

ウイルスとは？

- 1) 遺伝物質がDNAかRNAのいずれか
- 2) 成長・分裂しない
- 3) エネルギー產生系・リボソームをもたない
- 4) ウィルスの複製には宿主細胞の素材を利用する

ウイルス粒子(Virion)－核酸と蛋白性外被(CapsidProtein)

核酸を包むタンパク性外被
(nucleocapsid)

これら全部を包むエンベロープ
(envelope)

昆虫病原性ウイルス

| 科名 | 属名 | 和名 | 核酸の種類 |
|----------------|-----------------------|----------------------|-------|
| Baculoviridae | <i>Baculovirus</i> | 核多角体病ウイルス 顆粒病ウイルス | 複鎖DNA |
| Reoviridae | <i>Cyopivirus</i> | 細胞質多角体病ウイルス | 複鎖RNA |
| Poxviridae | <i>Entomopoxvirus</i> | 昆虫ポックスウイルス | 複鎖DNA |
| Iridovirus | <i>Iridovirus</i> | 虹色ウイルス | 複鎖DNA |
| Parvoviridae | <i>Densovirus</i> | 濃核病ウイルス | 单鎖DNA |
| Rhabdoviridae | <i>Sigmavirus</i> | σウイルス | 单鎖RNA |
| Picornaviridae | <i>Enterovirus</i> | 急性麻痺病ウイルス | 单鎖RNA |

「昆虫病理学」：福原敏彦編から転載

<http://hashi.agr.hokudai.ac.jp/IP/IP.html>

Viral Diseases of Insects in the Literature (VIDIL)
Literature, species, and subjects to 1985
Developed by Mauro E. Martignoni



Text Last Updated: June 13, 2001
Database Last Updated: March 12, 1999

<http://insectweb.inhs.uiuc.edu/Pathogens/VIDIL/>

The Illinois Agricultural Experiment Station, University of Illinois
The Center for Economic Entomology, Illinois Natural History Survey

Search the Ecological Database of the World's Insect Pathogens (EDWIP)

| | |
|---|--|
| <h1>EDWIP</h1> | |
| Welcome | |
| Guestbook NEW | |
| SEARCH NOW! | |
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| Indexes NEW | |
| Database Structure | |
| Submit Data to EDWIP | |
| NEW | |
| Data Entry Protocols | |
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| <p>NONVIRAL PATHOGEN</p> | |
| <p>ASSOCIATIONS i</p> | |
| <p>NEMATODE ASSOCIATIONS i</p> | |
| <p>VIRUS ASSOCIATIONS i</p> | |
| <p>NEGATIVE TEST RESULTS i</p> | |
| <p>HOST CLASSIFICATION AND</p> | |
| <p>ECOLOGY i</p> | |
| <p>NONVIRAL PATHOGEN</p> | |
| <p>CLASSIFICATION i</p> | |
| <p>CITATIONS i</p> | |
| <p>NOTE: Older browsers may be unable to read EDWIP database files. Please use the Guestbook to report problems. Be sure to include the browser name and version in your message.</p> | |

Families and Groups of Viruses Associated with Insect

DNA VIRUSES

Double-stranded DNA, enveloped

POXVIRIDAE, Poxviruses

ENTOMOPOXVIRINAE, Poxviruses of insects

Three probable genera (**poxvirus α , β , γ**)

BACULOVIRIDAE

Baculovirus

Subgroup A, nuclearpolyhedrovirus

Subgroup B, granulosis virus

Subgroup C, enveloped nonoccluded
rod-shaped nuclear virus

POLYDNAVIRIDAE

Polydnavirus

Subgroup A, polydnavirus with fusiform
nucleocapsids

Subgroup B, polydnavirus with rod-shaped
nucleocapsids

Double-stranded DNA, nonenveloped

IRIDOVIRIDAE, icosahedral cytoplasmic deoxyriboviruses

Iridovirus, small iridescent insect virus

Chloriridovirus, large iridescent insect virus

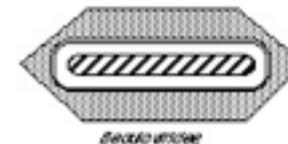
Single-stranded DNA, nonenveloped

PARVOVIRIDAE, parvoviruses

Densovirus, denonucleosis virus

Families of Viruses Infecting Invertebrates

dsDNA



100 nm

Families of Viruses
Infecting Invertebrates

ssDNA



100 nm

ポックスウイルス科 Poxviridae (Pox (痘症、天然痘) に由来する。)

コードポックスウイルス亜科 Chordopoxvirinae (脊椎動物に感染する)

オルトポックスウイルス属 Genus: Orthopoxvirus

主なウイルス： 痘瘍ウイルス Variola virus

牛痘ウイルス Cowpox virus

ワクシニアウイルス [Vaccinia virusem_poxvi.htm](#)

パラポックスウイルス属 Genus: Parapoxvirus

主なウイルス： オーフウイルス Orf virus

アヴィポックスウイルス属 Genus: Avipoxvirus

主なウイルス： 鶏痘ポックスウイルス Fowlpox virus

カプリポックスウイルス属 Genus: Capripoxvirusem_poxvi.htm

主なウイルス： ヒツジポックスウイルス Sheeppox virus

レポリポックスウイルス属 Genus: Leporipoxvirus

主なウイルス： 粘液腫ウイルス Myxoma virus

モラシポックスウイルス属 Genus: Molluscipoxvirus

主なウイルス： Molluscum contagiosum virus

ヤタポックスウイルス属 Genus: Yatapoxvirus

主なウイルス： ヤバサル腫瘍ウイルス Yaba monkey tumor virus

痘瘍ウイルス [Variola virus]

古代エジプトのミイラにも痘瘍の痕跡が見つかり、天然痘は古い病気であること、人類にとって死亡率の高い伝染病であった。種痘によるWHOの全世界痘瘍根絶計画が奏を功し、1977年10月東アフリカのソマリアにおける患者発生が最後だったことから、WHOは1979年この地球上から天然痘が撲滅されたと宣言しました。英國のエドワード・ジェンナーが生ワクチンを開発した。

エントモ(昆虫) ポックスウイルス亜科 Entomopoxvirinae (昆虫に感染する)

昆虫ポックスウイルス Alpha属 Genus: Alphaentomopoxvirus (鞘翅目昆虫)

主なウイルス： *Melolontha melolontha entomopoxvirus*

昆虫ポックスウイルス Beta属 Genus: Betaentomopoxvirus (直翅目、鱗翅目昆虫)

主なウイルス： *Amsacta moorei entomovirus*

昆虫ポックスウイルス Gamma 属 Genus: Gammaentomopoxvirus (双翅目昆虫)

主なウイルス： *Chironomus luridus entomopoxvirus*

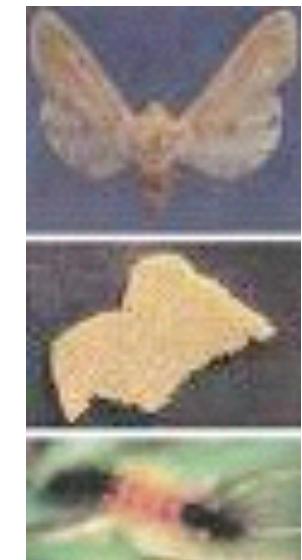
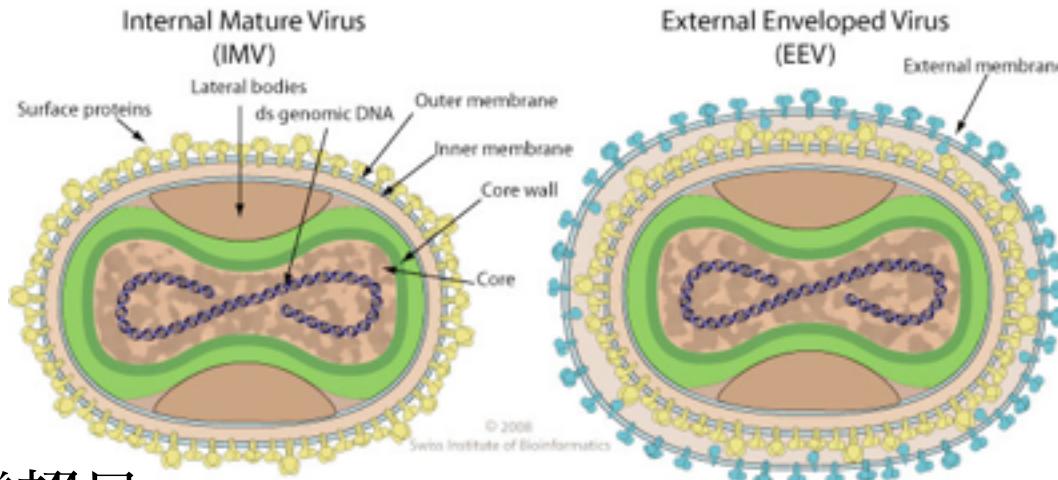
- ・感染と病気 脊椎動物、昆虫に感染。痘瘍ウイルスは、天然痘を引き起こす。
- ・粒子構造 レンガ状の大型ウイルス
- ・遺伝子 直鎖二本鎖DNA

Alpha : 鞘翅目

Melolontha melolontha entomopoxvirus (MMEV)

Anomala cuprea entomopoxvirus [AB005053] (ACEV)

Complete nucleotide sequence of spheroidin gene of *Anomala cuprea* entomopoxvirus Virus Res. 55 (1), 61-69 (1998)



Beta : 鱗翅目

Amsacta moorei entomopoxvirus 'L' (AMEV)

Complete genomic sequence of the *Amsacta moorei* entomopoxvirus: analysis and comparison with other poxviruses
Virology 274 (1), 120-139 (2000)

Arphia conspersa entomopoxvirus 'O' (ACOEV)

Gamma : 双翅目

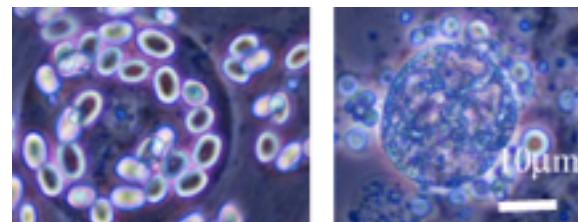
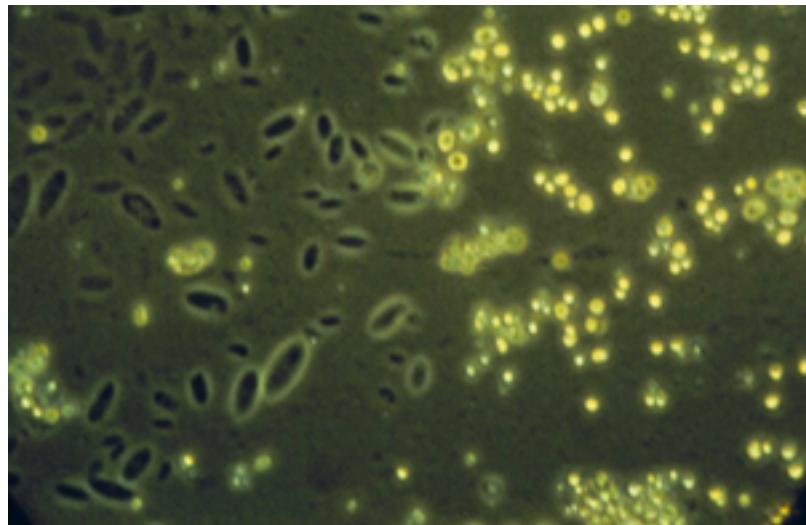
Chironomus luridus entomopoxvirus (CLEV)

Aedes aegypti entomopoxvirus (AAEV)



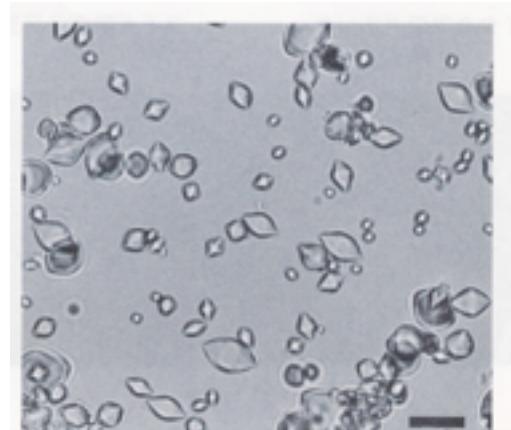
昆虫ポックスウイルス (Entomopoxvirus)

ポックスウイルス科に属する二本鎖DNAウイルス



昆虫ポックス感染虫の脂肪体細胞（左）。細胞質で包埋体が産生されている。右は、正常な脂肪体細胞。

種々の核多角体病ウイルスの感染力を大きく増強する
昆虫ポックスウイルスのスピンドル



精製したドウガネブイブイのEPVの
スピンドル。バーは10 μmを示す。

Sequence 1: gi|13876680|gb|AAK43568.1| 1210 aa-Lumpy skin disease virus=Homolog of Vac.J6R
 Sequence 2: gi|6681610|dbj|BAA88785.1| 1285 aa-Yaba monkey tumor virus=Homolog of Vac J6R
 Sequence 3: gi|9964535|ref|NP_065003.1| 1301 aa-AmEPV221 *Amsacta moorei* entomopoxvirus
 Sequence 4: gi|9631519|ref|NP_048113.1| 1319 aa-MsEPV043 *Melanoplus sanguinipes* entomopoxvirus
 Sequence 5: D1EPV36 772 aa-D1EPV RI-36-1 *Diachasmimorpha* entomopoxvirus

CLUSTAL W (1.81) multiple sequence alignment

| | |
|---------------------------|---|
| gi 13876680 gb AAK43568.1 | -----MDGALC 6 |
| gi 6681610 dbj BAA88785.1 | -MAVISKVTV <ins>SLYNQEEI</ins> NATDV <ins>LINHV</ins> KNDDD <ins>IGTVKD</ins> GRLGAMDGALC 49 |
| AmEPV221 | ---MTTFKYTL <ins>LD</ins> MSTIDA <ins>IPIVID</ins> SIGNDNEN-SVKSPKLGGTKFNVC 45 |
| MsEPV043 | <ins>HAKVNTEIILF</ins> FSMIPNSI <ins>IIDS</ins> IPIIINSI <ins>SNDADN</ins> -NVKSTKLGGTKFNVC 49 |
| D1EPV36 | -----EFSLVS <ins>NSEEIES</ins> IPIAIKNISGENKTEAAQLG--AIDSFQKC 40 |
| | * . |
| gi 13876680 gb AAK43568.1 | KTC GKT-E LQC FGHWGKVRLYETHI <ins>I</ins> KPEYIGEVIRILN--HICIRCGL 53 |
| gi 6681610 dbj BAA88785.1 | KTC EKT-E LQC FGHWGKVRLYETHI <ins>V</ins> KPEYIGEVVRILN--HICIRCGL 96 |
| AmEPV221 | STCNLTRENGDMGHPGRTPLRDMCIVKSGC <ins>I</ins> KNVLDTLNTLKL CNSCFMI 95 |
| MsEPV043 | STCR <ins>L</ins> TKDNGDLGHPGRTPLKKM <ins>AII</ins> IKP <ins>AFI</ins> KSVLDTLNALKIC <ins>SNCKMF</ins> 99 |
| D1EPV36 | ETC QLT--SACPGHFGKFHLTQP-LFKV <ins>AFKKLVEN</ins> IFK--YTCSFCGAL 85 |
| | .** * ** *; * . :.* * : ; : * * : . |
| gi 13876680 gb AAK43568.1 | RSRE PYIEDVTKMSHALRK-----LKDKILSKKK-----SCWNSK 89 |
| gi 6681610 dbj BAA88785.1 | RSRE PYMEDITRMPL <ins>I</ins> SLKK-----LKDKILSKKK-----SCWNSK 132 |
| AmEPV221 | KNNTIFSE <ins>IIE</ins> KYNSEYNIN-----LKKEILSLLKNNRQGGVKCNNEN 138 |
| MsEPV043 | RDNEALYK <ins>I</ins> LKKYNIDVQDNKID PPTELKKEILT <ins>L</ins> IKLN <ins>K</ins> QSASKCNNIN 149 |
| D1EPV36 | QNLEL <ins>LELI</ins> IKQIDER <ins>IT</ins> GIT-----VKDRAAFKKILEAT <ins>K</ins> QSKFKCIAPN 130 |
| | : : . : . * : . |
| gi 13876680 gb AAK43568.1 | C <ins>M</ins> Q <ins>Q</ins> YQKITFSKKK-VCFVN-KSDDITIPNALIYQKITSIYKRFWPLLEI 137 |
| gi 6681610 dbj BAA88785.1 | C <ins>M</ins> Q <ins>P</ins> YQKITFSKKK-VCFIN-KSDEITIPNALIYQKVT <ins>SIYKR</ins> FWPLLEI 180 |
| AmEPV221 | C <ins>Q</ins> NI <ins>IT</ins> GTYKYMQKKS <ins>YFYVK</ins> -KQ <ins>K</ins> KDEII <ins>L</ins> NKTVYTMLL <ins>G</ins> IPDI <ins>I</ins> YKCVTV 187 |
| MsEPV043 | C <ins>Q</ins> LP <ins>I</ins> ATYKYM <ins>TIKAQFYIK</ins> -VI <ins>KDKV</ins> I <ins>SNEQIYKMLIGI</ins> PHI <ins>VY</ins> KCIKS 198 |
| D1EPV36 | C <ins>Q</ins> Q <ins>Q</ins> V <ins>SPLQYSK</ins> NNMF <ins>IYNSGTTKGIVL</ins> DNRHVF <ins>NILQNL</ins> PQT <ins>FK</ins> LLLTP 180 |
| | * . : . : . : * : : . : . : . : |

a large gene that encodes a homolog of a DNA-directed RNA polymerase.

