

# VIRAL LIFE CYCLE

- **ATTACHMENT**
- **ENTRY**
- **UNCOATING**
  
- **GENOME REPLICATION**
  
- **ASSEMBLY**
- **RELEASE**

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On the surface membrane of all living cells are protein structures called "receptors". Some viral proteins "antireceptors" usually expressed on the enveloped or on the capsid are able to interact with them.

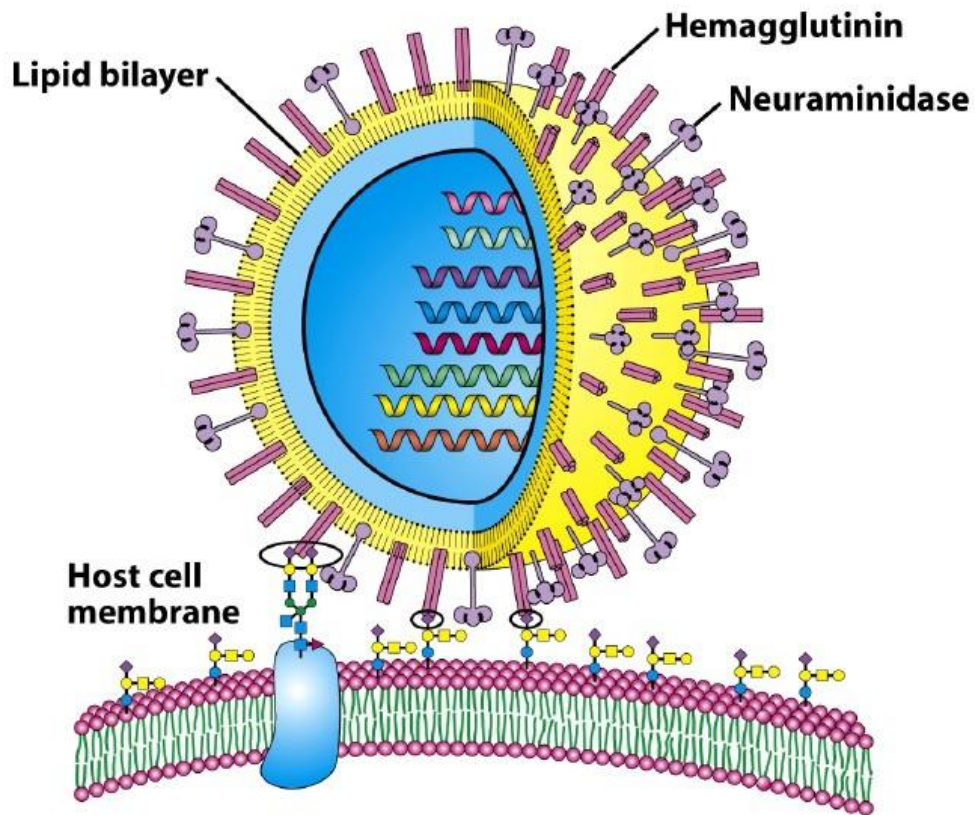
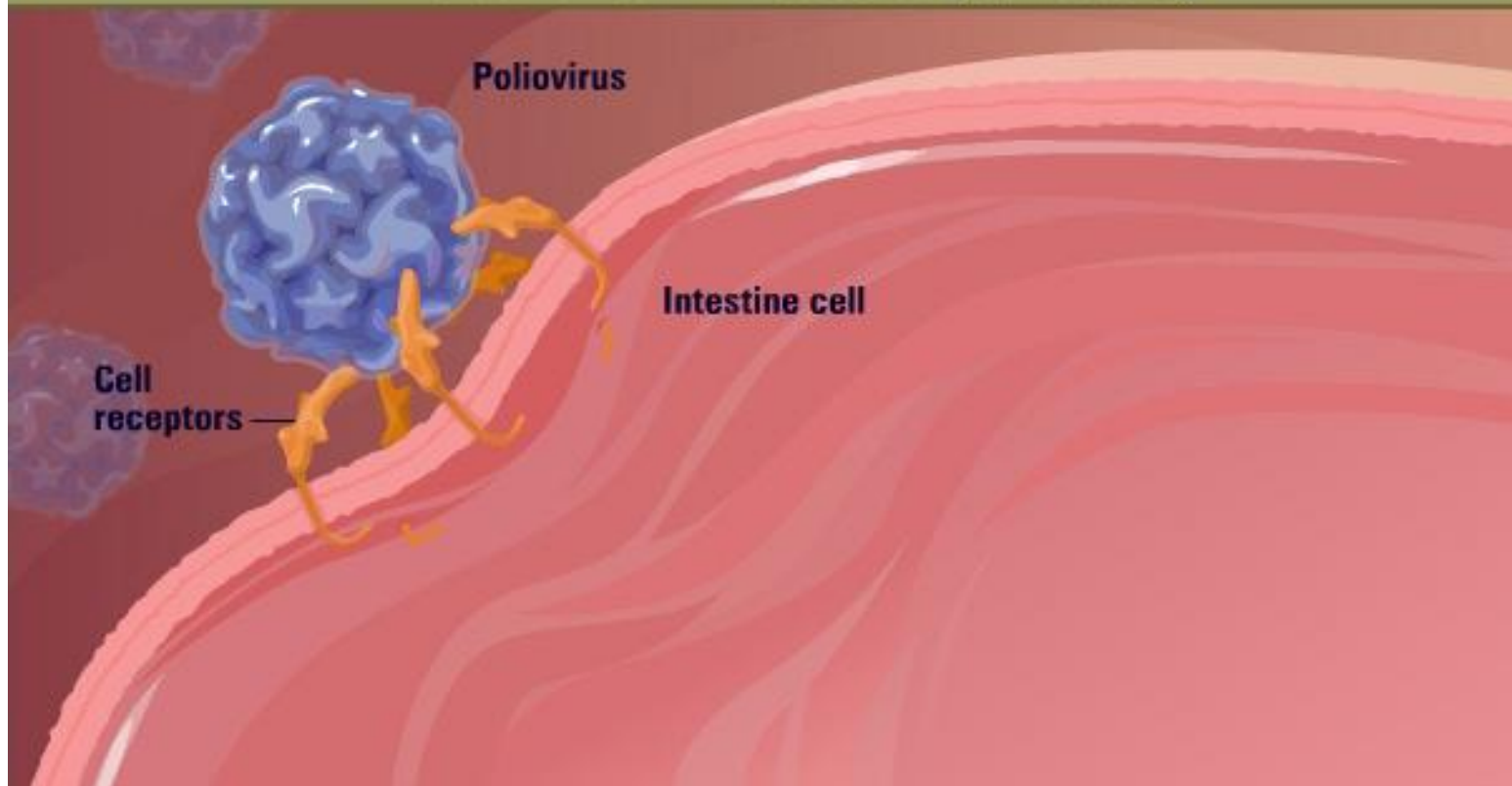
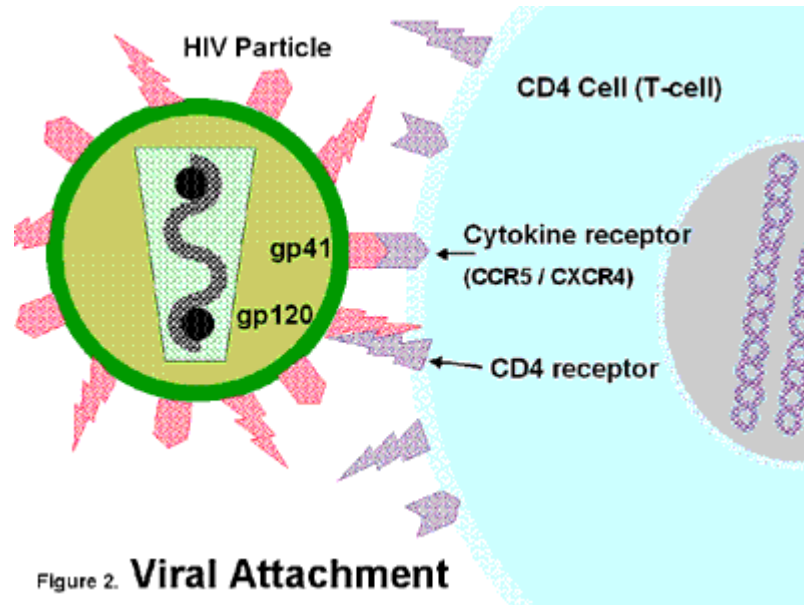


Figure 11-29  
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If the virus finds a cell with the correct receptor, infection begins.



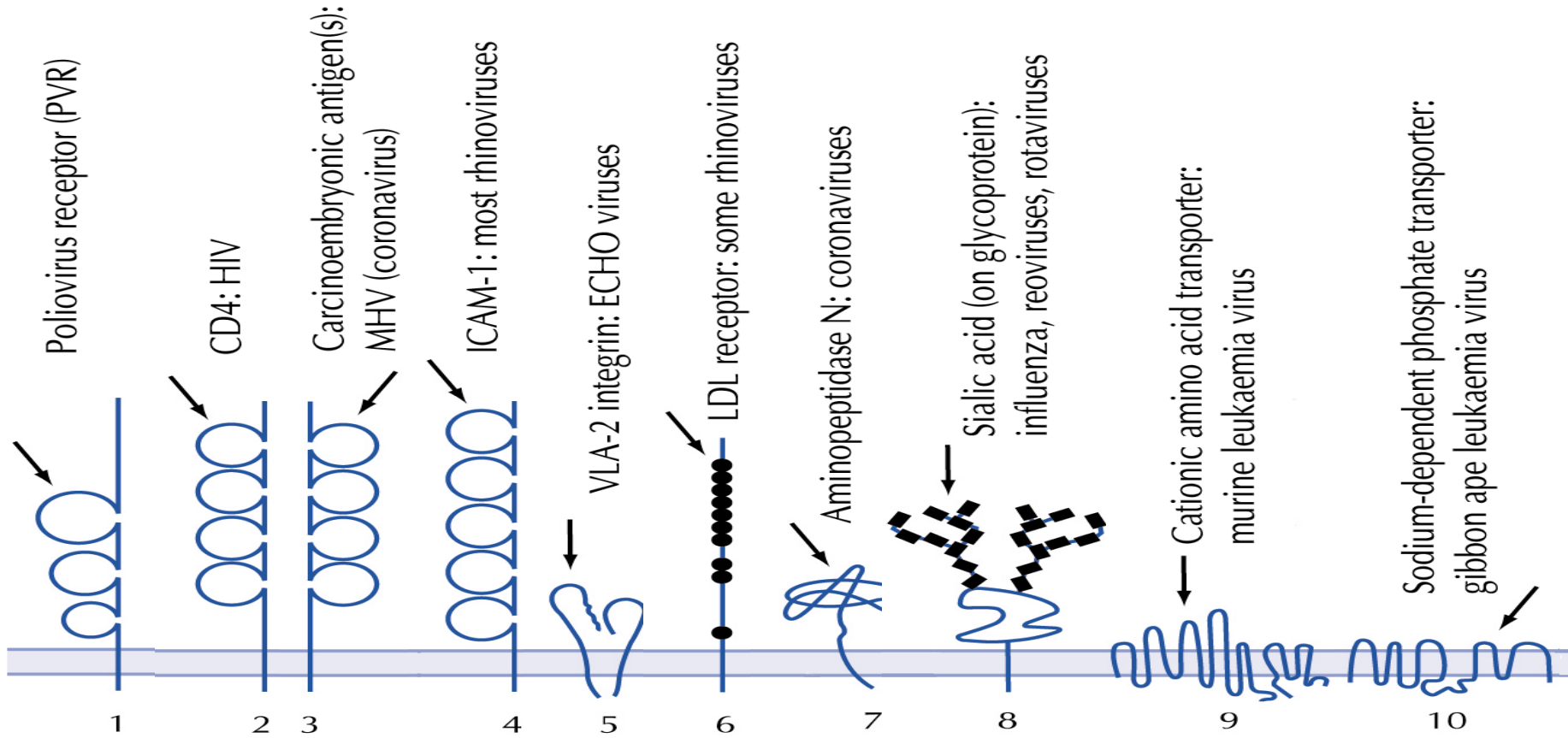


There are at least two receptors on T-lymphocytes to which the human immunodeficiency virus (HIV) sticks. The primary receptor, called "CD4" and a second receptor that loops through the cell membrane (chemokine receptor). Both are critical for infection to occur.

HIV infection of a lymphocyte requires attachment of the virus to the cell membrane through both of these "ligand-receptor" links. In cells whose the transmembrane receptor is different, the HIV "key" no longer matches the lymphocyte "lock" and attachment is incomplete.

Those cells may avoid infection by HIV.

# Virus Binding to the Cell Surface



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On the surface membrane of all living cells are protein structures called "receptors". Some viral proteins "antireceptors" usually expressed on the enveloped or on the capsid are able to interact with them.

**This event turns on the life replication cycle of the virus**

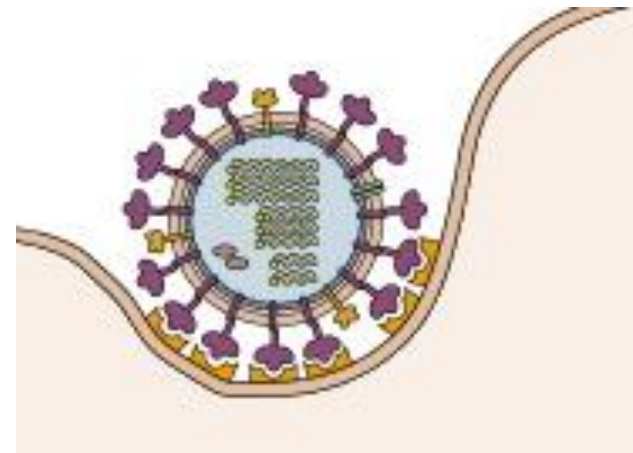
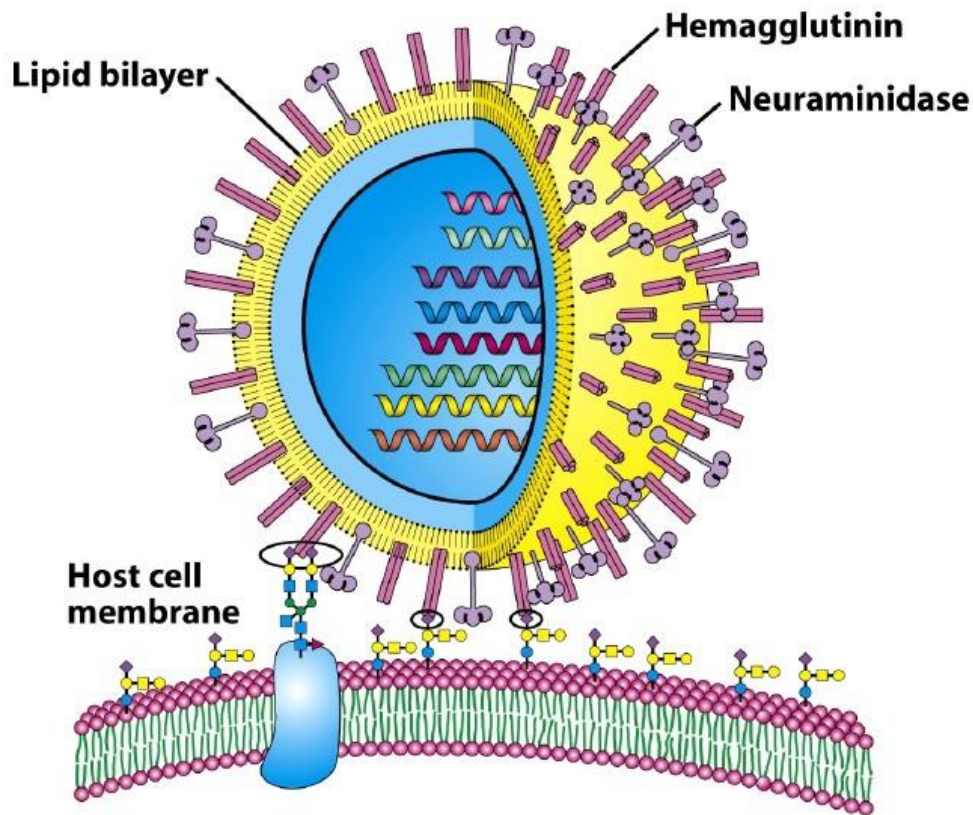
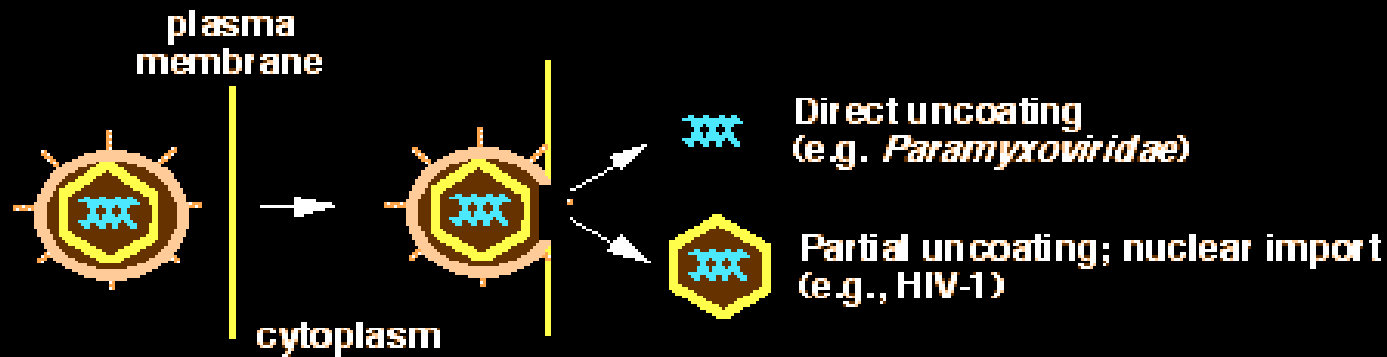


