TU 02 : Basic knowledge in Virology



Pr. Audrey Esclatine

audrey.esclatine@universite-paris-saclay.fr

Plan

Definition

• Virus Structure

- Genome
- Capside
- Envelope
- Classification
- Multiplication des virus (within a cell)
 - Initiation
 - Replication/ Transcription
 - maturation

Viral-associated Pathologies

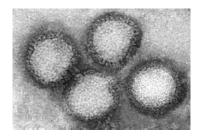
- Respiratory infections
 - Rhino pharyngitis, laryngitis, otitis...
 - Influenza
 - Bronchiolitis in infants
 - COVID19 (SARS-CoV2), SRAS (SARS-CoV1)
- Infections of the gastro intestinal tract
 - Viral gastroenteritis, rotavirus
- Poliomyelitis
- Viral hepatitis
 - Hepatitis A
 - Hepatitis B
 - Hepatitis C
 - Hepatitis D
 - Hepatitis E
- AIDS
 - HIV
 - Opportunistic infections

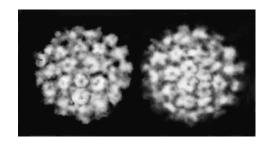
- Arboviruses (transmitted by arthropods)
 - ▲ Dengue
 - ▲ Yellow fever
 - ▲ Chikungunya, Zika...
- ▲ Hemorrhagic fevers
 - ▲ Ebola, Marburg
- ▲ Eruptive diseases
 - ▲ Measles
 - 🔺 Rubella
 - ▲ Mumps
 - ▲ Herpes, chicken pox
 - ▲ Papillomavirus...
- ▲ Zoonosis....
 - ▲ Rabies
 - 🔺 Мрох

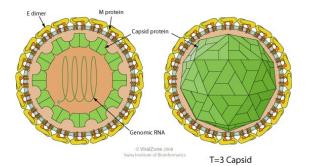
A virus A virosis

Definition of a virus

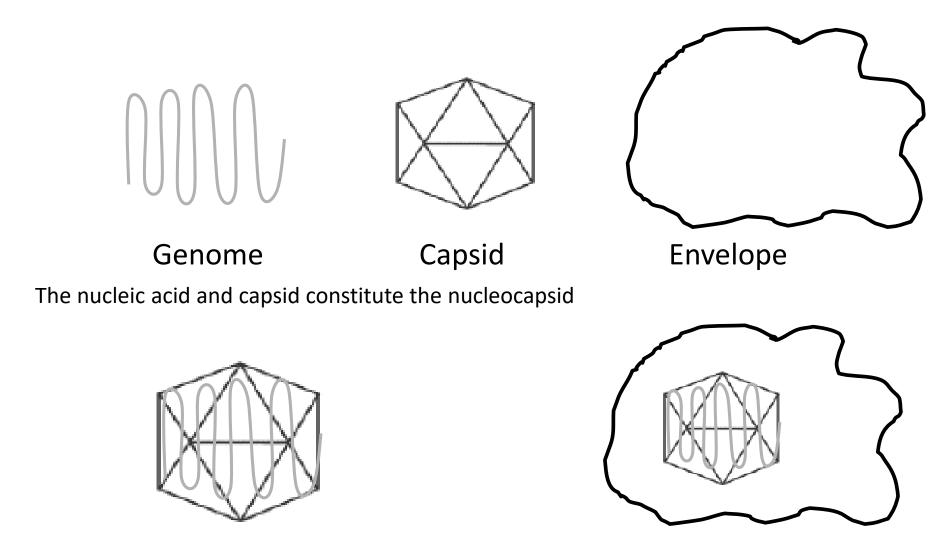
- 4 essential criterions :
- 1) Only one type of nucleic acid
 - either DNA or RNA viral genome, carries all the genetic information of the virus
- 2) Replication form genetic material: a virus cannot growth or divide
- 3) Obligate intracellular Parasitism
 - . Lack of enzymatic or energetic systems
 - hijacking of the cellular machinery
 - . Isolation of viruses by cell culture
 - . Difficult to find an antiviral chemotherapy
- 4) A define structure
 - . different from the one of cells or bacteria
 - . a characteristic type of symmetry
 - virion : mature, infectious, extracellular viral particle
 - virus : the infectious agent at every step of the viral cycle (intra and extra cellular)







Virion Structure



Naked Virus

Enveloped Virus

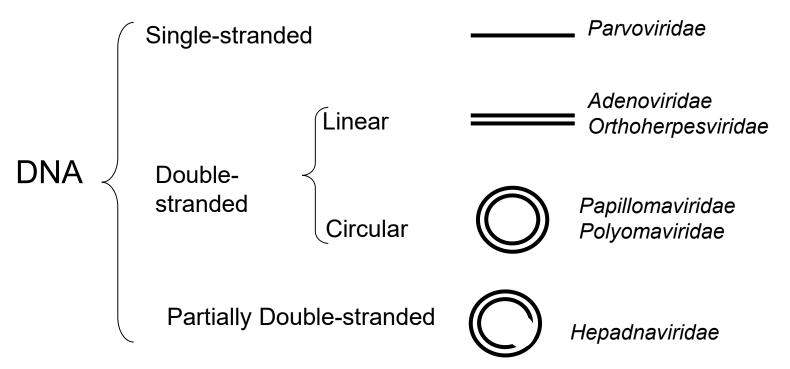
DNA genomes

▲ Variable genome size (in base or base pair)

- ▲ 3,2 kbp for Hepatitis B Virus (HBV)
- ▲ 235 kbp for human cytomegalovirus
- ▲ 1-5 Mb for bacteria and 3400 Mb for human genome
- Most often double-stranded and linear DNA



Classical three-dimensional double helix structure

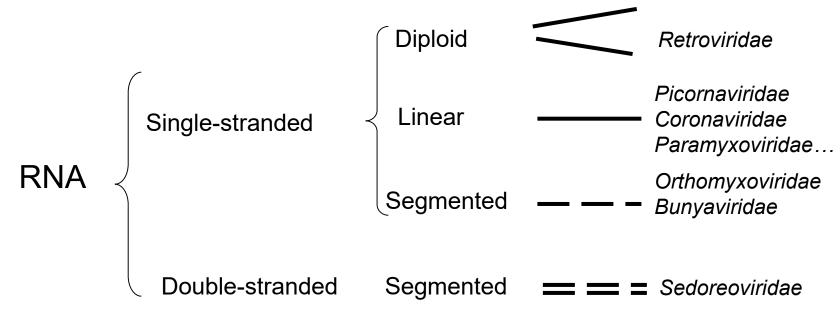


RNA genomes

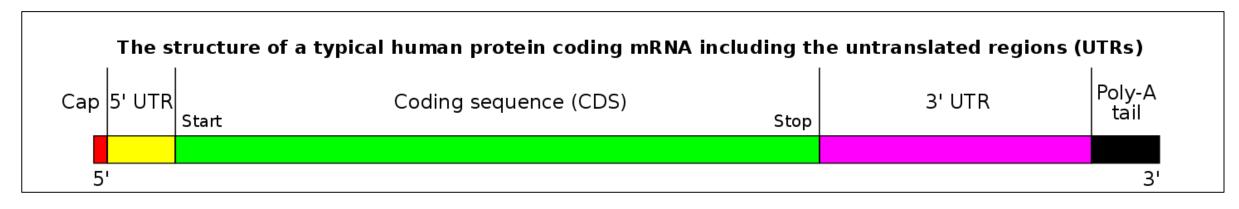
▲ Size (in bases)

From 7000 nucleotides for the smallest (enterovirus) to 30 kb for the biggest (Coronaviridae)

- A Needs a viral RNA dependent RNA polymerase to replicate
- Most often single-stranded and linear RNA
- Some genomes are double-stranded (Sedoreoviridae)
- Some genomes (segmented genomes) are organized into distinct subgenomic fragments (10 to 11 segments for Sedoreoviridae and 7 to 8 segments for influenza viruses)



RNA genomes

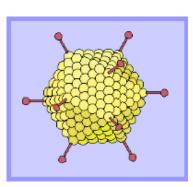


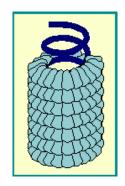
- Positive-strand RNA genomes can act as messenger RNAs (mRNAs), are able to be immediately translated into proteins
- A Negative-strand RNA genomes have to be transcribed first
- A Negative-strand RNA
 - ▲ Some viral polymerase proteins associated with the genome inside the virion
- Positive-strand or positive-sense RNA => Special characteristics
 - ▲ A cap at the 5' terminal end
 - ▲ A poly-A Sequence at the 3' terminal end
 - IRES internal ribosome entry site...

capsids

Viral capsid

- Three roles :
 - Protect the genome from the environment
 - For naked viruses, involved in the attachment of the virus the host cell
 - For DNA viruses and Retroviruses, transport of the viral genome inside the cell
- Resistant and very stable structure
- Limited viral genome coding capacity
- Capsid is formed by polymerization of one or a small number of proteins
 - Principe of self-assembly
 - Two main shapes
 - Tubular capsid with helical symmetry
 - Icosahedral capsid with cubic symmetry
 - Nature of the capsid constitutes a criterion for the virus classification





Helical capsids structure

- Animal viruses with helical capsid are always enveloped
- A flexible nucleocapsid folded inside the viral envelope
- Most of them are negative-sense RNA virus