Families and Groups of Viruses Associated with Insect

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Double-stranded DNA, enveloped

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ENTOMOPOXVIRINAE, Poxviruses of insects

Three probable genera

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Baculovirus

Subgroup A, nuclearpolyhedro virus

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rod-shaped nuclear virus

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Polydnavirus

Subgroup A, polydnavirus with fusiform
nucleocapsids

Subgroup B, polydnavirus with rod-shaped
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Double-stranded DNA, nonenveloped

IRIDOVIRIDAE, icosahedral cytoplasmic deoxyriboviruses

Iridovirus, small iridescent insect virus

Chloriridovirus, large iridescent insect virus

Single-stranded DNA, nonenveloped

PARVOVIRIDAE, parvoviruses

Densovirus, denonucleosis virus

Families of Viruses Infecting Invertebrates

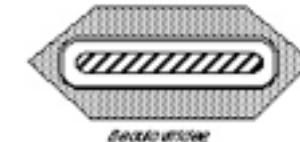
dsDNA



Poxviridae
Entomopoxvirinae



Iridoviridae
Chloririvirus



Baculoviridae



Polydnaviridae
Iridoavirinae



Polydnaviridae
Baculovirinae

100 nm

Families of Viruses
Infecting Invertebrates

ssDNA

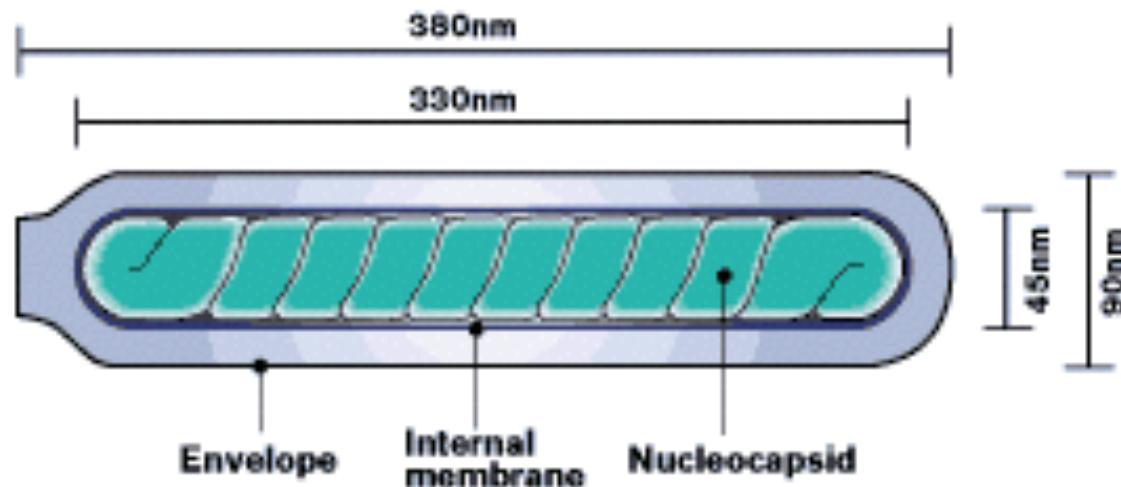


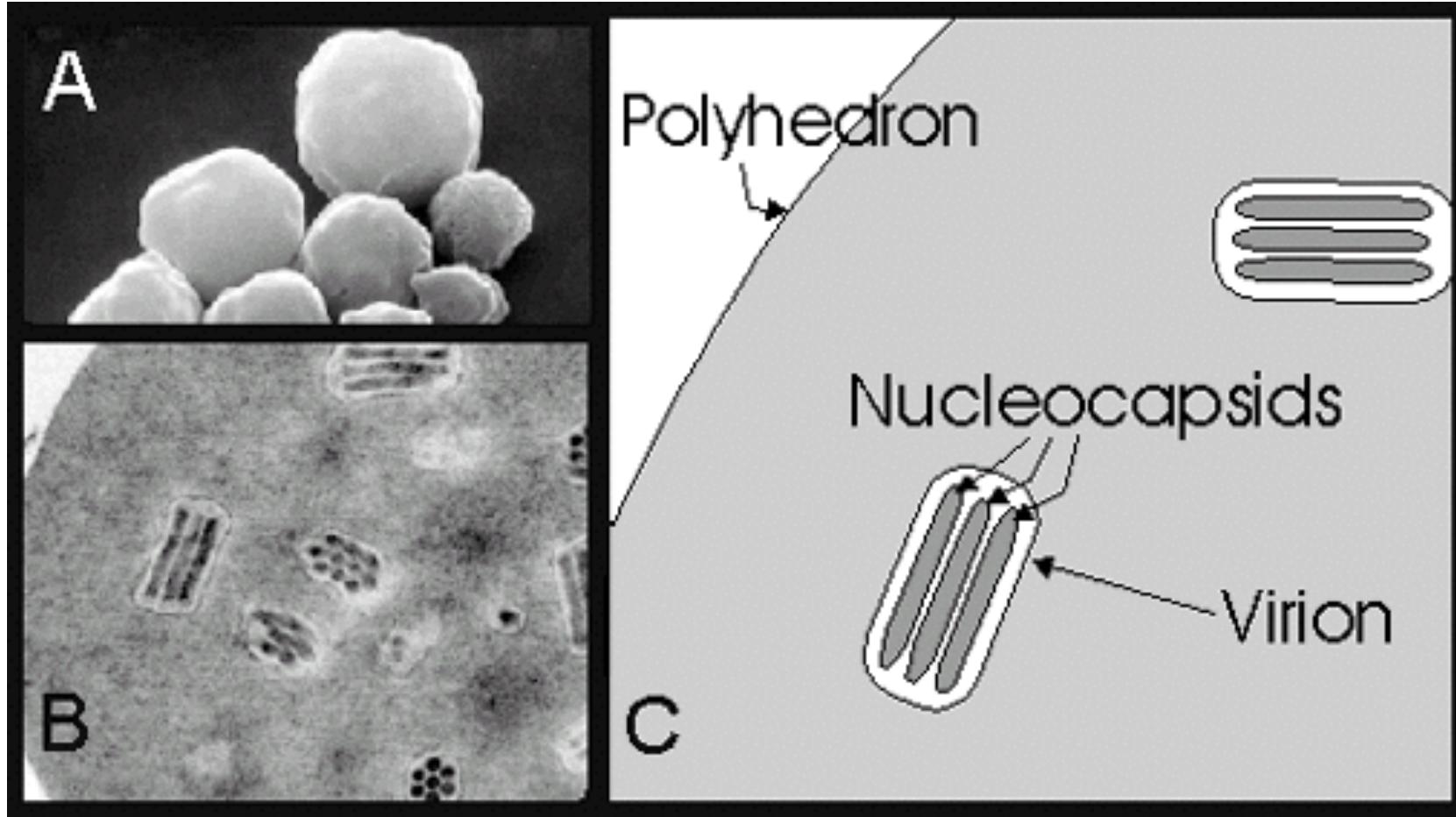
Parvoviridae
Entomoparvovirinae

100 nm

バキュロウイルス(Baculovirus)とは

昆虫を主な宿主として病気を引き起こすウイルスは、9つのグループに分類される。そのなかで、4つのグループは封入体を作るという特徴がある。この封入体はウイルス粒子を包埋するタンパク性の結晶構造物で、しかもその封入体はウイルス自身にコードされているタンパク質です。それらのウイルスの中で、環状2本鎖で約130Kbpの1断片のDNAをゲノムとするのが、バキュロウイルス (Baculovirus) 科に属する核多角体病ウイルス (nucleopolyhedrovirus: NPV) です。（注：NPVの英名は、 nuclear polyhedrosis virusから、 nucleopolyhedrovirusに変更になりました）。

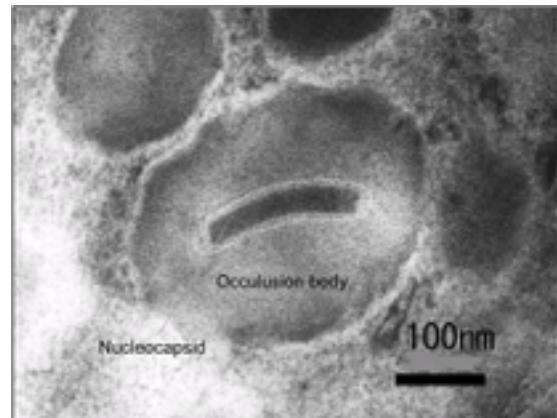




A) Baculovirus particles, or polyhedra; **B)** Cross-section of a polyhedron; **C)** Diagram of polyhedron cross-section.

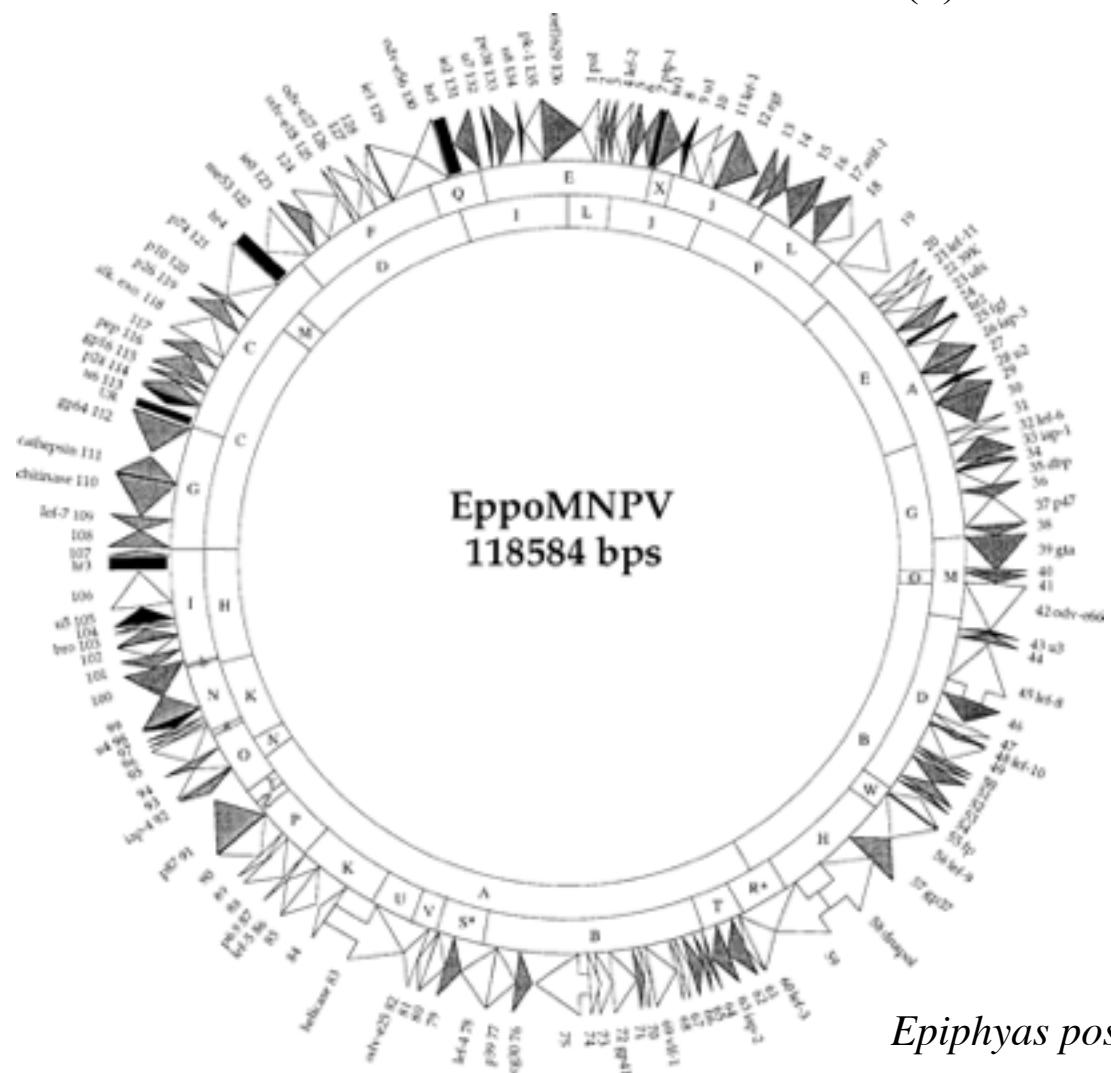
顆粒病ウイルス (granulovirus)

顆粒病ウイルスは、
核多角体病ウイルス
と同じバキュロウイ
ルス科に属す



Bar=5mm

沖縄に生息するチビカクモンハマキから新規に分離されたチビカクモンハマキ顆粒病ウ
イルス(上)
チャノコカクモンハマキ顆粒病ウイルスに感染したチャノコカクモンハマキ幼虫(下)



Circular map of the EppoMNPV genome. The sites for the restriction enzymes HindIII (outer circle) and EcoRI (inner circle) are shown with fragments named according to size. Fragments R and S (*) are reversed from the original map published by

