Structural Virology

Lecture 4

Pavel Plevka

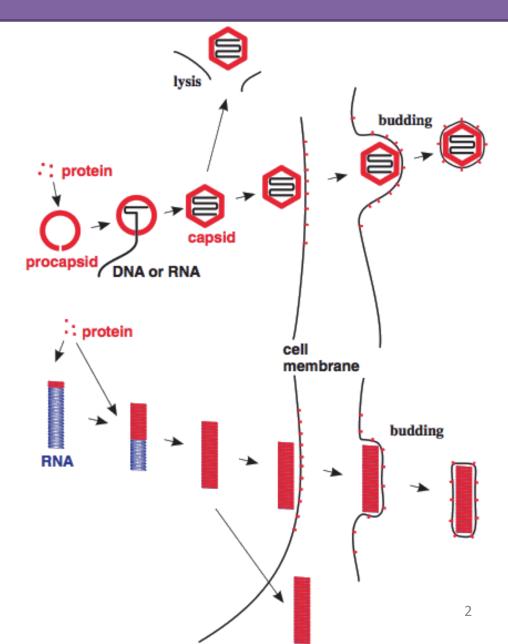


Financováno Evropskou unií NextGenerationEU

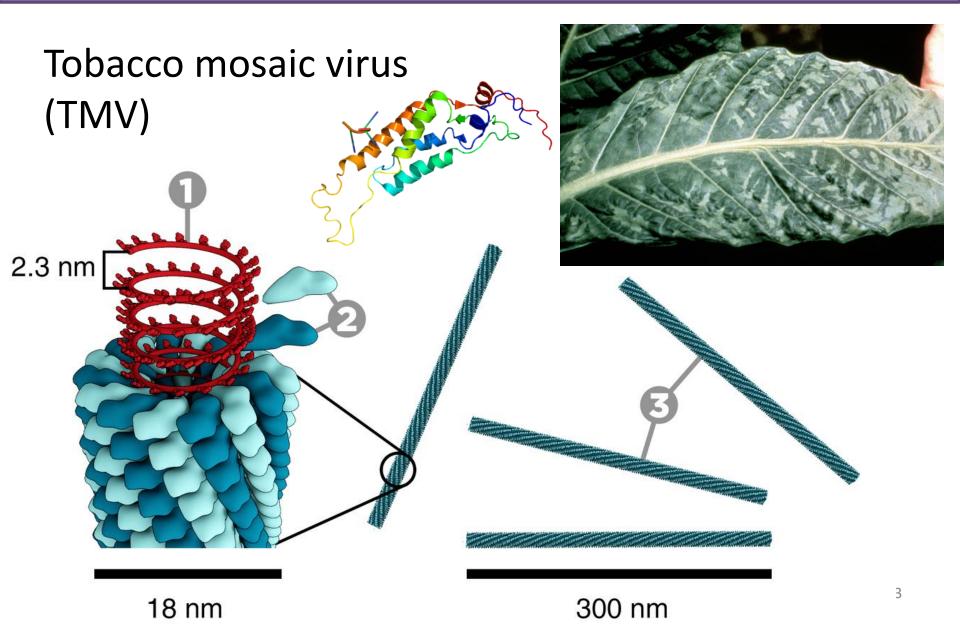


Assembly and exit of virions from cells

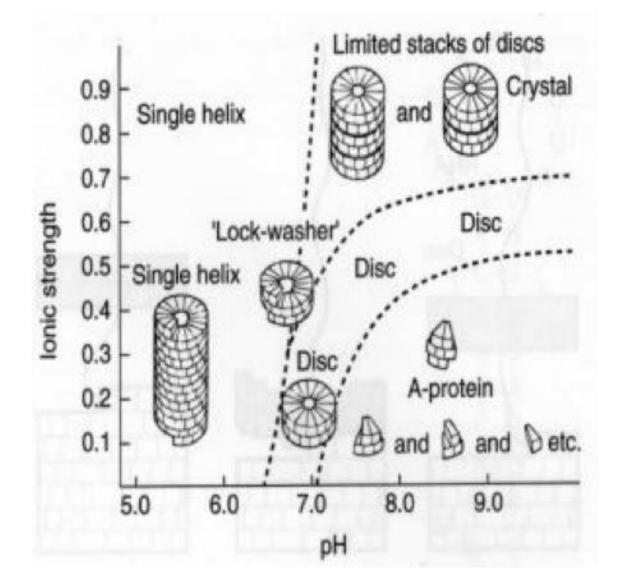
- 1. Attachment
- 2. Entry
- 3. Transcription
- 4. Translation
- 5. Genome replication
- 6. Assembly
- 7. **E**xit



Assembly of helical viruses

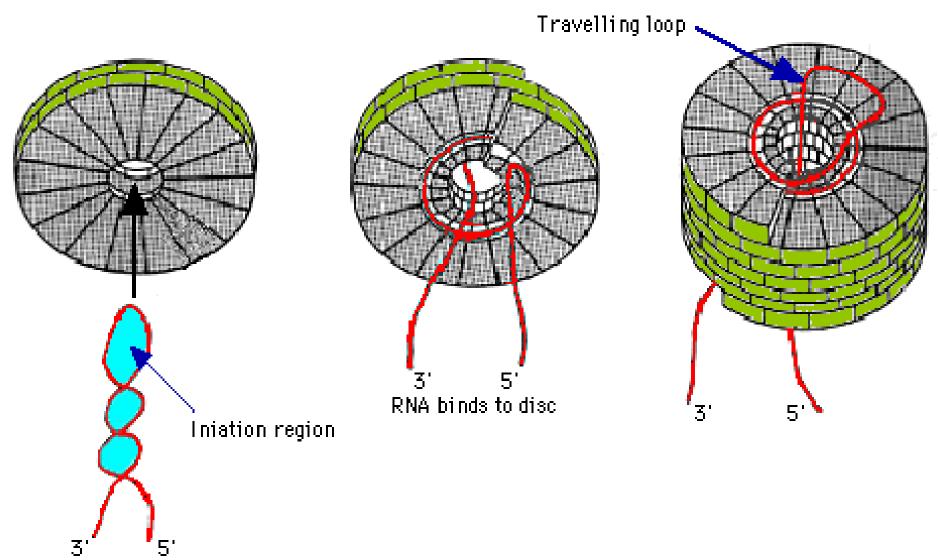


Effects of pH and ionic strength on formation of TMV capsid protein aggregates

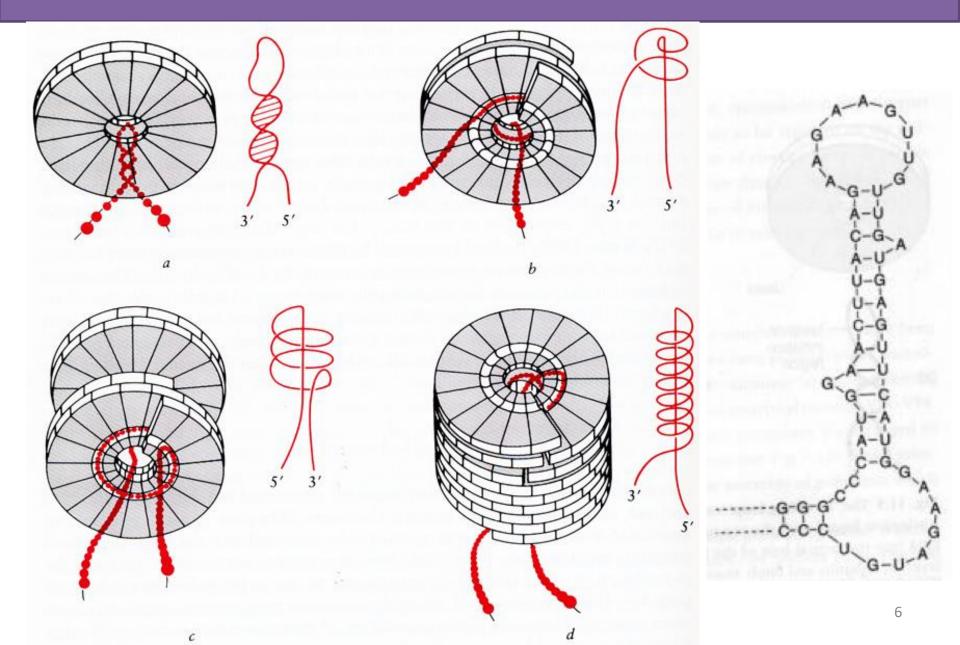


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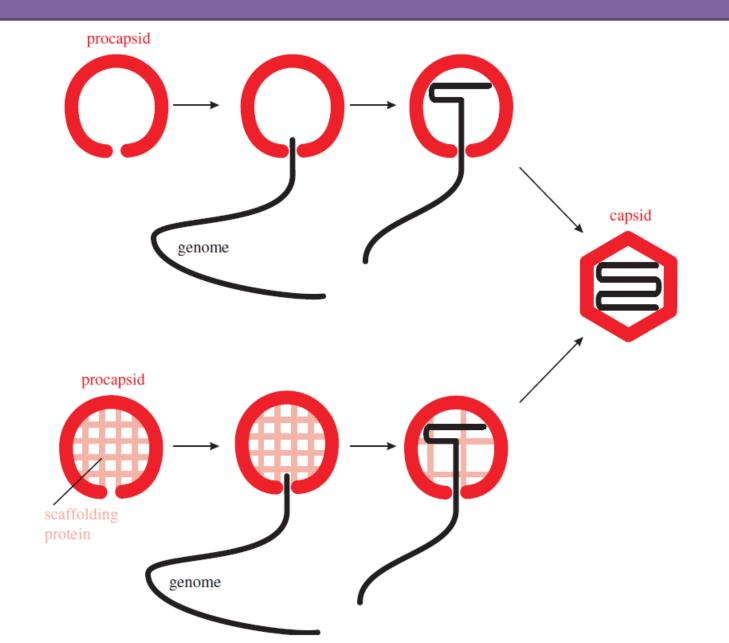
TMV initiation of assembly



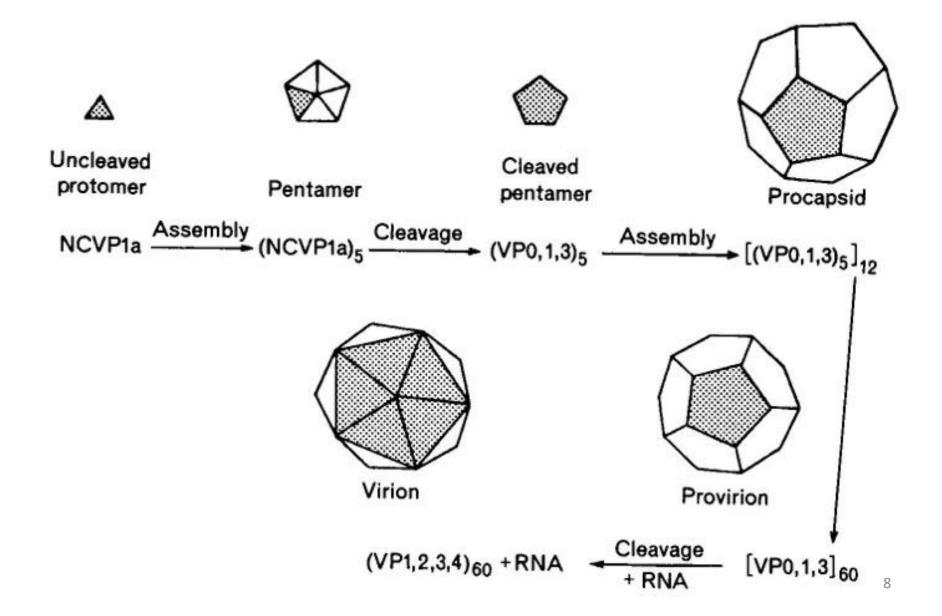
TMV assembly (GMO tobacco)

