slightly affecting the kinetics of BV production (Gandhi <i>et al.</i> , 2012)
PE38	153	Mut (Prikhod'ko and Miller, 1999), Del (Gandhi <i>et al.</i> , 2012; Milks <i>et al.</i> , 2003)	Del (Ono <i>et al.</i> , 2012)	Important for viral DNA synthesis, BV production, G-actin translocation to the nucleus and viral resistance of host (Gandhi <i>et al.</i> , 2012; Gebhardt <i>et al.</i> , 2014; Milks <i>et al.</i> , 2003) as a baculovirus trans activator (Wu <i>et al.</i> , 1993) which is involved in DNA replication (Kool <i>et al.</i> , 1994) and augments IE1-induced apoptosis but is unable to induce apoptosis (Prikhod'ko and Miller, 1999)
AC154	154	Del (Bai <i>et al.</i> , 2020; Tong <i>et al.</i> , 2021)	Del (Ono <i>et al.</i> , 2012)	Dispensable for BV and ODV production (Bai <i>et al.</i> , 2020; Ono <i>et al.</i> , 2012; Tong <i>et al.</i> , 2021); may function to be anti-apoptotic to provide a more favorable environment for progeny virus replication and packaging (Bai <i>et al.</i> , 2020)

∞: Mut: mutation; Del: Deletion, means "Knock out"; Ac: AcMNPV; Bm: BmNPV; Ha: HearNPV.

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