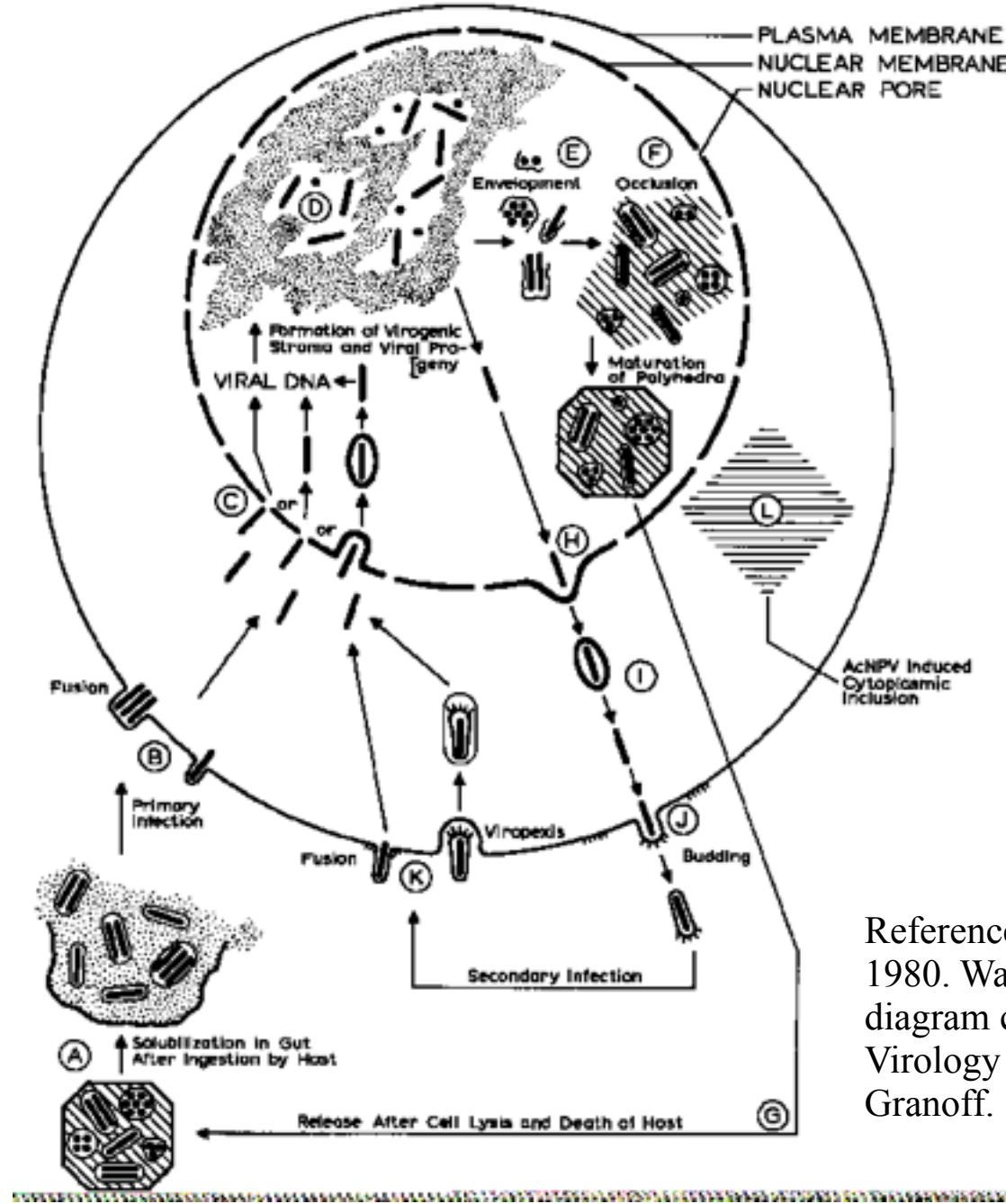
Hyink et al. (1998). Arrows indicate the size, location and direction of the 136 ORFs identified on the EppoMNPV genome.

Open arrows indicate ORFs conserved between all the sequenced baculoviruses shaded arrows indicate ORFs that have homologues in some but not all baculoviruses and black arrows indicate ORFs unique to EppoMNPV.

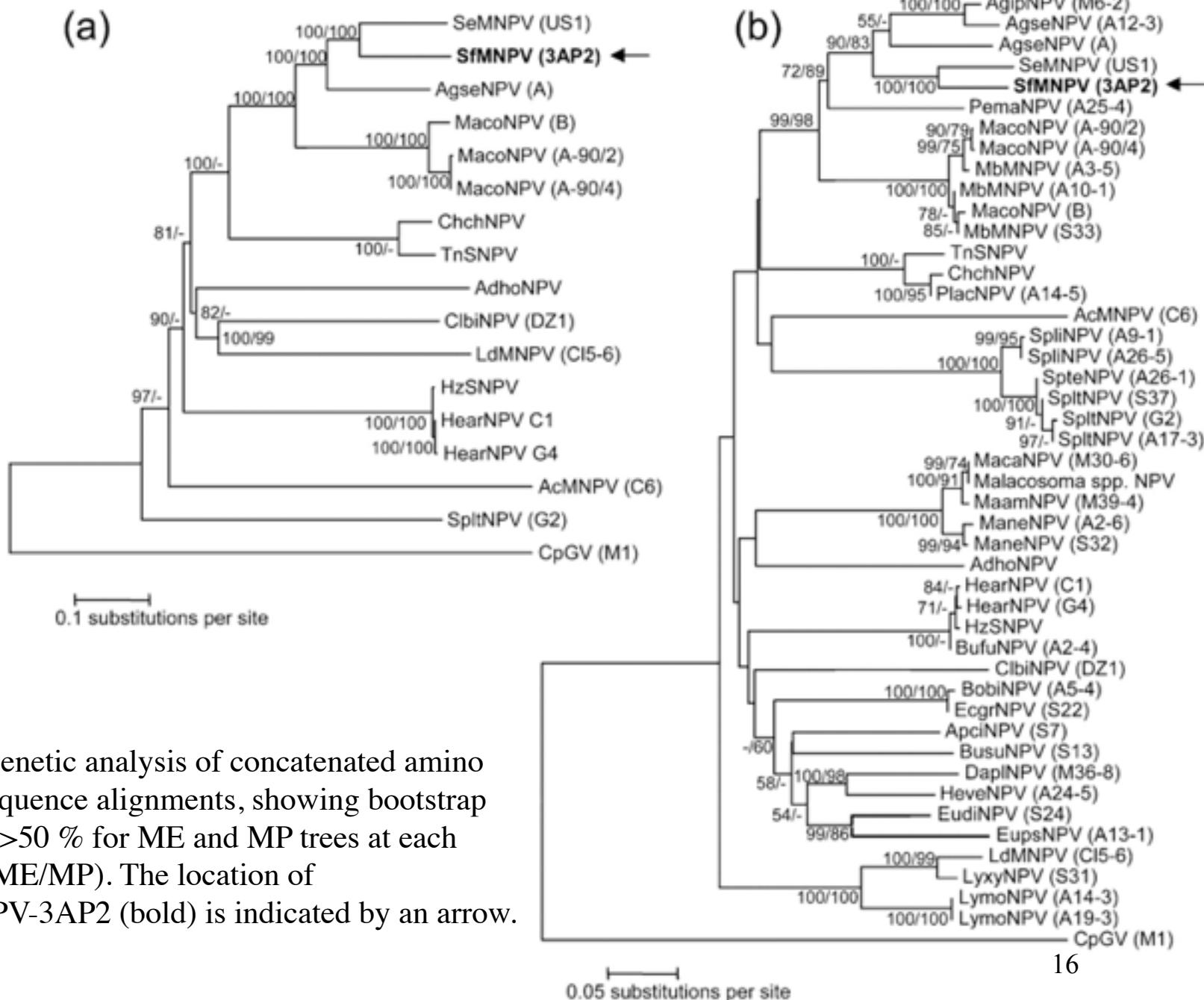
The positions of repeat regions are indicated by black boxes.

The unique repeat is identified as UR.

Epiphyas postvittana nucleopolyhedrovirus (EppoMNPV) genome



Reference: C.P. van der Beek. Doctoral Thesis 1980. Wageningen, The Netherlands a similar diagram can be found in the: Encyclopedia of Virology Vol. 1, page 134. eds. Webster and Granoff.



Phylogenetic analysis of concatenated amino acid sequence alignments, showing bootstrap values >50 % for ME and MP trees at each node (ME/MP). The location of SfMNPV-3AP2 (bold) is indicated by an arrow.

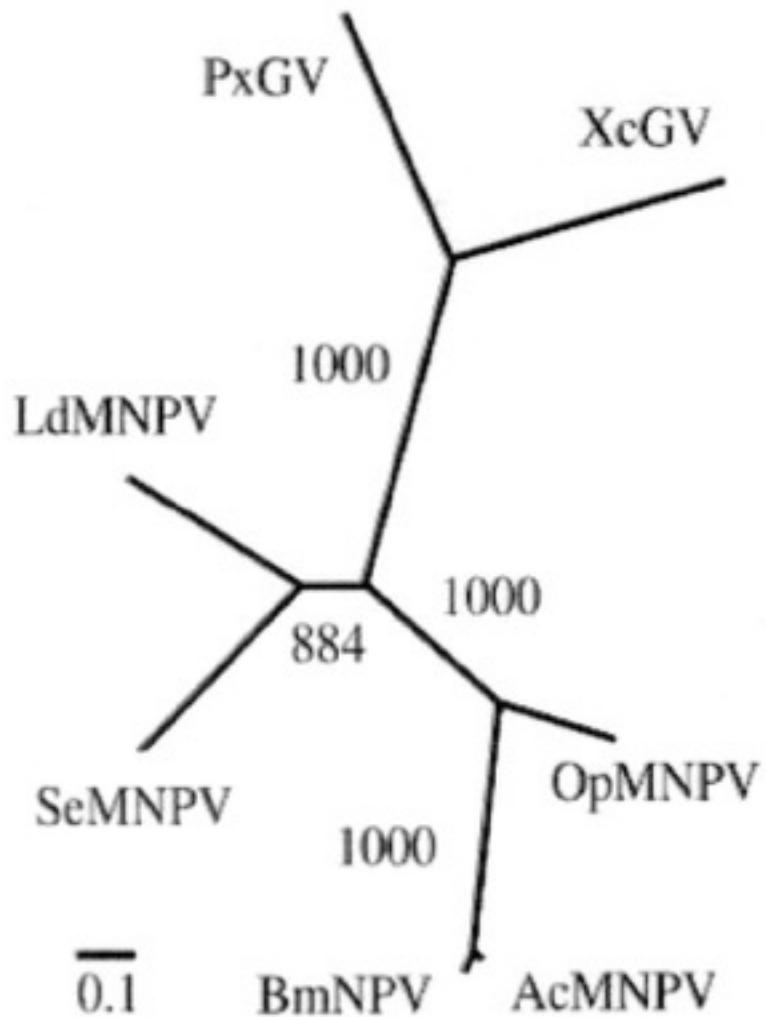


FIG. 4. Baculovirus phylogeny based on the major capsid protein. The sequences of Ac89 (vp39-capsid) and its homologs were determined using CLUSTAL W (Thompson *et al.*, 1994) and TREEVIEW (Page, 1996). Numbers are values determined by bootstrap analyses. For references, see Table 1.

Oryctes Virus:

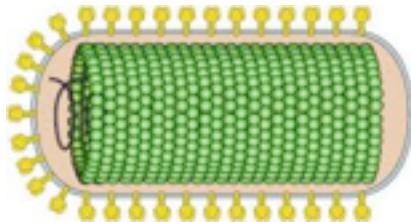
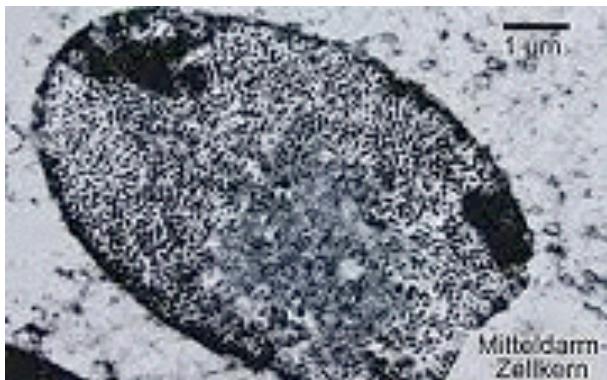
Family of INSECT VIRUSES containing two subfamilies: Eubaculovirinae (occluded baculoviruses) and Nudibaculovirinae (nonoccluded baculoviruses).

USE OF MICROBES FOR CONTROL AND ERADICATION OF INVASIVE ARTHROPODS

Progress in Biological Control, 2009, Volume 6, Part IV, 133-140, DOI: 10.1007/978-1-4020-8560-4_8

The Use of *Oryctes* Virus for Control of Rhinoceros Beetle in the Pacific Islands

Trevor A. Jackson



Cocoanut Rhinoceros Beetle
Pests and Diseases of American Samoa
Volume 6

Description: The coconut rhinoceros beetle, *Oryctes rhinoceros* L., has become one of the most serious pests of the coconut palm in several countries around the world from Africa to Asia to Oceania. This pest can damage a great amount of coconut and oil palm trees and attacks other palm species.

Habits: Adults emerge from the pupal cell 17–27 days before emerging, and during the adult stage, the beetles are active at night and hide or huddle or burrow during the day. After mating, adult females lay eggs among the fibrous roots of the coconut tree. Adults may live up to three years.

Damage: Coconut rhinoceros beetle adults damage palms by feeding on the leaves of young trees. They strip the young, green, young leaves and feed on the seedlings. As they approach the mature trees, they damage the developing fruits. When the fruits split open and fall onto the damage appears as V-shaped cuts on the fruits of trees throughout the island.

Egg: Cocoanut beetle eggs, brown, pointed, and white may be scattered by various predators, including frogs, bats, and other beetles. They also, when scattered by birds, damage the foliage. When laid on coconut and the fibrous roots, damage.

Life Cycle: Egg → larva → pupa → adult. In temperate regions, adults of decomposing vegetation or coconut, or other organic matter. Eggs hatch after 7 days, and larvae feed and grow 3–4 weeks. 10–15 days before reaching an 8.0–10.0 mg instar third-stage pupae. Pupal development is variable due to the season and the

18

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ENTOMOPOXVIRINAE, Poxviruses of insects

Three probable genera

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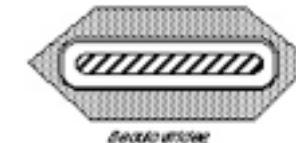
dsDNA



Poxviridae
Entomopoxvirinae



Iridoviridae
Chloriridovirus



Baculoviridae



Polydnaviridae
Iridoavirinae



Polydnaviridae
Baculovirinae

100 nm

Families of Viruses
Infecting Invertebrates

ssDNA



Parvoviridae
Entomoparvovirinae

100 nm

The **Polydnaviruses** (PDV) are a family of insect [viruses](#) that contain two genera: Ichnoviruses (IV) and Bracoviruses (BV). The [genome](#) of the virus is composed of multiple segments of double-stranded, superhelical DNA packaged in capsid proteins and a double layer (IV) or single layer (BV) envelope.