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# VIRUS ENTRY

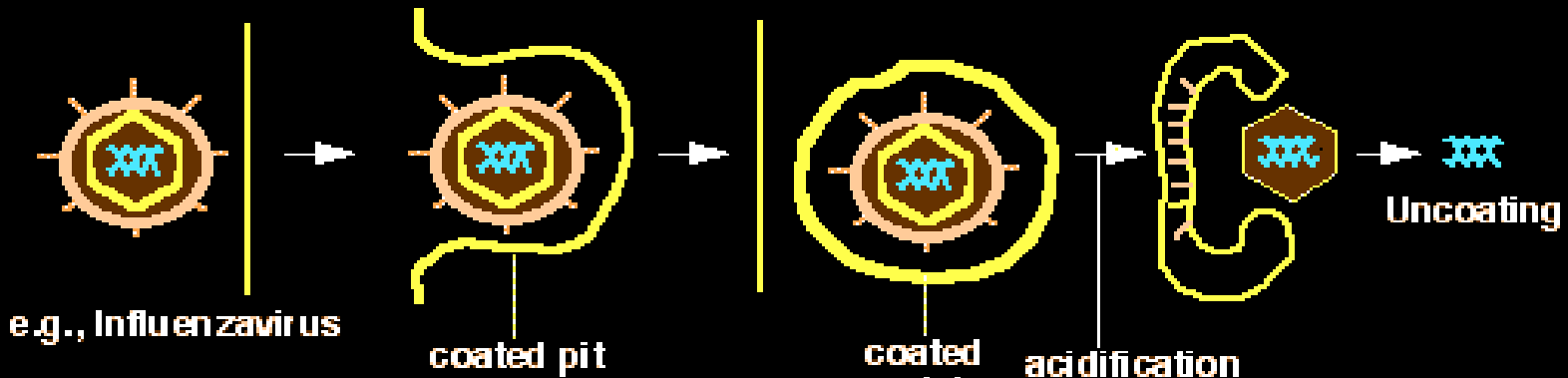
- Penetration by Membrane Fusion
- Endocytic Pathways
- Penetration of Nonenveloped Viruses
  - Membrane puncture
  - Perforation

## Surface Fusion

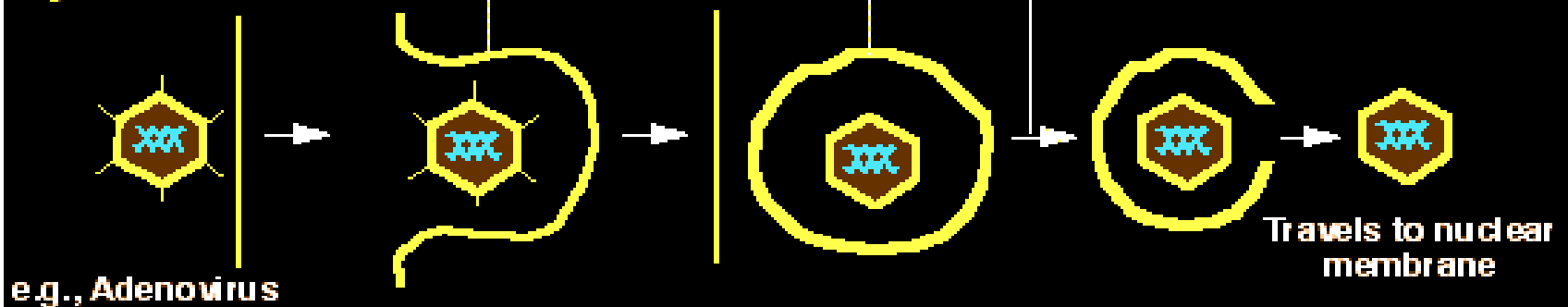


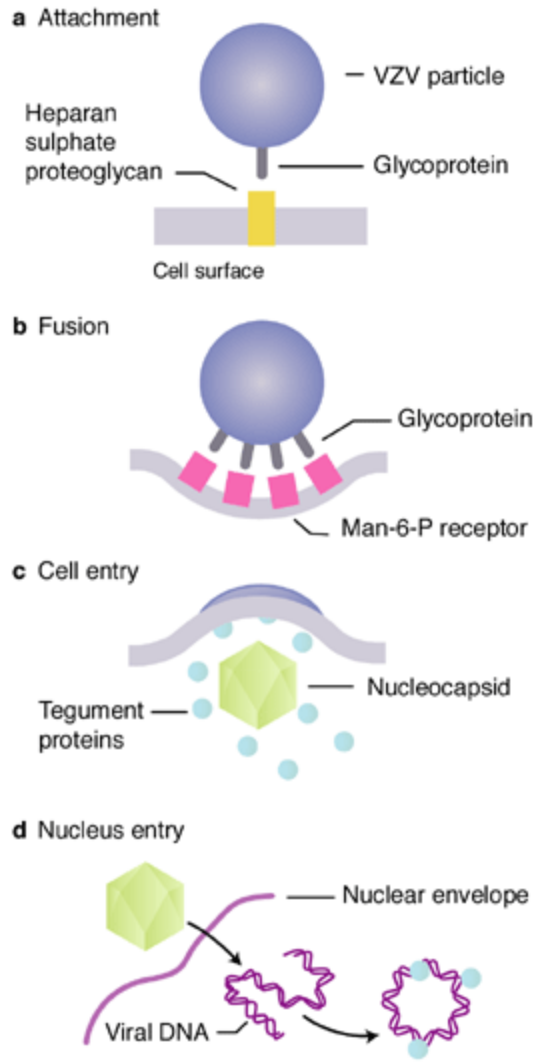
## Receptor-mediated endocytosis

### Fusion in endosome

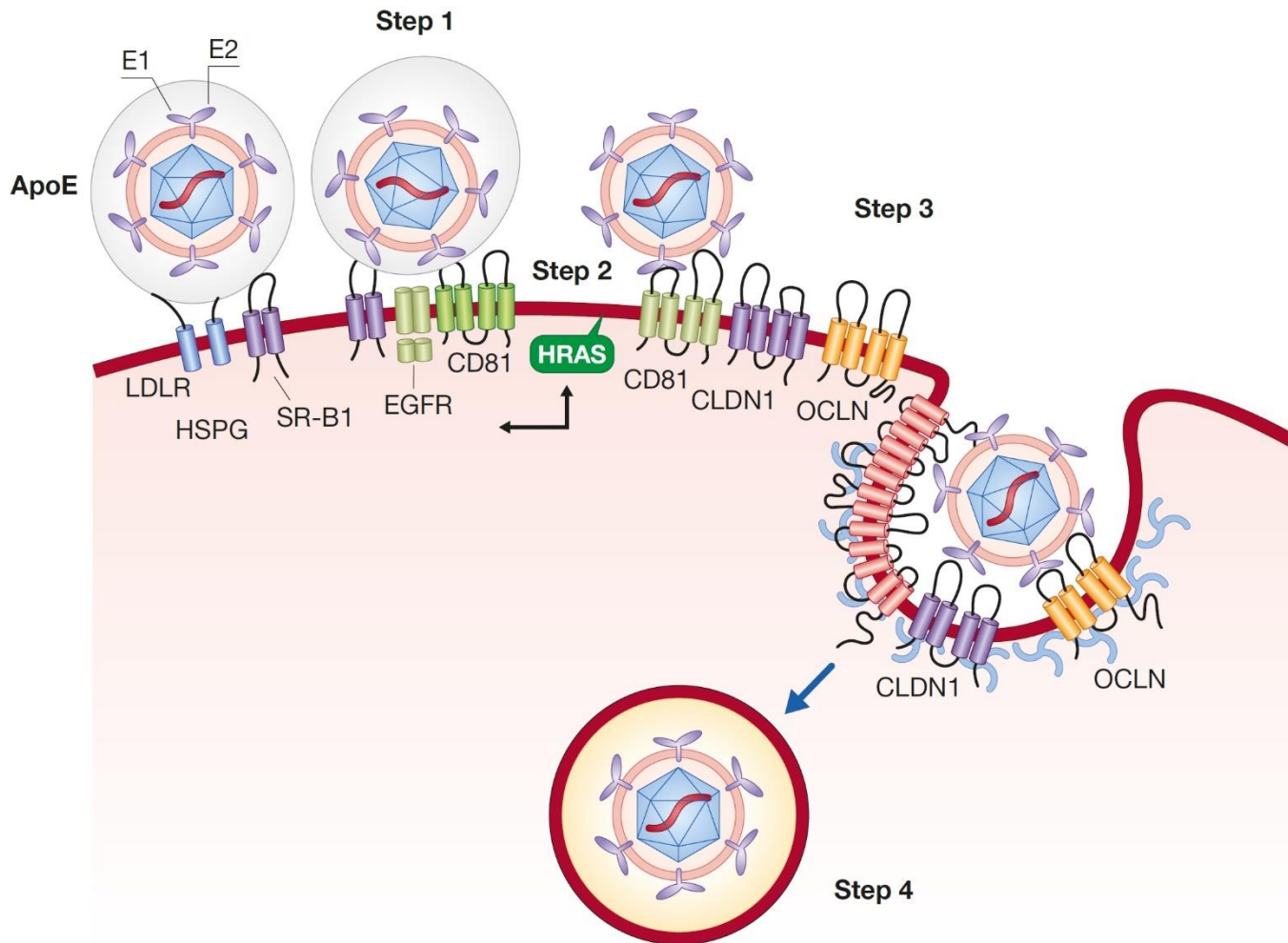


### Lysis of endosome

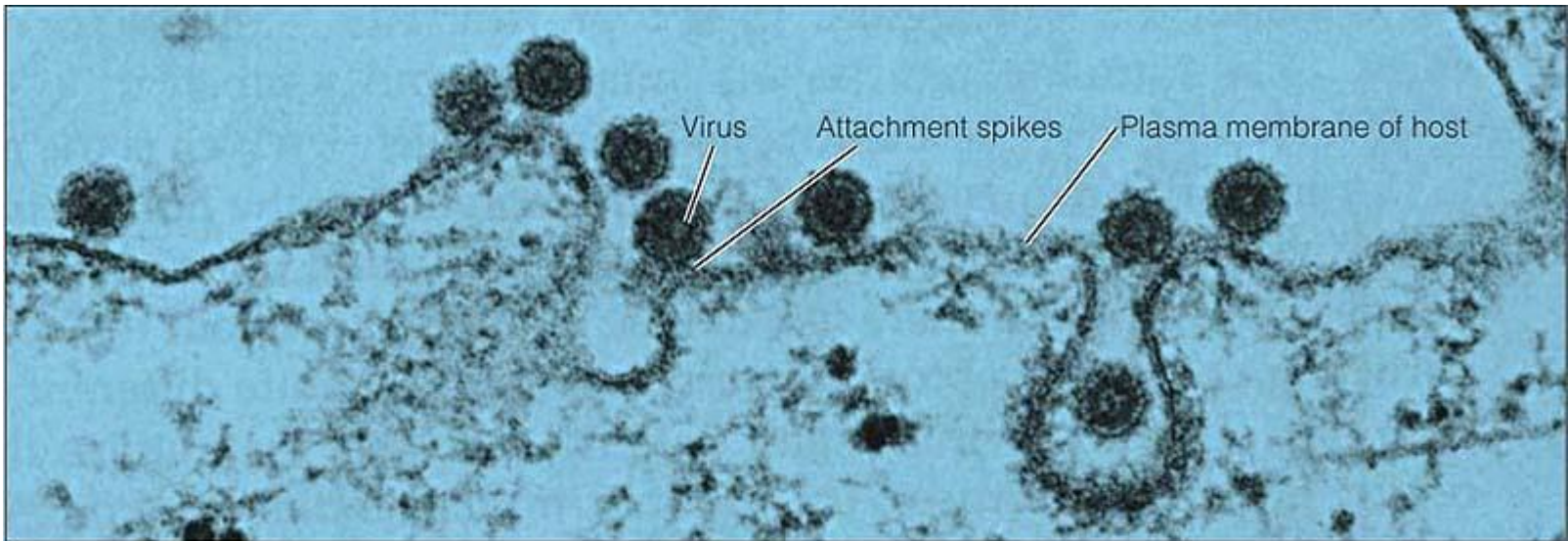




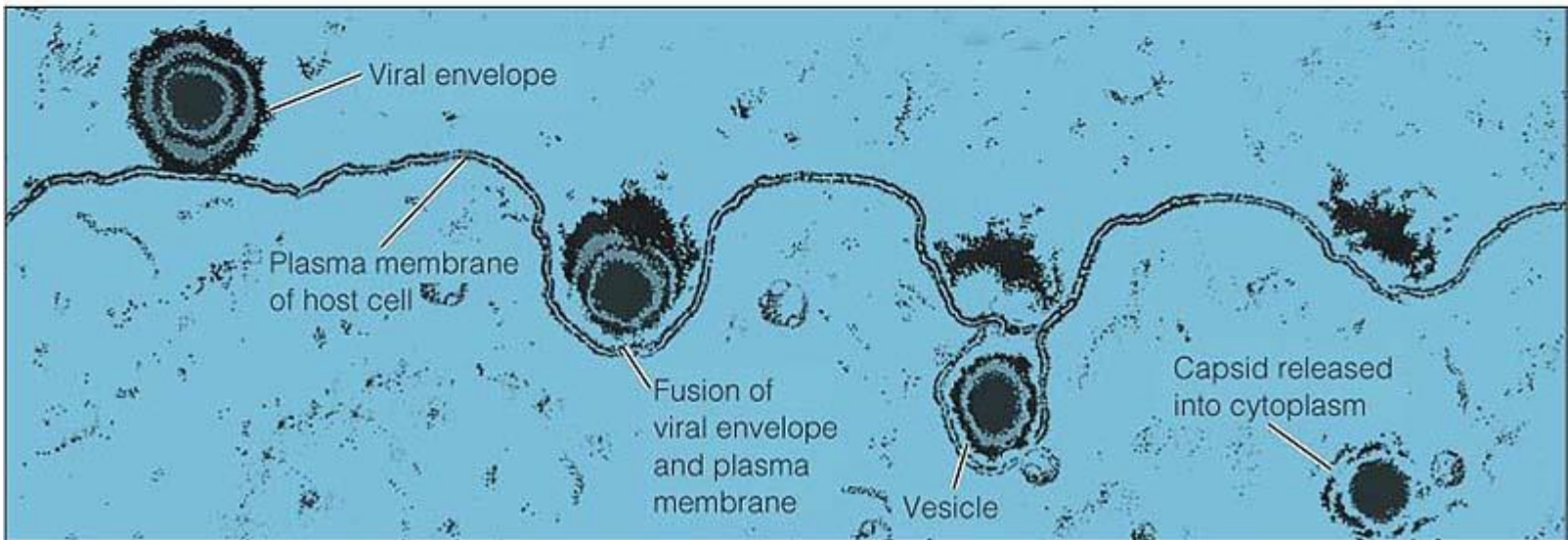
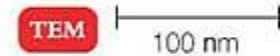
**Varicella-zoster virus cell attachment and fusion**



**Figura 36.1** Il capsid di HCV è circondato da lipidi, in cui sono incorporate le glicoproteine E1 ed E2, ed è associato a lipoproteine sieriche quali l'apolipoproteina E (ApoE). L'attacco alla superficie della cellula richiede l'interazione tra ApoE e una serie di molecole cellulari quali eparansolfati proteoglicani (HSPG), recettori per le proteine a bassa densità (LDLR) e il recettore scavenger di classe B di tipo 1 (SR-B1). Questa interazione induce un cambiamento conformazionale della glicoproteina E2 che gli permette di legarsi al CD81 (step 1). L'interazione tra E2 e CD81 porta all'attivazione delle vie di segnale mediante il recettore per il fattore di crescita epidermico (EGFR) e la via HRAS (step 2) che promuovono il movimento laterale del complesso CD81-HCV verso la *tight junctions protein claudina-1* (CLDN1) (step 3). Il CD81 interagisce con CLDN1 per formare un complesso co-recettore che insieme con la particella HCV viene internalizzato negli endosomi (step 4).