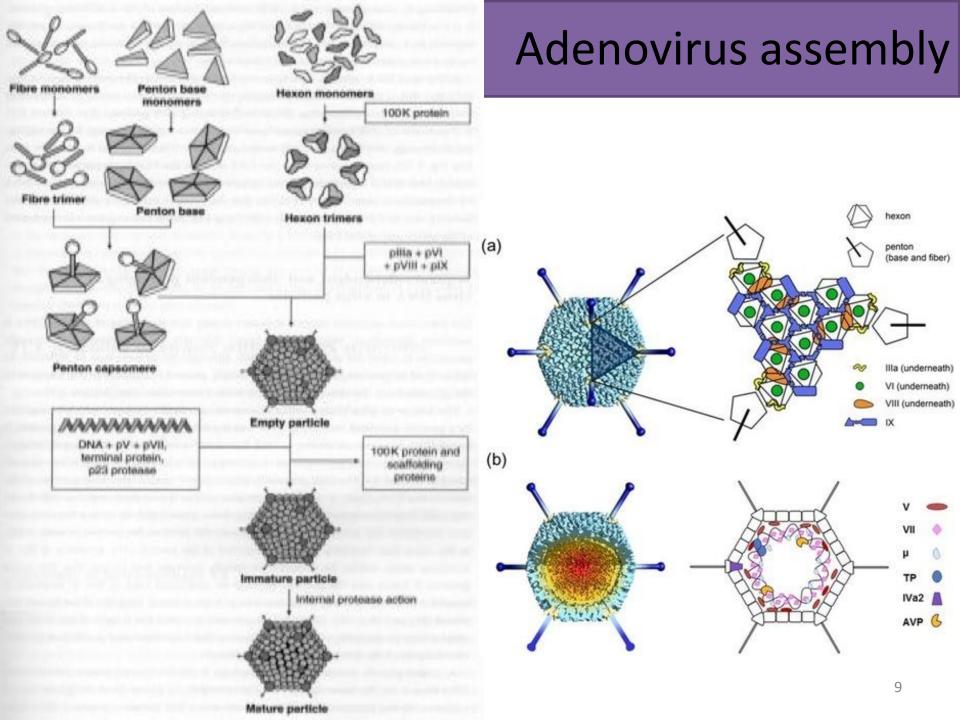


Assembly of icosahedral viruses

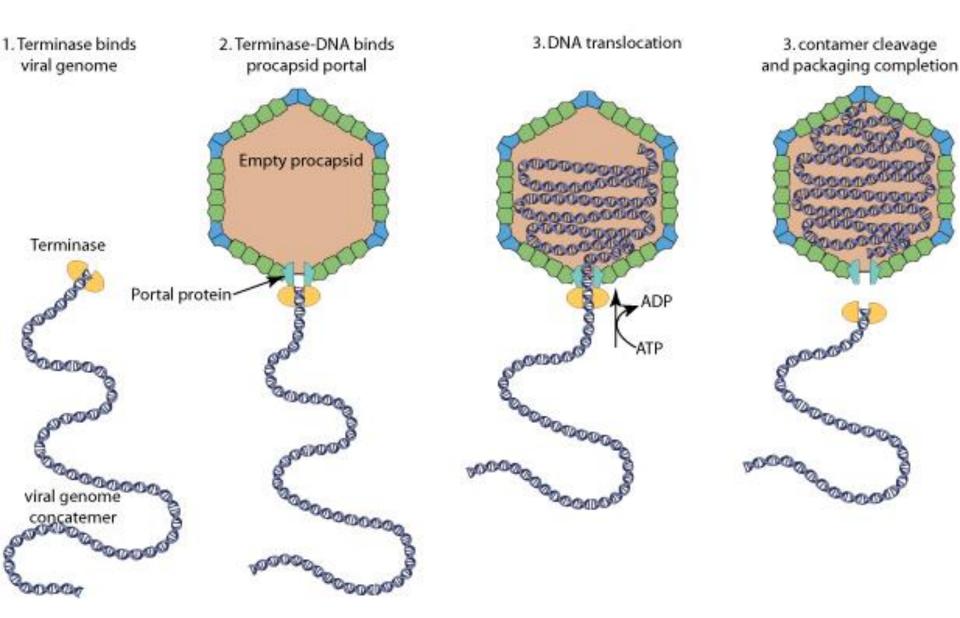


Picornavirus assembly

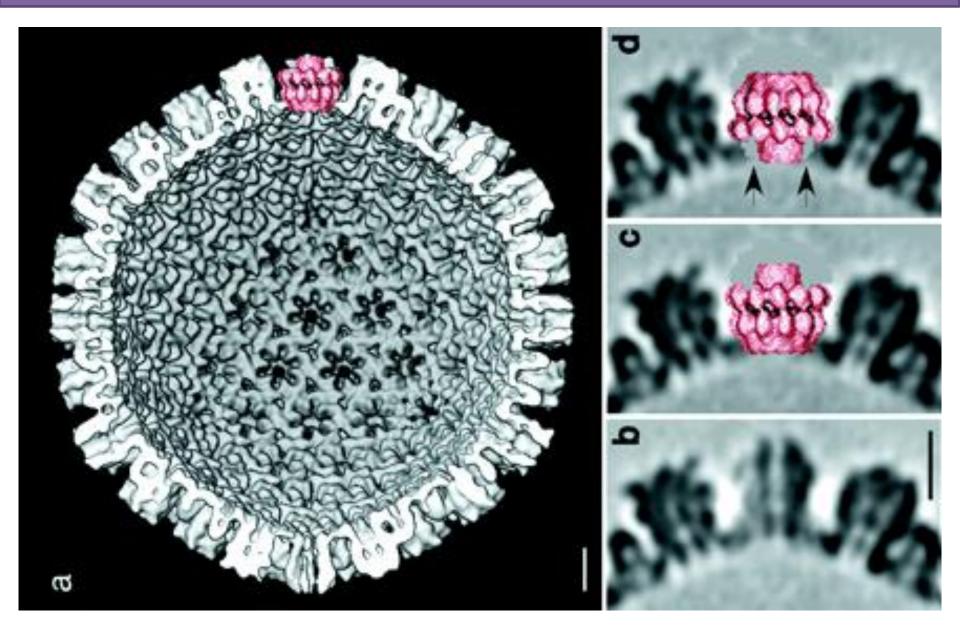




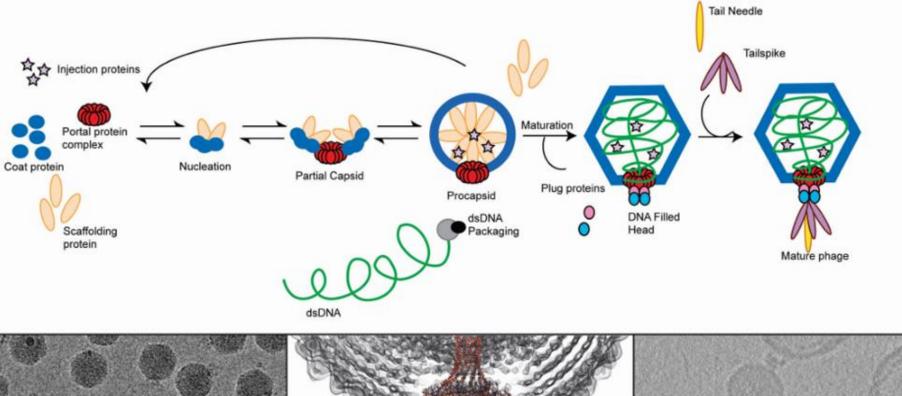
dsDNA virus genome packaging

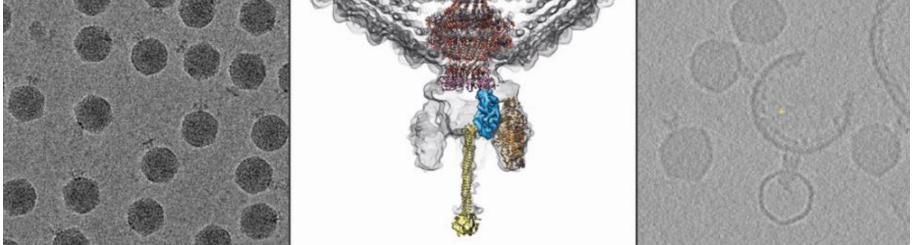


Herpesvirus portal structure

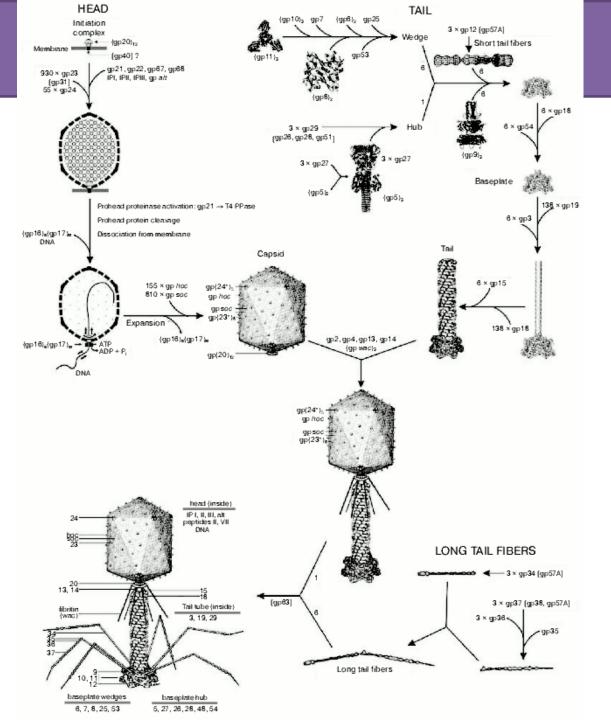


Podoviridae phage assembly

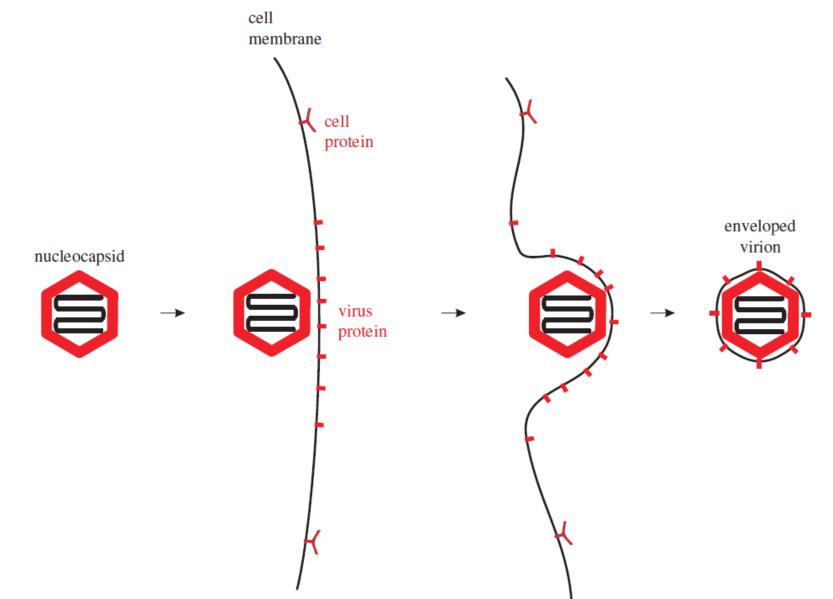




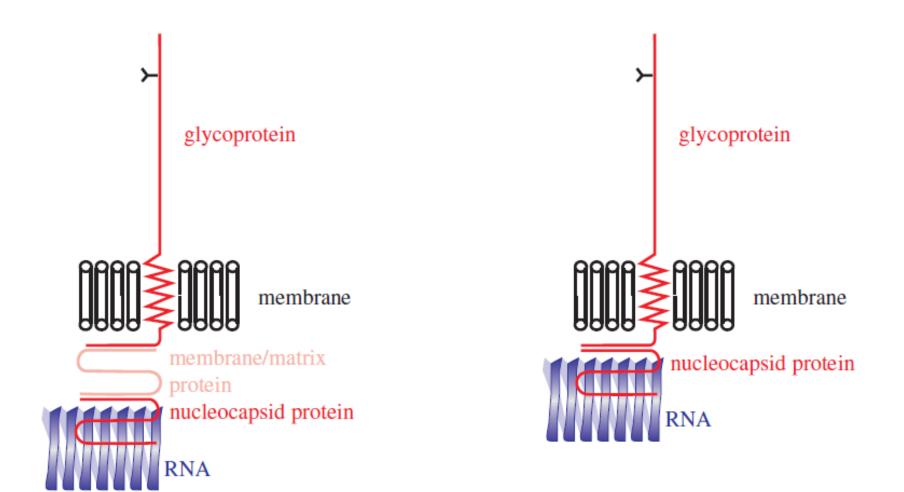




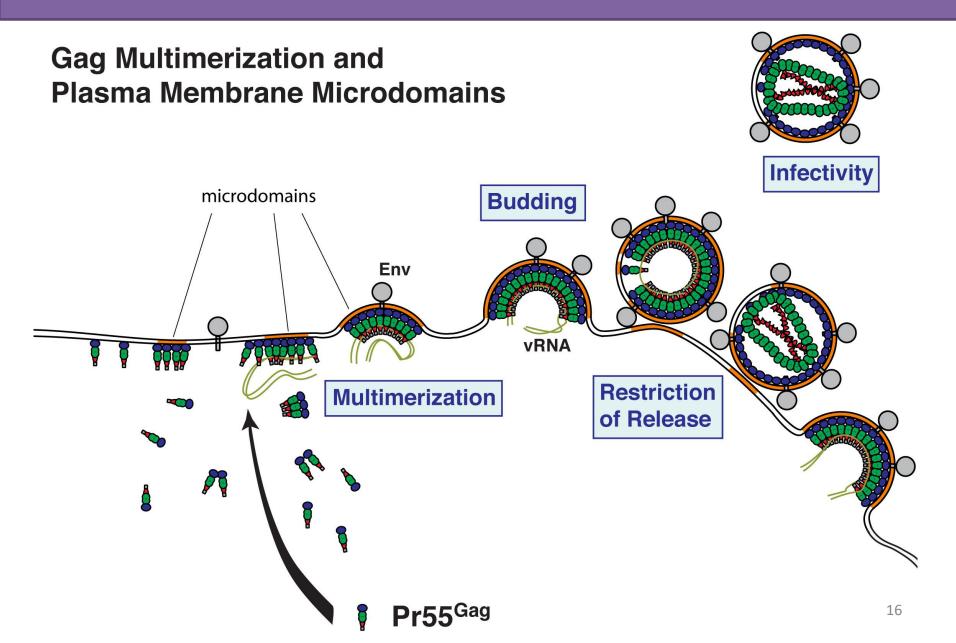
Acquisition of a virion envelope by budding