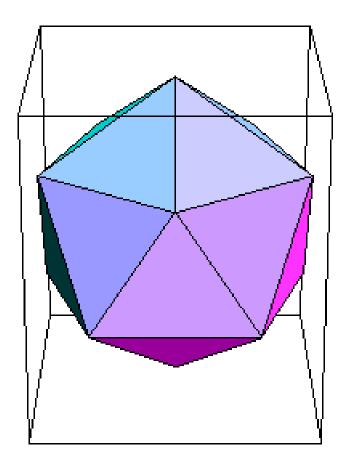
Some viral proteins which function as a RNA polymerase have to be inside the capsid

+RNA viruses
Coronavirus
-RNA viruses
Rabies Virus
Mumps Virus
Influenza Virus

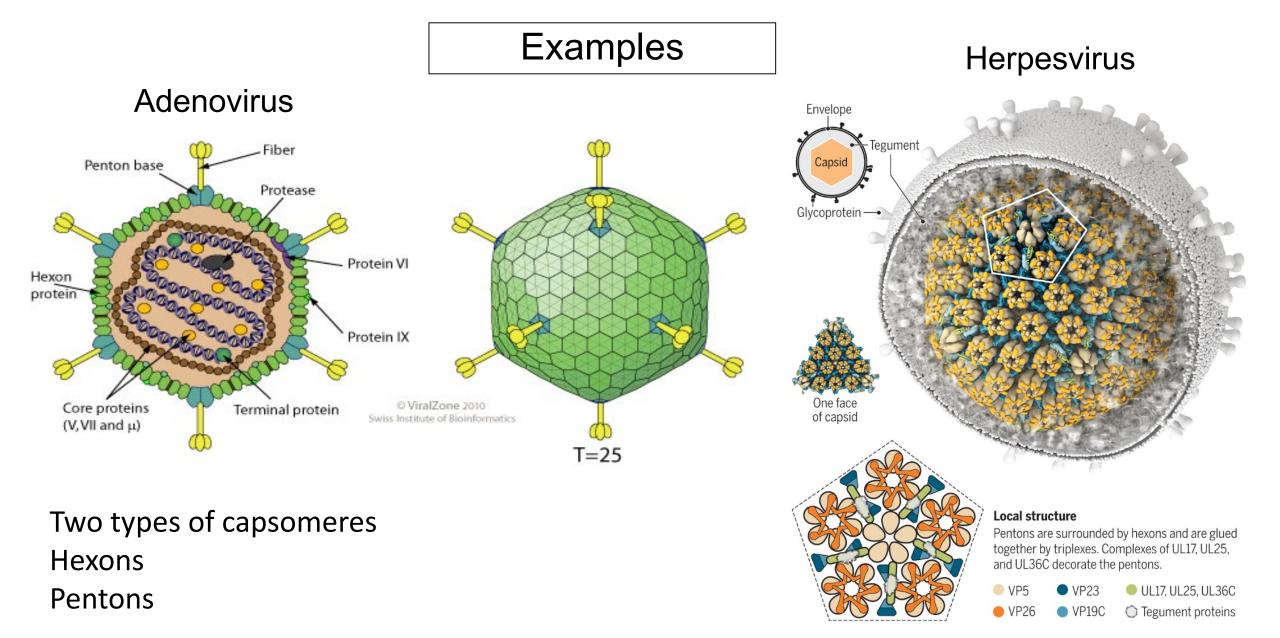
Icosahedral capsids with cubic symmetry

An icosahedron is a geometric shape with 20 sides, each composed of an equilateral triangle, 12 vertices and 30 edges

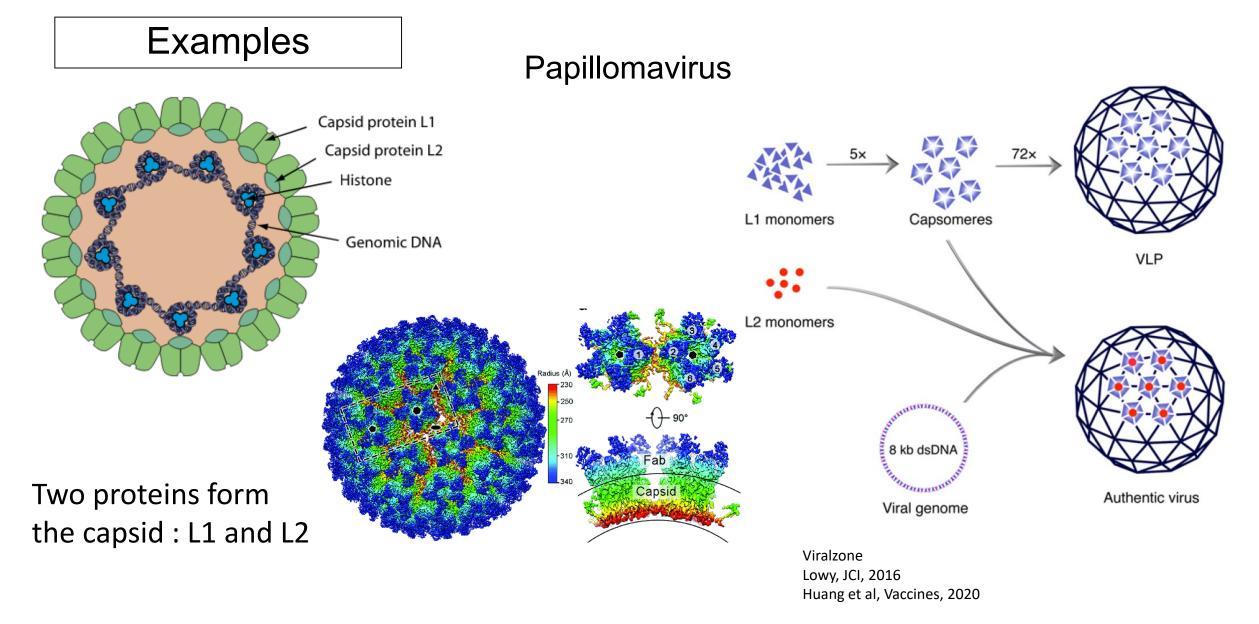
- Capsomeres = morphological units
 - Pentons (containing 5 structural units)
 - Hexons (containing 6 structural units)
- Every icosahedral virus has 12 pentons and a variable number of hexons



Icosahedral capsids with cubic symmetry



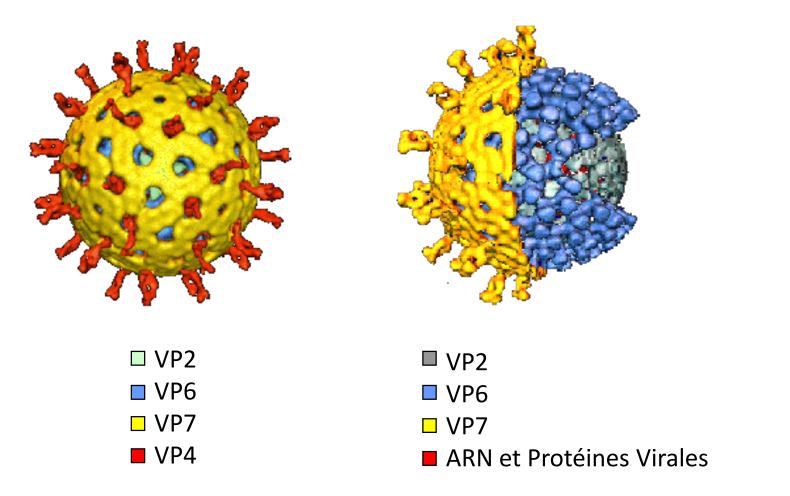
Icosahedral capsids with cubic symmetry



Sedoreoviridae

Virus with a triple capsid (double icosahedral capsid)

Rotavirus

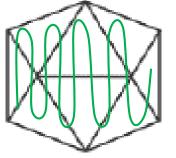


From Prasad et coll. 1994. Curr. Top. Microbiol. Immunol. 185: 9-29

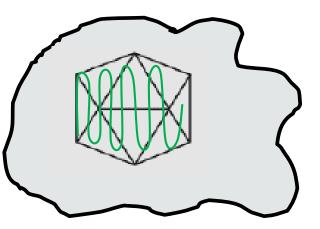
The viral envelope

- Some but not all viruses have an envelope surrounding the capsid
- All helical animal viruses are enveloped
- Complex composition with lipids, glycoproteins
 - Due to their dual viral and cellular origin
- Very sensitive to physico-chemical actions, thermo-sensitive
- More susceptible to environmental conditions
- Consequences for epidemiology and diagnosis