**(a)** Entry of togavirus



**(b)** Entry of herpesvirus

# VIRUS ENTRY

- Penetration by Membrane Fusion
- Endocytic Pathways
- Penetration of Nonenveloped Viruses
  - Membrane puncture
  - Perforation

# VIRUS ENTRY

- Penetration by Membrane Fusion
- Endocytic Pathways

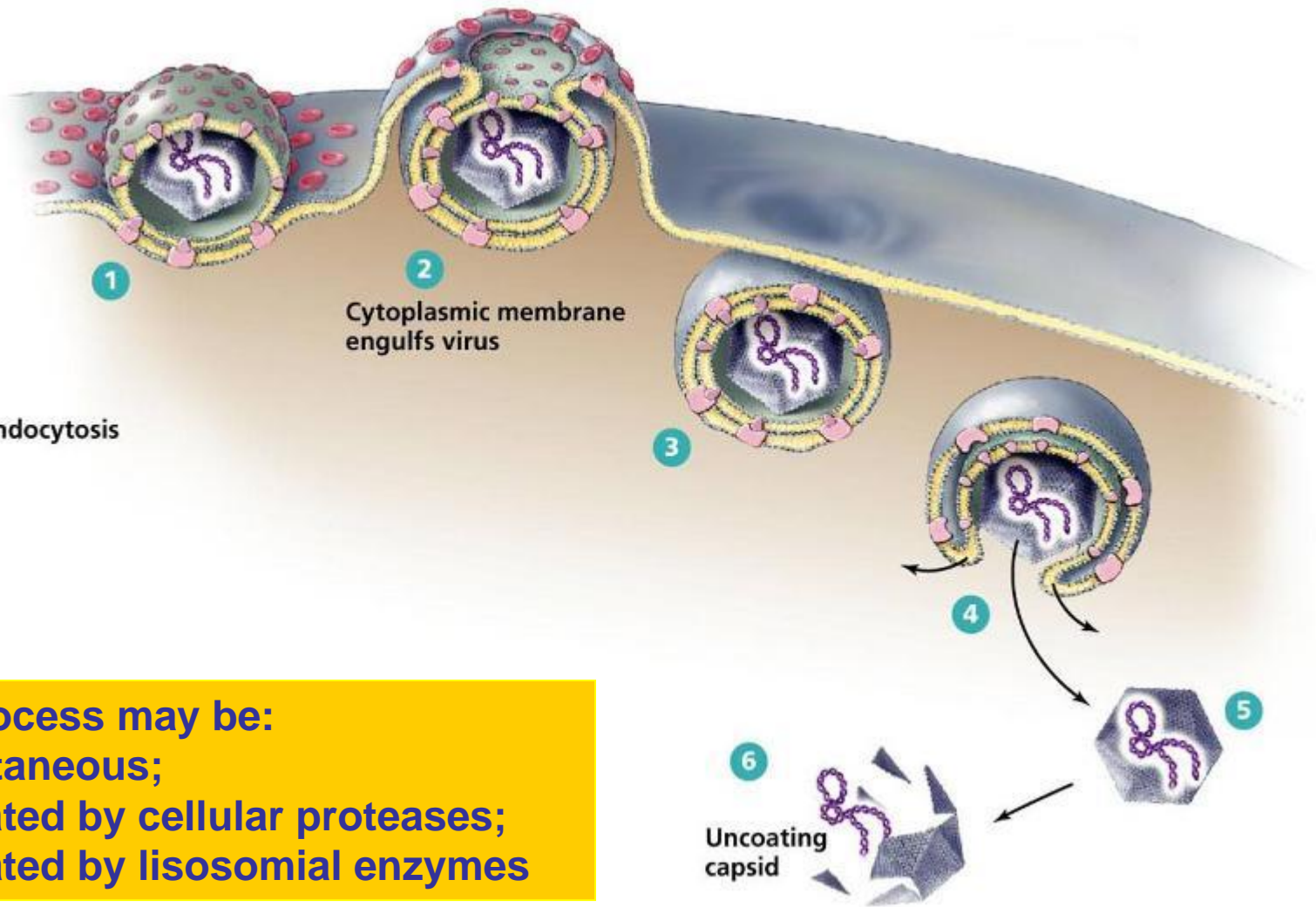
Nonenveloped viruses can use three distinct, general strategies:

- (a) Membrane puncture. The virus particle generates a pore in the membrane through which the genome is selectively released into the cytosol.
- (b) Perforation. The entire capsid is transferred through the membrane without major lysis of the membrane.



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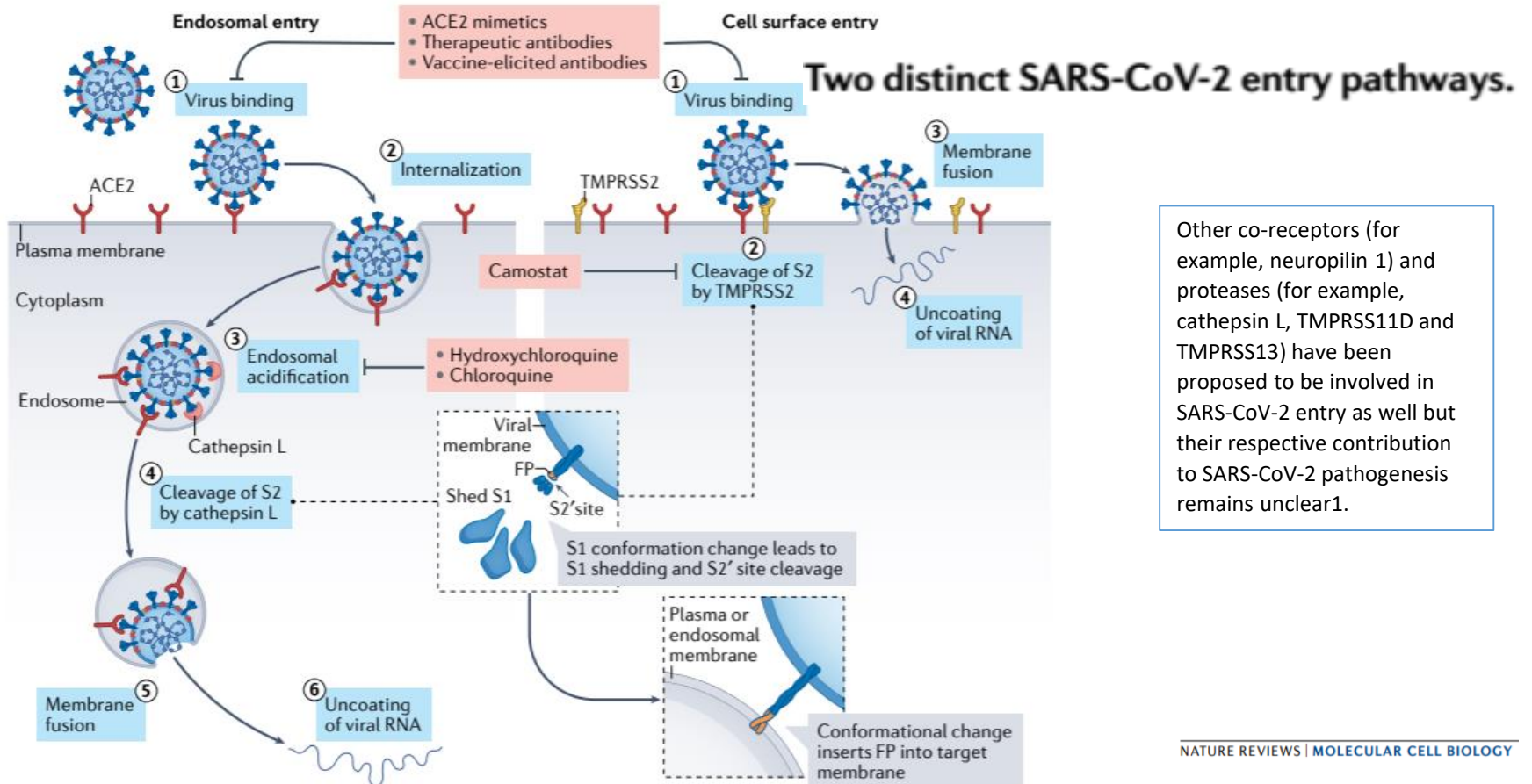


(c) Endocytosis

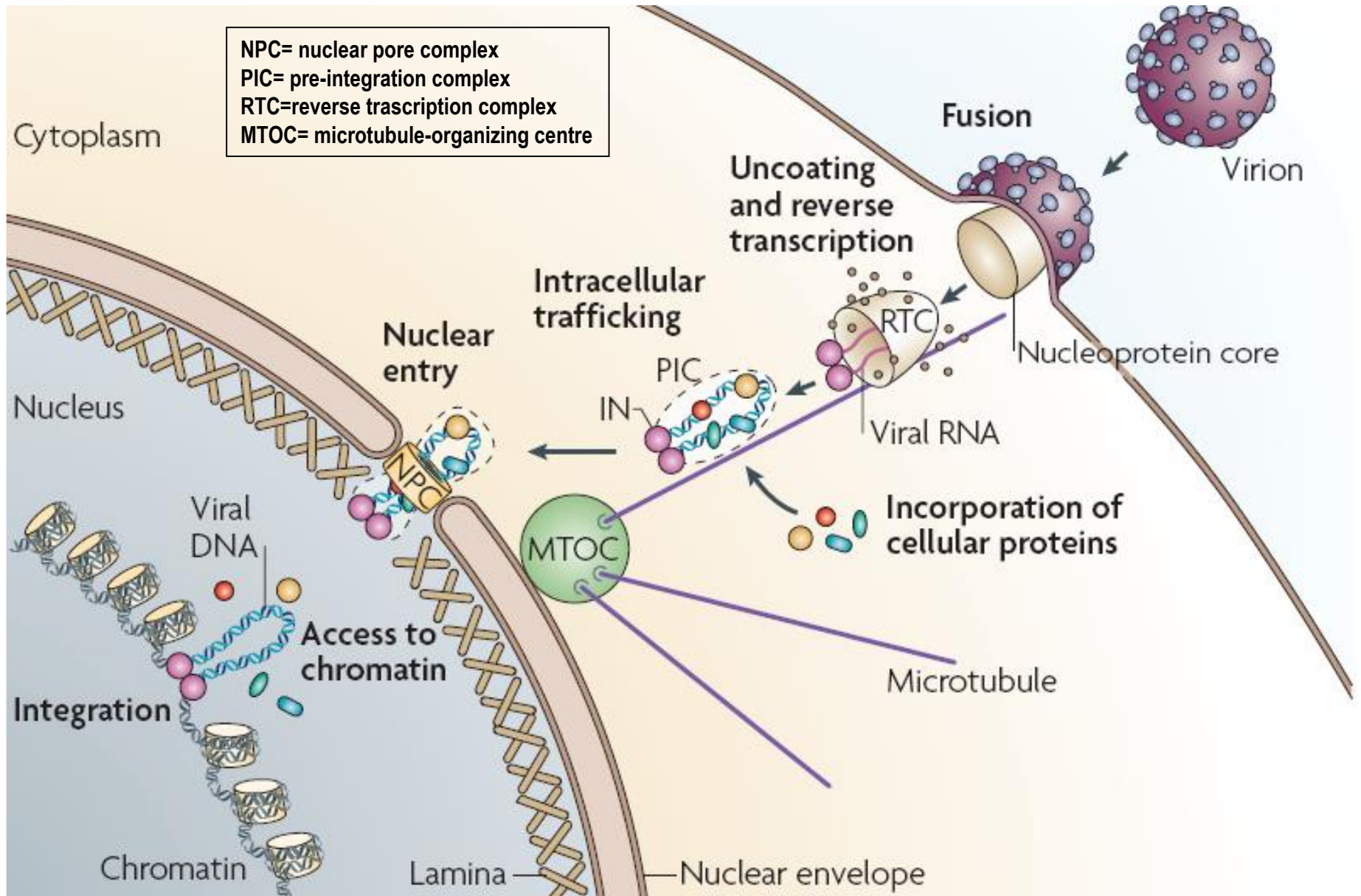
The process may be:

- Spontaneous;
- Mediated by cellular proteases;
- Mediated by lysosomal enzymes

# Why coronavirus and SARS-CoV-2 are so successful -2B- PRODUCTIVE VIRAL LIFE CYCLE



Other co-receptors (for example, neuropilin 1) and proteases (for example, cathepsin L, TMPRSS11D and TMPRSS13) have been proposed to be involved in SARS-CoV-2 entry as well but their respective contribution to SARS-CoV-2 pathogenesis remains unclear.

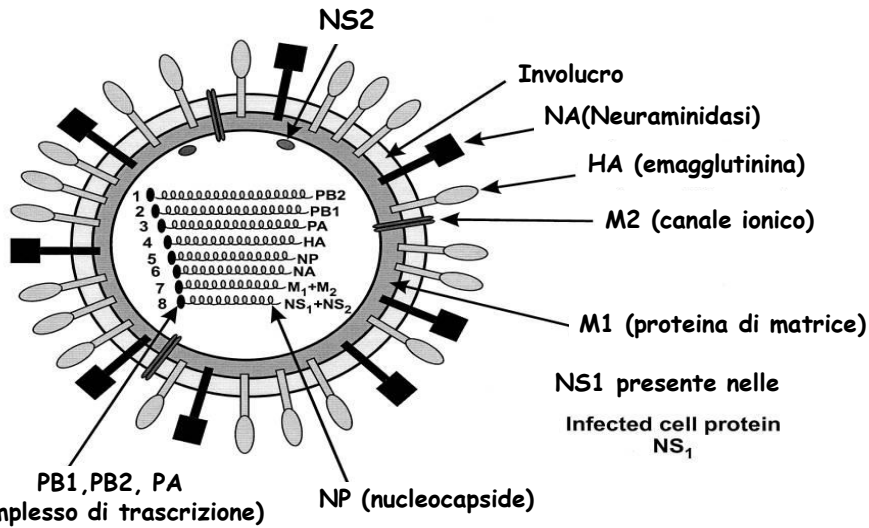


NPC= nuclear pore complex  
 PIC= pre-integration complex  
 RTC=reverse trascription complex  
 MTOC= microtubule-organizing centre