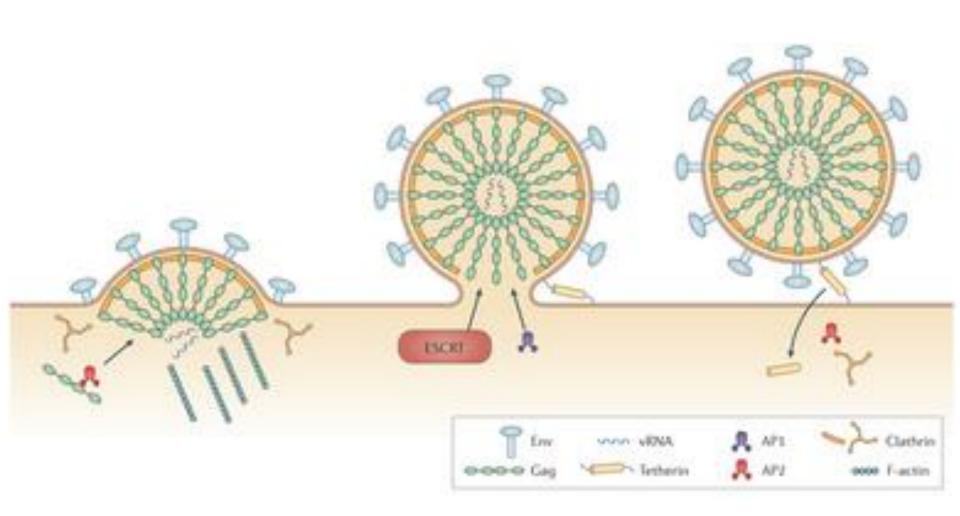
Complementation of enveloped viruses



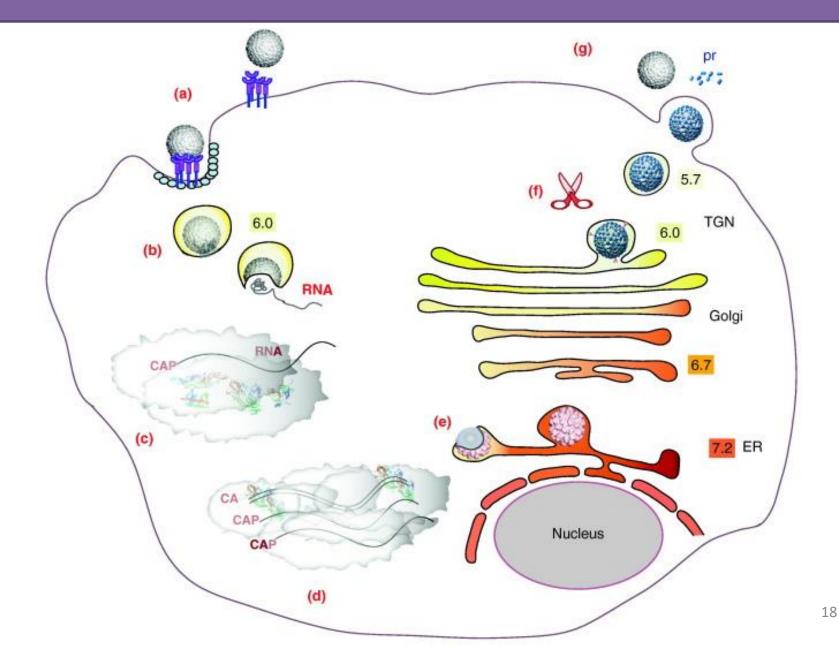
HIV assembly and maturation



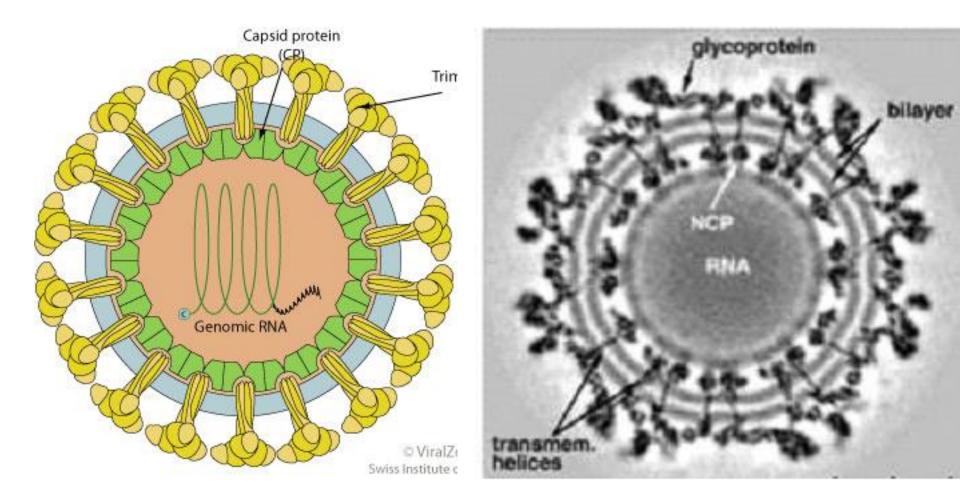
Retrovirus budding



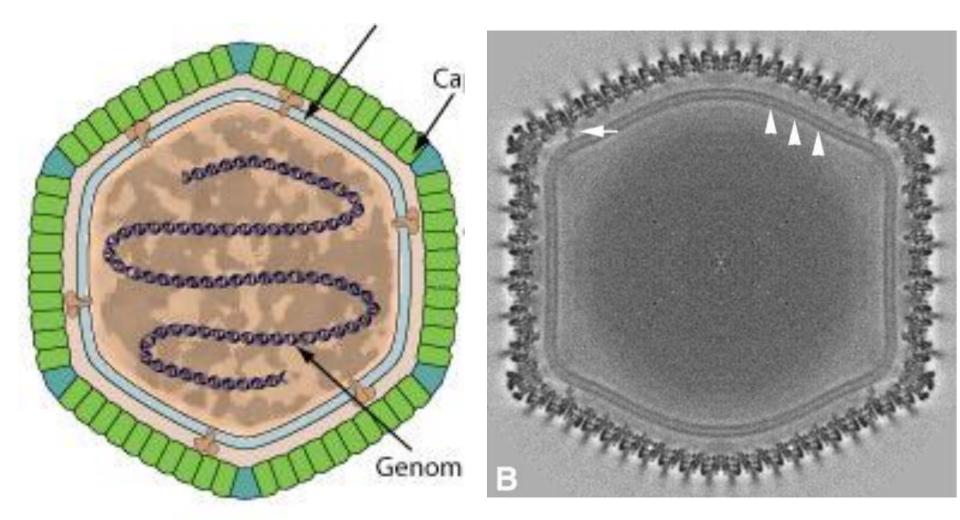
Assembly and maturation of dengue virus



Structure of alphavirus



Structure of iridovirus



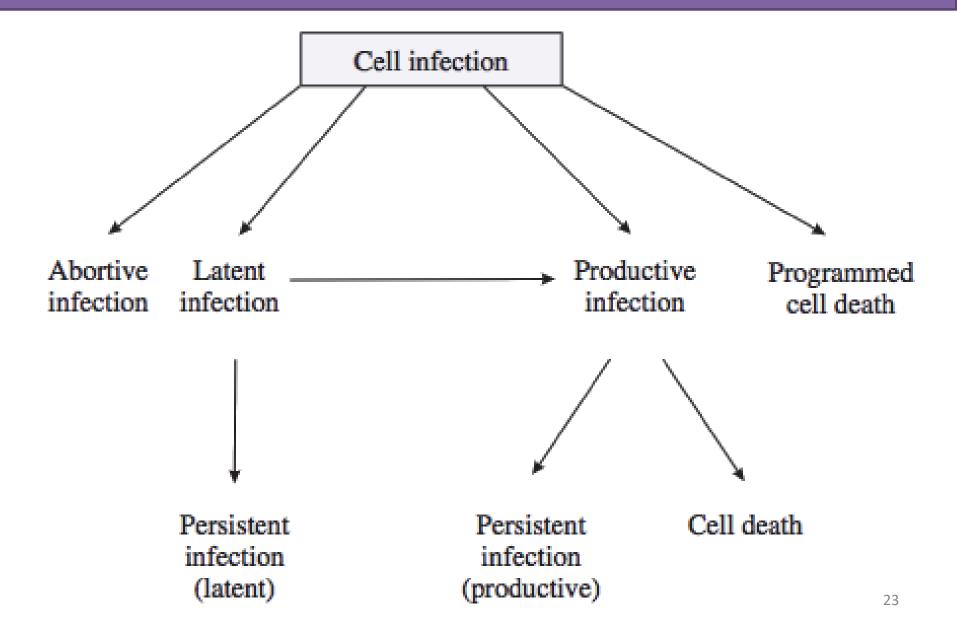
Summary of virus membrane acquisition

Site of virion assembly	Origin of virion membrane	Examples	
		Virion envelope	Internal virion membrane
Eukaryotic cell cytoplasm	Plasma membrane	Most retroviruses Most rhabdoviruses	
	Post-RER membrane	Hepadnaviruses	
	De novo synthesis	Poxviruses	Iridoviruses
Eukaryotic cell nucleus	Inner nuclear membrane	Nucleorhabdoviruses	
	De novo synthesis	Baculovirus virions that will be occluded	
Prokaryotic cell	Cell membrane	Acholeplasma laidlawii virus L2	
	De novo synthesis	Pseudomonas phage $\varphi 6$	Alteromonas phage PM2

Learning outcomes

- describe the assembly mechanisms for nucleocapsids with (a) helical symmetry and (b) icosahedral symmetry
- discuss the origins of internal virion membranes and of virion envelopes
- explain the roles played by membrane/matrix proteins in the budding of some enveloped viruses