Naked virus



Enveloped virus

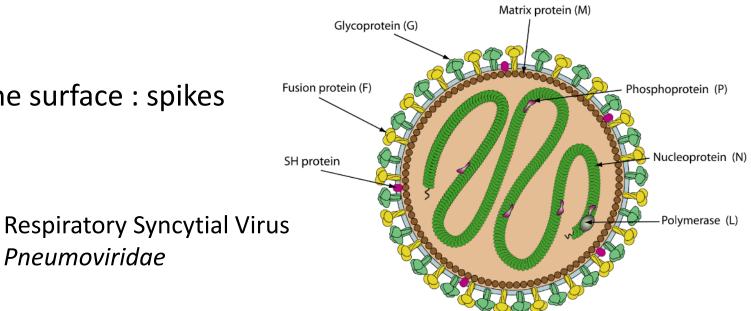


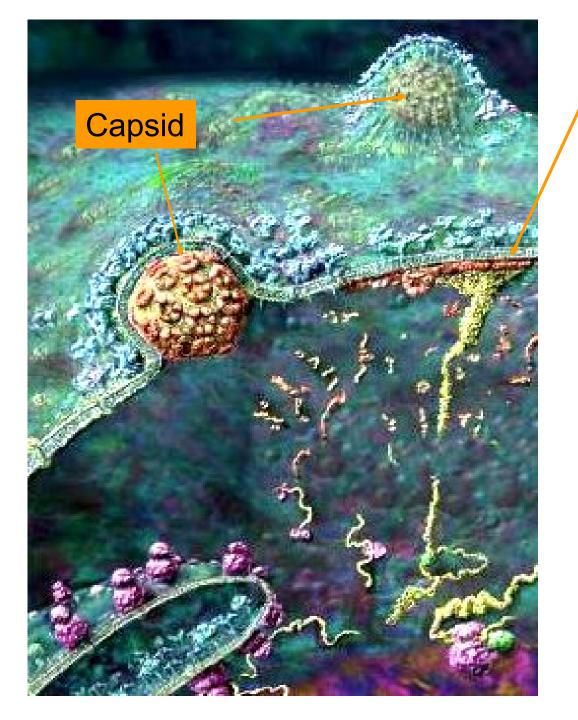
The viral envelope

- Enveloped viruses acquire their envelope by budding through a cellular membrane
 - Either through the plasma membrane (influenza virus, HIV, rabies virus)
 - or through intra-cytoplasmic membranes, Golgi apparatus, endoplasmic reticulum, intracellular vesicles...

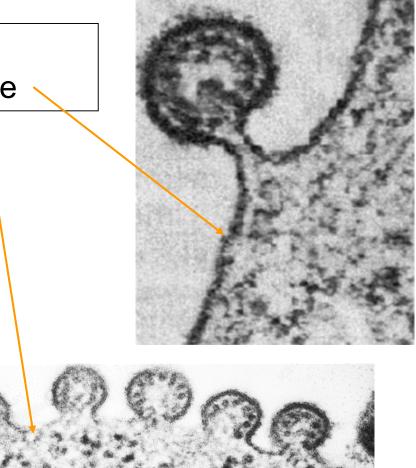
Pneumoviridae

- Transient envelopment of Herpes virus in the nuclear membrane
- Viral glycoproteins
 - Specific biological activities
 - Projections perpendicular to the surface : spikes





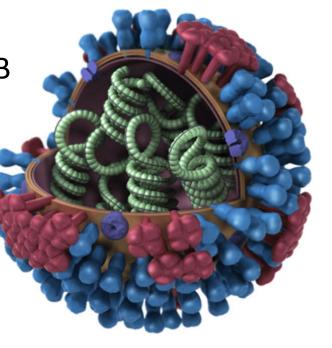
Plasma membrane <



Formation of the viral envelope by budding at the plasma membrane

Influenza virus

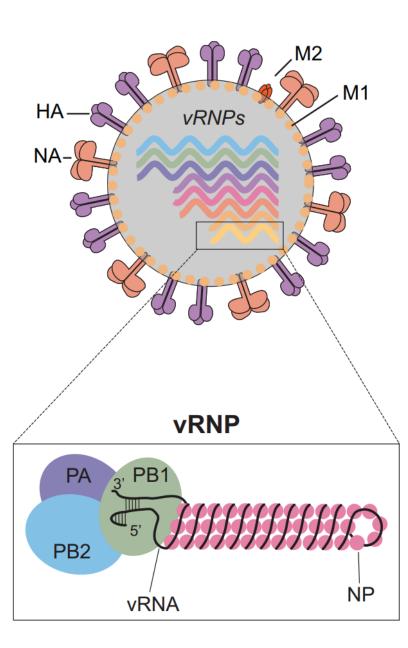
- Envelope + spikes hemagglutinin and neuraminidase
- Under the envelope the matrix (M1)
- Capsid with helical symmetry
- Single-stranded negative-sense linear RNA associated with polymerase PA/PB1/PB2
- Segmented genome
- => 8 fragments for A and B



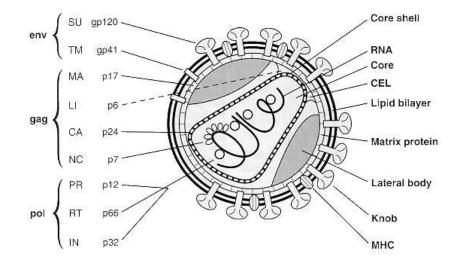






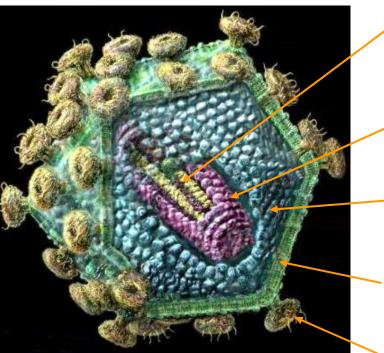






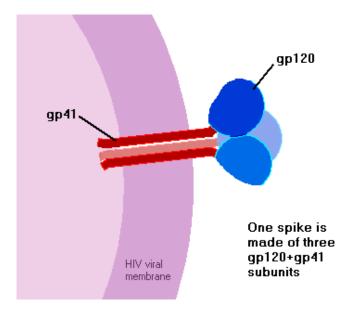
HIV-1

human immunodeficiency virus type 1



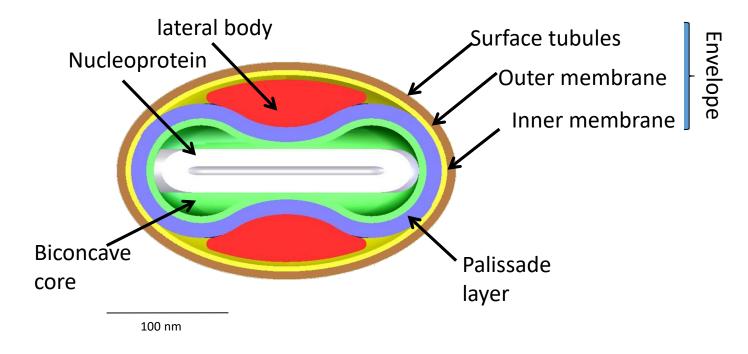
- Diploid RNA genome
- Cone-shaped Capsid
- Matrix
- Envelope

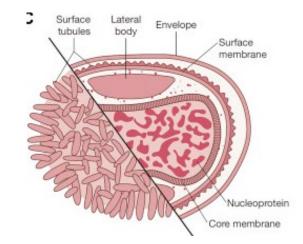
HIV1 spike : a trimer of gp120 and gp41

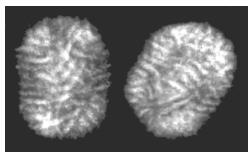


• Envelope (Env) spike

Poxvirus







Molluscum contagiosum



Orthopoxvirus : Mpox

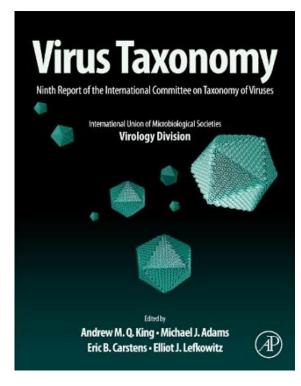
> Big size

- > Brick-shaped
- > Complexe internal structure
- Lateral bodies
- > A complex envelope

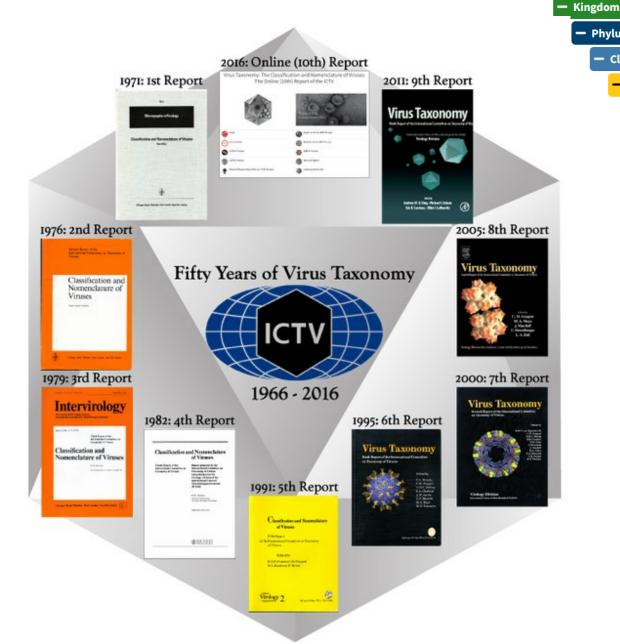
VIRUS CLASSIFICATION

- Classification determined by the International Committee on Taxonomy of Viruses (ICTV)
- Helps to define the evolutionary relationships between viruses and understand the consequences of virus diversity
- website ictvonline.org
- Based on
 - 1. type and organization of the viral genome
 - 2. viral replication strategy
 - 3. virion **structure**

(virion size, capsid shape, envelope or not)