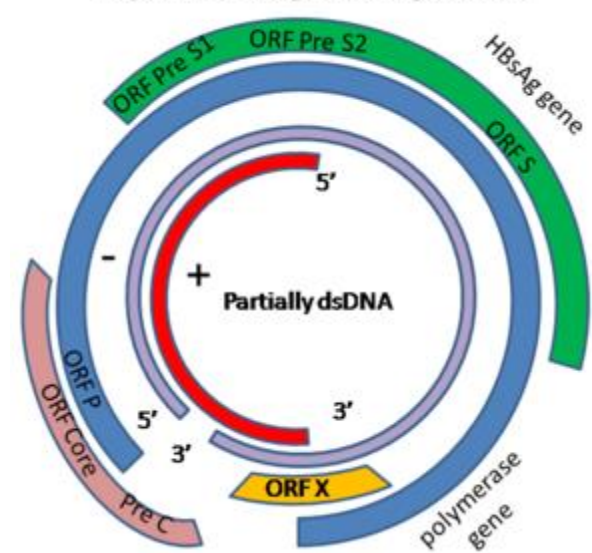
# VIRAL LIFE CYCLE

- **ATTACHMENT**
- **ENTRY**
- **UNCOATING**
- **GENOME REPLICATION (and Expression)**
- **ASSEMBLY**
- **RELEASE**

# Influenza



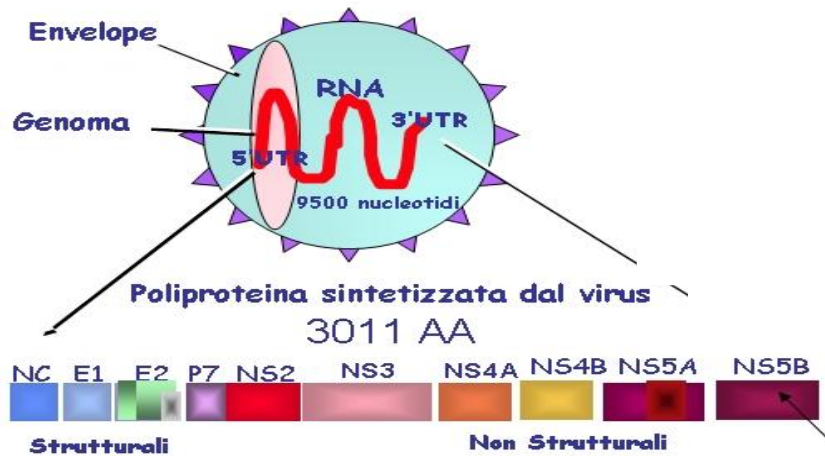
# HepatitisB virus genome organisation



DNA ds circolare

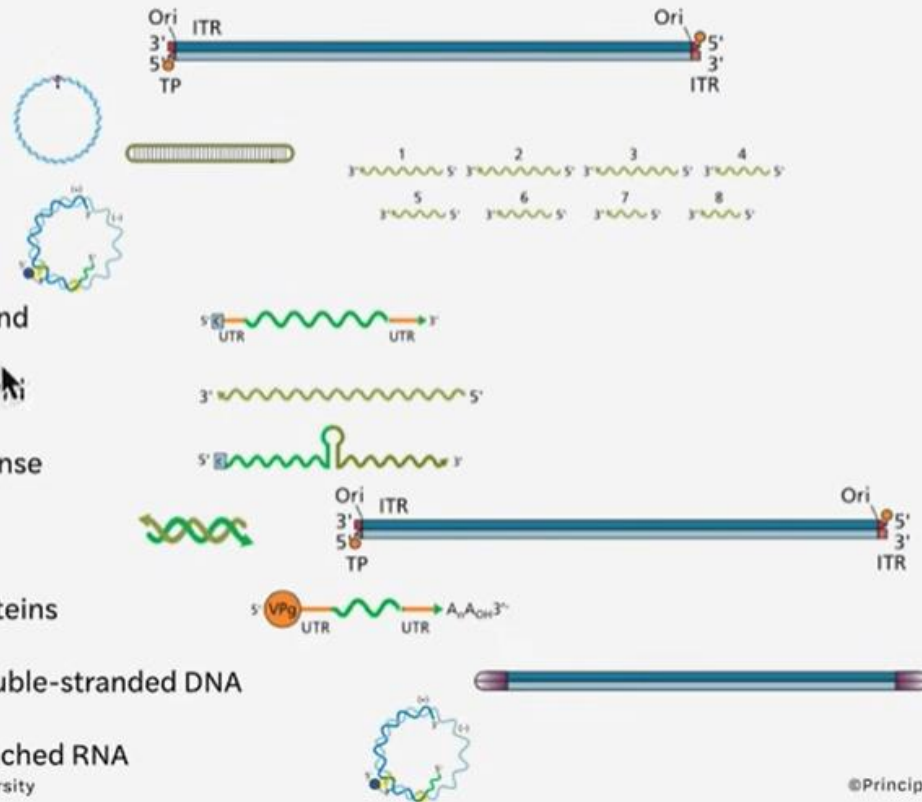
RNA (-) segmentato

HCV RNA ss +

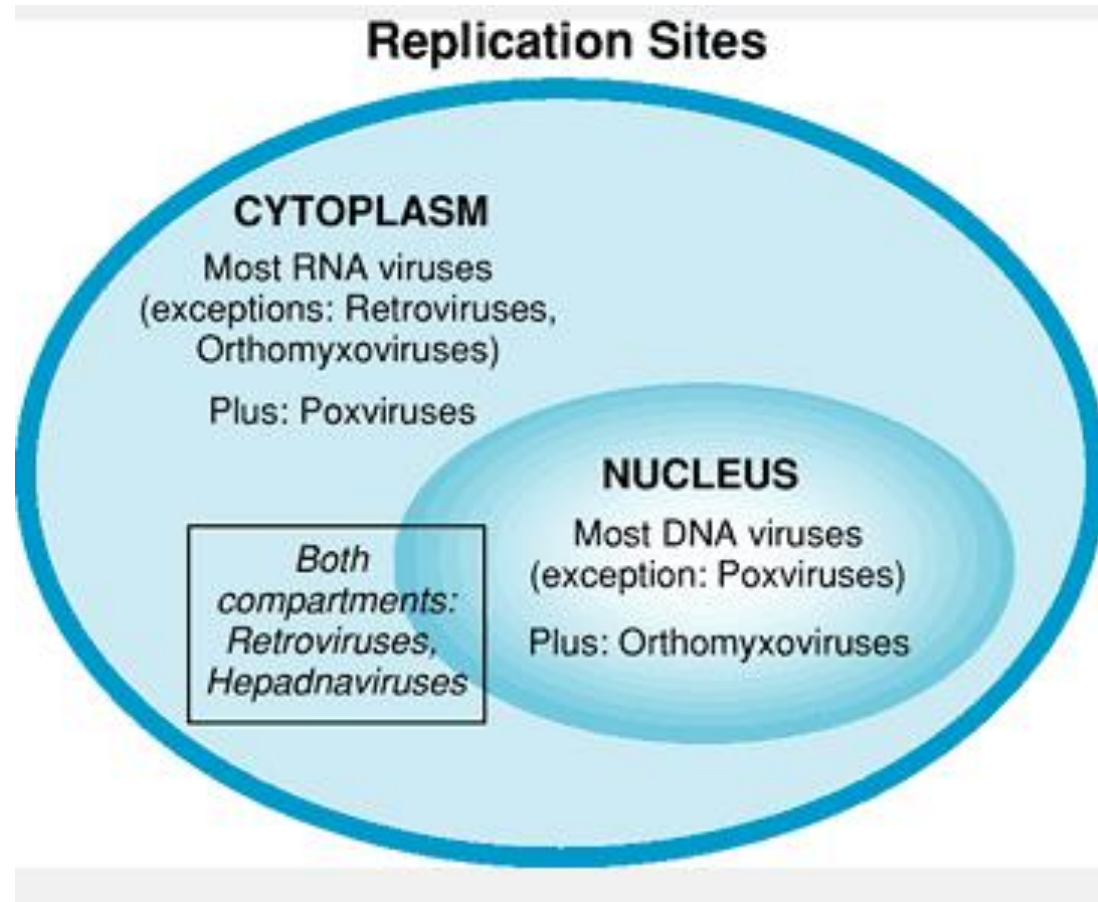


# Viral DNA or RNA genomes are structurally diverse

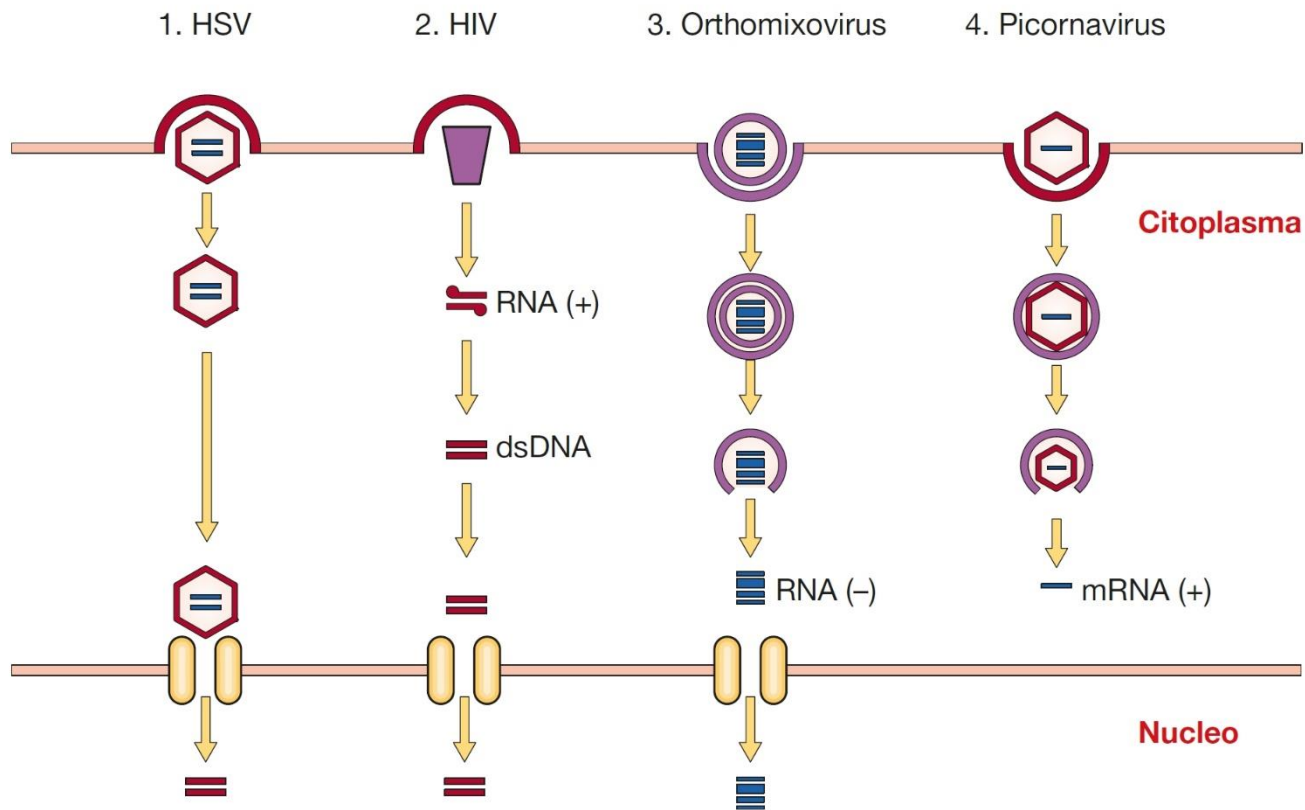
- Linear
- Circular
- Segmented
- Gapped
- Single-stranded (+) strand
- Single-stranded (-) strand
- Single stranded, ambisense
- Double-stranded
- Covalently attached proteins
- Cross-linked ends of double-stranded DNA
- DNA with covalently attached RNA



# Replication sites for major families of viruses





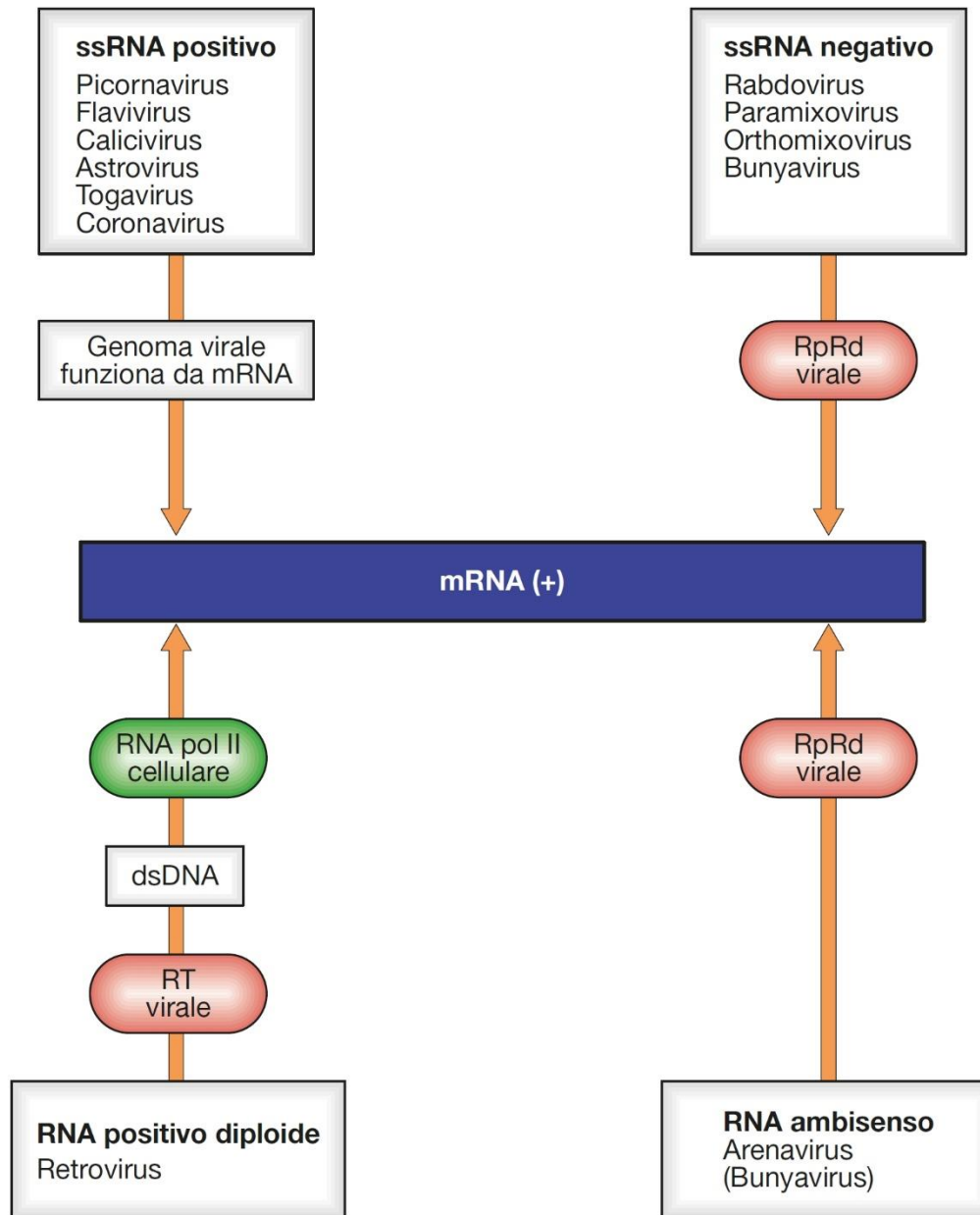


**Figura 36.2 Meccanismi di entrata per diverse classi di virus.** I virus rivestiti possono entrare o mediante fusione (1 e 2) oppure mediante endocitosi seguita da un processo di fusione con la membrana endosomiale (3). I virus nudi, come i picornavirus (4), penetrano all'interno della cellula mediante un processo di endocitosi.

# RNA Virus Genome Strategies

# RNA VIRUSES

- Most RNA viruses replicate **in the cytoplasm**
- **Notable exceptions** are the **orthomyxo-** and **bornaviruses**, whose linear negative-sense RNA genomes replicate in the nucleus, **retroviruses** (that integrate DNA copies of their genomes into cellular chromosomes) and the circular RNA genome of **hepatitis delta virus**
- RNA virus can derive several separate protein products from a single genome through **fragmentation** at the level of:
  - proteins (processing of polyprotein precursors)
  - mRNAs (production of several different monocistronic mRNAs from a single RNA template)
  - genes (multiple genome segments)

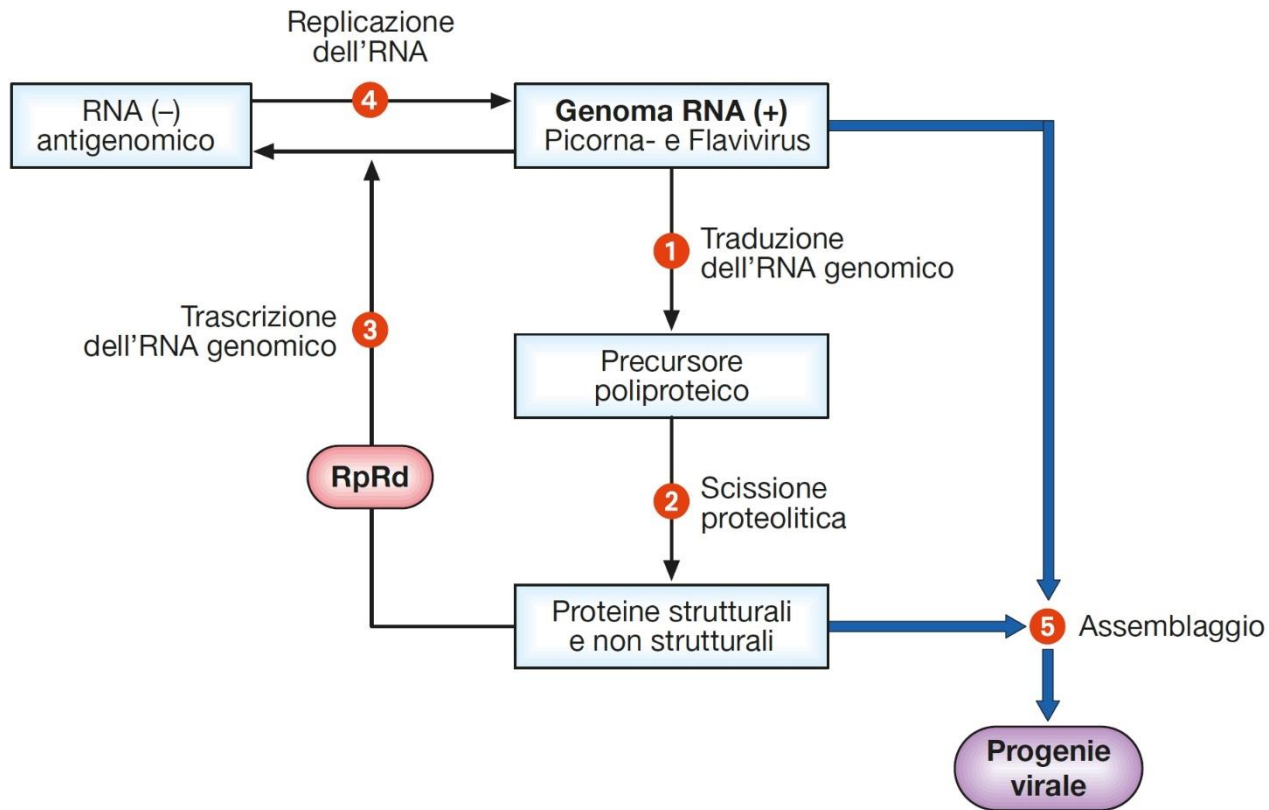


**Figura 36.3** Principali strategie replicative dei virus a RNA. L'RNA virale monofilamento (ssRNA) a polarità positiva viene immediatamente tradotto; l'RNA virale a polarità negativa necessita di essere trascritto da una RNA polimerasi-RNA dipendente (RpRd). Nei retrovirus la trascrittasi inversa (RT) retrotrascrive l'RNA virale in una doppia catena di DNA (dsDNA).

The type of RNA genome largely determines whether the first step of macromolecular synthesis is translation, transcription, or RNA replication.