- describe mechanisms used by viruses to exit from cells

Outcomes of virus infection for the host



Outcomes of virus infection for the host

Reasons for non-productive infection:

- latent infection
- abortive infection

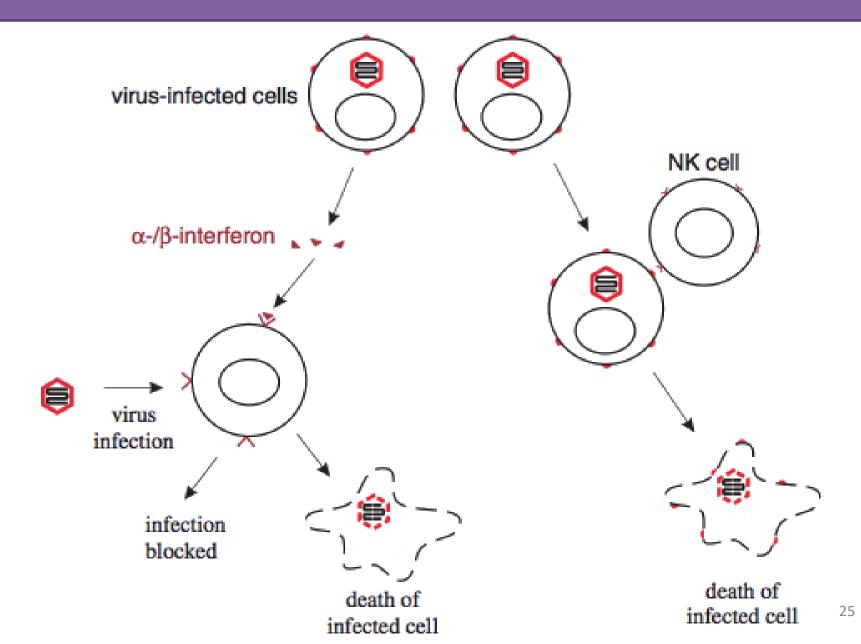
Persistent infections:

- productive (HIV)
- latent (herpesviruses)

Factors affecting outcomes of infection:

- host immune system
- "quality" of the virus (mutations, suitable host)

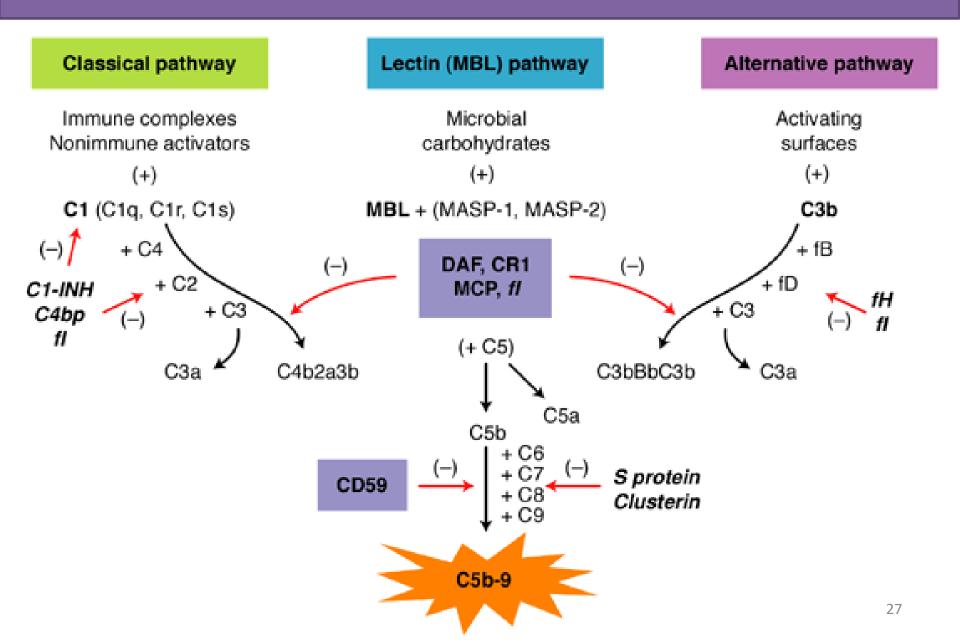
Innate immunity



Components of innate immunity

- Complement
- Interferons
- Natural Killer (NK) cells
- APOBEC3 protein complex
- tetherin

Activation and regulation of complement system



Complement effector system

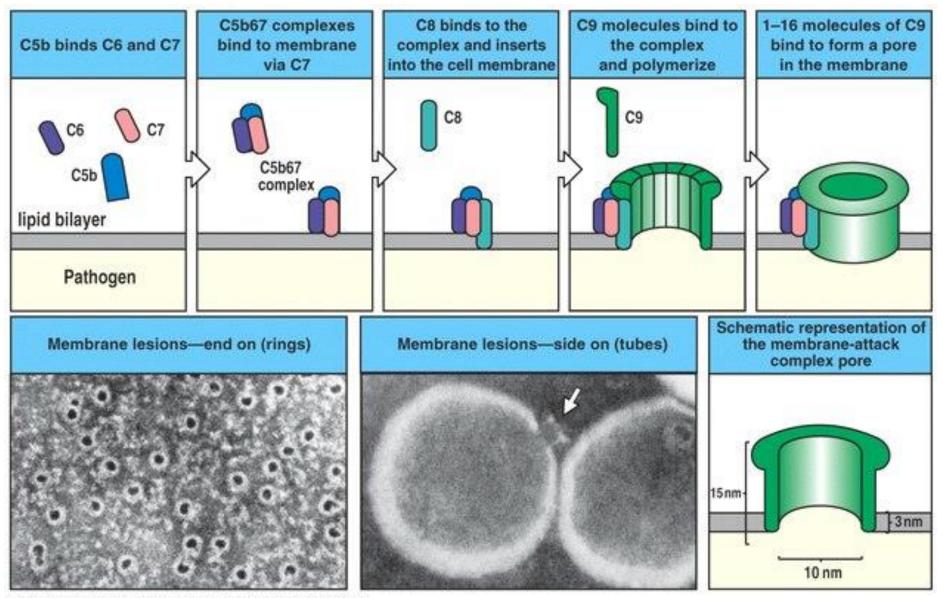
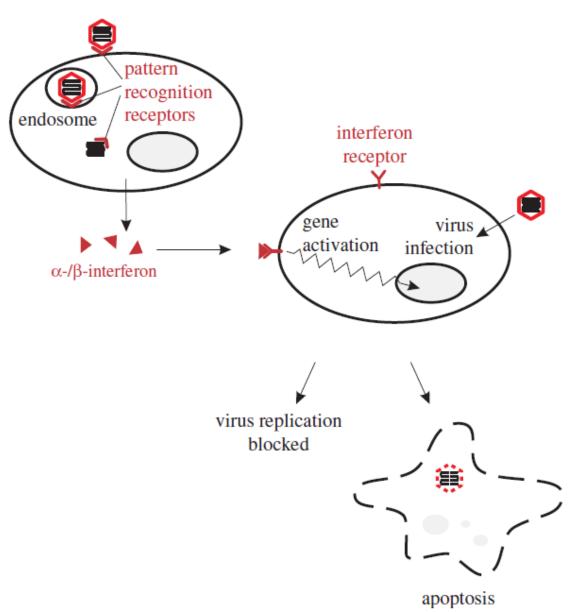


Figure 2-35 Immunobiology, 6/e. (© Garland Science 2005)

Interferon action



Interferon effects: Alpha and beta

- activation of expression of antiviral proteins: dsRNA dependent protein kinase R; tetherin

production of MHC I and proteasome components (presentation of peptides for control by T cells)

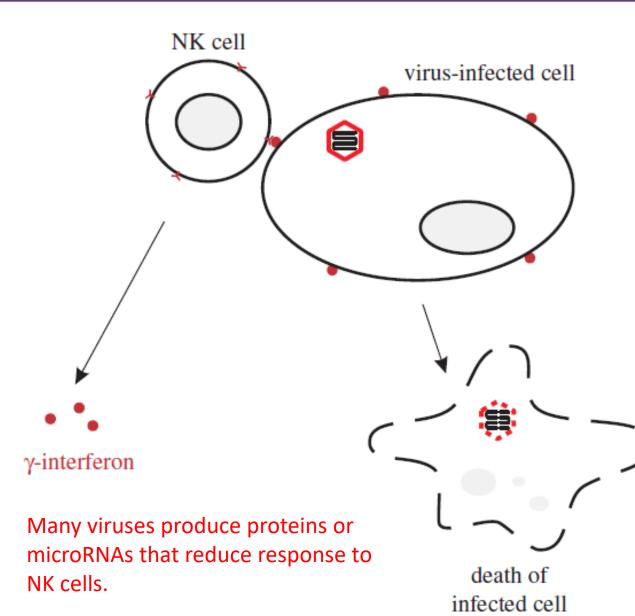
- activation of NK cells
- induction of apoptosis