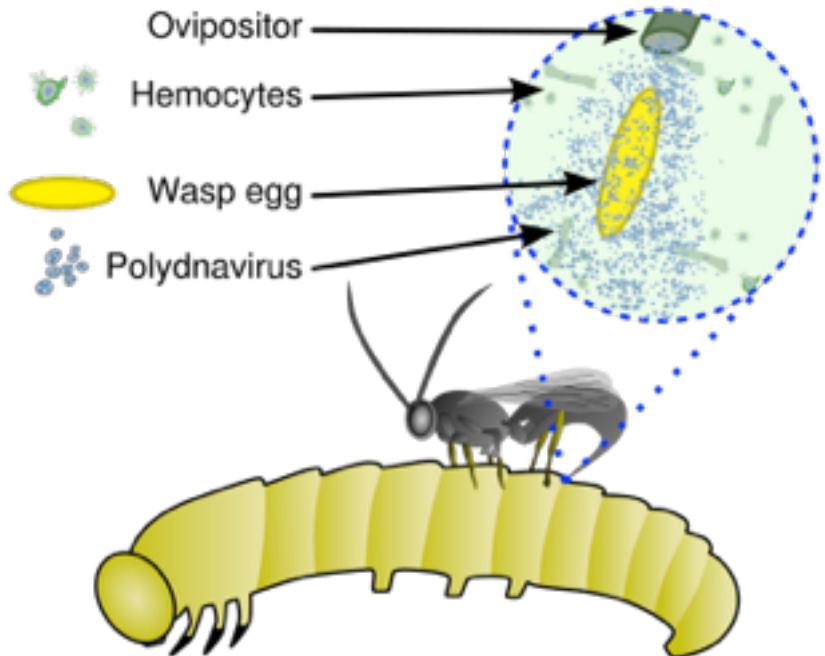


Diagram of a PDV host association

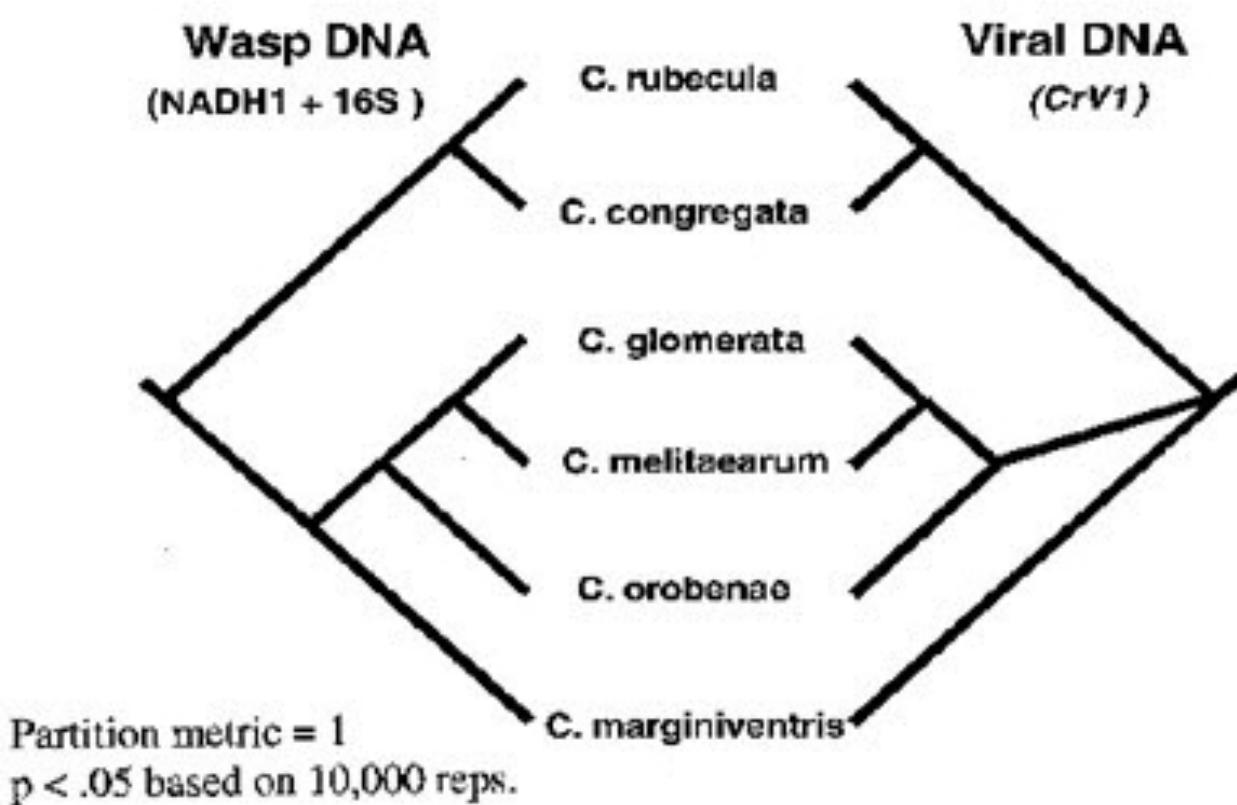
Both genera of PDV share certain characteristics:

- the virus particles of each contain **multiple segments of dsDNA** with each segment containing only part of the full genome (much like chromosomes in eukaryotic organisms);
- the genome of each is **integrated into the host wasp genome**;
- the virus particles are **only replicated (produced) in specific cell types in the female wasp reproductive organs**.

However, the morphology of the two genera are different when observed by electron microscopy. Ichnoviruses tend to be ovoid (egg-shaped) while bracoviruses are short rods.

Also, as their names suggest, the ichnoviruses occur in [ichneumonid wasps](#) (ヒメバチ科) species and bracoviruses in [braconid wasps](#) (コマユバチ科).

Coevolution of Polydnnaviruses with Braconid Wasps



C. rubecula cocoon

Polydnnaviruses are apparently mutualistic hereditary viruses that are injected into host caterpillars by some parasitoid wasps and aid in suppression of the host immune system and redirection of host physiology to aid the survival of developing wasp eggs and larvae.

Families and Groups of Viruses Associated with Insect

DNA VIRUSES

Double-stranded DNA, enveloped

POXVIRIDAE, Poxviruses

ENTOMOPOXVIRINAE, Poxviruses of insects

Three probable genera

BACULOVIRIDAE

Baculovirus

Subgroup A, nuclearpolyhedro virus

Subgroup B, granulosis virus

Subgroup C, enveloped nonoccluded
rod-shaped nuclear virus

POLYDNAVIRIDAE

Polydnavirus

Subgroup A, polydnavirus with fusiform
nucleocapsids

Subgroup B, polydnavirus with rod-shaped
nucleocapsids

Double-stranded DNA, nonenveloped

IRIDOVIRIDAE, icosahedral cytoplasmic deoxyriboviruses

Iridovirus, small iridescent insect virus

Chloriridovirus, large iridescent insect virus

Single-stranded DNA, nonenveloped

PARVOVIRIDAE, parvoviruses

Densovirus, denonucleosis virus

Families of Viruses Infecting Invertebrates

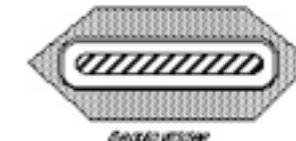
dsDNA



Poxviridae
Entomopoxvirinae



Iridoviridae
Insectivirinae
Chlorovirus



Baculoviridae



Polydnaviridae
Ichnavirinae



Polydnaviridae
Breniereinae

100 nm

Families of Viruses
Infecting Invertebrates

ssDNA



Parvoviridae
Entomoparvovirinae

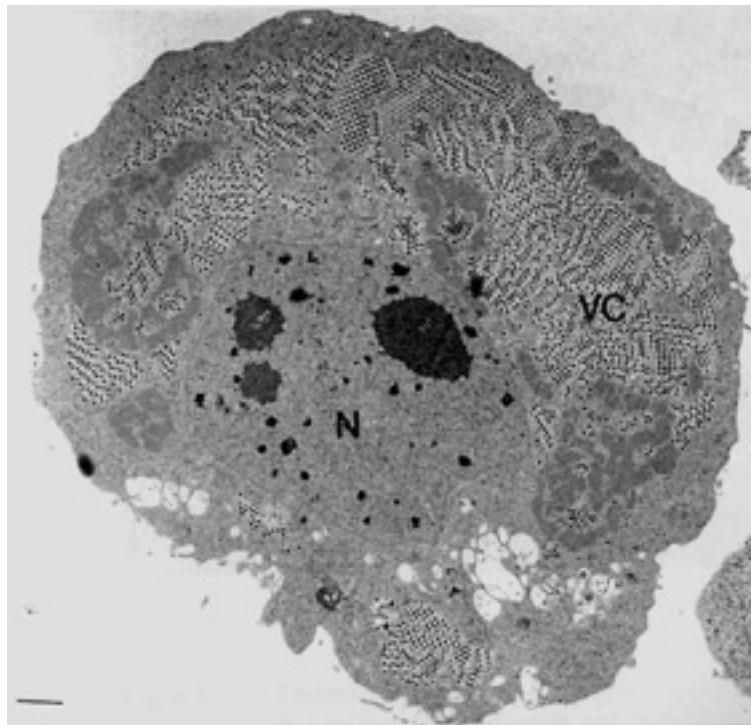
100 nm

The family **Iridoviridae** is comprised of four genera, two infecting vertebrates and two infecting invertebrates

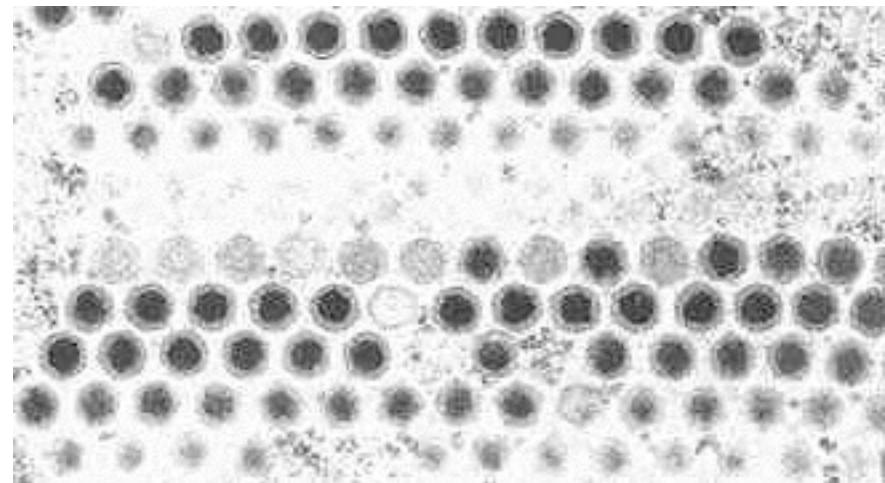
| Genus | Vernacular name | Host species | Type species |
|------------------------|---------------------------------|--------------------------------|----------------------------------|
| Iridovirus | Small iridescent insect virus | Invertebrates (mainly insects) | Chilo iridescent virus (IV6) |
| Chloriridovirus | Large iridescent insect viruses | Mosquitos | Mosquito iridescent virus (IV3) |
| Lymphocystivirus | Lymphocystis disease virus | Fish | Lymphocystivirus type 1 (LCDV-1) |
| Ranavirus | Frog virus | Amphibia | Frog Virus 3 (FV3) |



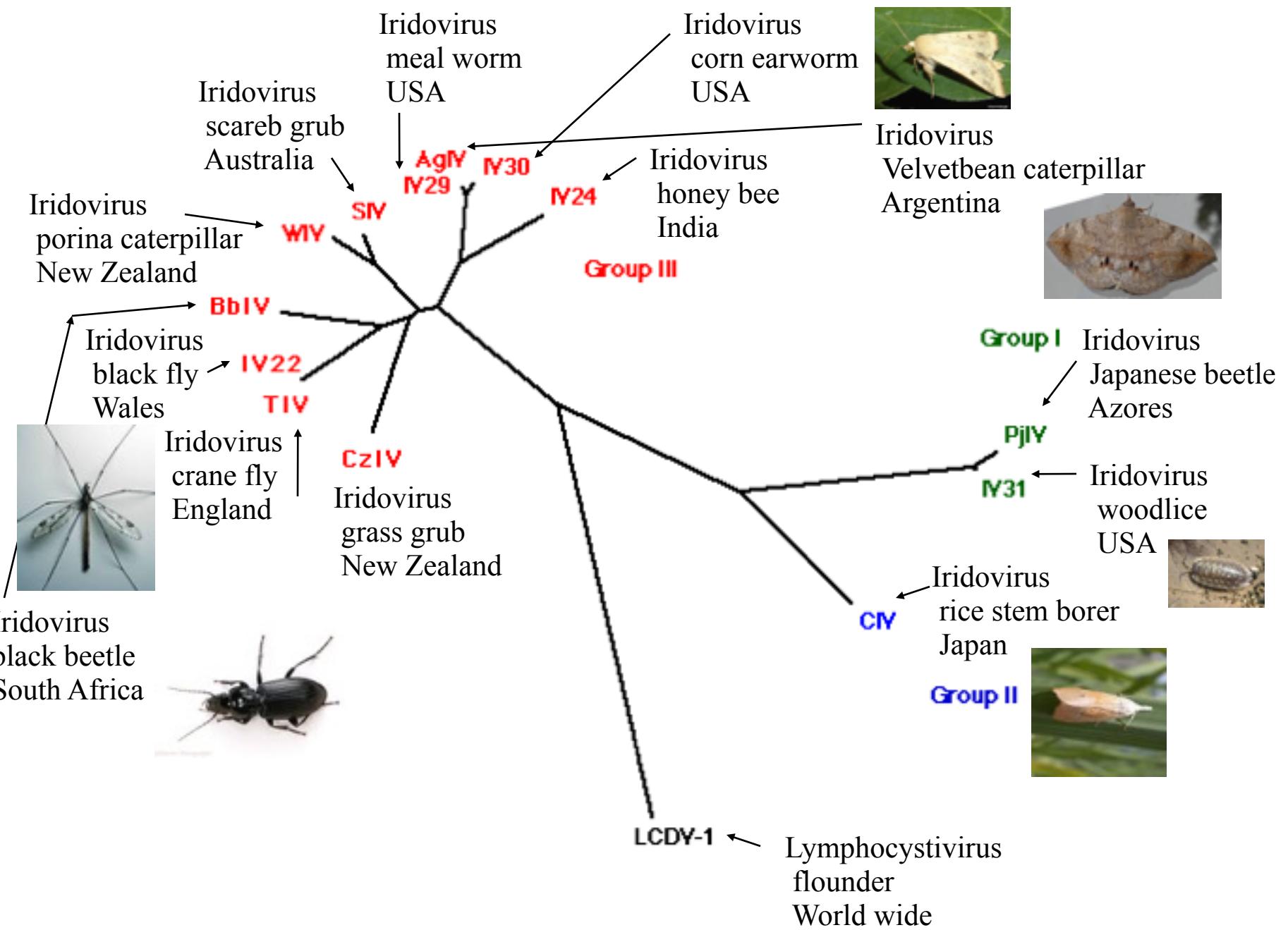
Larvae of the grass grub *Costelytra zealandica* displaying blue colouration of the hindgut due to iridovirus infection

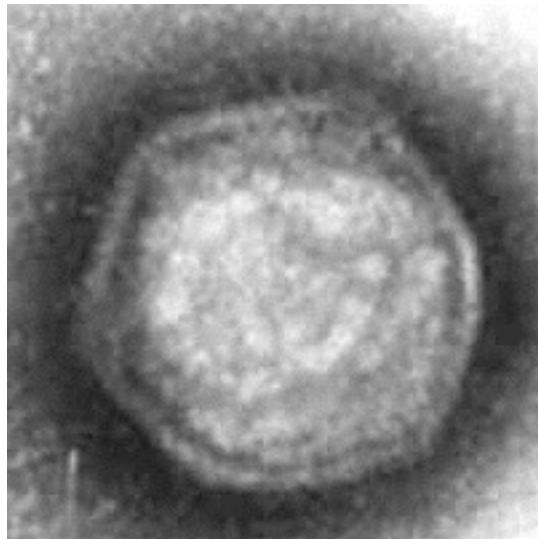


Sf21 tissue culture cell showing cytoplasmic localisation of assembling Wiseana iridescent virus particles. N= nucleus, VC = viroplasmic centre