# Positive-sense RNA genomes

- Viruses with positive-sense ssRNA genomes (**except the retroviruses**) deliver their genomic RNAs directly to cellular ribosomes and begin the infectious cycle with translation.
- Positive-sense RNA viruses fall into two groups: those that produce subgenomic mRNAs and those that do not.

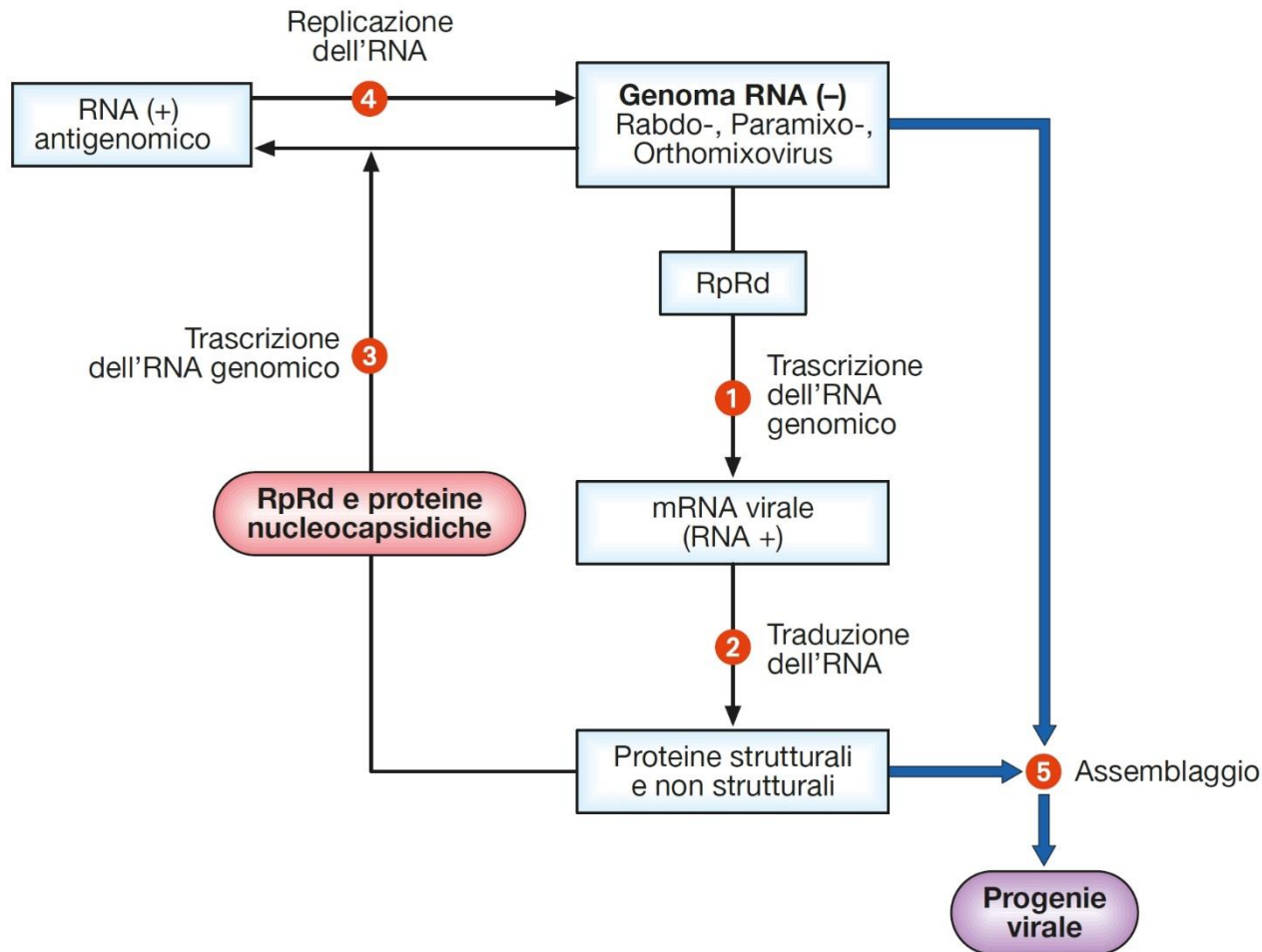


**Figura 36.4** Replicazione di alcuni virus con genoma a RNA (+). **1.** Traduzione dell'RNA genomico in un precursore poliproteico; **2.** scissione proteolitica e formazione delle proteine strutturali e funzionali; **3.** trascrizione dell'RNA genomico in RNA (-) (antigenoma) a opera dell'RNA polimerasi-RNA dipendente (RpRd) neoformata; **4.** sintesi di nuovo RNA (+); **5.** assemblaggio delle proteine strutturali e dei nuovi genomi.

# Negative-sense RNA genomes

- Viral genomes that consist of negative- or ambisense ssRNA or dsRNA are not infectious by themselves because they must begin the infectious cycle by transcribing viral mRNAs, and uninfected cells do not contain an appropriate RdRp. In general, primary transcription is catalyzed by enzymes that are carried into cells by the infecting virions, producing individual mRNAs for the viral proteins

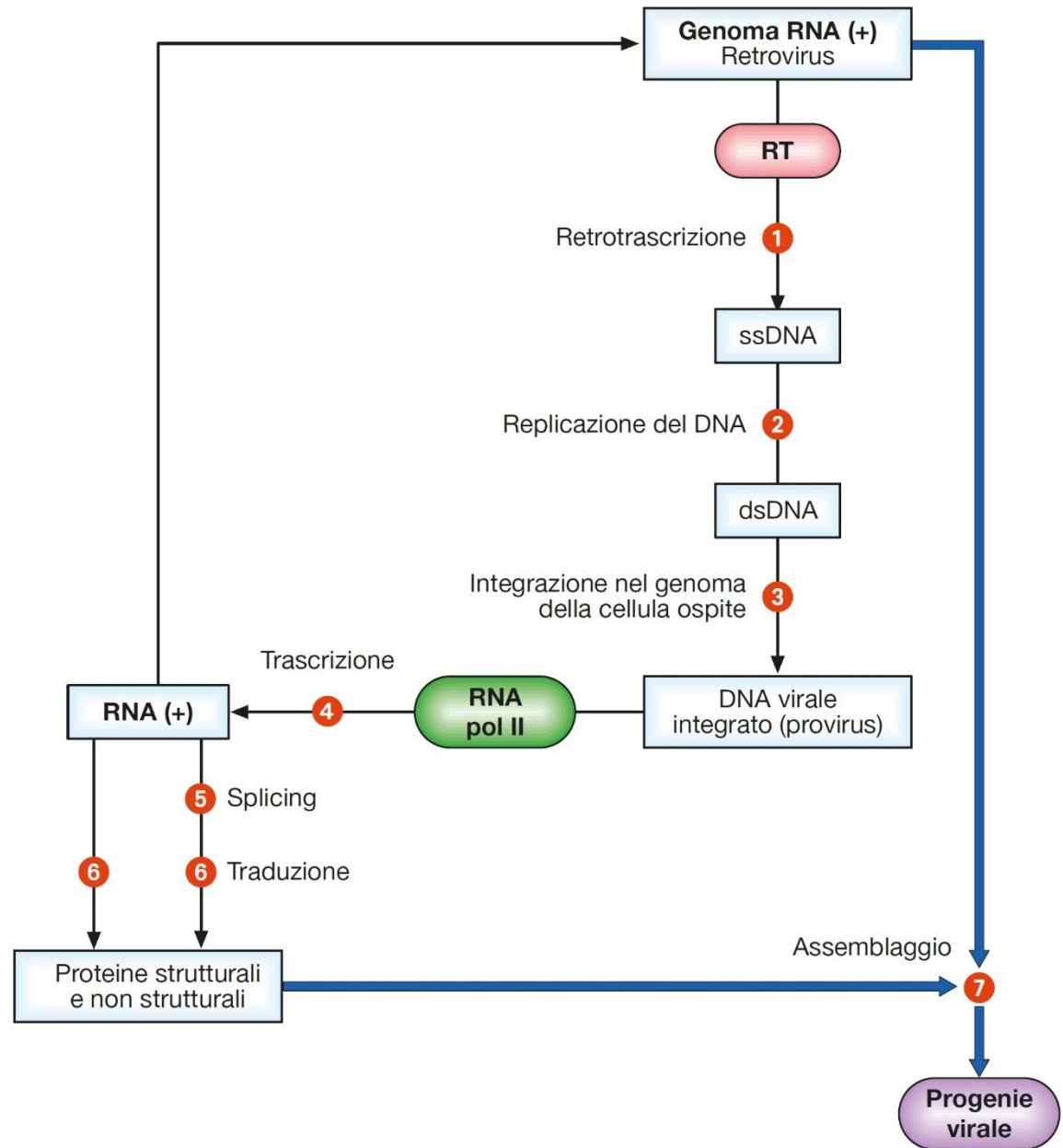


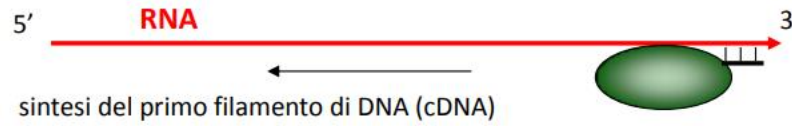


**Figura 36.5** Replicazione di alcuni virus con genoma a RNA (-). **1.** Trascrizione dell'RNA (-) in RNA (+) a opera della RNA polimerasi-RNA dipendente (RpRd) associata al virione; **2.** traduzione dell'RNA (+) neoformato; **3.** trascrizione dell'RNA genomico in RNA (+) (antigenoma) a opera della RpRd virale; **4.** sintesi del nuovo RNA di progenie; **5.** assemblaggio delle proteine strutturali e dei nuovi genomi.

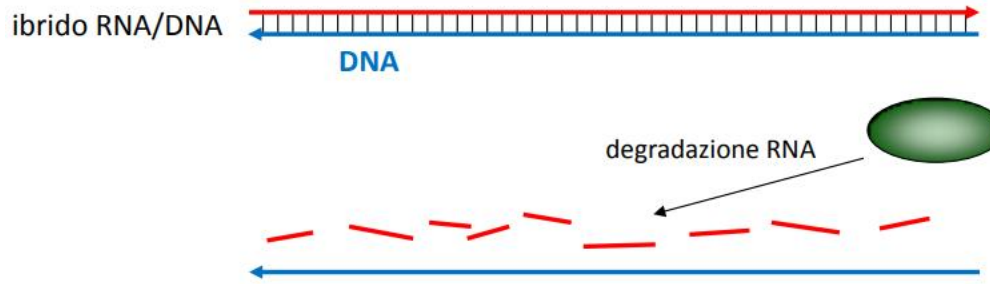
Positive-sense RNA genomes  
which are transcribed into DNA  
(Retroviridae)

**Figura 36.6** Replicazione dei retrovirus. **1.** e **2.** Retrotrascrizione dell'RNA genomico in DNA ad opera della trascrittasi inversa (RT) virale; **3.** integrazione del genoma virale nel DNA cellulare; **4.** trascrizione del genoma virale ad opera dell'RNA polimerasi II; **5.** splicing dell'RNA; **6.** traduzione dell'mRNA (sottoposto e non a splicing); **7.** assemblaggio delle proteine strutturali e dei nuovi genomi.