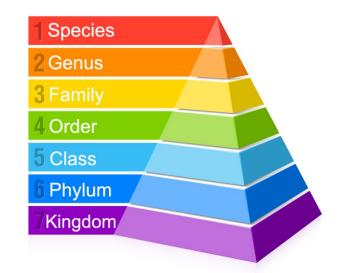
VIRUS CLASSIFICATION



- Realm: Dupl

– Cl

odnaviria		
: Heunggongvirae Realm: Duplodnaviria		
um: Peploviricota Kingdom: Heunggongvirae		
lass: Herviviricetes Phylum: Peploviricota	3 families, 3 subfamilies, 23 genera, 133 species	
- Order: Herpesvirales Class: Herviviricetes		
+ Family: Alloherpesviridae Order: Herpesvirales		
+ Family: Malacoherpesviridae Order: Herpesvirales		
- Family: Orthoherpesviridae Order: Herpesvirales		
— Subfamily: Alphaherpesvirinae Family: Orthoherpesviridae		
+ Genus: Iltovirus Subfamily: Alphaherpesvirinae		
+ Genus: Mardivirus Subfamily: Alphaherpesvirinae		
+ Genus: Scutavirus Subfamily: Alphaherpesvirinae		
+ Genus: Simplexvirus Subfamily: Alphaherpesvirinae		
- Genus: Varicellovirus Subfamily: Alphaherpesvirinae		
Species: Varicellovirus humanalpha3 Genus: Varicellovirus		



Classification

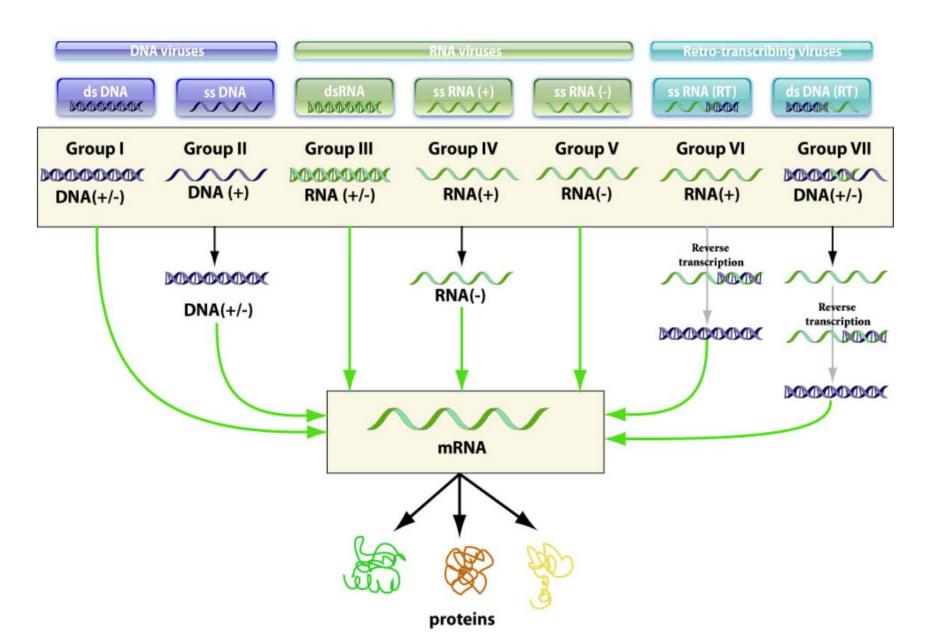
- the same taxonomical hierarchy than other organisms
- Viruses are classified using domain, kingdom, phylum, class, order, family, genus, and species **taxa** (singular: **taxon**)
- Open access up-date available online

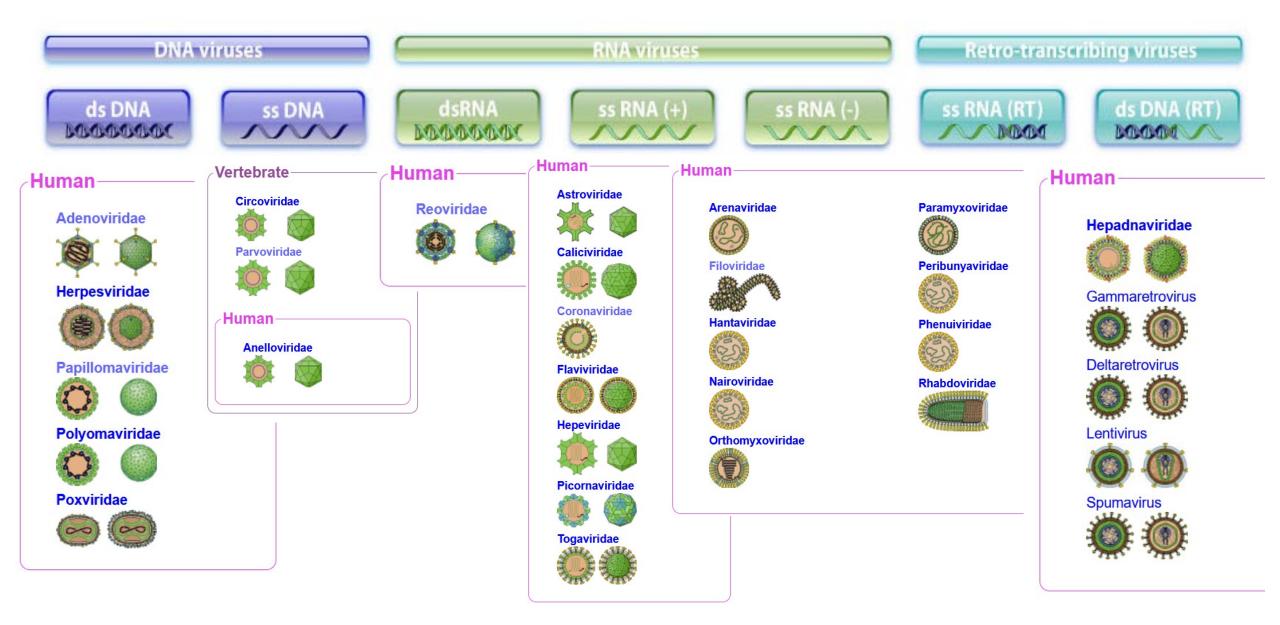
Taxon	Suffix	Examples
order	virales	Mononegavirales
family	viridae	Paramyxoviridae
Sub-family	virinae	Paramyxovirinae
genus	virus	Morbilivirus
species	(virus name)	Measles Virus

constantly evolving

In 2024, 81 orders, 314 families, 200 subfamilies, 3522 genus and 14690 species

Baltimore classification



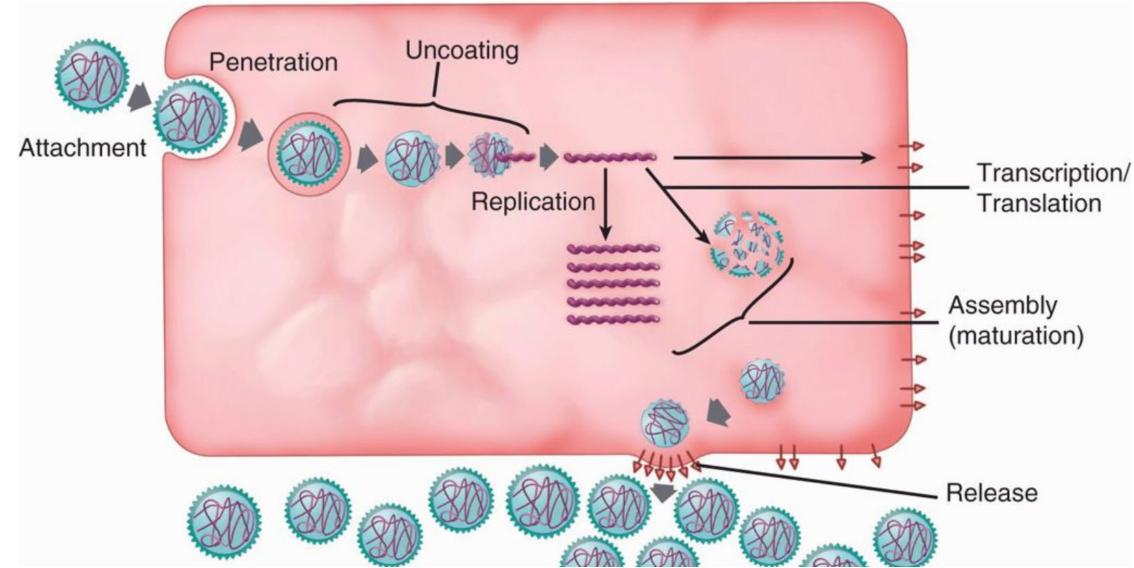


MULTIPLICATION

OF VIRUSES

Different steps of the viral replication cycle

general scheme



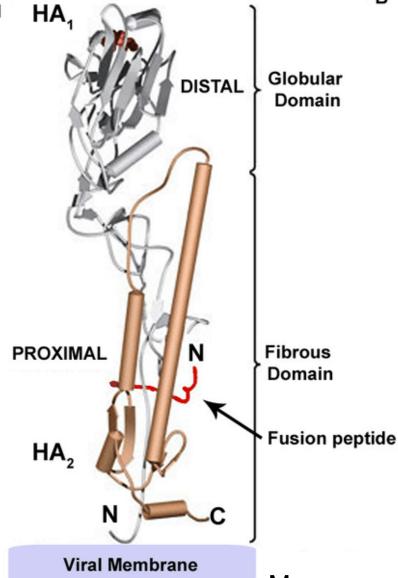
Viral attachment

- First step of the infection, mandatory
- Interactions between viral glycoproteins, proteins and cellular receptors
- Viral proteins are called « ligands » or receptor-binding protein
- Integrity of ligand is essential for attachment of the virus (mutations, neutralizing antibodies)