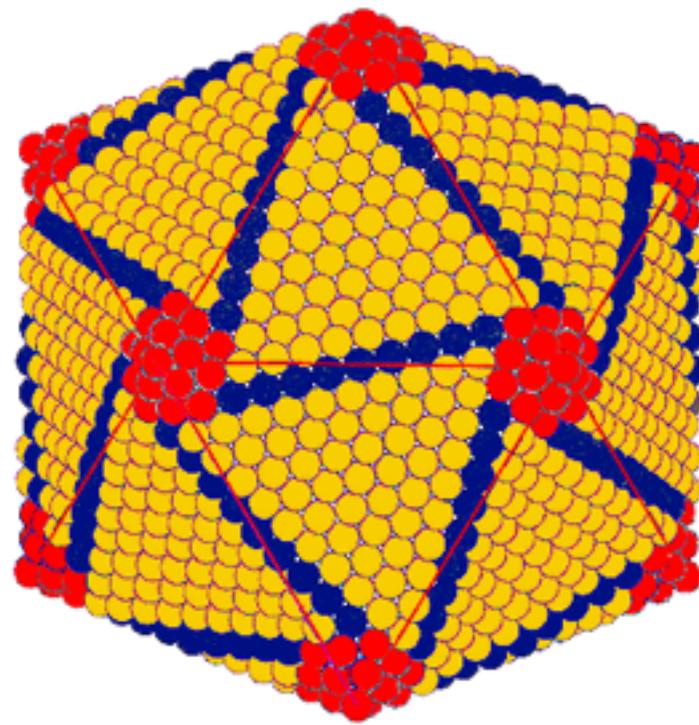


Paracrystalline array of virus particles within infected cell



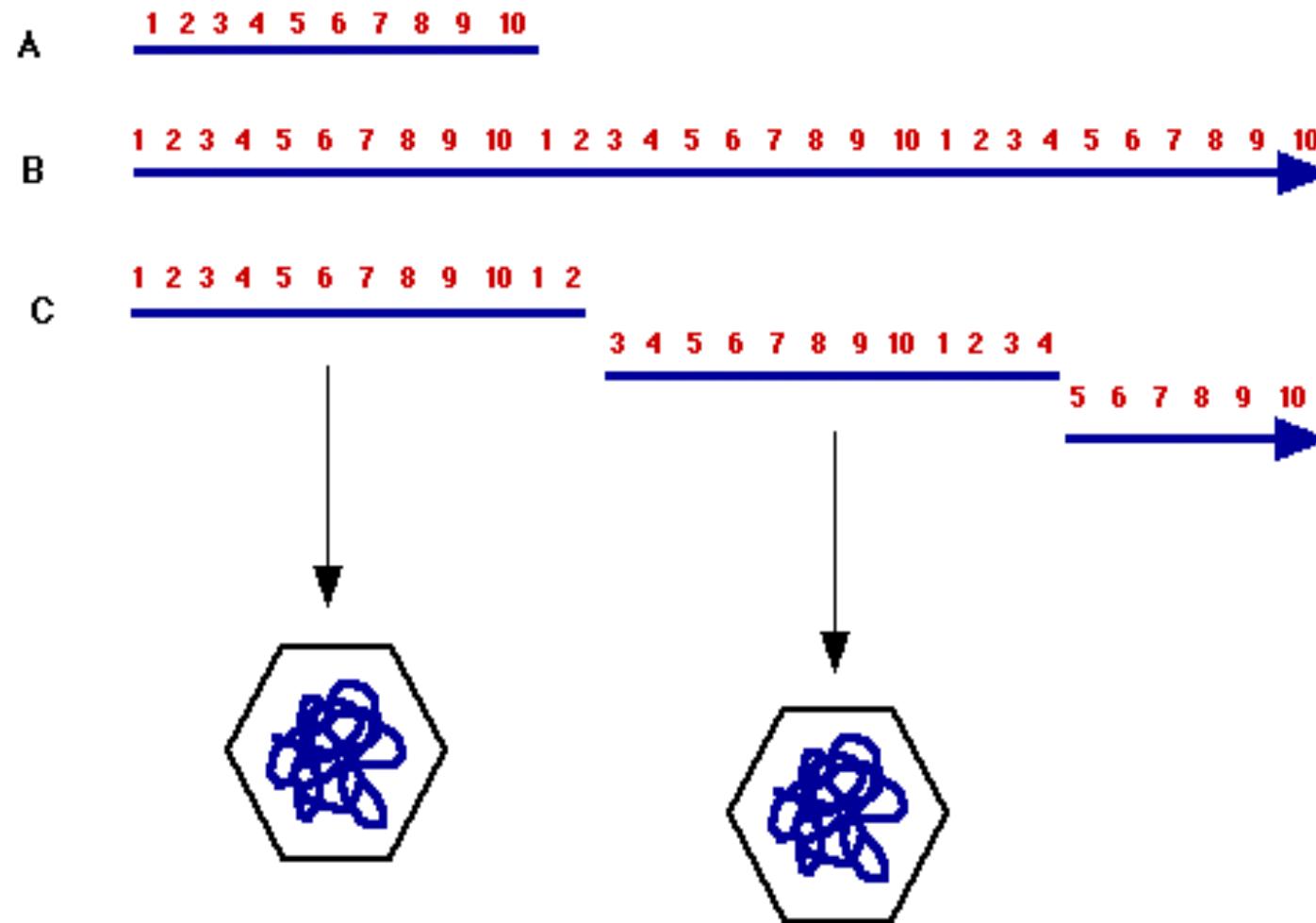


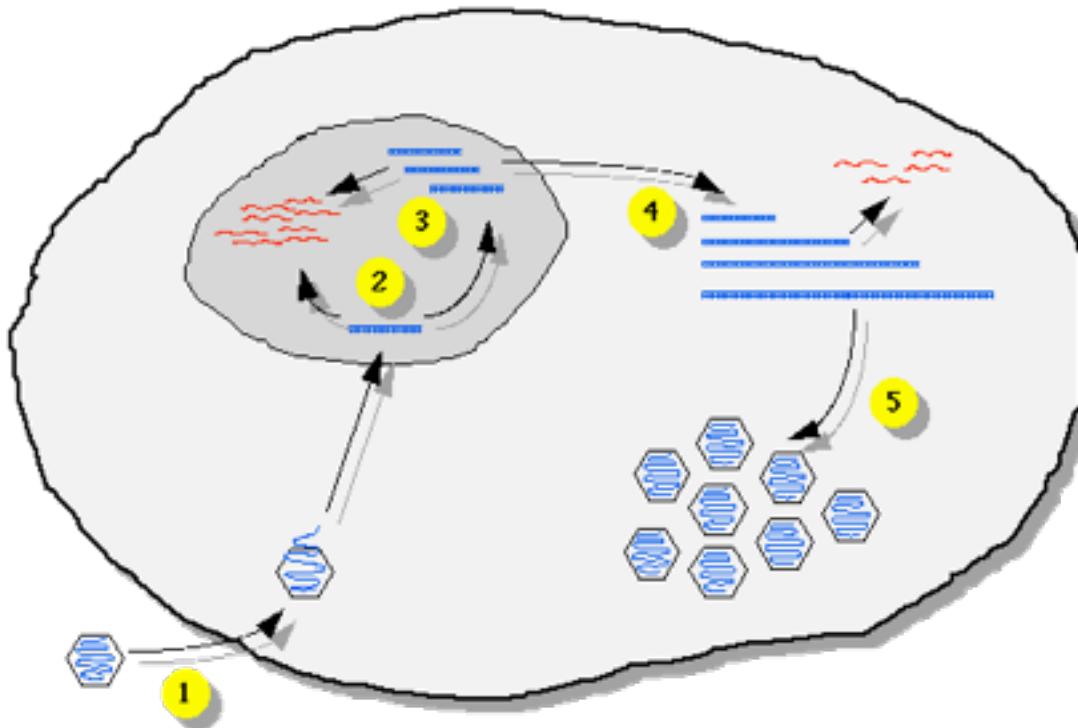
Electron micrograph of a typical iridovirus



Schematic diagram adapted from N.G. Wrigley 1969 (J. Gen. Virol. 5:123-134)

Iridoviruses contain a single copy linear dsDNA genome that ranges in size from 150 to 280 kbp depending on viral species. The genomes appear unique within the eucaryotic viruses in that they are terminally redundant and cyclically permuted. This structure is a result of the resolution of genome concatamers during DNA replication (see replication).





FV3 replication model adapted from R. Goorha 1982 (J. Virol. 43(2):519-528)

- 1) Virus particles enter the cell by pinocytosis and uncoating occurs.
- 2) Viral DNA is transported to the cell nucleus where host macromolecular synthesis is rapidly shutdown. Transcription is initiated by virally modified host RNA polymerase II.
- 3) Parental DNA is used to produce genome and greater than genome length DNA. This becomes the template for cytoplasmic replication.
- 4) Progeny DNA is transported into the cytoplasm where large concatamers of viral DNA are formed by recombination. Transcription of very late transcripts may also take place in the cytoplasm.
- 5) Concatamers are resolved into packaged lengths, possibly by a headful packaging approach. Virions exit the cell by budding or cell lysis.

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Single-stranded DNA, nonenveloped

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Densovirus, denonucleosis virus

Families of Viruses Infecting Invertebrates

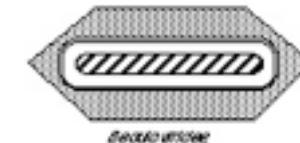
dsDNA



Poxviridae
Entomopoxvirinae



Iridoviridae
Insectivirinae
Chlorovirus



Baculoviridae



Polydnaviridae
Ichnavirinae



Polydnaviridae
Brecavirinae

100 nm

Families of Viruses
Infecting Invertebrates

ssDNA



Parvoviridae
Entomoparvovirinae

100 nm

Parvoviruses Sub-Family Densovirinae

| | |
|-------------|----------------------------------|
| Densovirus | <i>Junonia coenia</i> densovirus |
| Iteravirus | <i>Bombyx mori</i> virus |
| Contravirus | <i>Aedes aegypti</i> densovirus |

One genus in the family Parvoviridae has been recovered from insects, Densovirus (DNV). First isolated from the waxworm, *Galleria mellonella*, these viruses have now been recovered from Diptera, Orthoptera, Blattodea, Odonata, and other Lepidoptera (Kawase, 1985). This ssDNA, nonenveloped virus is characterized by its effects on the nucleus of host cells, and it attacks most host tissues. Infection results in extreme hypertrophy of cell nuclei and the filling of the nuclei with small isometric particles. The nuclei become strongly eosinophilic (dense nuclei, thus the name densovirus). Discoloration and paralysis of the host often occur. These viruses can be very virulent and infectious, and the type species, GmDNV, has caused serious problems for the waxworm bait industry in the Midwest in recent years. Despite their high virulence, densoviruses do not replicate in vertebrates (Bergoin and Tijssen, 1998).

Biological Control:

High virulence and the presence in insect pests have precipitated some interest in use of densoviruses in biological control (Bergoin and Tijssen, 1998). Infected waxworms were placed in beehives to successfully eliminate waxworms from the hives (Lavie et al., 1961). Other pest species have also been successfully treated, for example, the palm oil defoliator, *Sibine fusca* (Genty and Mariau, 1975). The sequence homologies between densoviruses and vertebrate parvoviruses, however, have presented concerns about use, despite lack of evidence that this genus can infect vertebrates. These viruses have insecticidal potential, therefore specificity and tissue tropism should be studied more thoroughly (Bergoin and Tijssen, 1998).

Subfamily. Densovirinae

Genus 00.050.2.01. Densovirus

Type Species 00.050.2.01.001. *Junonia coenia* densovirus (JcDNV)

List of Species in the Genus

Species, their serotypes, strains and isolates

- 00.050.2.01.002. *Galleria mellonella* densovirus [L32896] (GmDNV)
- 00.050.2.01.003. *Junonia coenia* densovirus [S17265] (JcDNV)
- 00.050.2.01.006. *Mythimna loreyi* densovirus (MIDNV)
- 00.050.2.01.007. *Pseudoplusia includens* densovirus (PiDNV)

Tentative Species in the Genus

- 00.050.2.81.014. *Acheta domestica* densovirus (AdDNV)
- 00.050.2.81.008. *Culex pipiens* densovirus (CpDNV)
- 00.050.2.81.005. *Diatraea saccharalis* densovirus (DsDNV)
- 00.050.2.81.009. *Periplanata fuliginosa* densovirus (PfDNV)

Genus 00.050.2.02. Iteravirus

Type Species 00.050.2.02.001. *Bombyx mori* densovirus (BmDNV)

List of Species in the Genus

Species, their serotypes, strains and isolates

- 00.050.2.02.001. *Bombyx mori* densovirus [M15123] (BmDNV)
- 00.050.2.02.001. *Bombyx mori* densovirus [M60583]
- 00.050.2.02.001. *Bombyx mori* densovirus [M60584]

Tentative Species in the Genus

- 00.050.2.83.004. *Casphalia extranea* densovirus (CeDNV)

Genus 00.050.2.03. Brevidensovirus

Type Species 00.050.2.03.001. *Aedes aegypti* densovirus (AaDNV)

List of Species in the Genus

00.050.2.03.001. *Aedes aegypti* densovirus (AaDNV)

00.050.2.03.002. *Aedes albopictus* densovirus (AIDNV)

Tentative Species in the Genus

00.050.2.83.014. *Acheta domestica* densovirus (AdDNV)

00.050.2.83.015. *Aedes pseudoscutellaris* densovirus (ApDNV)

00.050.2.83.003. *Agraulis vanillae* densovirus (AvDNV)

00.050.2.83.004. *Casphalia extranea* densovirus (CeDNV)

00.050.2.83.005. *Diatraea saccharalis* densovirus (DsDNV)

00.050.2.83.006. *Euxoa auxiliaris* densovirus (EaDNV)

00.050.2.83.007. *Leucorrhinia dubia* densovirus (LdDNV)

00.050.2.83.008. *Lymantria dubia* densovirus (LdDNV)

00.050.2.83.009. *Periplanata fuliginosa* densovirus (PfDNV)

00.050.2.83.010. *Pieris rapae* densovirus (PrDNV)

00.050.2.83.011. *Pseudalaetia includens* densovirus (PiDNV)

00.050.2.83.012. *Sibine fusca* densovirus (SfDNV)

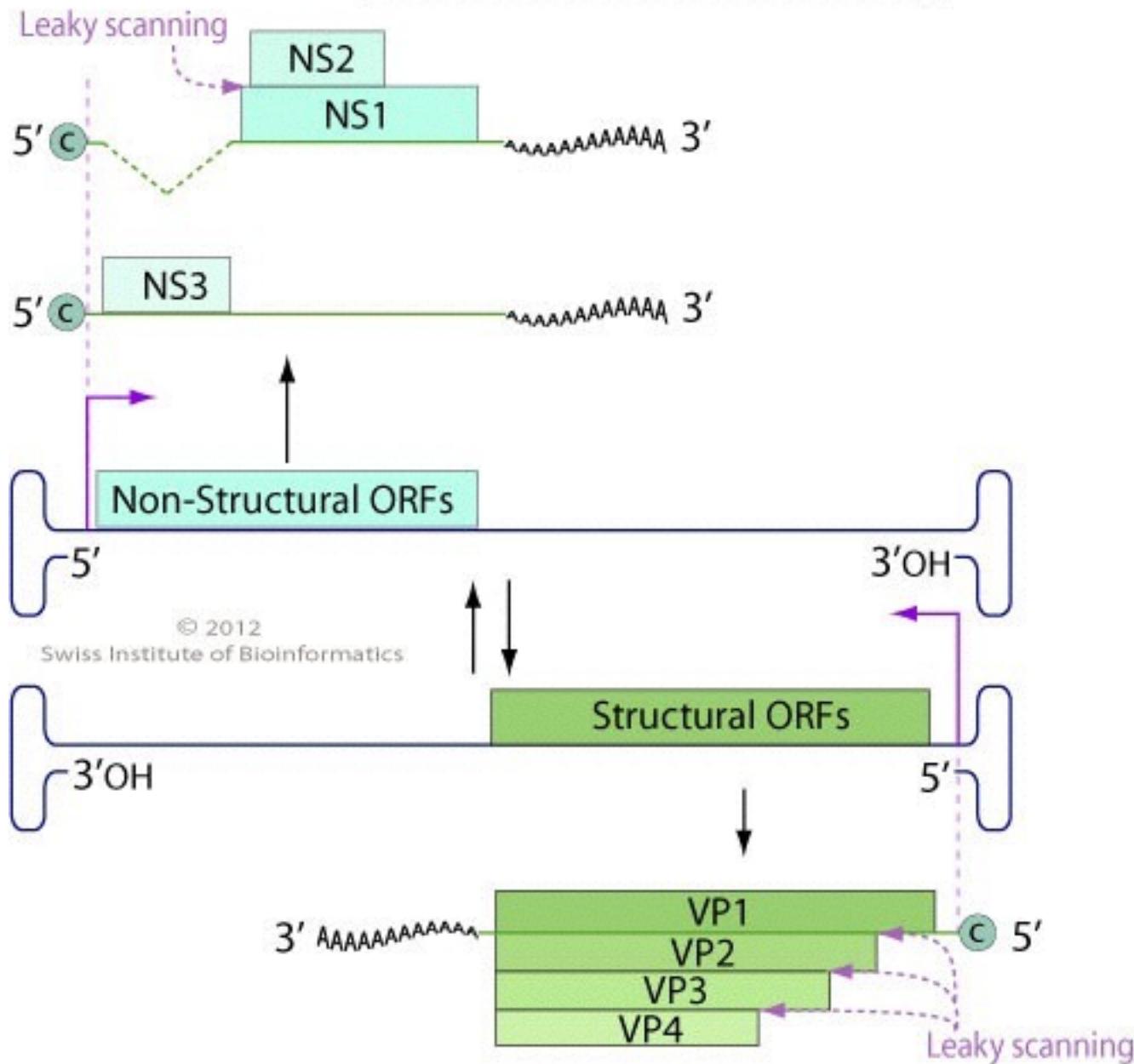
00.050.2.83.013. *Simulium vittatum* densovirus (SvDNV)