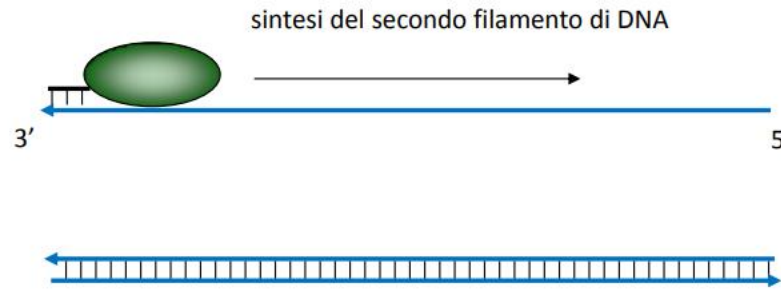




Attività DNA-polimerasi  
RNA-dipendente

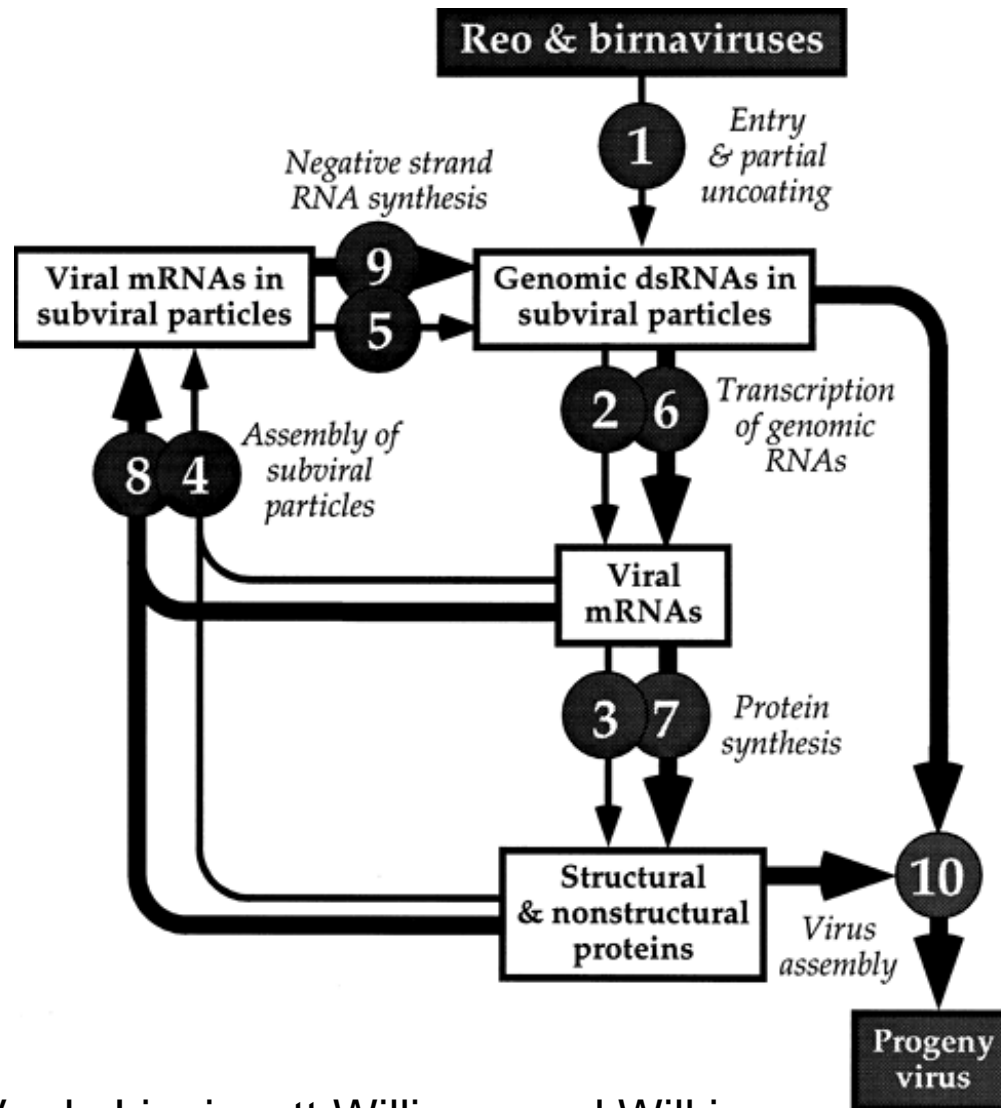


Attività RNasi H



Attività DNA-polimerasi  
DNA-dipendente

# Double-stranded RNA viruses



# DNA Virus Genome Strategies

# DNA VIRUSES (1)

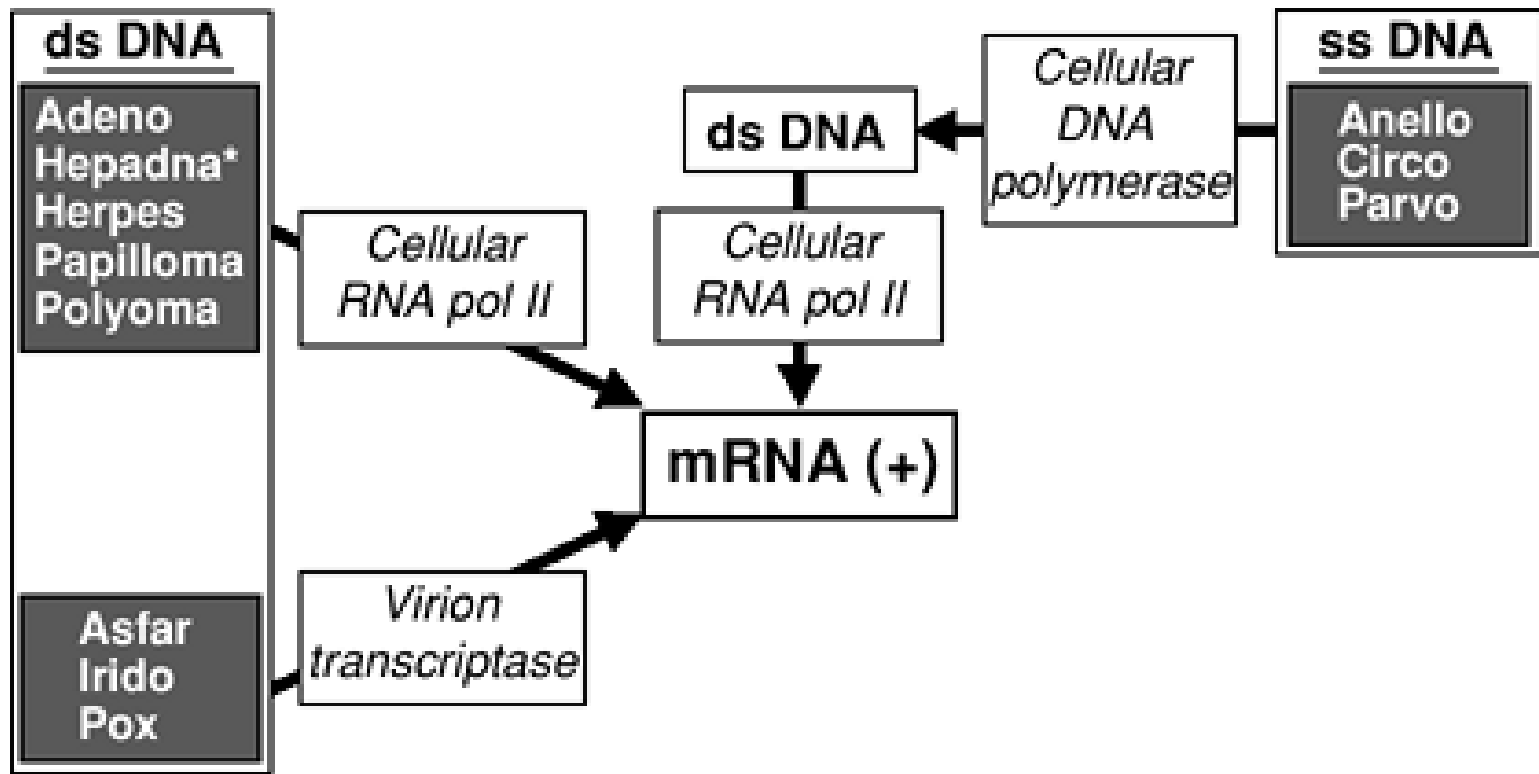
- Most DNA viruses of eukaryotes transcribe and replicate their genomes and assemble progeny **in the nucleus**, the site of cellular DNA transcription and replication. The exceptions are the poxviruses, iridoviruses, and African swine fever virus, which replicate their DNA genomes partly or completely in the cytoplasm.
- DNA viruses use **host cell enzymes for transcription**
- **Early in infection**, a subset of so-called immediate early viral genes is expressed to produce mostly catalytic amounts of the **nonstructural proteins** required for DNA replication and modulation of the intracellular environment
- Sometimes these early events can lead to **neoplastic transformation**
- Most viruses with small DNA genomes use host cell enzymes for **DNA replication**; those with intermediate and large size genomes have sufficient genetic capacity to encode DNA polymerases
- After genome replication, a different subset of genes is expressed (**late genes**) that directs synthesis of stoichiometric amounts of the **structural proteins** required for progeny virus assembly

# DNA VIRUSES (2)

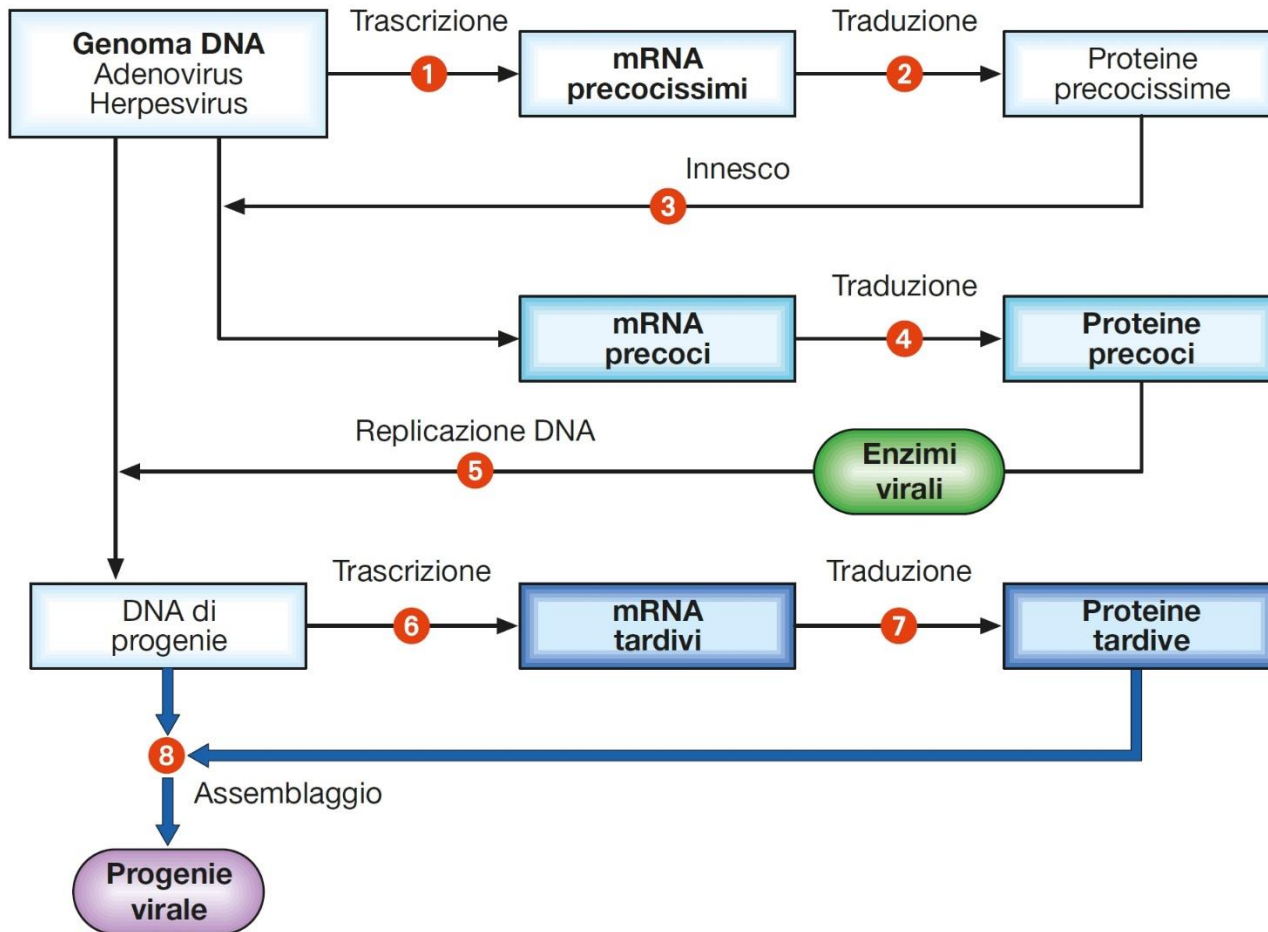
- Because cellular DNA synthesis occurs only **during the S-phase of the cell cycle** and not at all in terminally differentiated G0 cells, **viruses that depend on the DNA polymerases of the host must either wait** for infected cells to enter S-phase spontaneously, **as do parvoviruses**, or early in infection **they must express one or more viral oncogenes to** override the regulation imposed by the cell cycle control proteins p53 and pRb and thereby **stimulate infected cells to enter S-phase** prematurely, **as do polyoma- and papillomaviruses**, among others.
- **Viruses with large DNA genomes** (e.g., **herpes- and poxviruses**) **encode some of these enzymes themselves, and can thus replicate in nondividing cells** and other environments that would otherwise be inhospitable for DNA replication, such as terminally differentiated cells of the nervous system (some herpesviruses) or even the cytoplasm (poxviruses).



Pathways of primary mRNA synthesis by DNA viruses of animals.

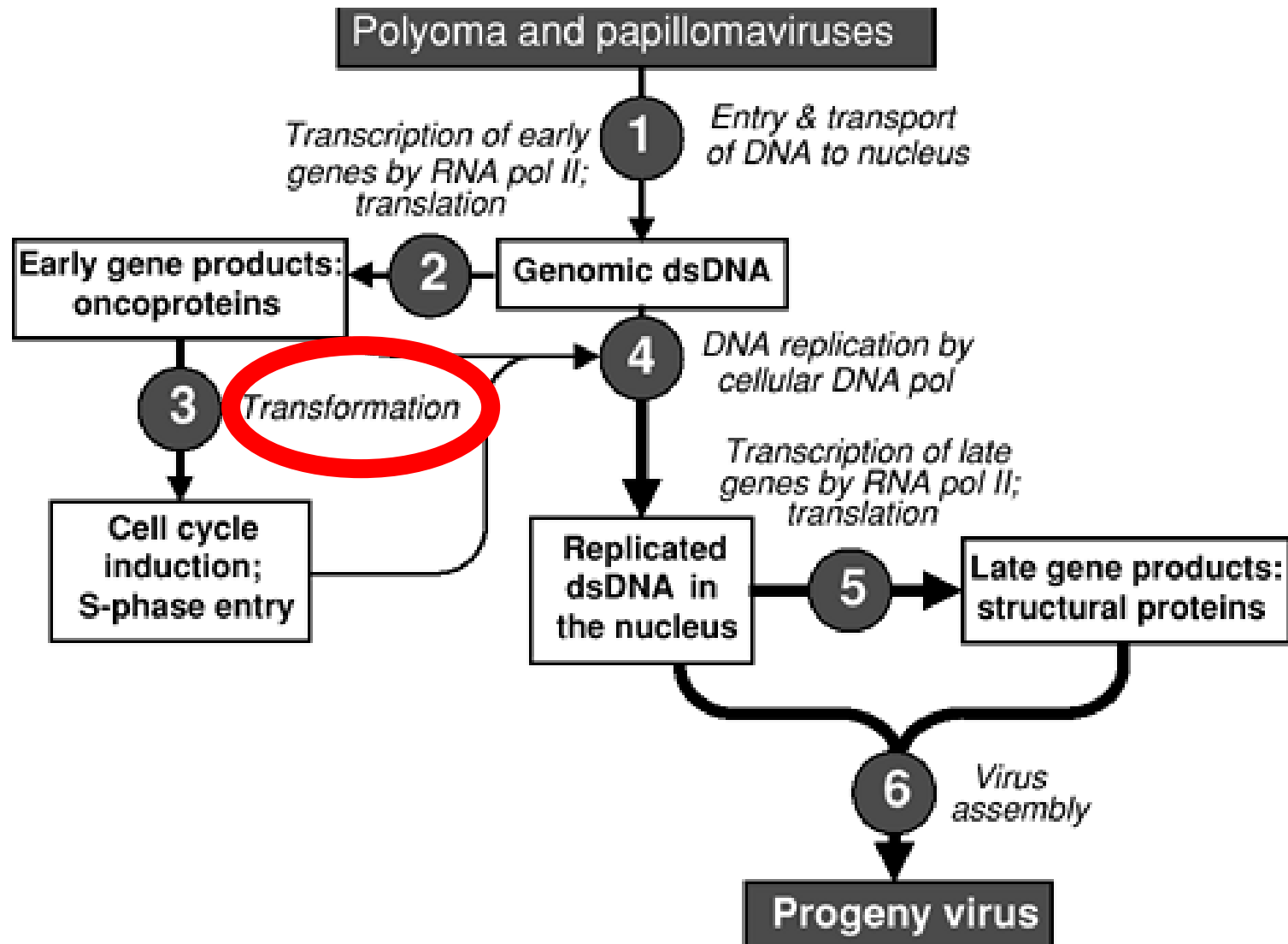


\*Hepadnaviruses replicate via reverse transcription of an ssRNA intermediate

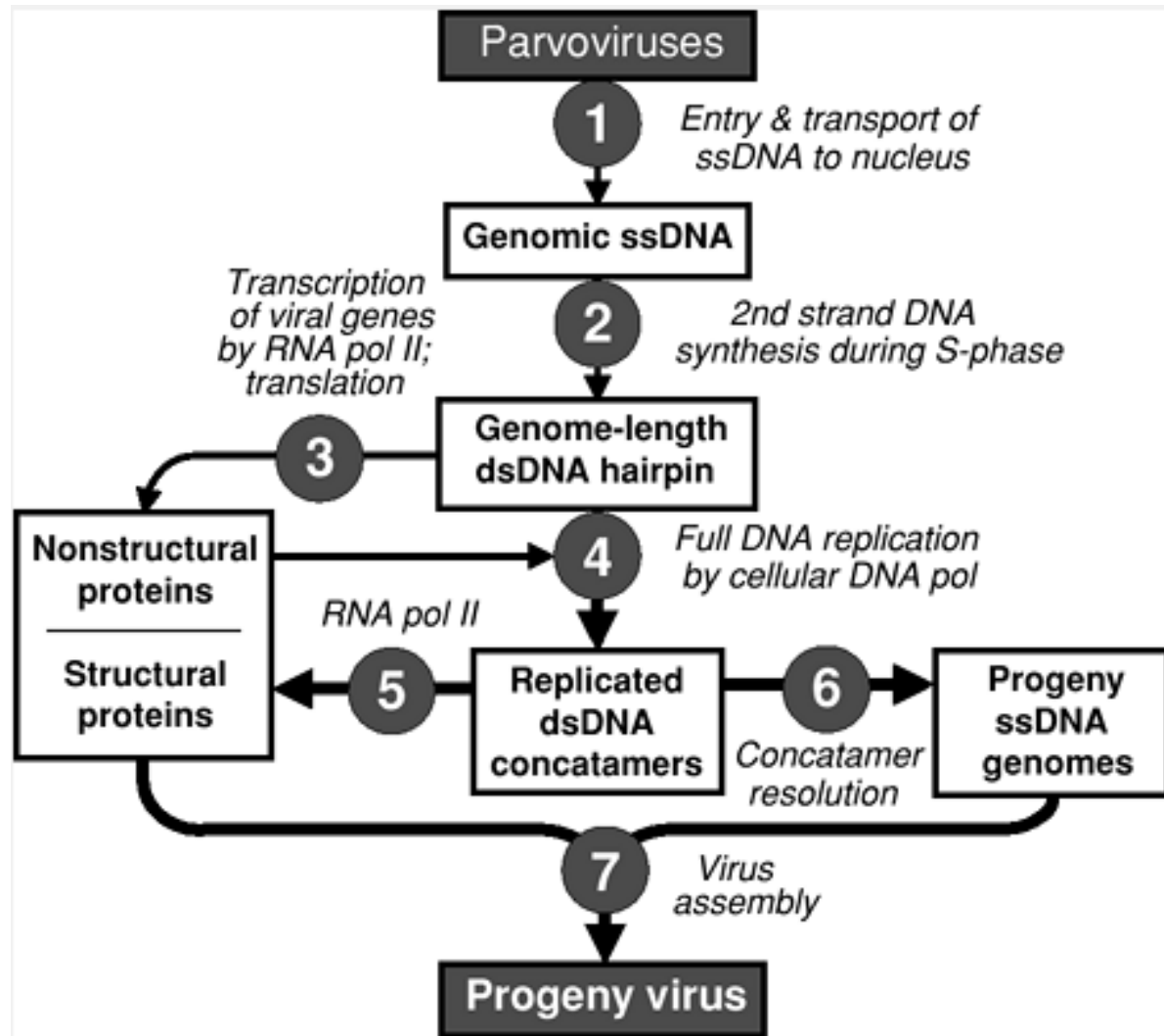


**Figura 36.7** Replicazione di alcuni virus con genoma dsDNA. **1.** Trascrizione del DNA in mRNA precocissimi; **2.** traduzione dei messaggeri precocissimi; **3.** innesco della sintesi dei messaggeri precoci da parte delle proteine precocissime; **4.** traduzione degli mRNA precoci; **5.** replicazione del DNA virale a opera di enzimi virali neoformati; **6.** e **7.** trascrizione e successiva traduzione degli mRNA tardivi dal DNA neoformato; **8.** assemblaggio delle proteine strutturali e dei nuovi genomi.