Viral ligands

- Enveloped viruses : glycoproteins
- Naked viruses : proteins

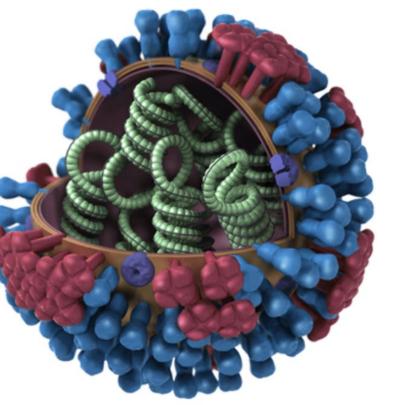
Ligands of enveloped virus



Influenza virus

- Trimer
 composed of 2 subunits
 HA1 and HA2
- >membrane-anchored HA2
- linked to HA1 by a disulfure bond
- HA1 contains the receptor site to sialic acid
- receptor site = conserved
 AA in different HA sub-types

Monomer of hemagglutinin





Hemagglutinin



uraminidase

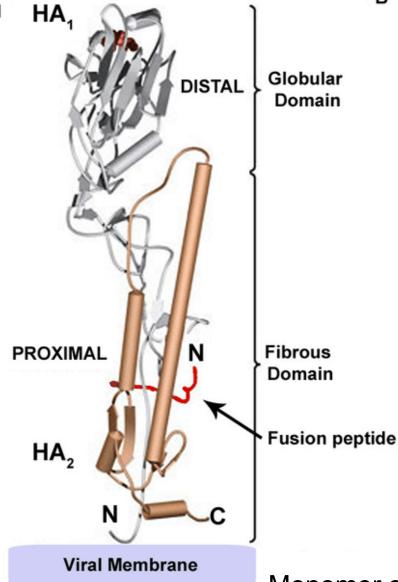


M2 Ion Channel





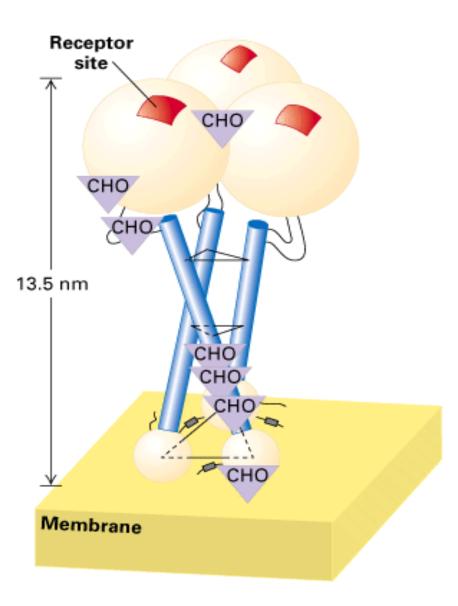
Ligands of enveloped virus



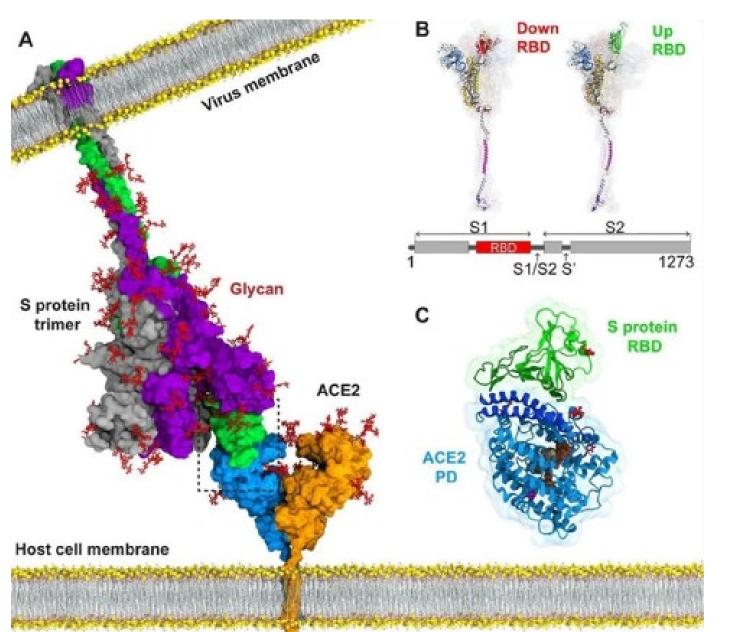
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- HA1 contains the receptor site to sialic acid
- receptor site = conservedAA in different HA sub-types

Monomer of hemagglutinin



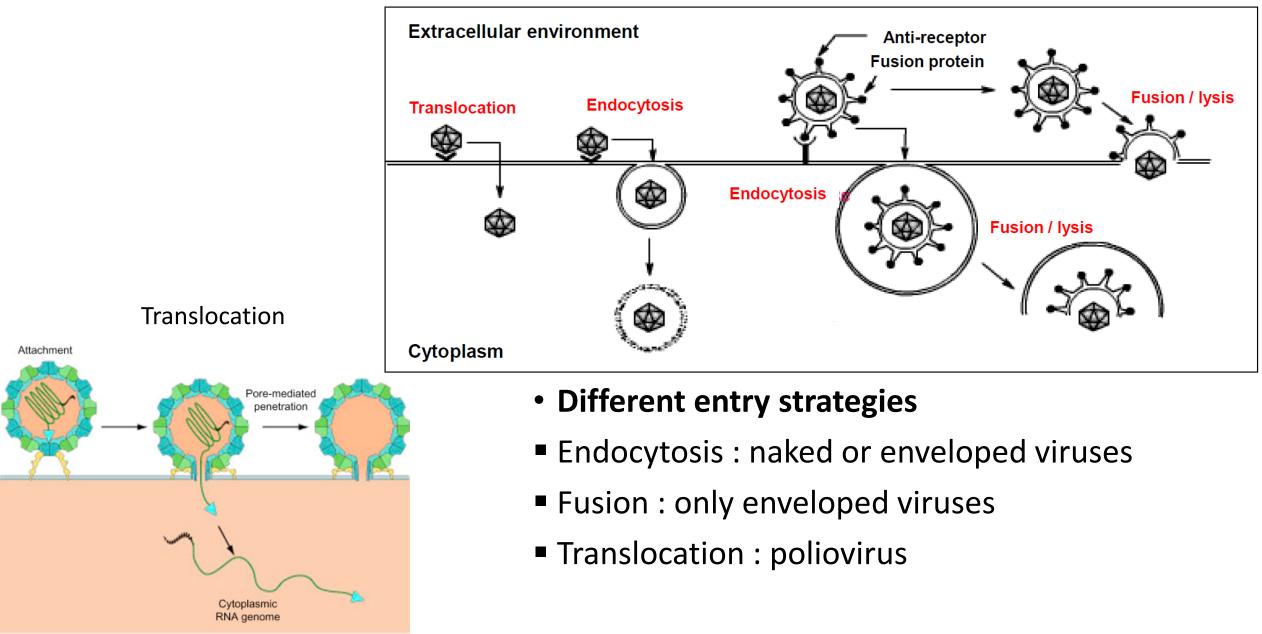
Ligands of enveloped virus



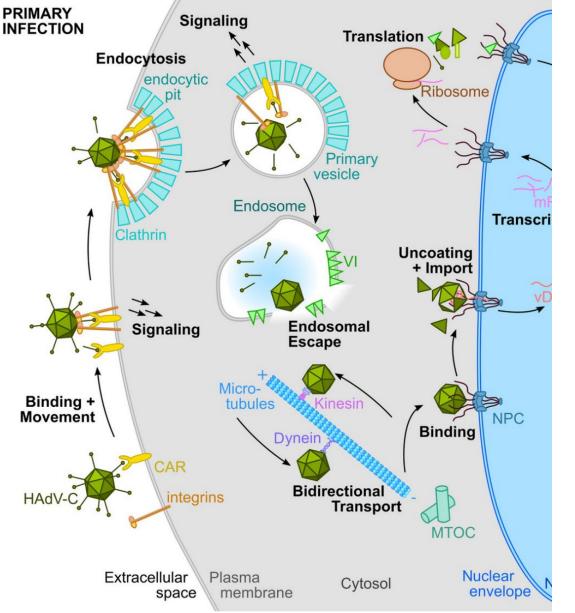
SARS-CoV-2

- Spike (S) Glycoprotein trimer of S (S1 and S2)
- binds on ACE2 (angiotensin Iconverting enzyme-2)
- RBD = receptor binding domain
- Triggers the entry of the virus