00.050.2.83.016. *Aedes Thai* densovirus (AThDNV)

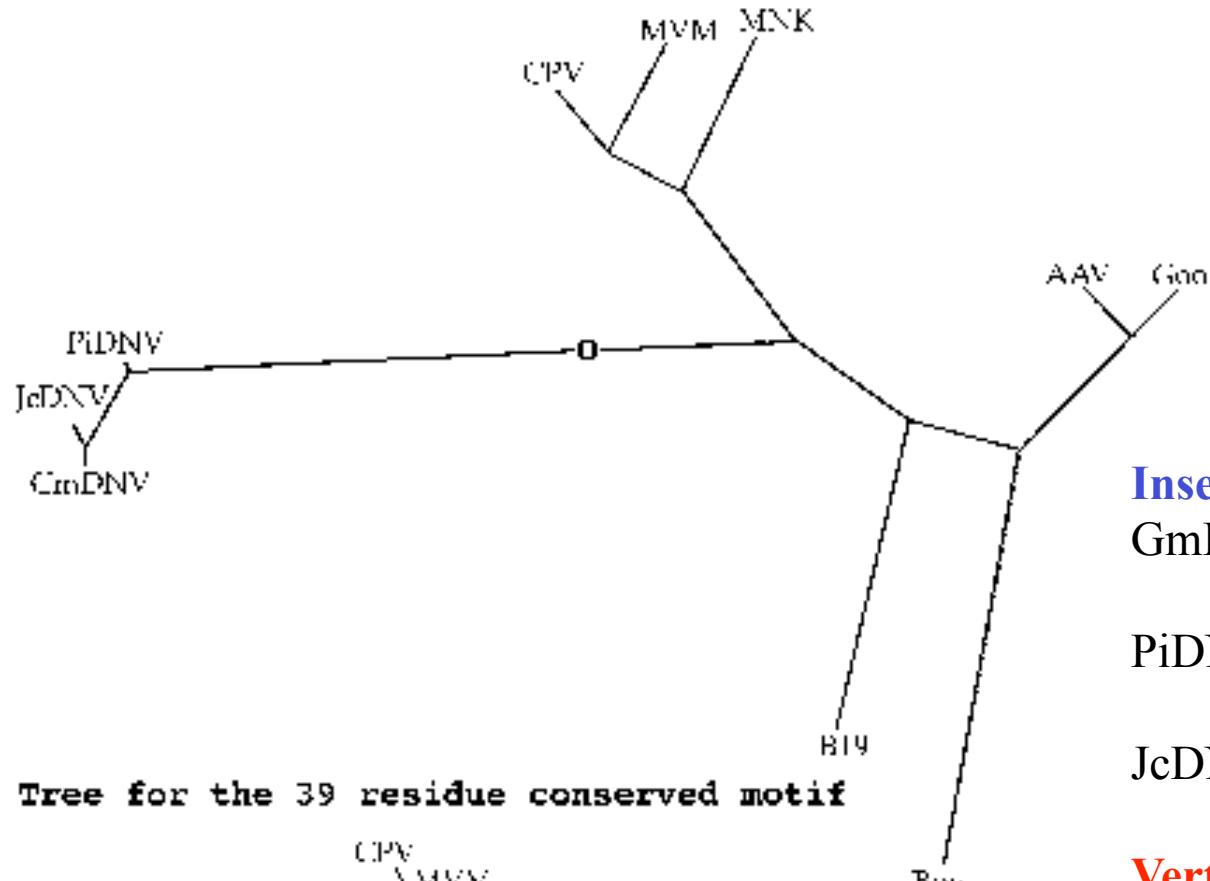
00.050.2.83.017. *Haemagogus equinus* densovirus (HeDNV)

00.050.2.83.018. *Toxorhynchites amboinensis* densovirus (TaDNV)

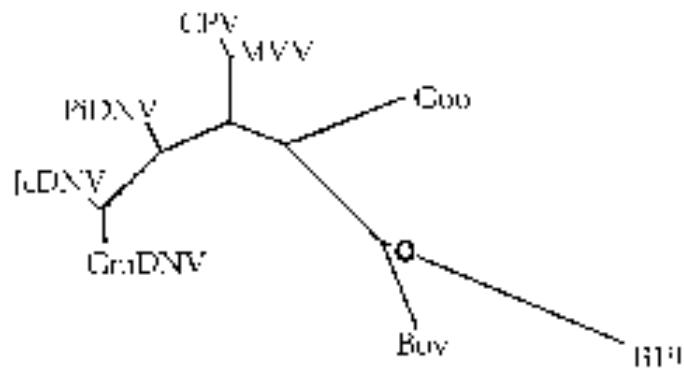
Galleria mellonella densovirus



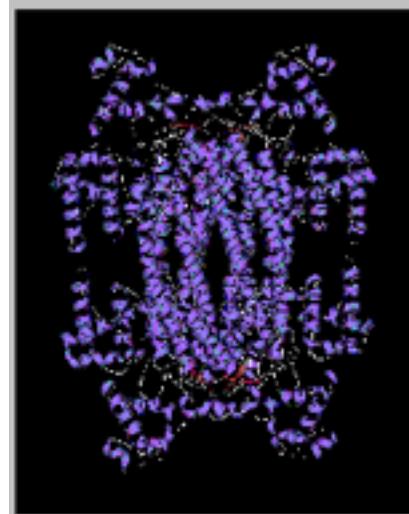
Tree for the structurally conserved regions



Tree for the 39 residue conserved motif



DELTA CRYSTALLIN



Insect parvoviruses (densovirus):

GmDNV :Galleria mellonella:
wax moth densovirus.

PiDNV :Pseudoplusia includens:
oybean looper densovirus

JcDNV :Junonia coenia densovirus

Vertebrate parvoviruses:

CPV :Canine parvovirus

MVM :Minute virus of mouse.

Bov :Bovine parvovirus

AAV :Adeno-associated virus

Mnk :Aleutian virus of mink

B19 :B19 (humnan) parvovirus

Goo :Goose parvovirus

RNA VIRUSES

Double-stranded RNA, nonenveloped

REOVIRIDAE, reoviruses

Cytoplasmic polyhedrosis virus

BIRNAVIRIDAE, bisegmented dsRNA viruses

Birnavirus, *Drosophila X* virus

Single-stranded RNA, enveloped

RHABDOVIRIDAE, bullet-shaped viruses

Sigmavirus, CO₂ sensitivity virus

Single-stranded RNA, nonenveloped

PICORNAVIRIDAE, picornaviruses

Cricket paralysis virus

Drosophila C virus

Gonometa virus

Infectious flacherie virus

Rhopalosiphum padi virus

CALICIVIRIDAE, caliciviruses

Amyelosis chronic stunt virus

Nudaurelia virus group

Nudaurelia virus

Trichoplusia RNA virus

Related viruses isolated from other Lepidoptera

NODAVIRIDAE

Nodavirus

Nodamura virus

Black beetle virus

Families of Viruses Infecting Invertebrates

dsRNA



Reoviridae
Cytopivirus



Birnaviridae
Entomabirnavirus

100 nm

Families of Viruses Infecting Invertebrates

ssRNA(-)



Anelloviridae

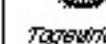


Bunyaviridae

ssRNA(+)



Picornaviridae



Togaviridae

Tetrviridae



Flaviviridae

Nodaviridae

100 nm

Species in the Genus

Cytopivirus 1 (CPV-1)
Bombyx mori cytopivirus 1 (BmCPV-1)
Dendrolimus spectabilis cytopivirus 1 (DsCPV-1)
Lymantria dispar cytopivirus 1 (LdCPV-1)
Cytopivirus 2 (CPV-2)
Aglais urticae cytopivirus 2 (AuCPV-2)
Agraulis vanillae cytopivirus 2 (AvaCPV-2)
Arctia caja cytopivirus 2 (AcCPV-2)
Arctia villica cytopivirus 2 (AviCPV-2)
Boloria dia cytopivirus 2 (BdCPV-2)
Dasychira pudibunda cytopivirus 2 (DpCPV-2)
Eriogaster lanestris cytopivirus 2 (ElCPV-2)
Hyloicus pinastri cytopivirus 2 (HpCPV-2)
Inachis io cytopivirus 2 (IiCPV-2)
Lacanobia oleracea cytopivirus 2 (LoCPV-2)
Malacosoma neustria cytopivirus 2 (MnCPV-2)
Mamestra brassicae cytopivirus 2 (MbCPV-2)
Operophtera brumata cytopivirus 2 (ObCPV-2)
Papilio machaon cytopivirus 2 (PmCPV-2)
Phalera bucephala cytopivirus 2 (PbCPV-2)
Pieris rapae cytopivirus 2 (PrCPV-2)
Cytopivirus 3 (CPV-3)
Anaitis plagiata cytopivirus 3 (ApCPV-3)
Arctia caja cytopivirus 3 (AcCPV-3)
Danaus plexippus cytopivirus 3 (DpCPV-3)
Gonometa rufibrunnea cytopivirus 3 (GrCPV-3)
Malacosoma neustria cytopivirus 3 (MnCPV-3)
Operophtera brumata cytopivirus 3 (ObCPV-3)
Phlogophora meticulosa cytopivirus 3 (PmCPV-3)
Pieris rapae cytopivirus 3 (PrCPV-3)
Spodoptera exempta cytopivirus 3 (SexmCPV-3)
Cytopivirus 4 (CPV-4)
Actias selene cytopivirus 4 (AsCPV-4)
Antheraea mylitta cytopivirus 4 (AmCPV-4)
Antheraea pernyi cytopivirus 4 (ApCPV-4)

Cytopivirus 5 (CPV-5)

Euxoa scandens cytopivirus 5 [J04338] Seg 10 (EsCPV-5)
Heliothis armigera cytopivirus 5 (HaCPV-5)
Orgyia pseudosugata cytopivirus 5 (OpCPV-5)
Spodoptera exempta cytopivirus 5 (SexmCPV-5)
Trichoplusia ni cytopivirus 5 (TnCPV-5)
Cytopivirus 6 (CPV-6)
Aglais urticae cytopivirus 6 (AuCPV-6)
Agrochola helvolva cytopivirus 6 (AhCPV-6)
Agrochola lychnidis cytopivirus 6 (AlCPV-6)
Anaitis plagiata cytopivirus 6 (ApCPV-6)
Anti xanthomista cytopivirus 6 (AxCPV-6)
Biston betularia cytopivirus 6 (BbCPV-6)
Eriogaster lanestris cytopivirus 6 (ElCPV-6)
Lasiocampa quercus cytopivirus 6 (LqCPV-6)
Cytopivirus 7 (CPV-7)
Mamestra brassicae cytopivirus 7 (MbCPV-7)
Noctua pronuba cytopivirus 7 (NpCPV-7)
Cytopivirus 8 (CPV-8)
Abrajas grossulariata cytopivirus 8 (AgCPV-8)
Heliothis armigera cytopivirus 8 (HaCPV-8)
Malacosoma disstria cytopivirus 8 (MdCPV-8)
Nudaurelia cytherea cytopivirus 8 (NcCPV-8)
Phlogophora meticulosa cytopivirus 8 (PmCPV-8)
Spodoptera exempta cytopivirus 8 (SexmCPV-8)
Cytopivirus 9 (CPV-9)
Agrotis segetum cytopivirus 9 (AsCPV-9)
Cytopivirus 10 (CPV-10)
Aporophyla lutulenta cytopivirus 10 (AlCPV-10)

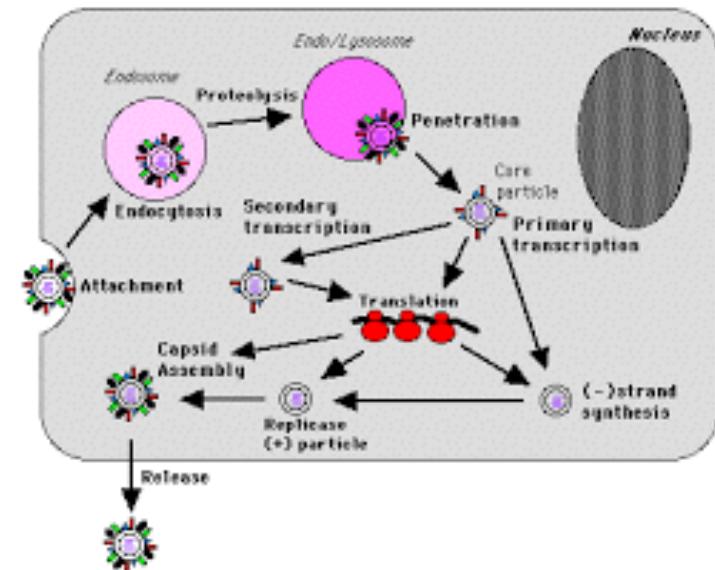
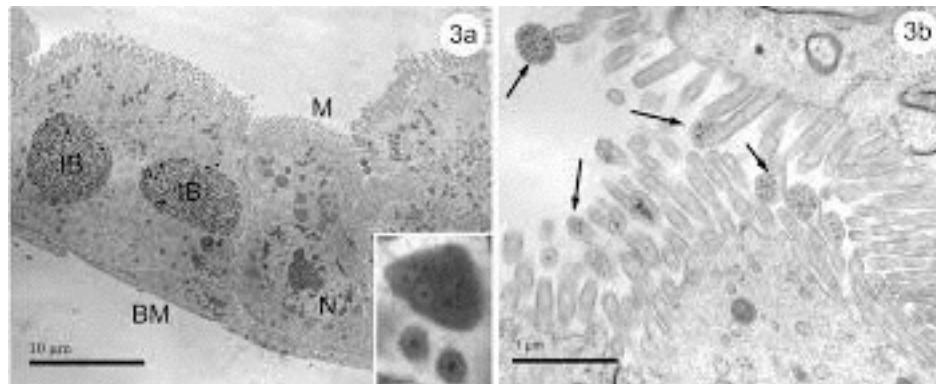
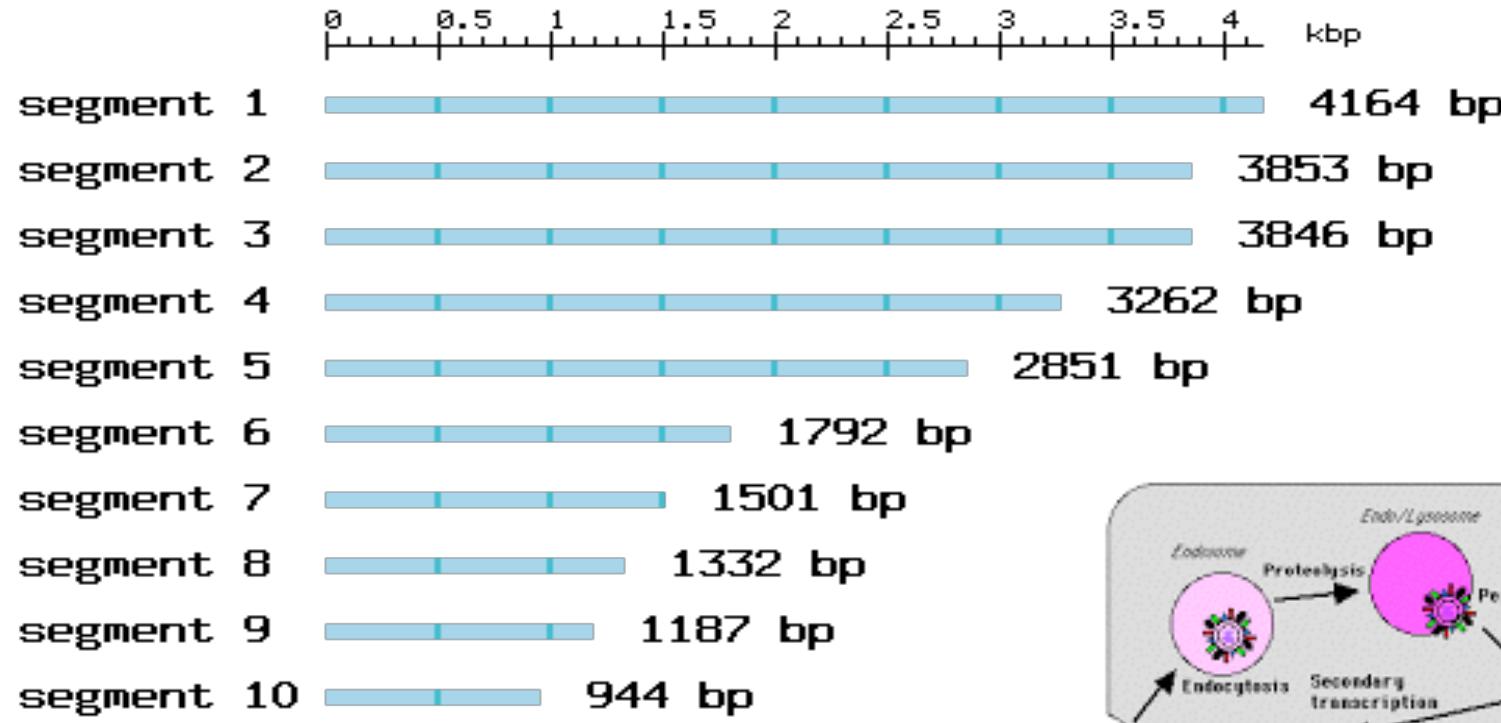
Cypovirus 11 (CPV-11)