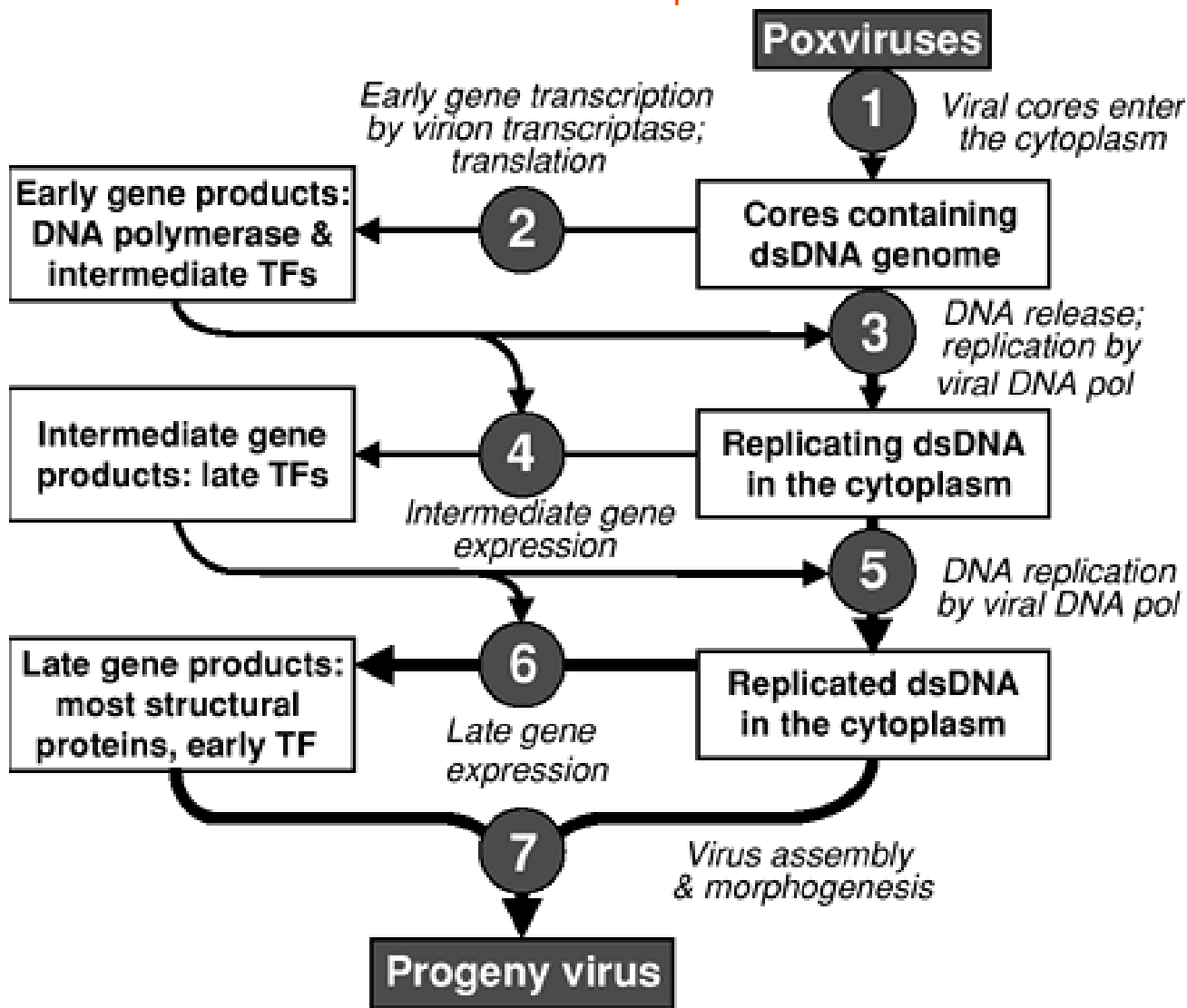
# Small DNA genomes use host cell enzymes for both transcription and DNA replication

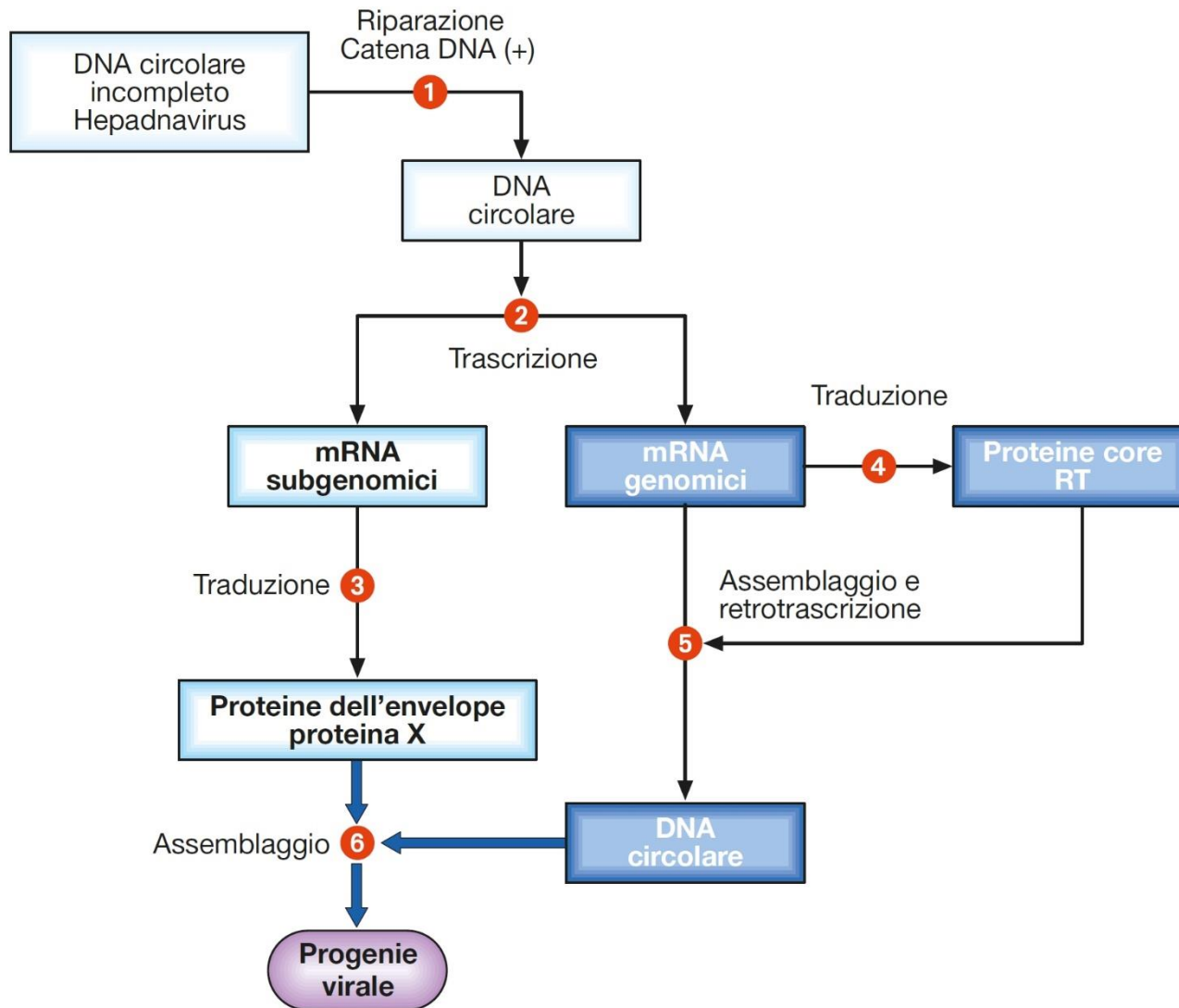


# Small DNA genomes use host cell enzymes for both transcription and DNA replication



Virus with large (130-350 kb) genomes have sufficient genetic capacity to encode DNA polymerases and binding proteins, and have entire multisubunit virus-specific RNA polymerases that mediate all viral transcription



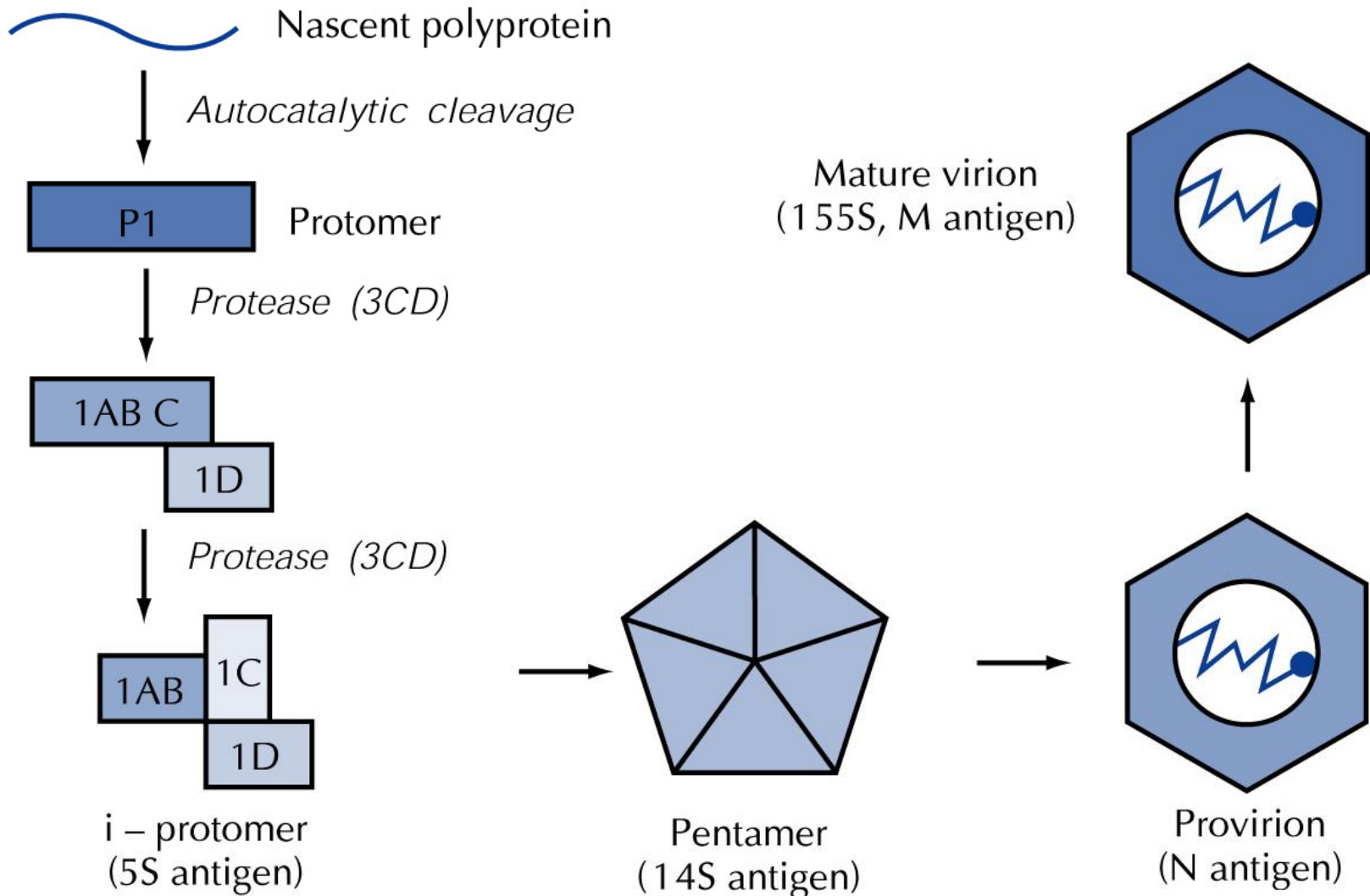


**Figura 36.8 Replicazione degli hepadnavirus.** 1. Riparazione della catena di DNA (+) interrotta; 2. trascrizione del DNA in RNA genomici e subgenomici; 3. traduzione degli mRNA subgenomici; 4. traduzione degli mRNA genomici; 5. retrotrascrizione dell'RNA genomico in DNA a opera della trascrittasi inversa (RT) virale; 6. assemblaggio delle proteine strutturali e dei nuovi genomi.

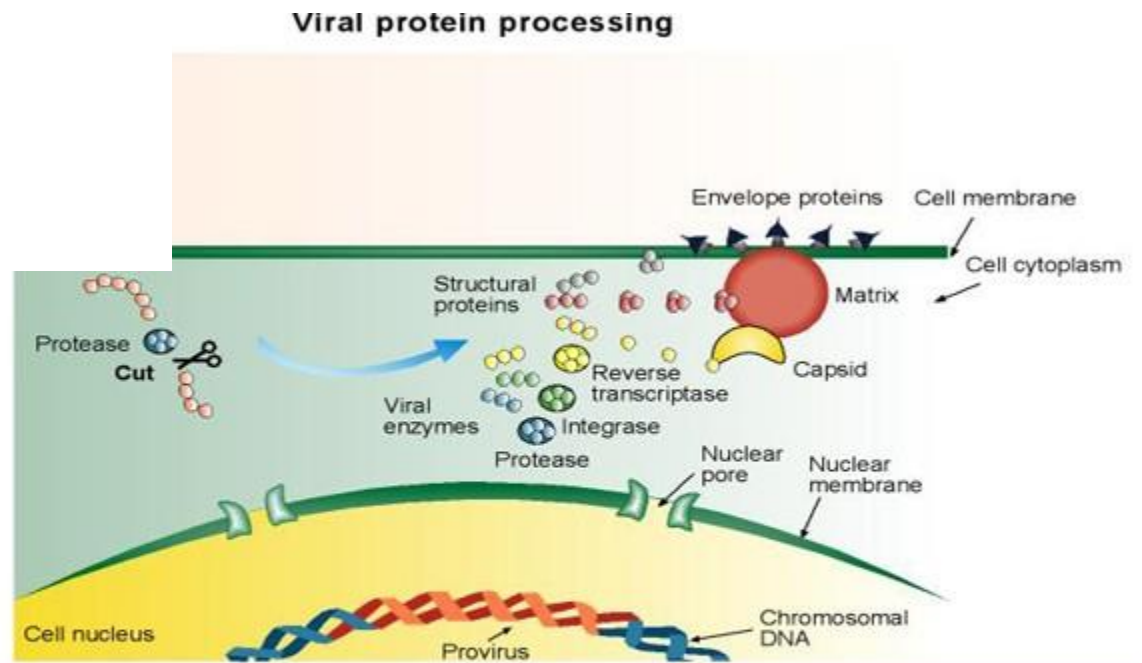
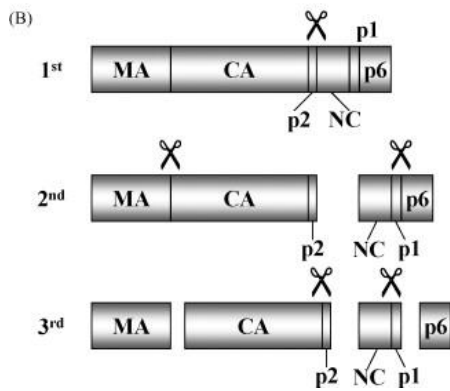
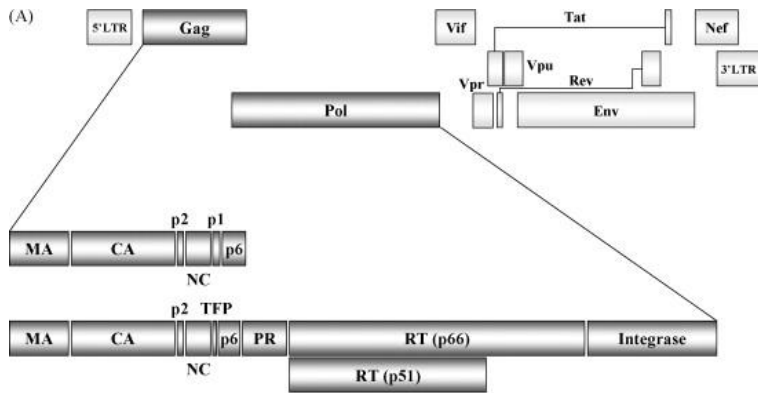
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# Assembly of Picornavirus