Gamma

- produced by cells of immune system
- activation of phagocytes and NK cells

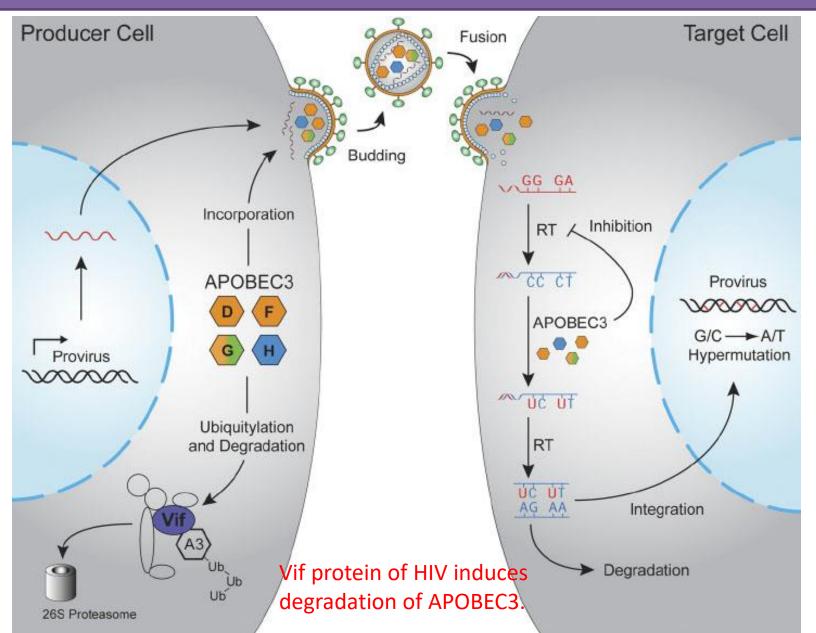
Many viruses inhibit production of interferons in infected cells. ²⁹

Activities of Natural Killer (NK) cells



Recognize infected cells Use perforins to kill cells Induce apoptosis

APOBEC3 proteins



Tetherin

In the absence of Vpu, CD317 expression causes tethering of buds to the cell surface and decreased infectivity.

8888c

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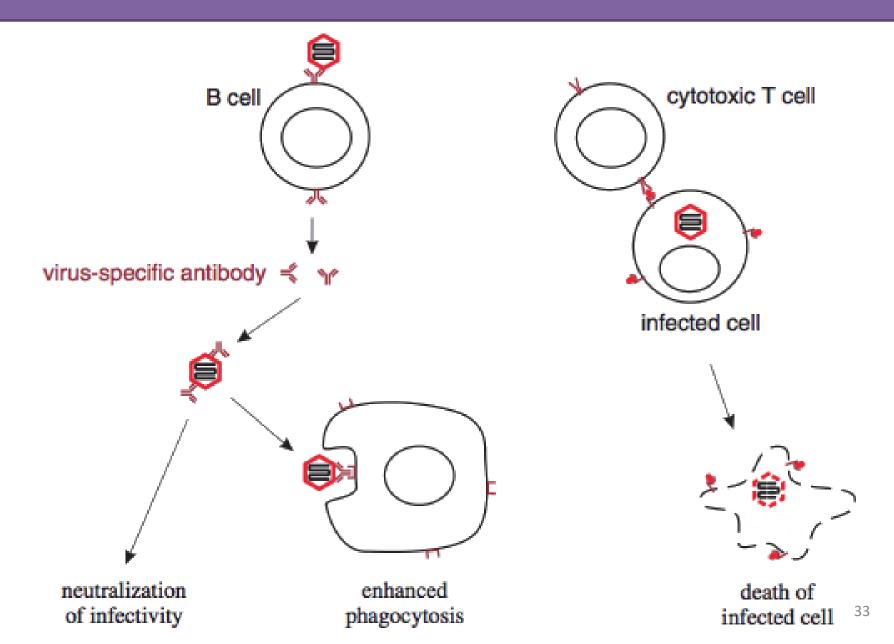
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CD4 also interferes with viral budding through interactions with the viral envelope.

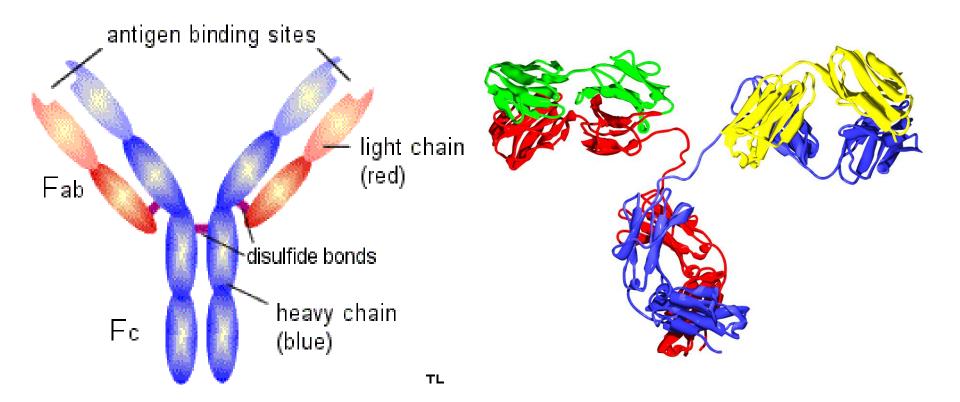


Vpu targets and degrades CD317 and CD4 to allow efficient budding.