Penetration

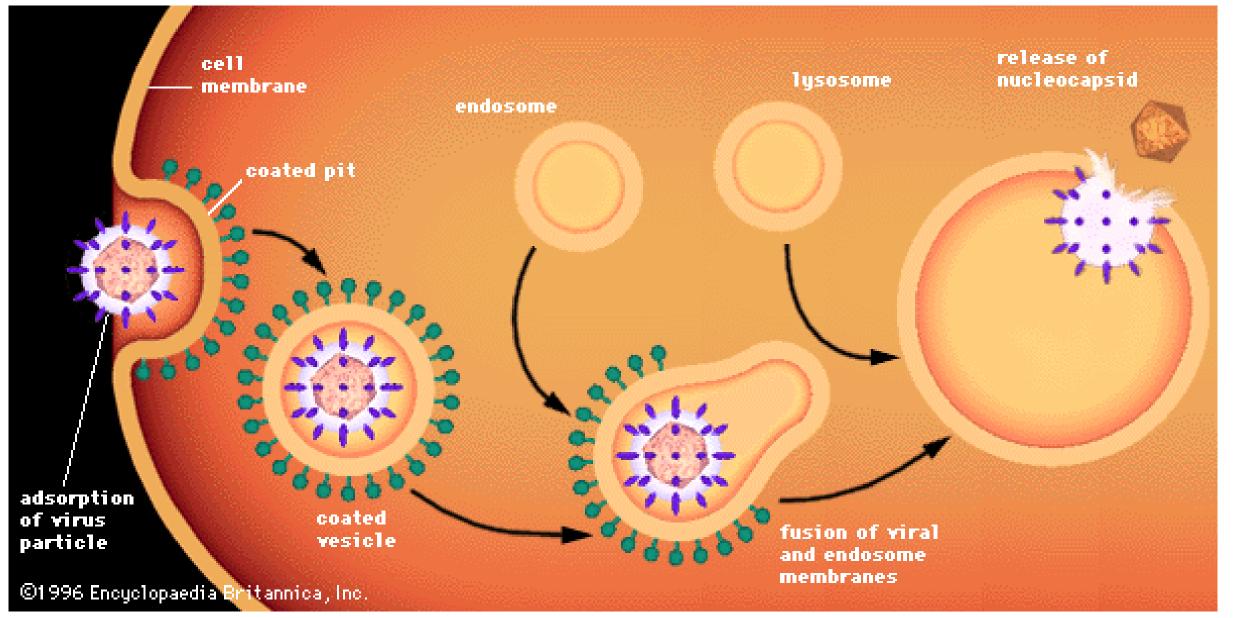


Adenovirus enters into the cell by endocytosis



Virus attaches to **CAR** (Coxsackie adenovirus receptor) via fiber tip (terminal button) Penton base interacts with integrins: triggers entry by endocytosis Acidification of the endosome: partial degradation of the capsid and lysis of the endocytosis vesicle, releasing the capsid into the cytoplasm **Decapsidation:** the capsid is delivered to the nucleus, and the genome penetrates the nuclear pores.

Entry of an enveloped virus by endocytosis

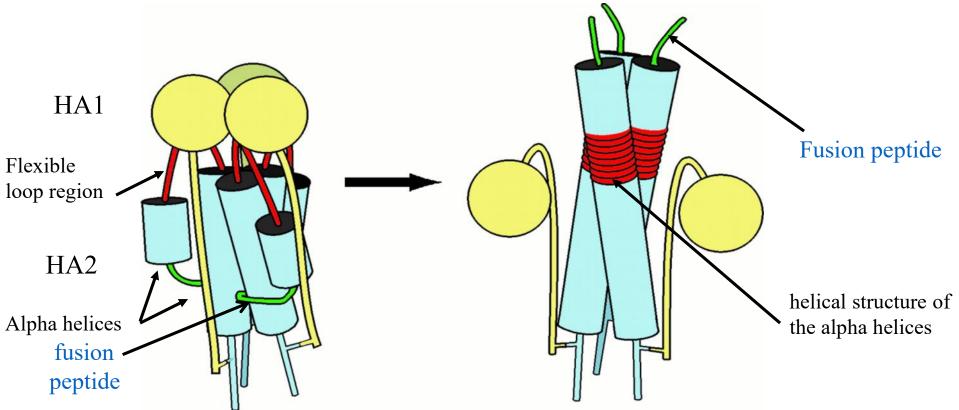


Myxovirus, Rhabdovirus, Bunyavirus, Togavirus, Flavivirus, Coronavirus

Enveloped virus

Influenza virus

- structural alterations at low pH
- hydrophobic fusion peptide released and projected
- Anchoring in the endosomal membrane

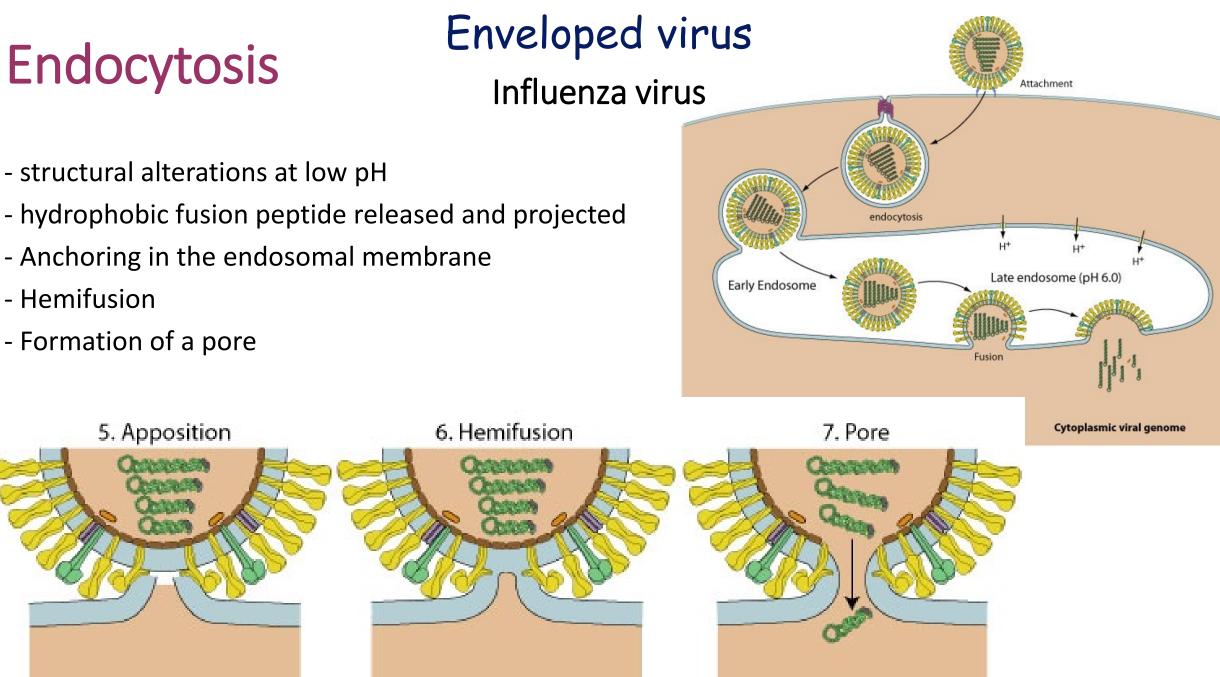


native form at neutral pH post

post fusion structure at acid pH

Endocytosis

- Hemifusion



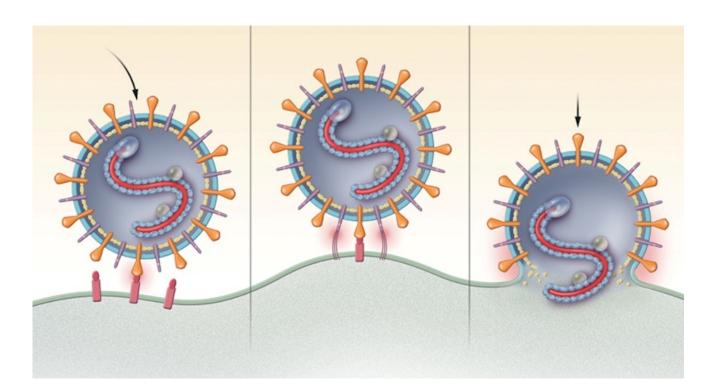
Fusion

Only for enveloped virus

• Fusion of the viral envelope with the plasma membrane

penetration of the nucleocapsid into the cytoplasm

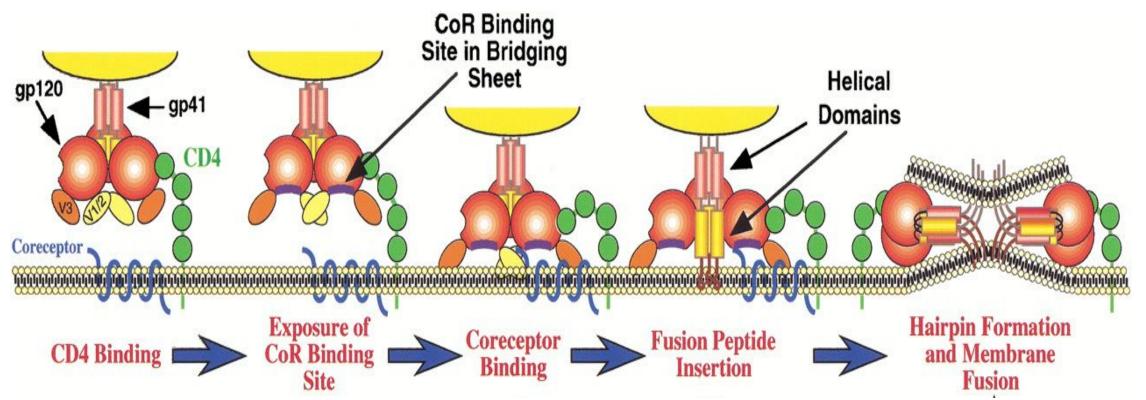
- Fusion peptide is active at physiological pH
- Some examples :
 - HIV
 - Paramyxovirus
 - Herpesvirus



Attachment / fusion

HIV entry

- Interaction between gp120 and CD4, the HIV receptor
- Binding of gp120 to a coreceptor (CCR5 or CXCR4)
- Anchoring of the fusion peptide (gp41) in the membrane
- Hairpin structure of gp41 and membrane fusion, formation of a pore



Expression and replication of the viral genome

- Synthesis of the viral proteins and replication of the viral genome
 - ✓ Transcription (mRNA)
 - ✓ Translation

-non structural proteins (present in the infected cell, absente in viral particles) : enzymes, proteins which regulate the virus and the cellular metabolism