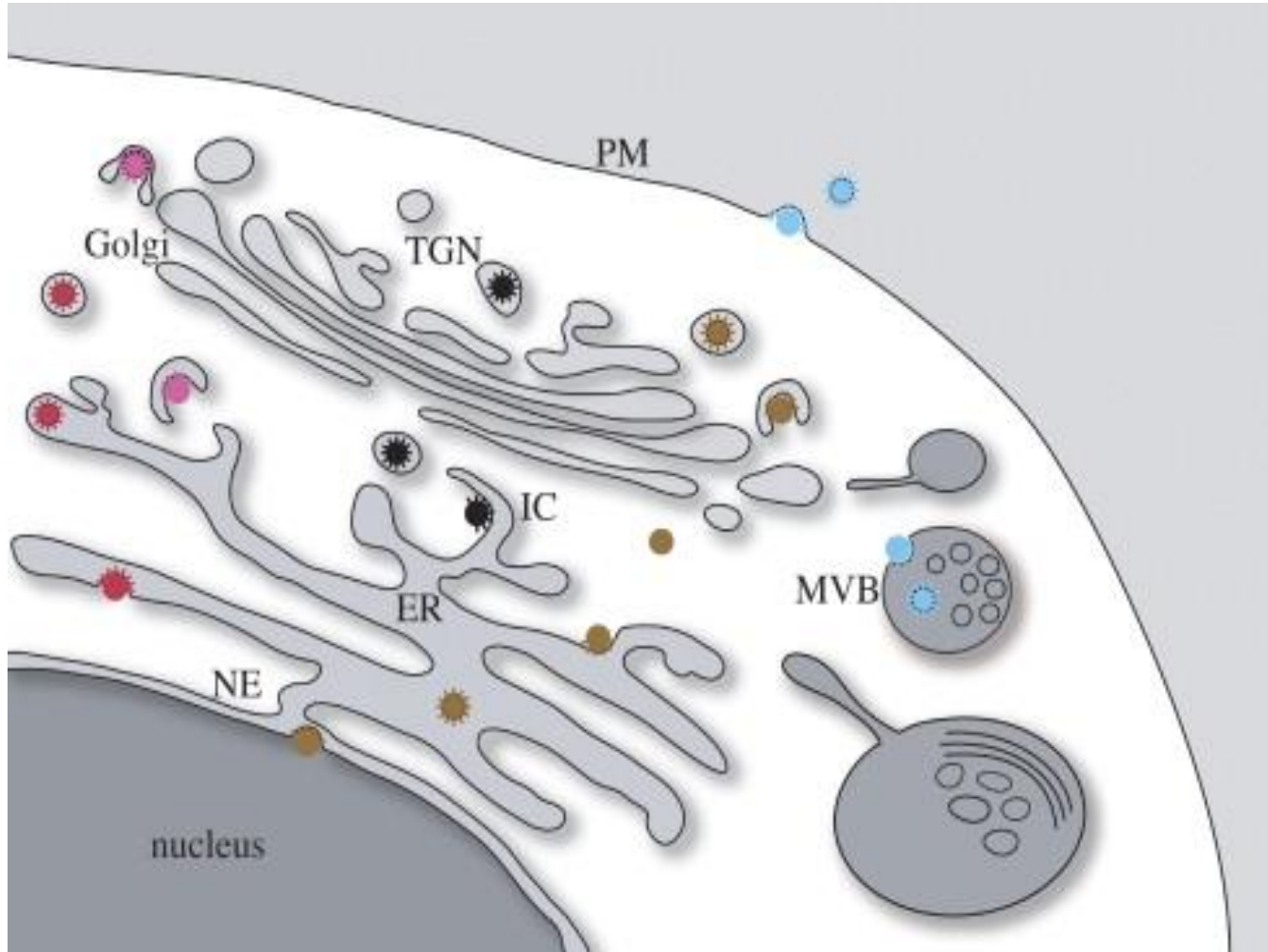


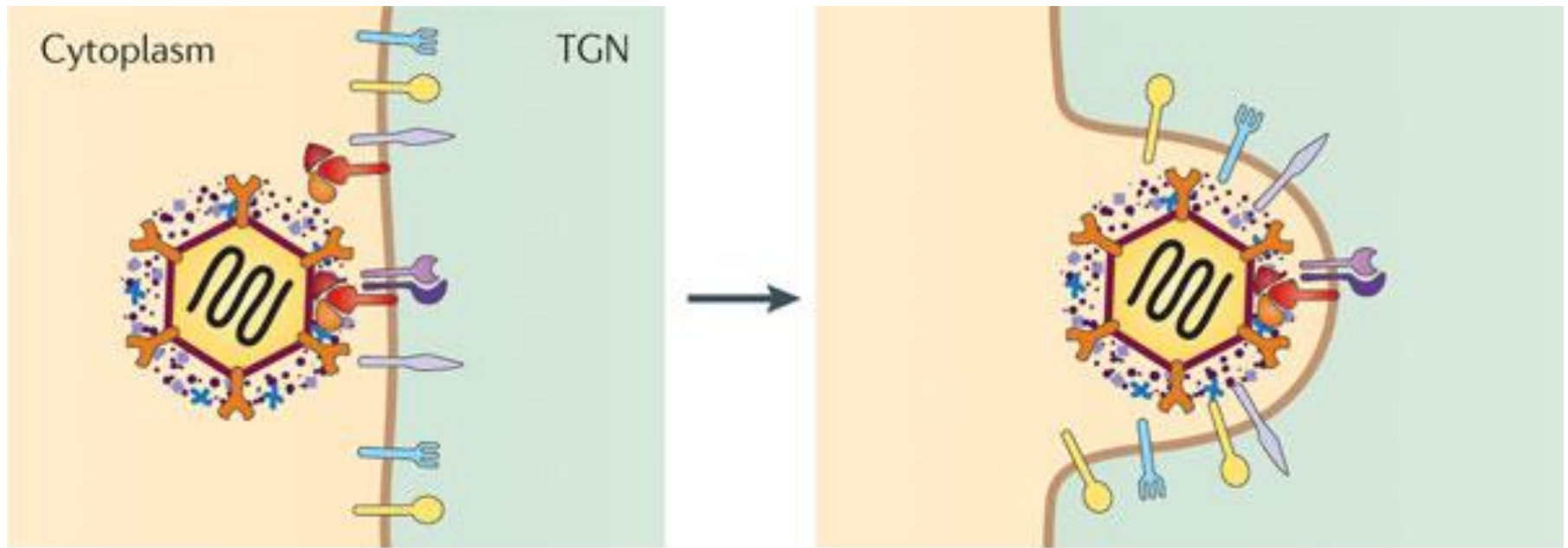


Virus particles are assembled at the cell membrane by recruitment of the newly synthesised viral proteins. Viral genomic RNA is also transcribed for incorporation into the virus core.

# Various membrane systems are implicated in budding of enveloped viruses

nuclear envelope (NE), endoplasmic reticulum (ER), endosomes, intermediate or pre-Golgi compartment (IC), Golgi cisternae and the *trans*-Golgi-network (TGN) multivesicular body (MVB) have been proposed to serve as platforms for virus budding





Tegument proteins   VHS-VP16   VP22   gE-gI

Herpes simplex virus

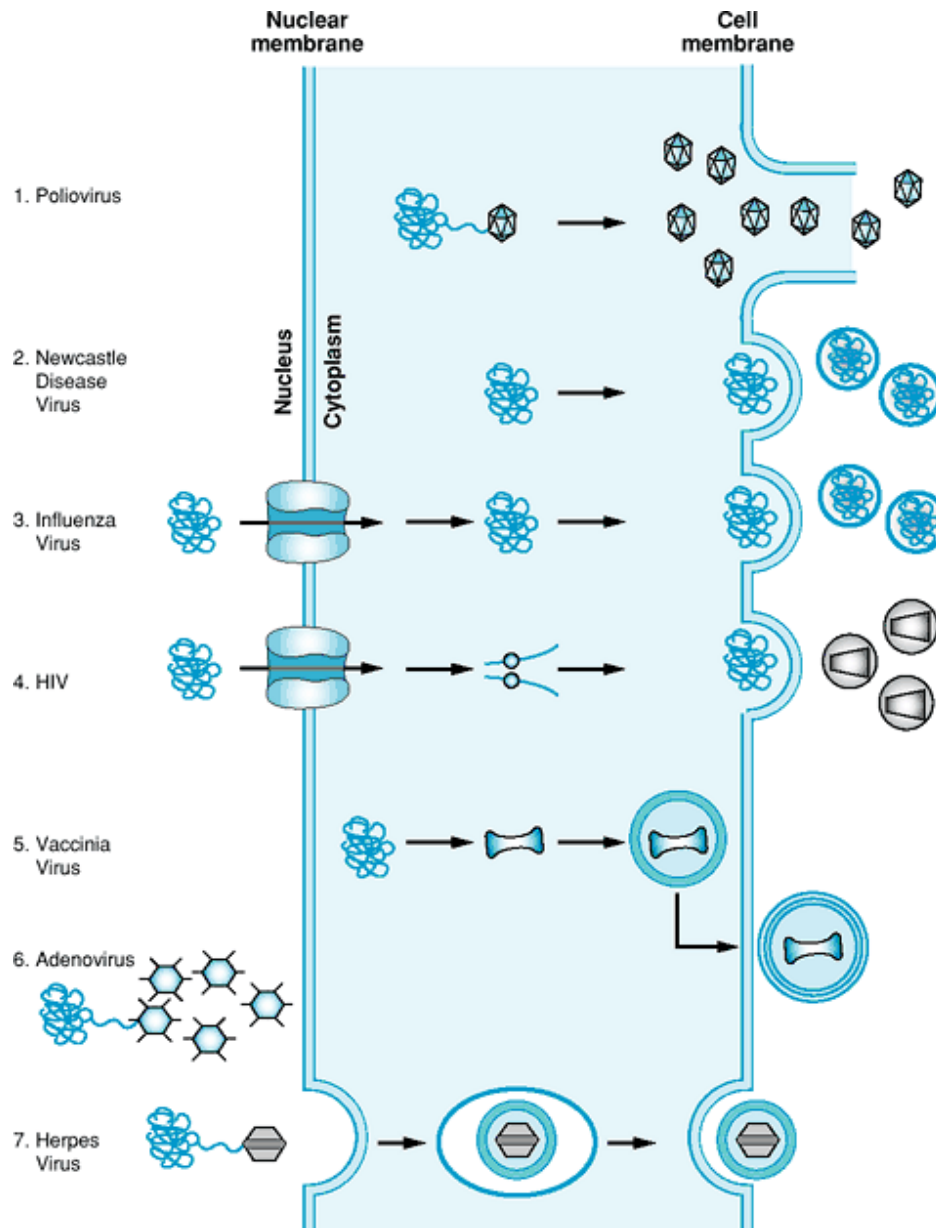
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# VIRUS RELEASE

- CELL LYSIS
- BUDDING

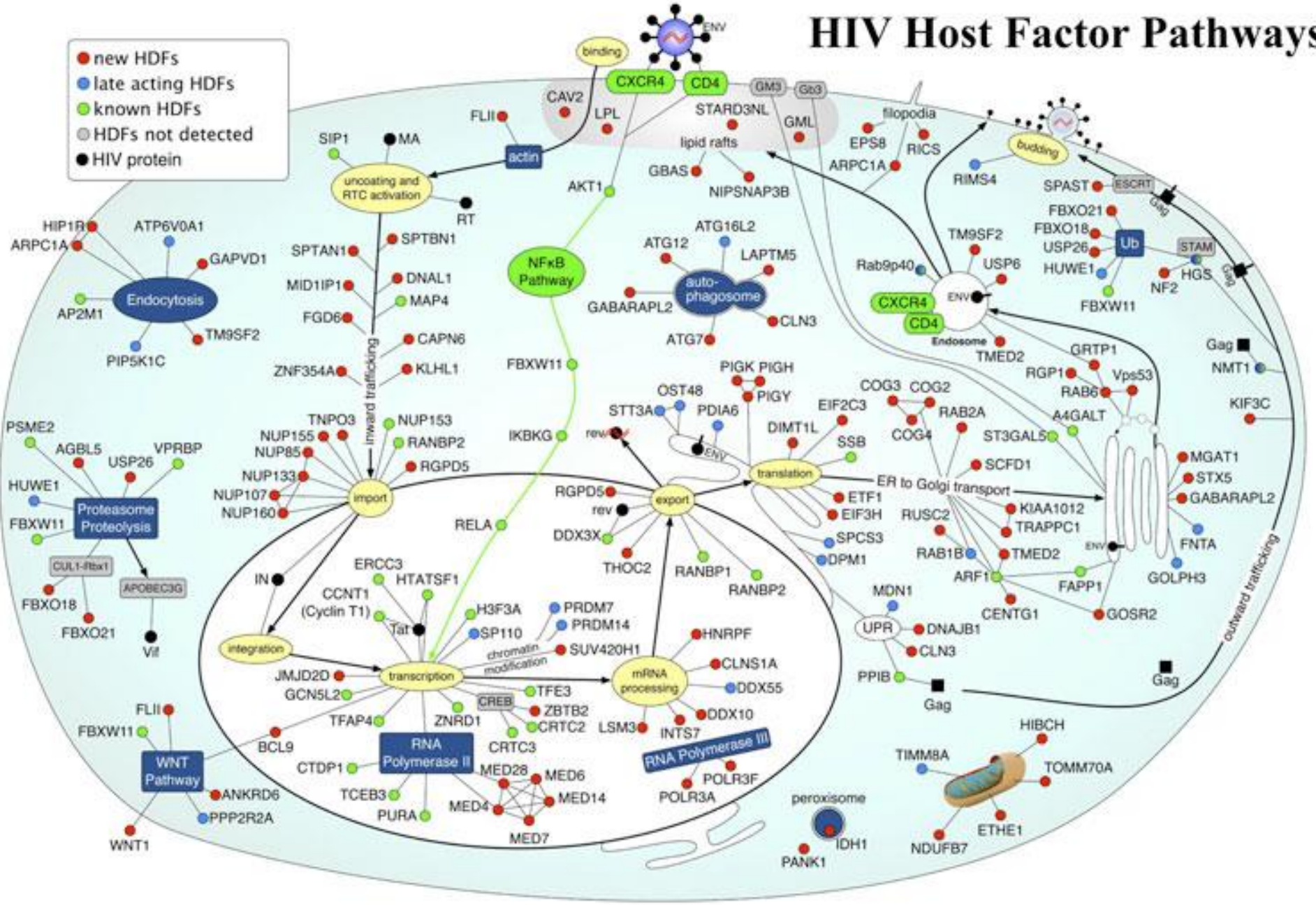
# Pathways of exit from cells for different viruses



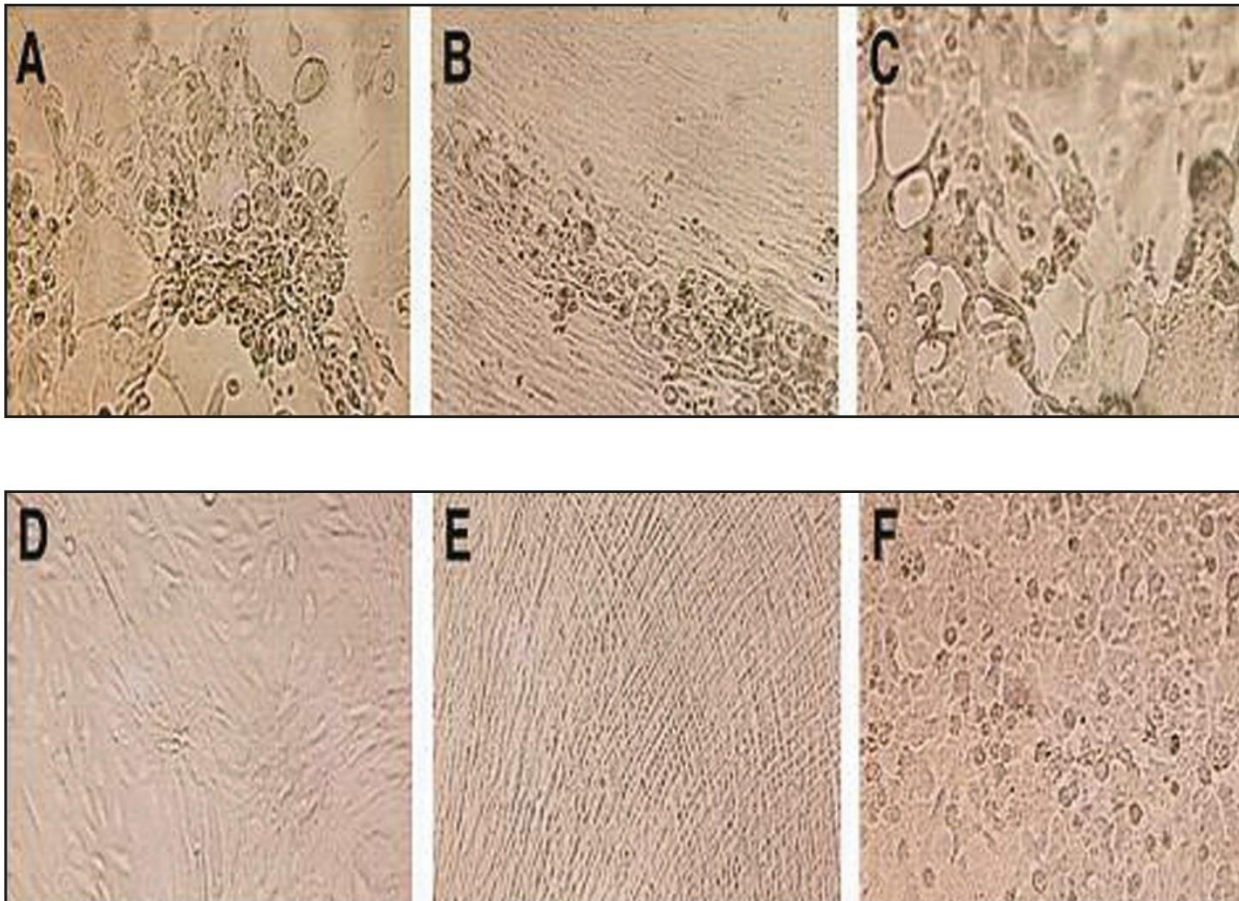


Electron micrograph of enveloped virus particles budding from the infected cell surface

# HIV Host Factor Pathways







**Figura 36.9 Effetto citopatico causato da diversi virus.** **A.** Effetto della replicazione del virus herpes simplex in cellule di rene di scimmia (in **D**, cellule di rene di scimmia non infette); **B.** effetto della replicazione di citomegalovirus in fibroblasti embrionali (in **E**, fibroblasti embrionali umani non infetti); **C.** effetto della replicazione del virus respiratorio sinciziale in una linea cellulare di carcinoma laringeo umano (in **F**, cellule di carcinoma laringeo non infette).