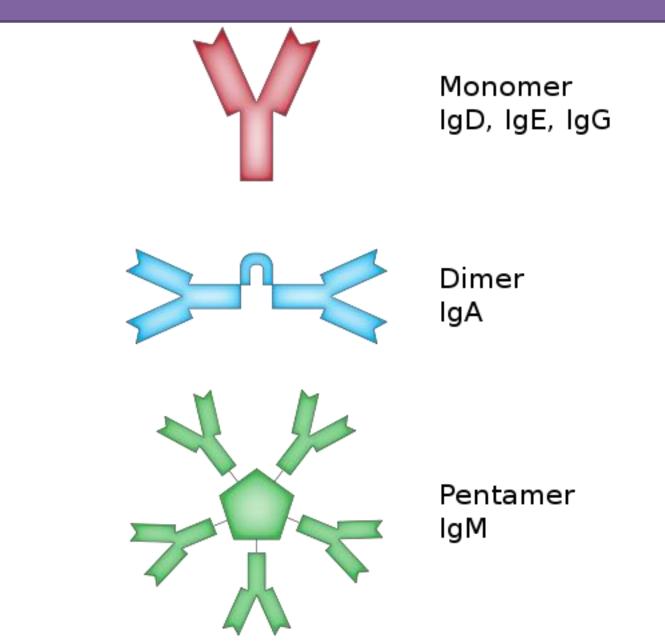
Adaptive immunity



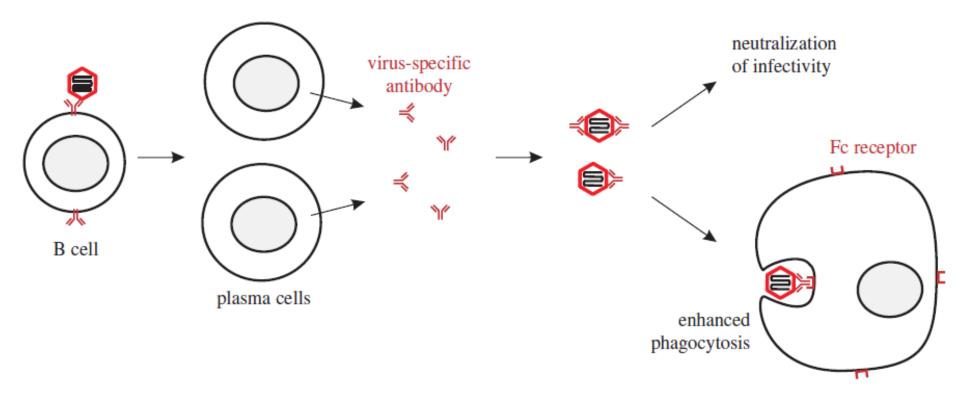
Antibody structure



Antibody structures



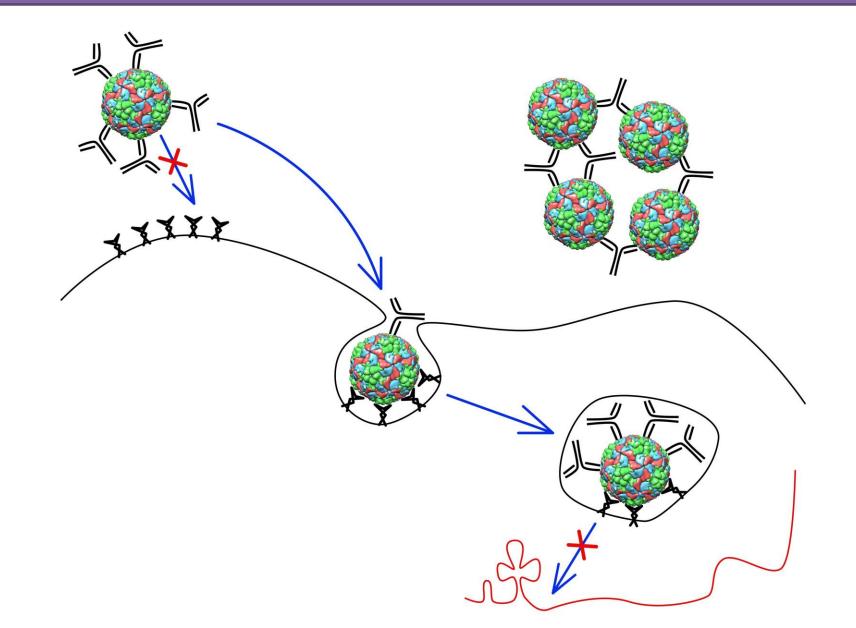
Production of virus-specific antibodies

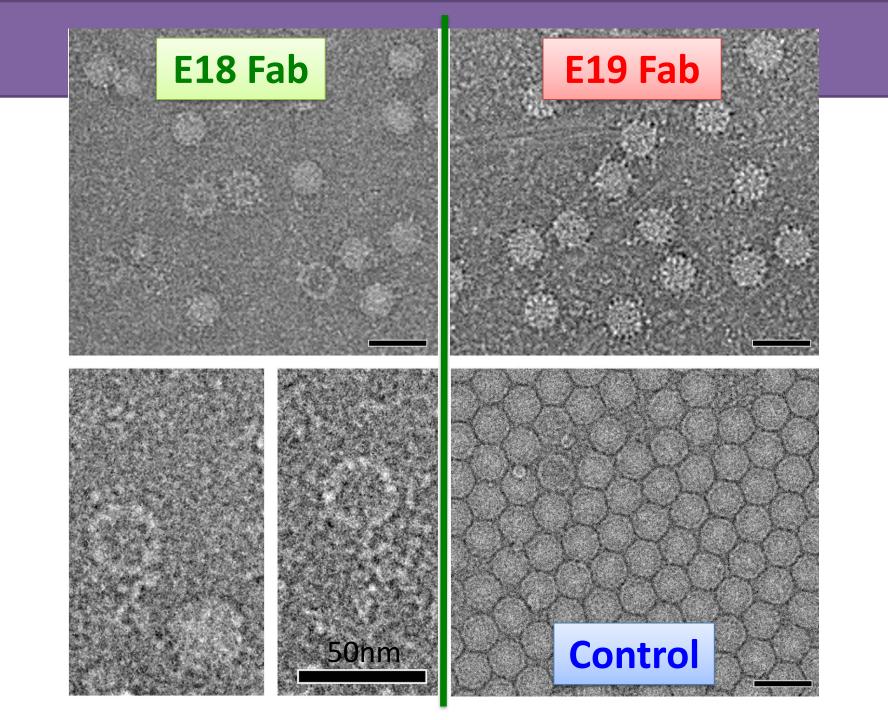


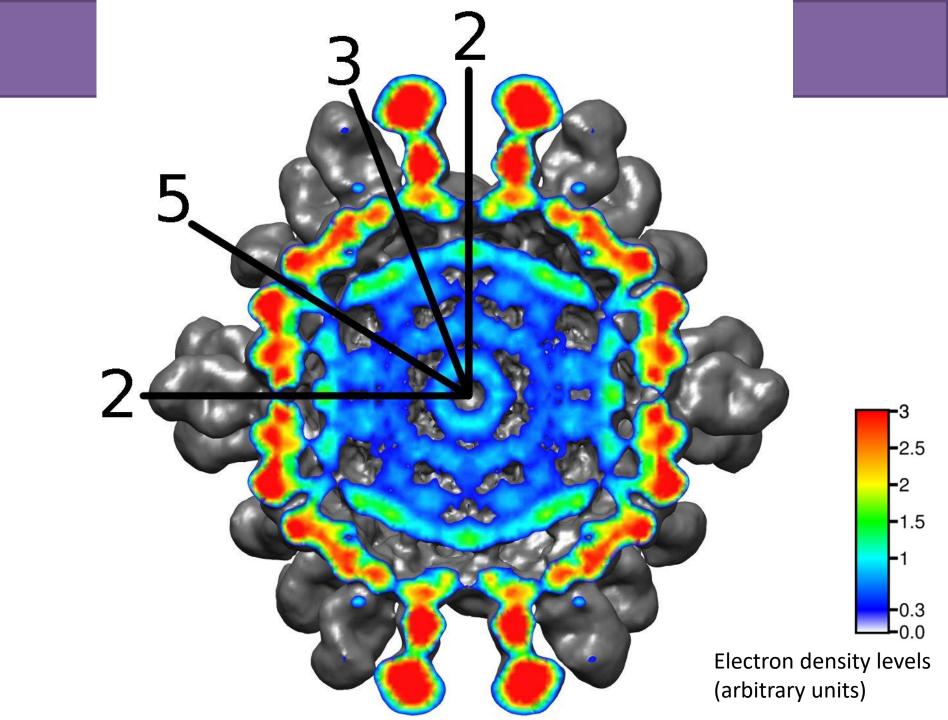
Antiviral effects of antibodies

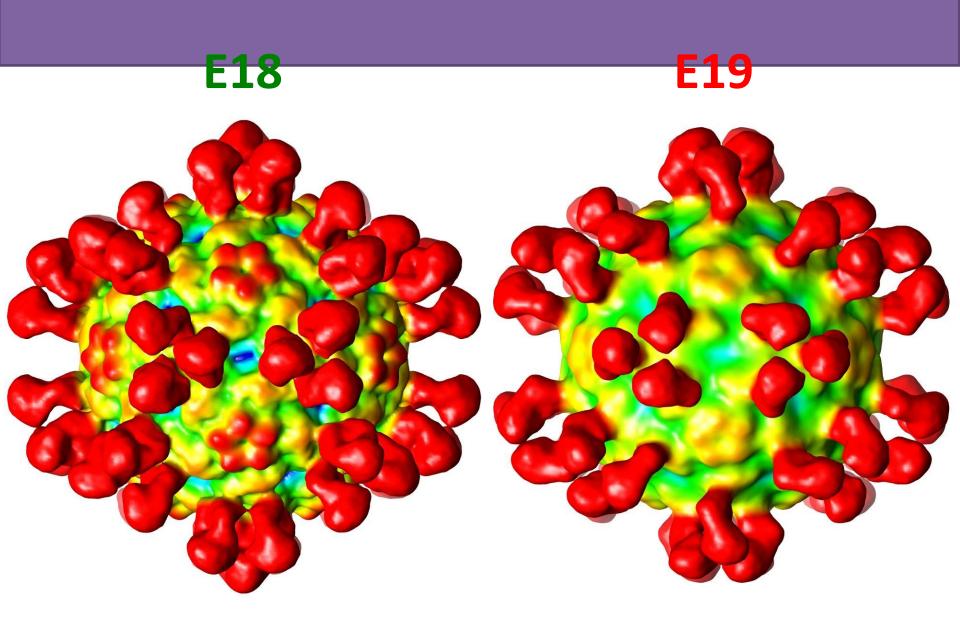
- Neutrophils and macrophages have IgG Fc receptors -> phagocytosis of infected cells and induction of apoptosis
- NK cells have IgG Fc receptors -> killing of infected cells.
- Antibodies can induce genome release from virions (poliovirus, EV71).
- Prevention of receptor binding
- Release of virions attached to cells
- Inhibition of cell entry (fusion proteins)
- Inhibition of genome uncoating
- Activation of complement

Neutralization of viruses by antibodies



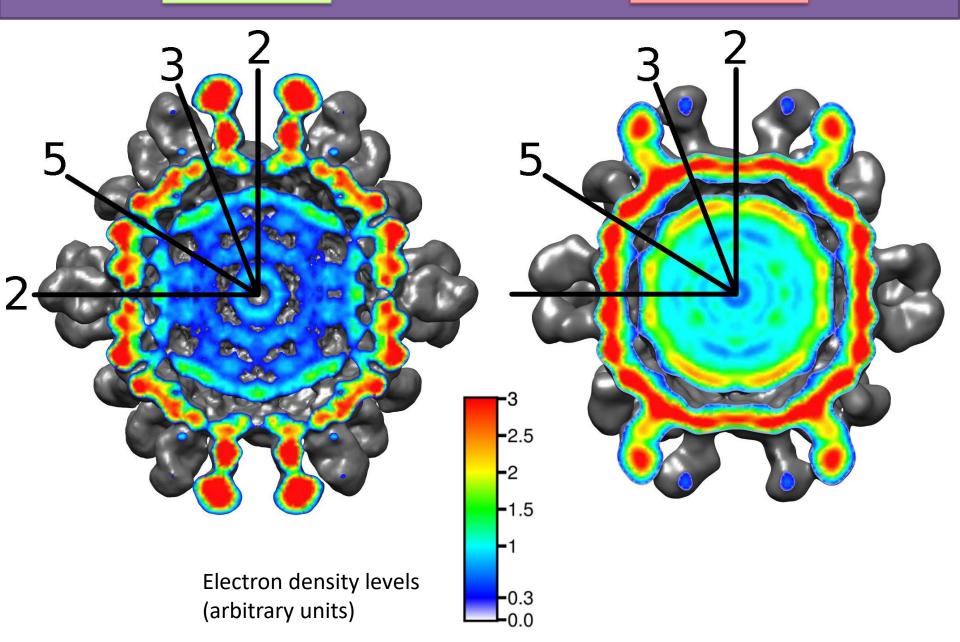




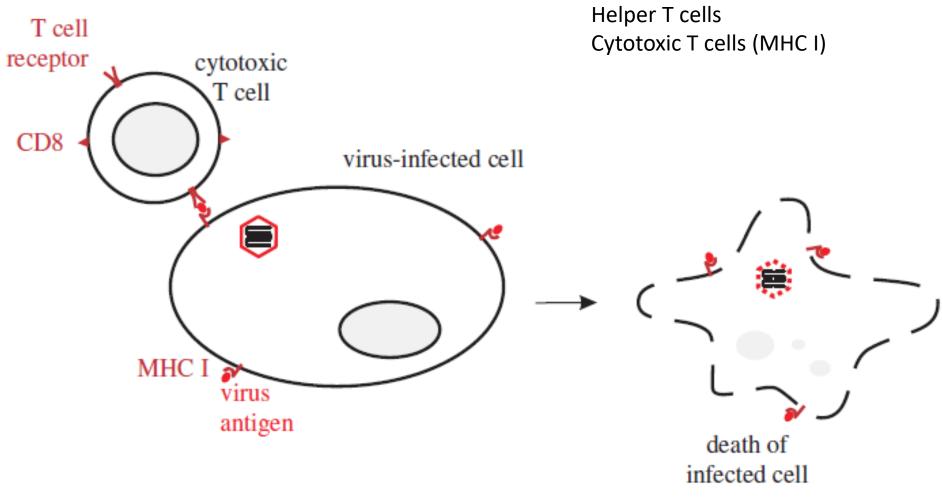


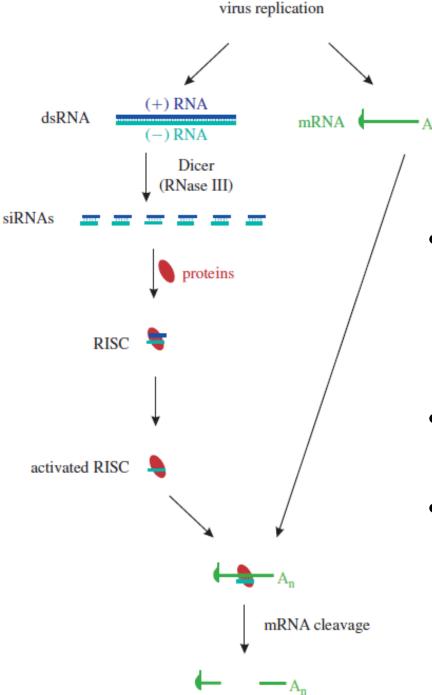






Recognition and killing of infected cell by cytotoxic T-cell



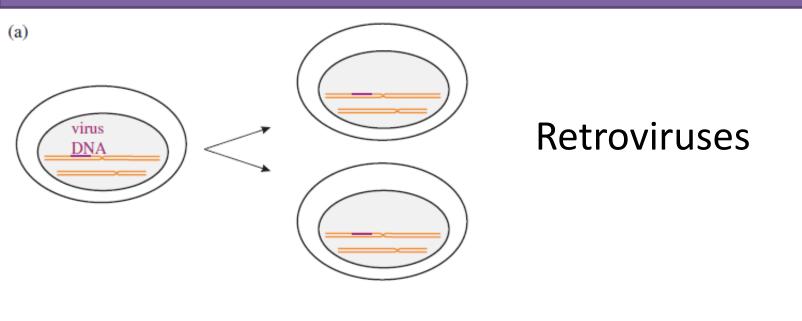


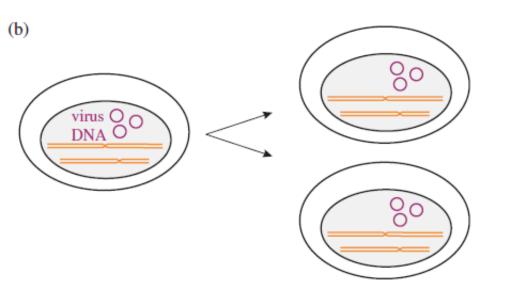
RNA silencing

- Fragmenting of dsRNA by DICER (production of 21-25bp fragments with 2-3nt 3' overhangs)
- Activation of RISC (RNA induced silencing complex)
- Destruction of mRNAs

Many plant viruses produce proteins that inhibit RNA silencing.