Heliothis armigera cypovirus 11 (HaCPV-11)
Heliothis zea cypovirus 11 (HzCPV-11)
Lymantria dispar cypovirus 11 (LdCPV-11)
Mamestra brassicae cypovirus 11 (MbCPV-11)
Pectinophora gossypiella cypovirus 11 (PgCPV-11)
Pseudaletia unipuncta cypovirus 11 (PuCPV-11)
Spodoptera exempta cypovirus 11 (SexmCPV-11)
Spodoptera exigua cypovirus 11 (SexgCPV-11)
Cypovirus 12 (CPV-12)
Autographa gamma cypovirus 12 (AgCPV-12)
Mamestra brassicae cypovirus 12 (MbCPV-12)
Pieris rapae cypovirus 12 (PrCPV-12)
Spodoptera exempta cypovirus 12 (SexmCPV-12)
Cypovirus 13 (CPV-13)
Polistes hebraeus cypovirus 13 (PhCPV-13)
Cypovirus 14 (CPV-14)
Heliothis armigera cypovirus 14 ('A' strain) (HaCPV-14)

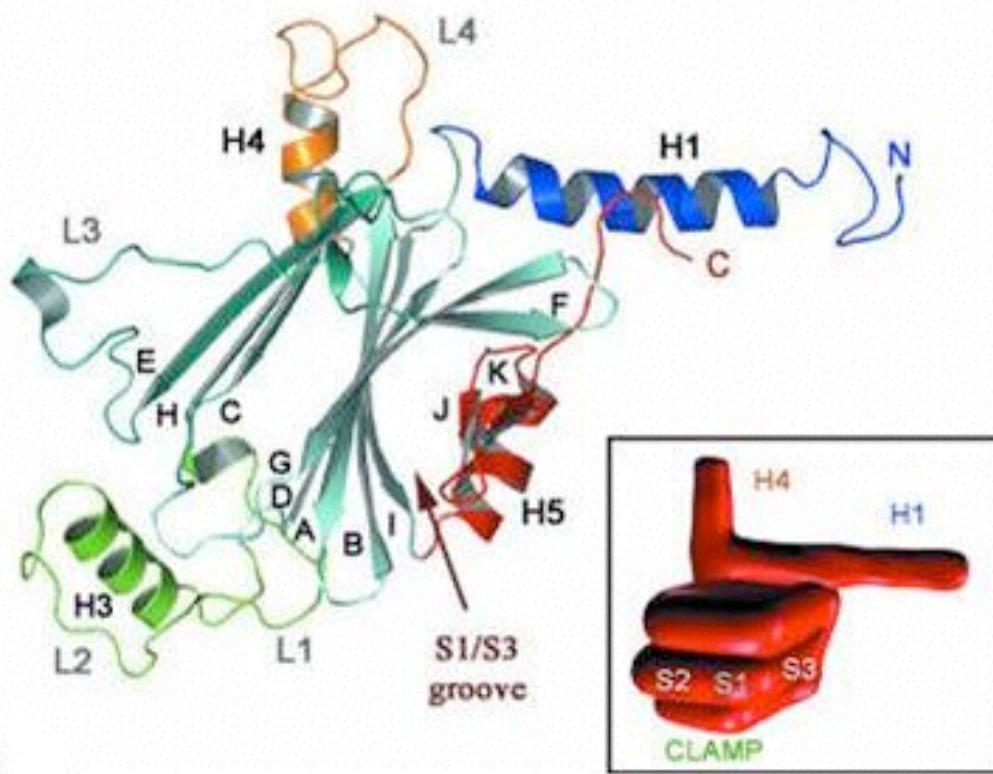
Taxonomic Structure of the Genus

Cypoviruses are currently classified within **14 species** that were initially characterised by their **distinctive dsRNA electropherotype patterns**. Cross-hybridisation analyses of dsRNA, limited comparative RNA sequence data and serological comparisons of cypovirus proteins have confirm the validity of this classification. However, only a few cypoviruses have been analysed in this way.

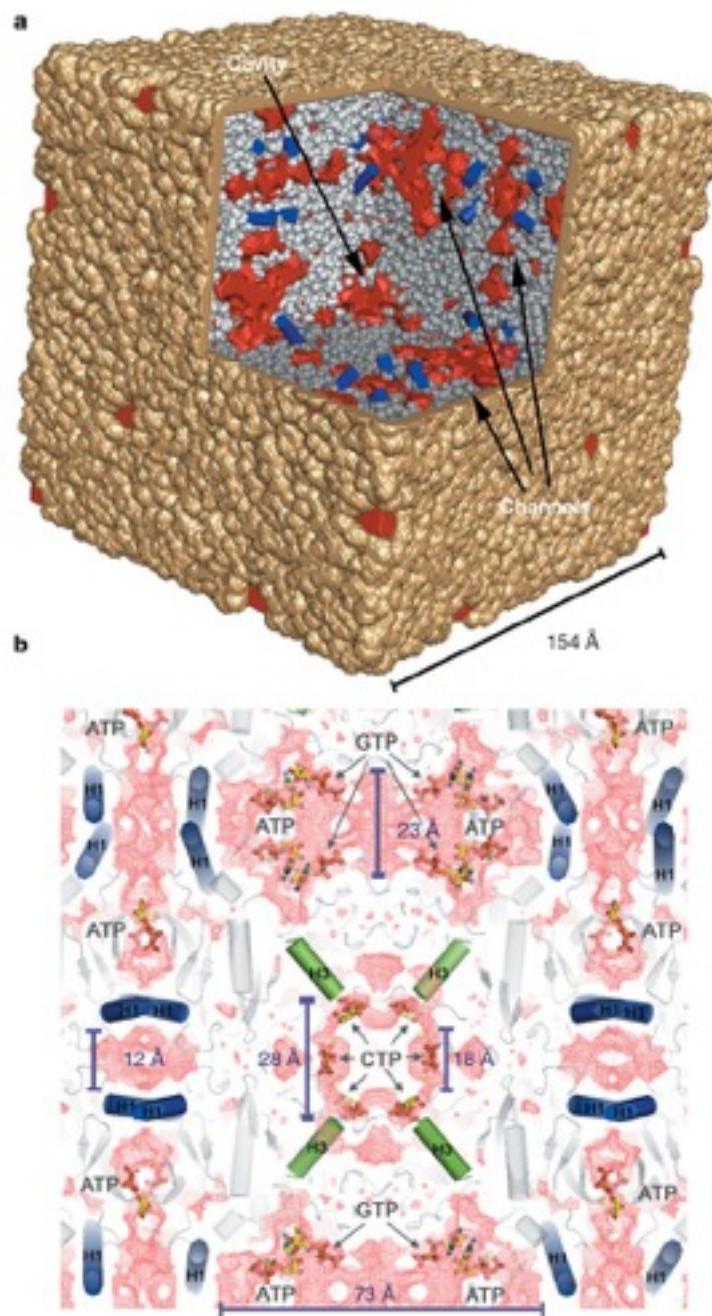
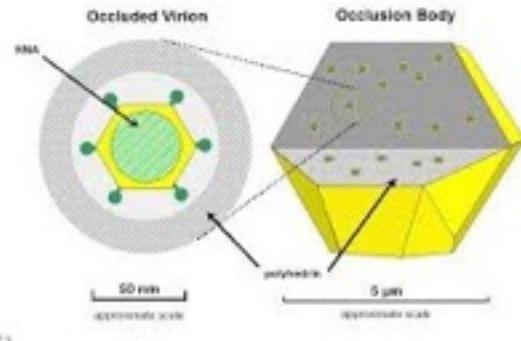
Lymantria dispar cypovirus 1 complete genome



Lineage:Cypovirus; Reoviridae; dsRNA viruses



Cypovirus
Cytoplasmic polyhedrovirus virus



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Gonometa virus

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Rhopalosiphum padi virus

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Amyelosis chronic stunt virus

Nudaurelia virus group

Nudaurelia virus

Trichoplusia RNA virus

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Nodavirus

Nodamura virus

Black beetle virus

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dsRNA



Reoviridae
Cytopivirus



Bimaviridae
Entomobimavirus

100 nm

Families of Viruses Infecting Invertebrates

ssRNA(-)



Anellovirus



Bunyaviridae

ssRNA(+)



Picornaviridae



Togaviridae

Tetrviridae



Flaviviridae

Nodaviridae

100 nm

Family 00.009. Birnaviridae

Genus 00.009.0.01. Aquabirnavirus

Genus 00.009.0.02. Avibirnavirus

Genus 00.009.0.03. **Entomobirnavirus**

Species, their serotypes, strains and isolates

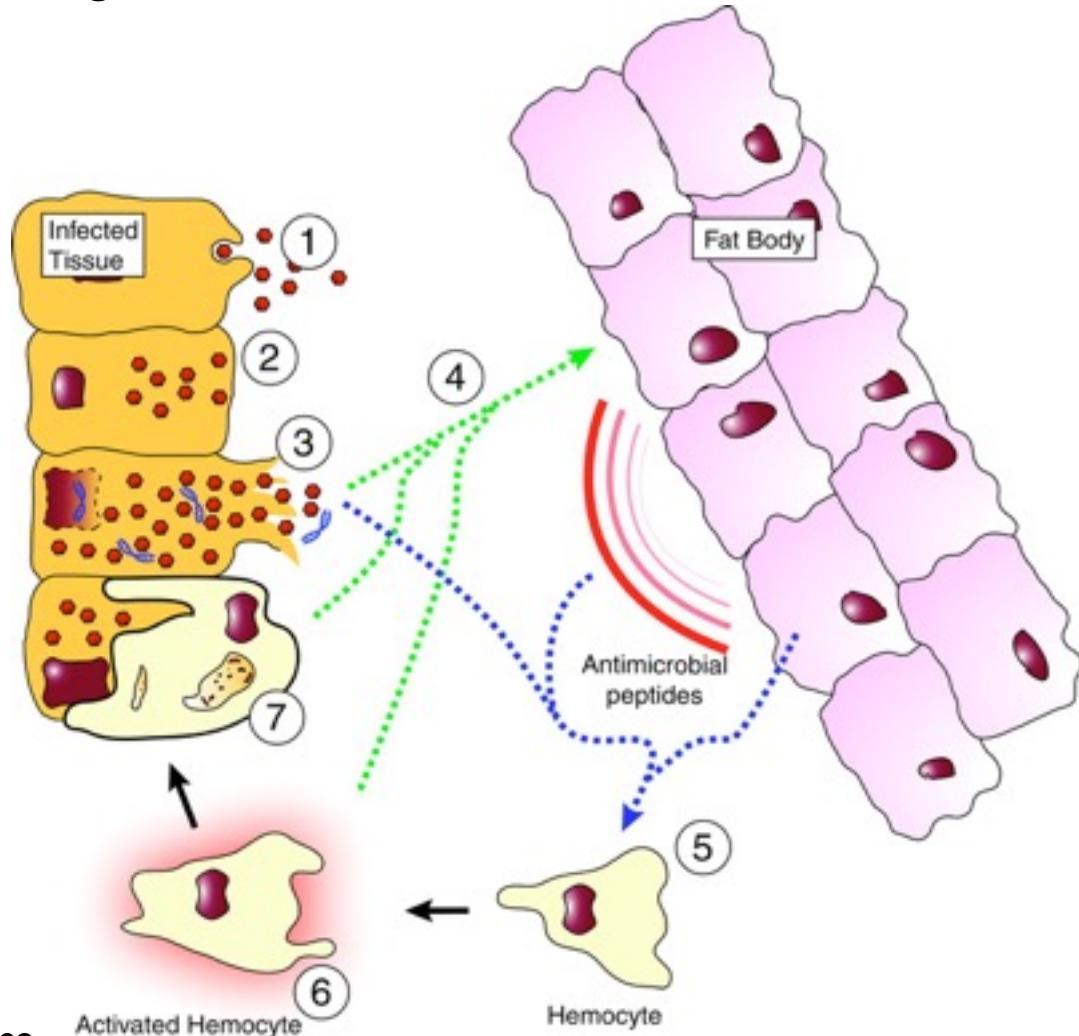
00.009.0.03.001. **Drosophila X virus** [U60650] RNA-A (DXV)

Sequence analysis of the bicistronic Drosophila X virus genome segment A and its encoded polypeptides

Tentative Species in the Genus

None reported.

Model for *Drosophila* antiviral response. 1, Virus enters system and infects the cell. 2, Virus replication. 3, Lysis of infected cell, releasing internal cellular compounds and virus. 4, Released materials activate the IMD and Toll humoral immune pathways and cause local activation of the cellular response. 5, Global activation of hemocytes via Toll pathway signaling. 6, Activated hemocytes signal to the fat body enhancing Toll and IMD pathways activation. 7, Hemocytes recognize aberrant infected cells and engulf and eliminate these cells



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Double-stranded RNA, nonenveloped

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BIRNAVIRIDAE, bisegmented dsRNA viruses

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PICORNAVIRIDAE, picornaviruses

Cricket paralysis virus

Drosophila C virus

Gonometa virus

Infectious flacherie virus

Rhopalosiphum padi virus

CALICIVIRIDAE, caliciviruses

Amyelosis chronic stunt virus

Nudaurelia virus group

Nudaurelia virus

Trichoplusia RNA virus

Related viruses isolated from other Lepidoptera

NODAVIRIDAE

Nodavirus

Nodamura virus

Black beetle virus

Families of Viruses Infecting Invertebrates

dsRNA



Reoviridae
Cytopivirus



Birnaviridae
Entomabirnavirus

100 nm

Families of Viruses Infecting Invertebrates

ssRNA(-)



Anelloviridae

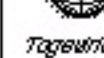


Bunyaviridae

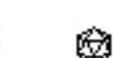
ssRNA(+)



Picornaviridae



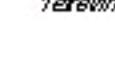
Togaviridae



Tetrviridae



Flaviviridae



Nodaviridae

100 nm

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ARTICLES

Carbon dioxide sensitivity in mosquitoes infected with sigma, vesicular stomatitis, and other rhabdoviruses

L Rosen

Carbon dioxide, usually an innocuous narcotic for insects, kills mosquitoes infected with rhabdoviruses. Such toxicity was originally observed in *Drosophila* harboring a hereditary virus, sigma, and has been considered unique to *Drosophila*. The new findings support the possibility that insects with piercing and sucking mouthparts harbor similar hereditary viruses and transmit some of them to vertebrates or plants.