- structural proteins : capsid, envelope, enzymes
- \checkmark Replication of the genome Hepatitis C virus Viral particle ipids Apolipoproteins non structural structural 5 NS2 NS4A NS4B NS5A NS5B Core EI E2 D7 NS3 HCV genome Protéase et son cofacteur **ARN** polymérase

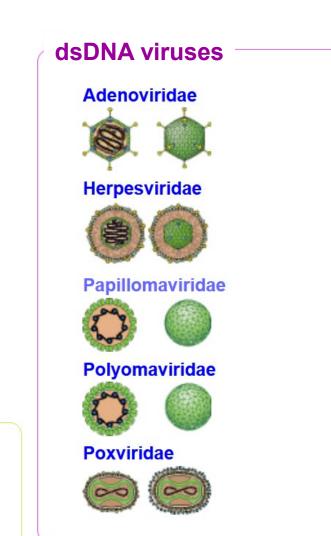
enveloppe glycoprotein

DNA VIRUSES REPLICATION

DNA viruses

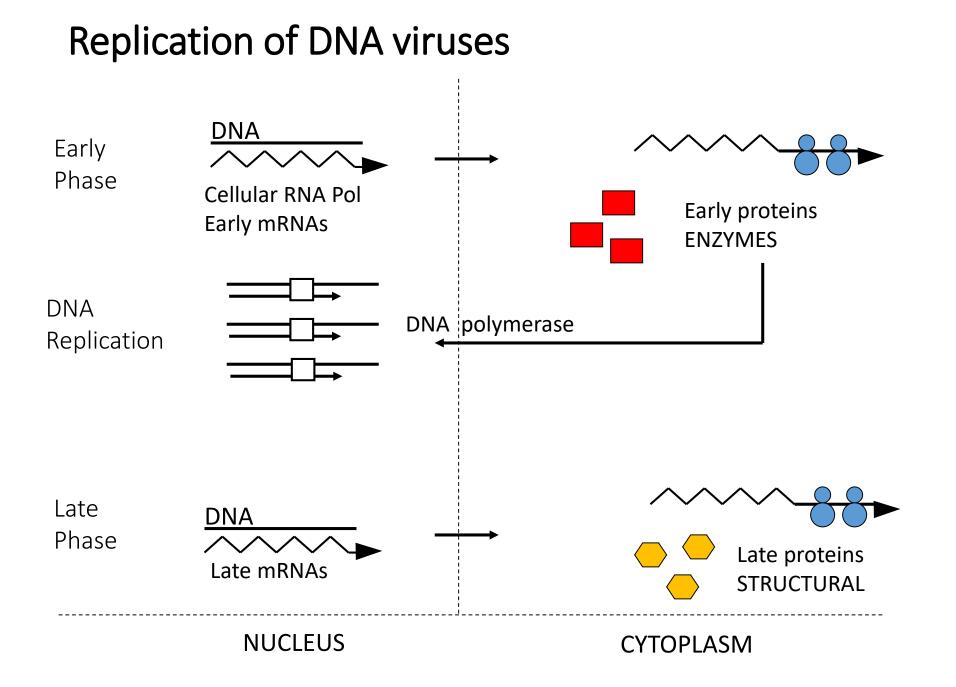
Parvovirus, Adenovirus, Herpesvirus, Papillomavirus...

- Intranuclear replication (except Poxvirus)
- Transcription : cellular RNA polymerase II-DNA dependent
- single-stranded or double-stranded genomes

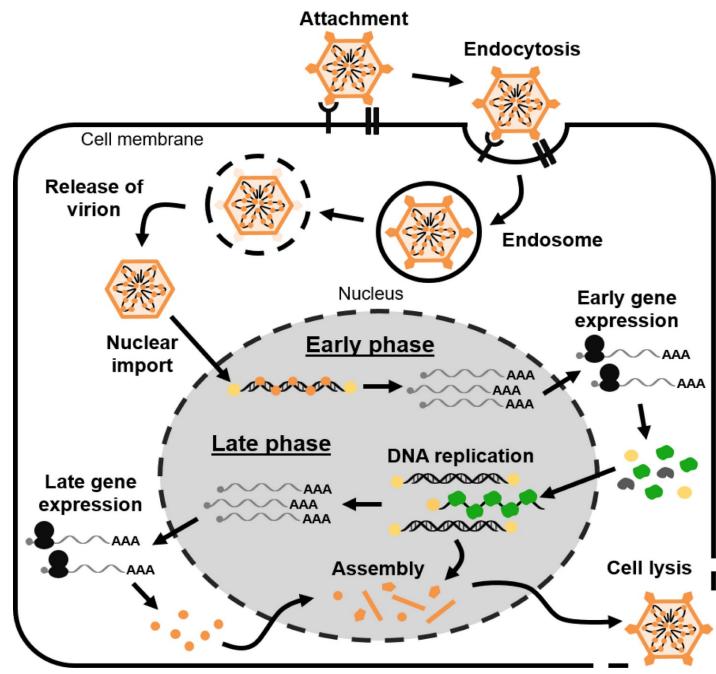


ssDNA viruses





Adenovirus DNA replication



RNA VIRUSES REPLICATION

Replication of RNA viruses

- Intracytoplasmic multiplication (except nuclear phase of Orthomyxovirus)
- need to code a viral RNA-dependent RNA polymerase
- They have a single-stranded genome (except Reovirus)
- Positive or negative sense ssRNA

RNA+

• +ssRNA, same sense as cellular mRNA:

cap in 5', polyA tails in 3'

(Picornavirus, Coronavirus, Togavirus)

• - ssRNA, have to be transcribed in mRNA

(Orthomyxovirus, Paramyxovirus, Rhabdovirus)

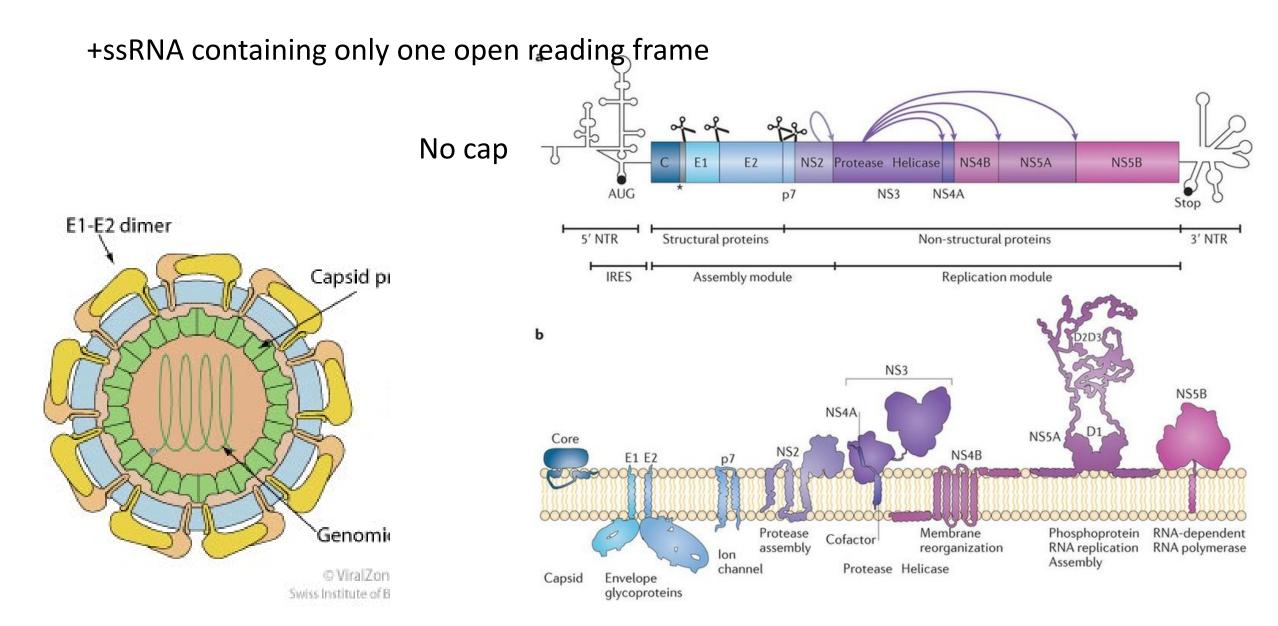
• Retrovirus, constitutive reverse transcriptase

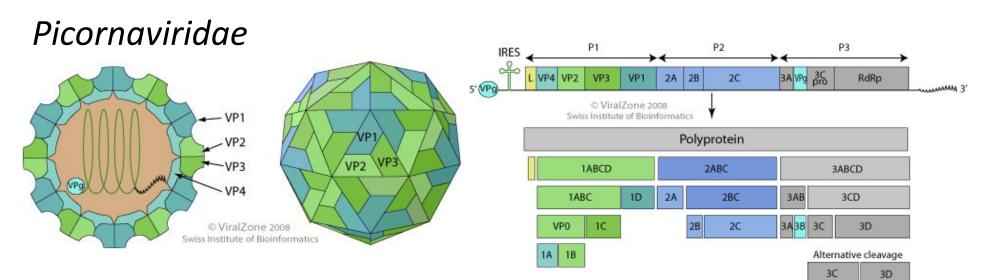
DNA, DNA integration

+ssRNA viruses

- Two roles for <u>genomic RNA</u>:
 - mRNA, directly translated
 - template for the synthesis of a complementary strand -RNA +RNA
- Viral RNA polymerase is synthetized
- Two roles for <u>neosynthesized RNA</u>:
 - mRNA, which will be translated
 - genomic RNA, which will be incorporated in the capsid to form new virions