Maintenance of genomes during latent infection





Polyoviruses, Papillomaviruses

Signals that induce activation of latent infections

- A eukaryotic host cell moves into another phase of the cell cycle
- The host cell is irradiated with ultra-violet light. (phages, herpes simplex virus)
- A host organism becomes immunocompromised. (herpes simplex virus)
- The host cell becomes infected with a second virus that provides a function that the first virus lacks. (satellite virus and helper virus)

Productive infections in plants

Mode of transport through plasmodesmata	Virus examples
Virus RNA-MP	Tobacco mosaic
complexes transported	virus
	Cowpea chlorotic mottle virus
Virus RNA-coat	Cucumber mosaic
protein-MP	virus
complexes transported	
Virions transported	Cowpea mosaic
through tubules	virus
composed of MP	

Disease

Virulence of a virus strain Dose of virus

Effectiveness of immune system (age, nutritional status, previous training) Human interventions

Virus elimination x latent infections

Learning outcomes

- describe the major components of innate and adaptive immunity in vertebrates
- outline the process of RNA silencing
- explain programmed cell death
- explain the terms
 - productive infection
 - non-productive infection
 - latent infection
 - abortive infection
 - defective virus
- discuss the spread of virus infections within animal bodies and within plants
- discuss the factors that determine whether virus infection results in disease

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Virus classification

Authority: International Committee on Taxonomy of Viruses

http://www.ncbi.nlm.nih.gov/ICTVdb/index.htm

Taxonomic groups: Order

Family

Subfamily

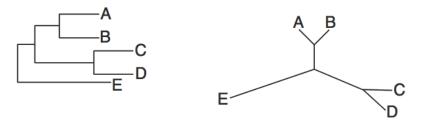
Genus

Species

Phylogenetic trees

- based on genome sequences

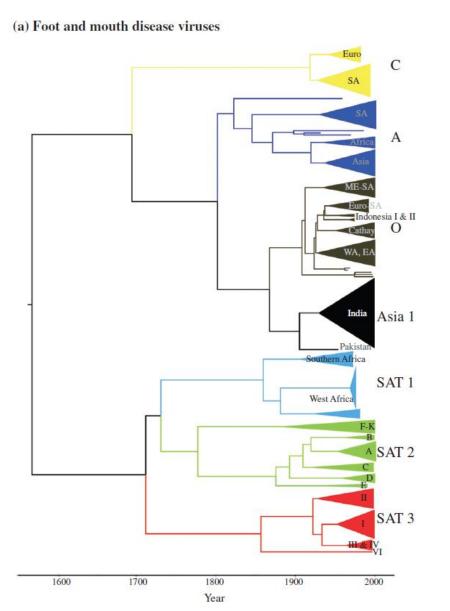
- indicate relationships between viruses:



Baltimore Classification: - seven classes of viruses

- based on genome type and transcription.

Phyogenetics trees



(b) Family Reoviridae

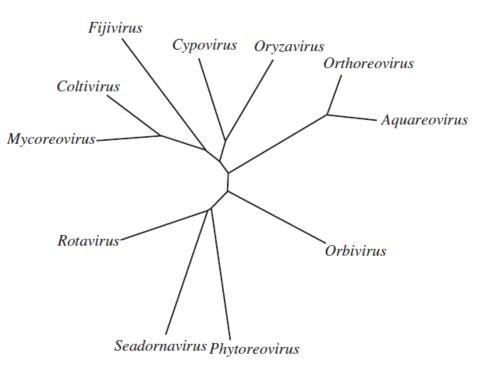
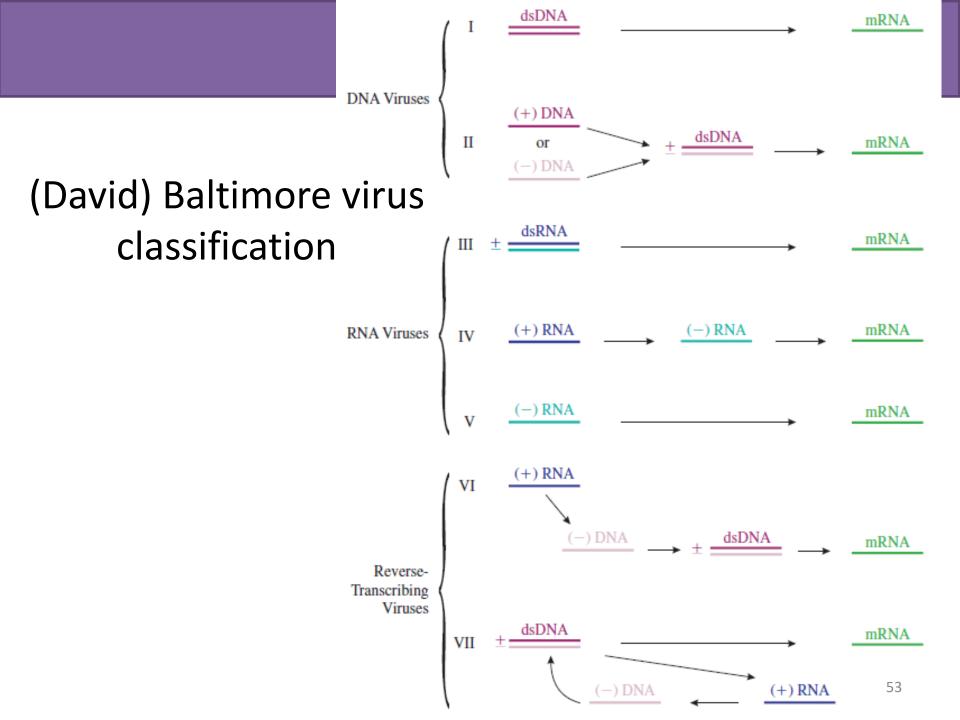


Figure 10.1 Phylogenetic trees. (a) Rooted tree showing relationships between foot and mouth disease virus serotypes based on *VP1* sequences. The serotypes evolved in different regions of the world. (b) Unrooted tree showing relationships between genera in the family *Reoviridae* based on *VP1* sequences.

Source: (a) Tully and Fares (2008) *Virology*, 382, 250. Reproduced by permission of Elsevier.

Nomenclature

Taxonomic group	Suffix	Example 1	Example 2	Example 3
Order	-virales	Caudovirales	Mononegavirales	Nidovirales
Family	-viridae	Myoviridae	Paramyxoviridae	Coronaviridae
Subfamily	-virinae	-	Paramyxovirinae	-
Genus	-virus	T4-like viruses	Morbillivirus	Coronavirus
Species	-	Enterobacteria phage T4	Measles virus	Severe acute respiratory
				syndrome virus



Learning outcomes

- evaluate the traditional criteria used to classify viruses into families and genera
- write family and genus names in the correct format
- explain how genome sequence data are used to classify viruses
- evaluate phylogenetic trees
- explain the basis of the Baltimore classification of viruses

Herpesviruses

Figures Chapter 11

2nd Edition

Principles and Applications

Virology

John B. Carter and Venetia A. Saunders

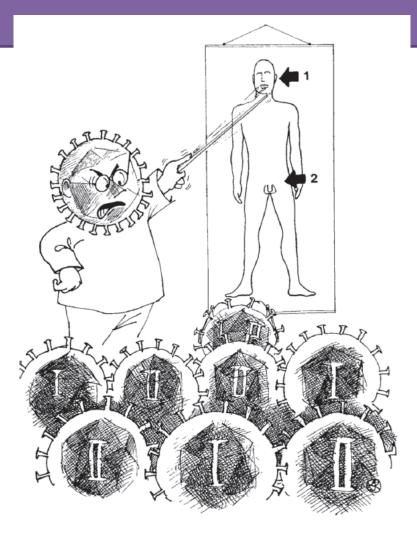
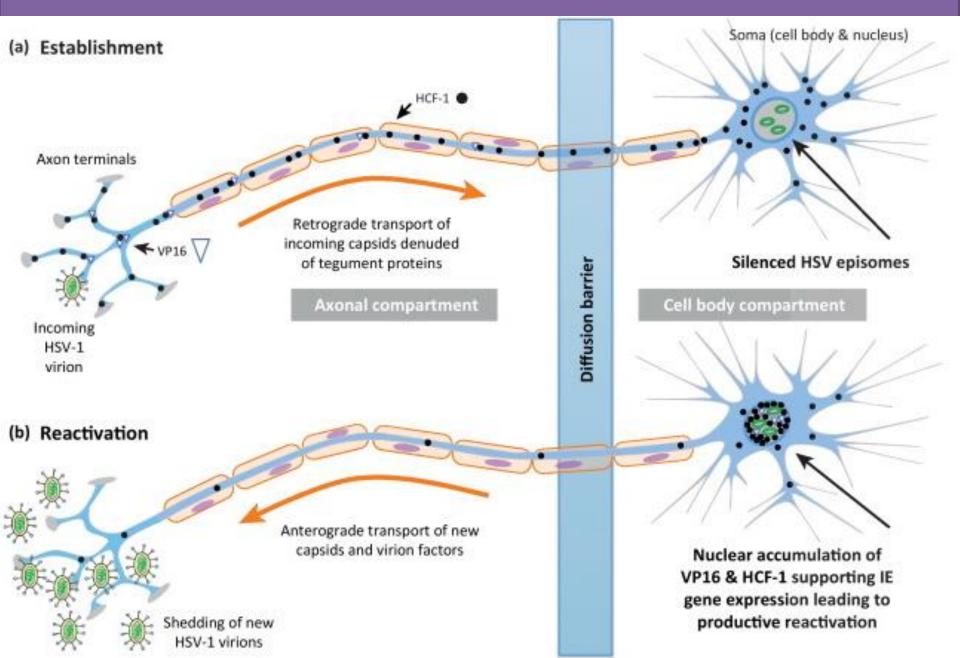
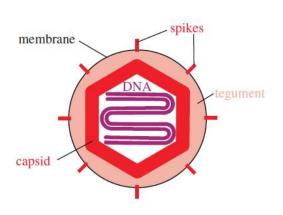


Figure 11.1 "Get your priorities right."

Source: Haaheim, Pattison, and Whiteley (2002) *A Practical Guide to Clinical Virology*, 2nd edition. Reproduced by permission of John Wiley & Sons.

Latent infection





(a) Virion components

Electron cryo-tomographic visualizations

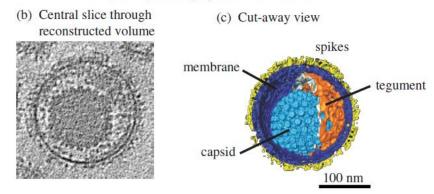


Figure 11.2 The herpesvirus virion.

Source: (b) and (c) are images of HSV-1, from Grünewald and Cyrklaff (2006) *Current Opinion in Microbiology*, 9, 437. Reproduced by permission of the authors and Elsevier Limited.