RNA VIRUSES

Double-stranded RNA, nonenveloped

REOVIRIDAE, reoviruses

Cytoplasmic polyhedrosis virus

BIRNAVIRIDAE, bisegmented dsRNA viruses

Birnavirus, *Drosophila X* virus

Single-stranded RNA, enveloped

RHABDOVIRIDAE, bullet-shaped viruses

Sigmavirus, CO₂ sensitivity virus

Single-stranded RNA, nonenveloped

PICORNAVIRIDAE, picornaviruses

Cricket paralysis virus

Drosophila C virus

Gonometa virus

Infectious flacherie virus

Rhopalosiphum padi virus

CALICIVIRIDAE, caliciviruses

Amyelosis chronic stunt virus

Nudaurelia virus group

Nudaurelia virus

Trichoplusia RNA virus

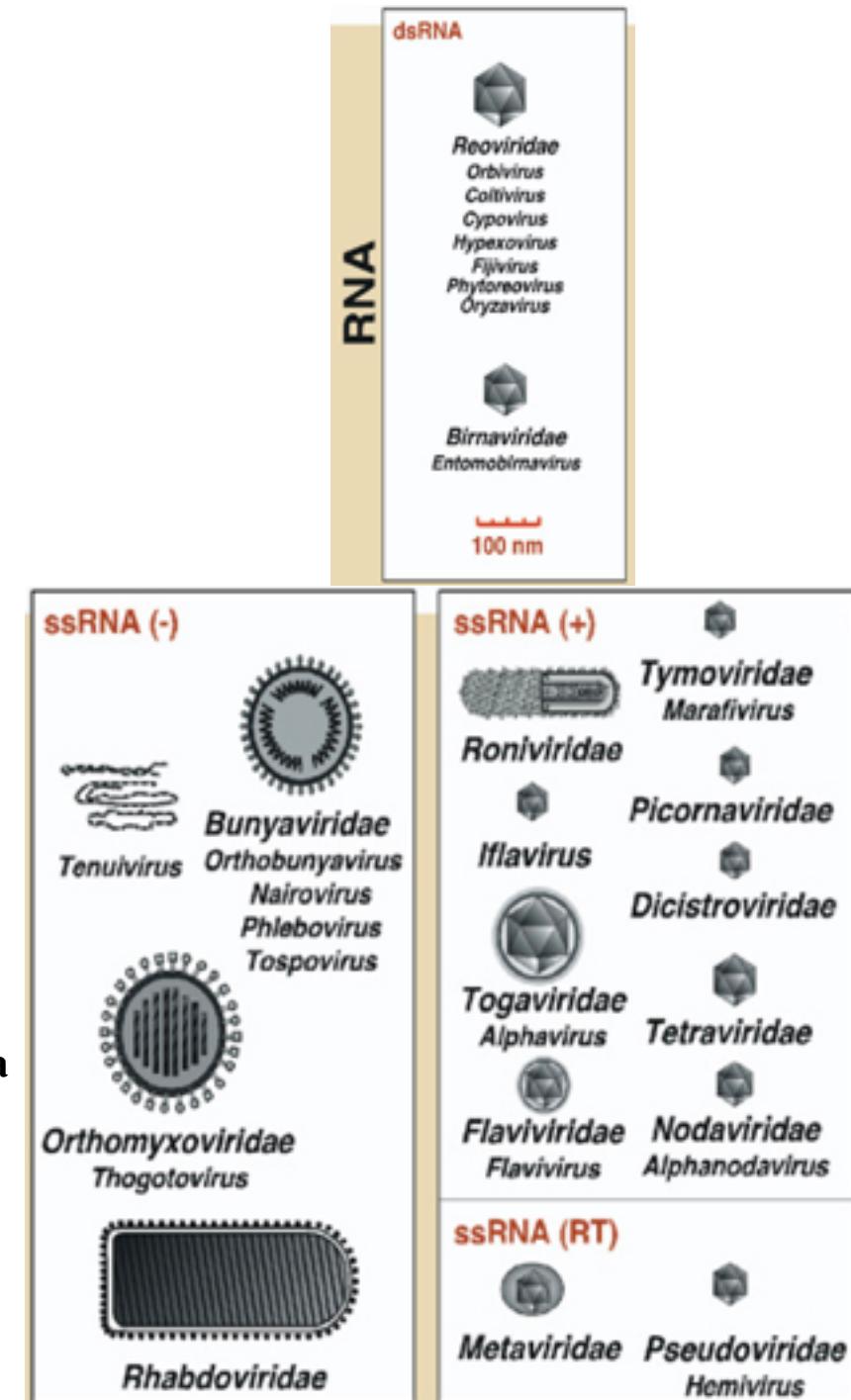
Related viruses isolated from other Lepidoptera

NODAVIRIDAE

Nodavirus

Nodamura virus

Black beetle virus



List of Invertebrate Picorna Viruses -listed by N.J. Knowles

Viruses isolated from **Orthoptera** (直翅目：コオロギ)

Cricket paralysis virus (CrPV) field crickets, *Teleogryllus oceanicus* and *T. commodus* Reinganum et al., 1970

Viruses isolated from **Diptera** (双翅目：カ・ハエ)

Drosophila C virus (DCV) Fruit fly, *Drosophila melanogaster* Jousset et al., 1972

Drosophila P virus (DPV) Fruit fly, *Drosophila melanogaster* Plus and Duthoit, 1969; Tenniges and Plus, 1972

Drosophila A virus (DAV) Fruit fly, *Drosophila melanogaster* ?

Queensland fruit fly virus (QFFV) Fruit fly *Dacus tryoni* Bashiruddin et al., 1988

Aedes taeniorhynchus virus ? Mosquito, *Aedes taeniorhynchus* Wagner et al., 1974

Kawino virus (KV) mosquito, *Mansonia uniformis* Pudney et al., 1978

Viruses isolated from **Lepidoptera** (鱗翅目：チョウ・ガ)

Gonometa virus ? caterpillar, *Gonometa podocarpi* Harrap et al., 1966

Infectious flacherie virus (IFV) silkworm, *Bombyx mori* Kawase et al., 1980

Latoia viridissima virus ? oil palm pest, *Latoia viridissima* Fediere et al., 1990

Pectinophora gossypiella virus (PgV) cotton pink bollworm, *Pectinophora gossypiella* Monsarrat et al., 1995

Perina nuda picornavirus (PnPV) Ficus transparent wing moth, Lymantriidae *Perina nuda* Wang et al., 1999

Viruses isolated from **Hymenoptera** (膜翅目 : ハチ)

| | |
|--|------------------------|
| Bee slow paralysis virus (BSPV) honey bee, <i>Apis mellifera</i> | Bailey and Woods, 1974 |
| Bee acute paralysis virus (BAPV) honey bee, <i>Apis mellifera</i> | Bailey et al., 1963 |
| bee virus X (BVX) honey bee, <i>Apis mellifera</i> | Bailey and Woods, 1974 |
| sacbrood virus (SBV) honey bee, <i>Apis mellifera</i> | Bailey et al., 1964 |
| Thai sacbrood virus ? Eastern hive bee, <i>Apis cerana</i> | Bailey et al., 1982 |
| Egypt bee virus (EBV) honey bee, <i>Apis mellifera</i> | Bailey et al., 1979 |
| Kashmir bee virus (KBV) Eastern hive bee, <i>Apis cerana</i> | Bailey et al., 1979 |
| South Australia Kashmir bee virus ? honey bee, <i>Apis mellifera</i> | Bailey et al., 1979 |
| black queen-cell virus (BQCV) honey bee, <i>Apis mellifera</i> | Bailey and Woods, 1977 |

Viruses isolated from **Hemiptera** (半翅目 : アブラムシ・カメムシ)

| | |
|--|--|
| Rhopalosiphum padi virus (RhPV) Oat bird cherry aphid, <i>Rhopalosiphum padi</i> | D'Arcy et al., 1981 |
| aphid lethal paralysis virus (ALPV) Oat bird cherry aphid, <i>Rhopalosiphum padi</i> | Williamson et al., 1988 |
| Acyrthosiphon pisum virus (APV) pea aphid, <i>Acyrthosiphon pisum</i> | van den Heuvel et al., 1997; van der Wilk et al., 1997 |
| Triatoma virus (TrV) Kissing bug, <i>Triatoma infestans</i> | Muscio et al., 1987, 1988 |
| Plautia stali intestine virus (PSIV) Brown-winged green bug, <i>Plautia stali</i> | Nakashima et al., 1998 |

Viruses isolated from **Isoptera** (等翅目 : シロアリ)

| | |
|---------------------------------------|---------------|
| termite paralysis virus (TPV) Termite | Gibbs et al., |
|---------------------------------------|---------------|

タンパク質合成開始の新機構

中島信彦



真核生物の普通の伝令RNA(mRNA)の先頭にはキャップ構造があり、この構造を翻訳開始因子類が認識し、そこにリボソームが結合する事によって翻訳が始まるとしてされています。これに対して、
ポリオウイルスやC型肝炎ウイルスなど一部のRNAウイルスは、mRNAのコード領域の上流にリボソーム内部進入部位(Internal Ribosome Entry Site、略してIRES)というRNAの高次構造を形成し、この構造を足掛かりにリボソームがmRNAに結合します。そのため翻訳開始の際にキャップ構造も、翻訳開始因子類のいくつかも必要ありません。しかし、どちらの場合にも翻訳開始メチオニンtRNAという翻訳開始の時にだけ使用される特殊なtRNAが、既にリボソームに組み込まれた状態でmRNAに結合します。したがって、蛋白質の合成は必ずメチオニンから始まります。つまり、これまでメチオニン以外のアミノ酸からのタンパク質

合成開始は不可能と考えられていました。

しかし、チャバネアオカメムシ腸管ウイルス(PSIV)の外被タンパク質遺伝子の翻訳開始では、翻訳開始メチオニンtRNAと翻訳開始コドンとの塩基対合は不要です。図に示したRNAの高次構造が自分で翻訳開始部位を決める働きをしているからです。これまでの研究結果から、

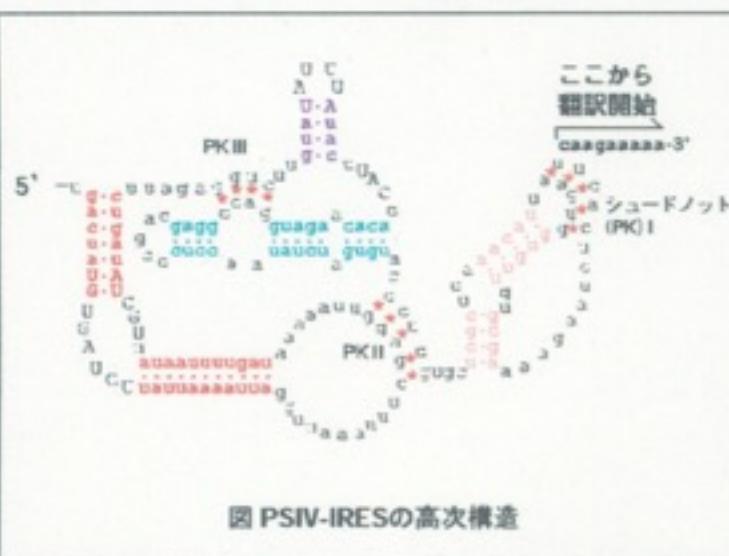


図 PSIV-IRESの高次構造

PSIVのIRESでは約180塩基があれば翻訳開始を行うことができ、その中には、赤、紫、緑、ピンクで示した4本のステム構造と、赤のアスタリスクで表した3カ所のシードノット構造が形成されます。このようなRNAの高次構造が形成されれば、このシードノットの直ぐ下流から翻訳が始まります。また、外被タンパク質はウイルスにとって最

も大量に合成する必要のあるタンパク質で、このIRESはタンパク質を大量合成するために特殊化したと考えられます。今後さらに解析が進めば、任意のアミノ酸からの蛋白質合成が可能になるかもしれません。

(昆虫適応遺伝研究グループ 昆虫共生媒介機構研究チーム主任研究官)

Picornavirus-like viruses of invertebrates

FAMILY I:

| | | |
|---|-------------------|------------------------------|
| Cricket paralysis virus (CrPV) | CG 9185 AF218039 | Wilson et al., 2000 |
| Drosophila C virus (DCV) EB 1998 | CG 9264 AF014388 | Johnson and Christian, |
| Plautia stali intestine virus (PSIV) | CG 8797 AB006531 | Sasaki et al., 1998 |
| Rhopalosiphum padi virus (RhPV) | CG 10011 AF022937 | Moon et al., 1998 |
| Himetobi P virus (HiPV) | CG 9275 AB017037 | Nakashima et al., 1999 |
| South African | CG 8550 AF183905 | Leat et al., 2000 |
| Triatoma virus (TrV) | CG 9010 AF178440 | Czibener et al., 2000 |
| Taura syndrome virus (TSV) capsid-coding region | 3278 AF277378 | Robles-Sikisaka et al., 2001 |

FAMILY 2:

| | | |
|---------------------------------|-------------------|---------------------------|
| Acyrthosiphon pisum virus (APV) | CG 10035 AF024514 | van der Wilk et al., 1997 |
|---------------------------------|-------------------|---------------------------|

FAMILY 3:

| | | |
|----------------------------------|------------------|--------------------|
| Infectious flacherie virus (IFV) | CG 9650 AB000906 | Isawa et al., 1998 |
| Sacbrood virus (SBV) Rothamstead | CG 8832 AF092924 | Ghosh et al., 1999 |

CG = complete genome; (p) = partial gene sequence;
N/A = not available on the EMBL/GenBank/DDBJ databases

Unclassified small spherical RNA viruses

| | | |
|--|-------|-----------------------------|
| Arkansas bee virus | APV | Acute Paralysis Virus |
| Bee acute paralysis virus | ArkBV | Arkansas Bee Virus |
| Bee chronic paralysis virus associate | BQCV | Black Queen Cell Virus |
| Bee slow paralysis virus | BVX | Bee Virus X |
| Bee virus X | BVY | Bee Virus Y |
| Bee virus Y | CBPV | Chronic Bee Paralysis Virus |
| Black queen cell virus | CWV | Cloudy Wing Virus |
| Cloudy wing virus | DWV | Deform Wing Virus |
| Crystalline array virus | EBV | Egypt Bee Virus |
| Drosophila A virus | FBV | Filamentous Virus |
| Drosophila P virus | JEBV | Japon strain of Egypt Virus |
| Kashmir bee virus | KBV | Kashmir Bee Virus |
| Kelp fly virus | SBV | Sacbrood Bee Virus |
| Sacbrood virus | SPV | Slow Paralysis Virus |

Unclassified small ovoid RNA viruses

- Bee chronic paralysis virus**
- Drosophila RS virus**

OTHER UNCLASSIFIED VIRUSES

Rod-shaped, enveloped

Cricket macronucleosis virus

Hypera virus

Long flexuous rod, enveloped

Bee filamentous virus

Drosophila (ショウジョウバエ)のRNAウイルス

Drosophila ananassae Tom virus (DanTomV) Metaviridae

Drosophila C virus (DCV) "CrPV-like viruses"

Drosophila F virus (DFV) Reoviridae

Drosophila line virus (DLV) Nodaviridae

Drosophila melanogaster 17.6 virus (Dme176V) Metaviridae

Drosophila melanogaster 1731virus (Dme1731V) Pseudoviridae

Drosophila melanogaster 297 virus (Dme297V) Metaviridae

Drosophila melanogaster 412 virus (Dme412V) Metaviridae

Drosophila melanogaster copia virus (DmeCopV) Pseudoviridae

Drosophilamelanogaster gypsy virus (DmeGypV) Metaviridae

Drosophila melanogaster mdg1virus (DmeMdg1V) Metaviridae

Drosophila melanogaster mdg4 virus (DmeMdg4V) Metaviridae

Drosophila melanogaster micropia virus (DmeMicV) Metaviridae

Drosophila P virus (DPV) Unassigned

Drosophila S virus (DSV) Reoviridae

Drosophila virilis Ulysses virus (DviU11V) Metaviridae

Drosophila X virus (DXV) Birnaviridae

Drosophilia A virus (DAV) Unassigned