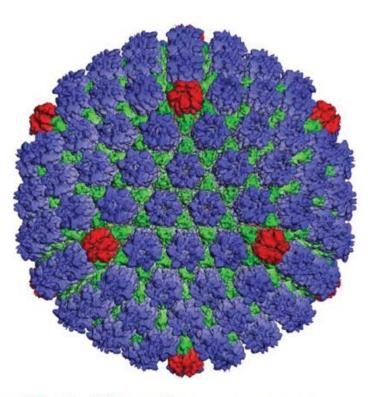


Figure 11.3 The HSV-1 capsid. Reconstructed image from cryoelectron microscopy.

Source: Courtesy of Professor Wah Chiu, Baylor College of Medicine, Houston, TX. Reinterpretation of data from Zhou *et al.* (2000) *Science*, 288, 877, with permission of the American Association for the Advancement of Science.

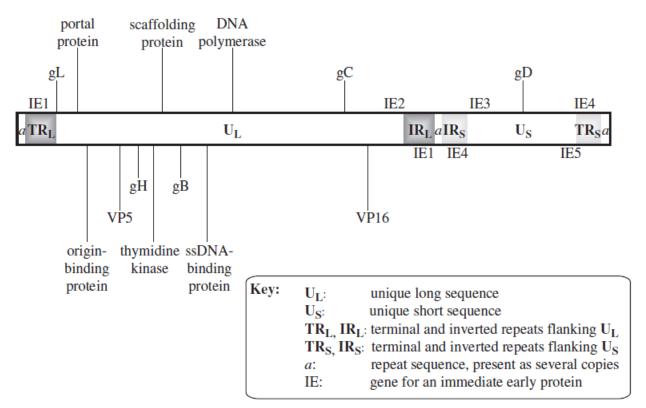


Figure 11.4 The HSV-1 genome.

The locations of some genes are indicated. Those shown above the genome are in ORFs read left to right; those shown below the genome are in ORFs read right to left.

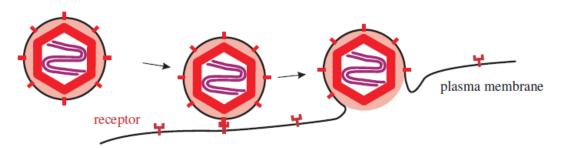


Figure 11.5 HSV-1 attachment and entry into the cell. Virus glycoproteins bind to receptors on the cell, then the virion envelope fuses with the plasma membrane.

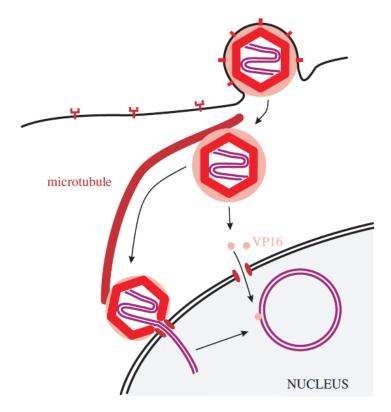


Figure 11.6 Transport of HSV-1 DNA and VP16 into the nucleus. The nucleocapsid is transported along a microtubule to a site close to the nucleus. Docking of the nucleocapsid at a nuclear pore is followed by release of the genome into the nucleus. Molecules of VP16 released from the tegument are also transported into the nucleus.

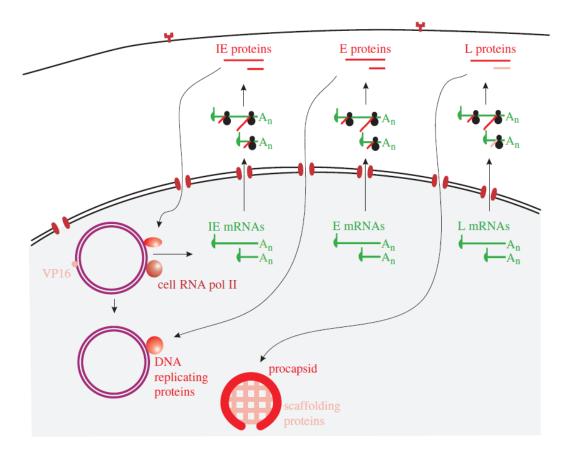
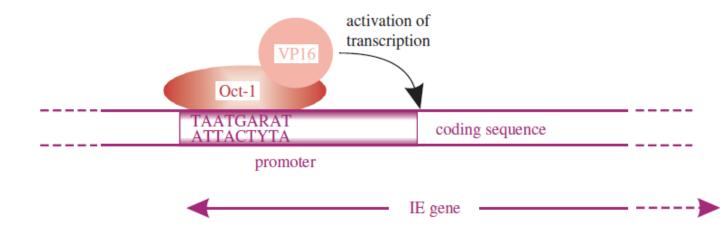


Figure 11.7 HSV-1 transcription and translation.

There are three phases of transcription and translation:

IE: immediate early E: early

L: late.



- Figure 11.8 Activation of transcription of HSV-1 IE genes by VP16.
- R: purine
- Y: pyrimidine

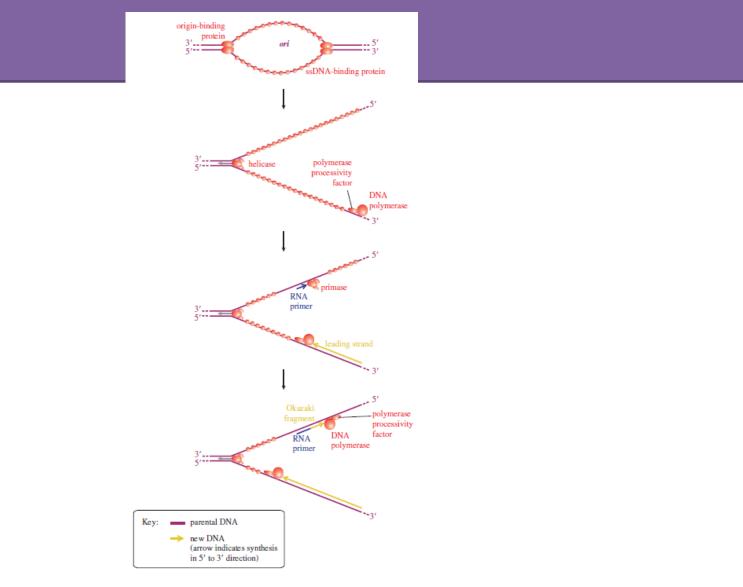


Figure 11.9 Roles of HSV-1 proteins in DNA replication. The roles of the seven virus proteins essential for DNA replication are shown. The helicase and the primase are complexes of the same three proteins. Please see text for details.

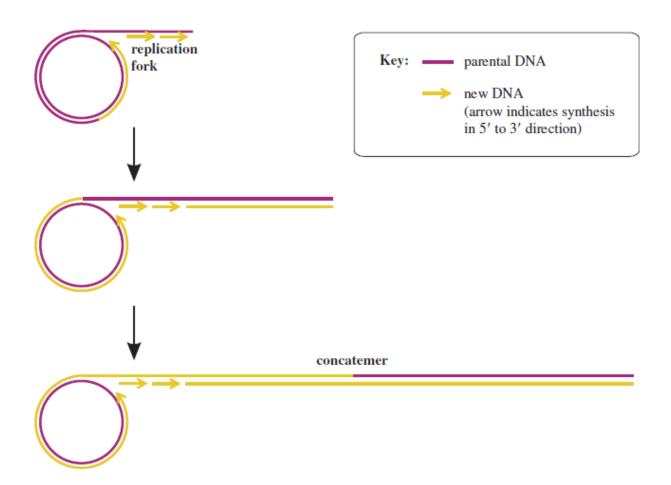


Figure 11.10 Formation of a concatemer. See Figure 7.5 for earlier stages of rolling circle replication of DNA. Color coding indicates the fates of the two parental strands of DNA.

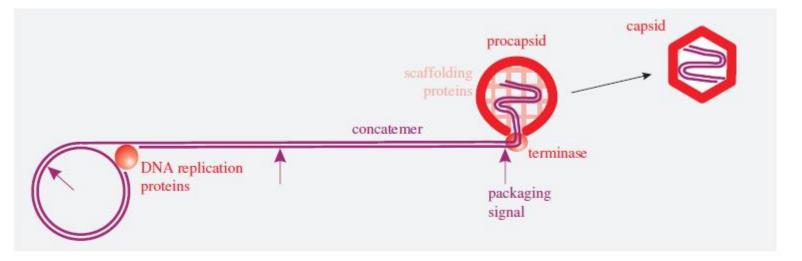
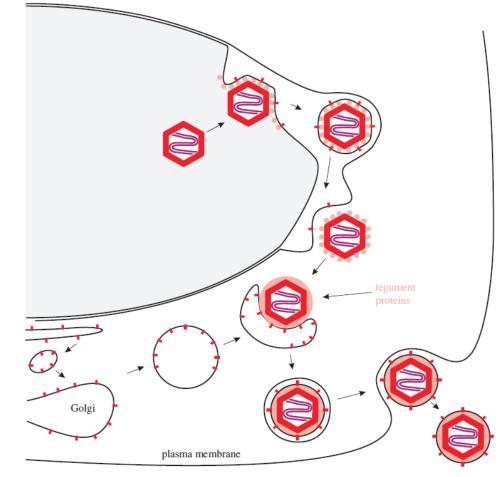
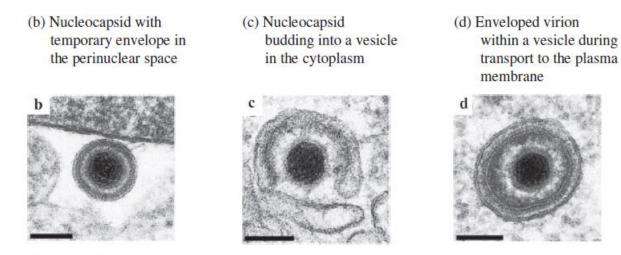


Figure 11.11 Packaging HSV-1 DNA into a procapsid. A genome-length of DNA enters a procapsid and is cleaved by a terminase at a packaging signal.



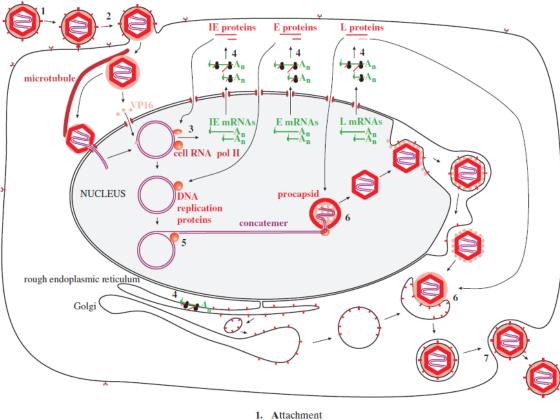
(a) Transport of nucleocapsids to the cytoplasm, envelopment, and exit from the cell

Figure 11.12 Final stages of HSV-1 assembly.





Source: (b)-(d) from Mettenleiter, Klupp, and Granzow (2009) Virus Research, 143, 222. Scale bar: 100 nm. Reproduced by permission of Elsevier and the authors.



- 2. Entry
- 3. Transcription
- 4. Translation
- 5. Genome replication
- 5. Genome replicatio
- 6. Assembly
- 7. Exit