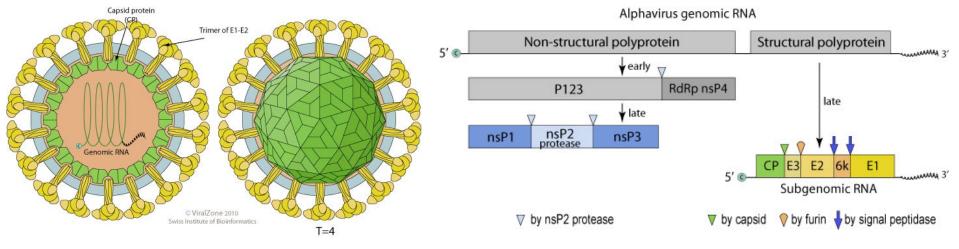
+ssRNA virus: Flaviviridae - Hepatitis C virus





Ex: Poliovirus, Hepatitis A virus

Togaviridae



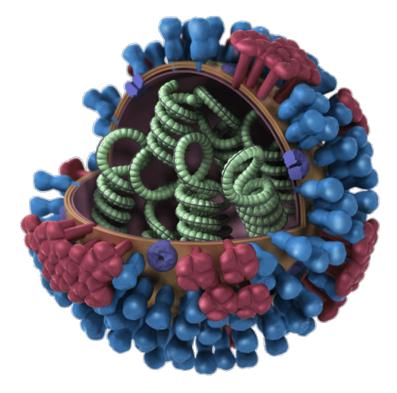
Ex: Rubella virus, Chikungunya virus

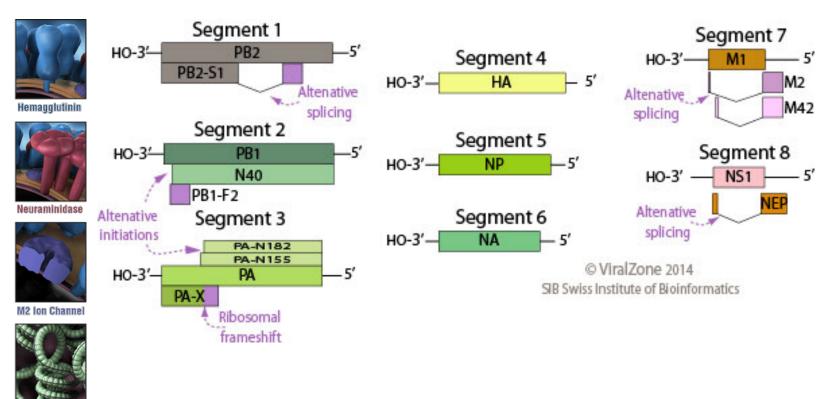
-ssRNA and dsRNA VIRUSES

- <u>Non messenger genomic RNA</u>
 - needs to be transcribed in mRNA
- Constitutive viral RNA polymerase (transcriptase)
- Two roles for <u>neosynthetized RNA</u>:
 - mRNA
 - Template for synthesis of complementary genomic -RNA, incorporated in the capsid

Orthomyxoviridae

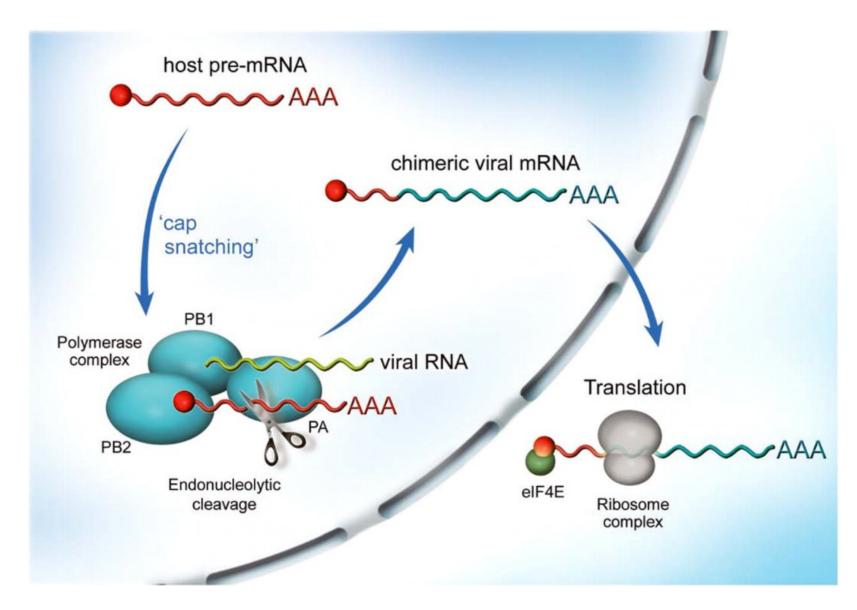
BNP



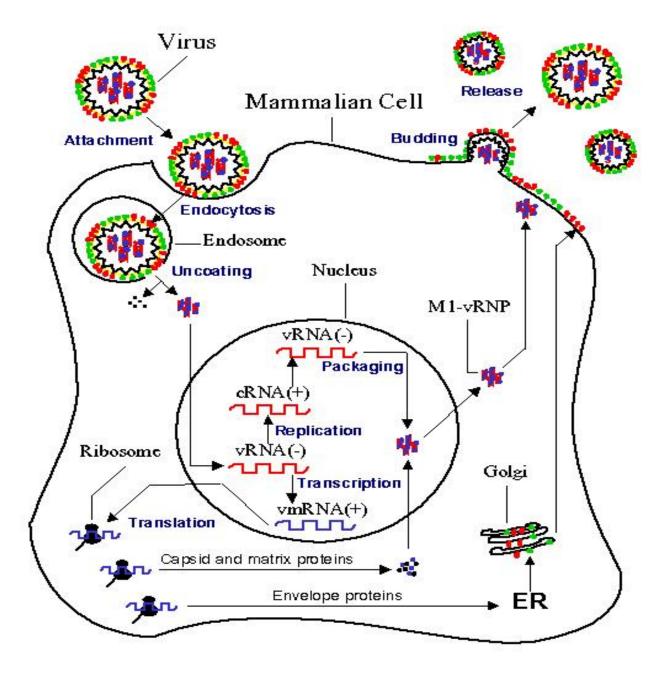


Ex: Influenza Virus

Cap snatching *Orthomyxoviridae* (influenza virus)



Influenza virus Cycle

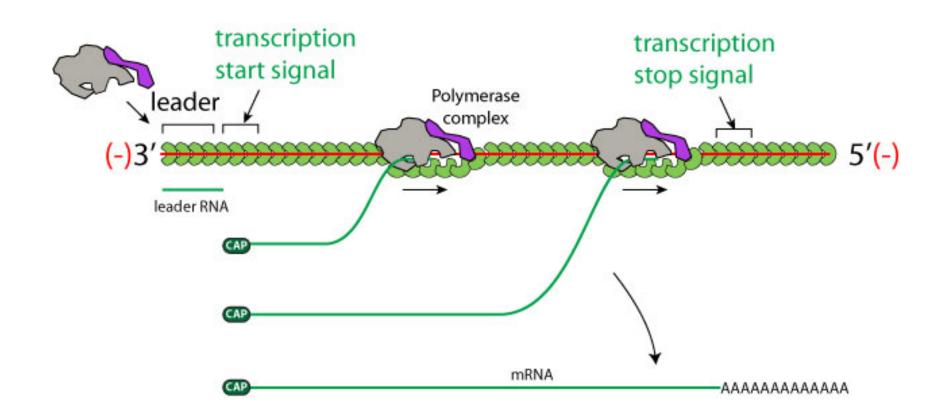


Example of a Negative RNA Virus : Measles virus

Non segmented genome

Filoviridae, Paramyxoviridae, Rhabdoviridae

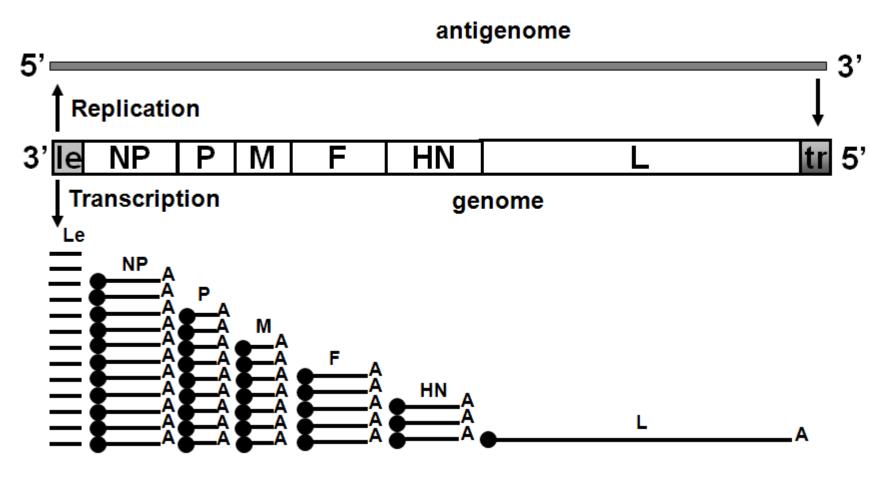
- Transcription and replication by the viral RNA polymerase
- Each gene is framed by a starting sequence and a stop sequence



Example of a Negative RNA Virus : Measles virus

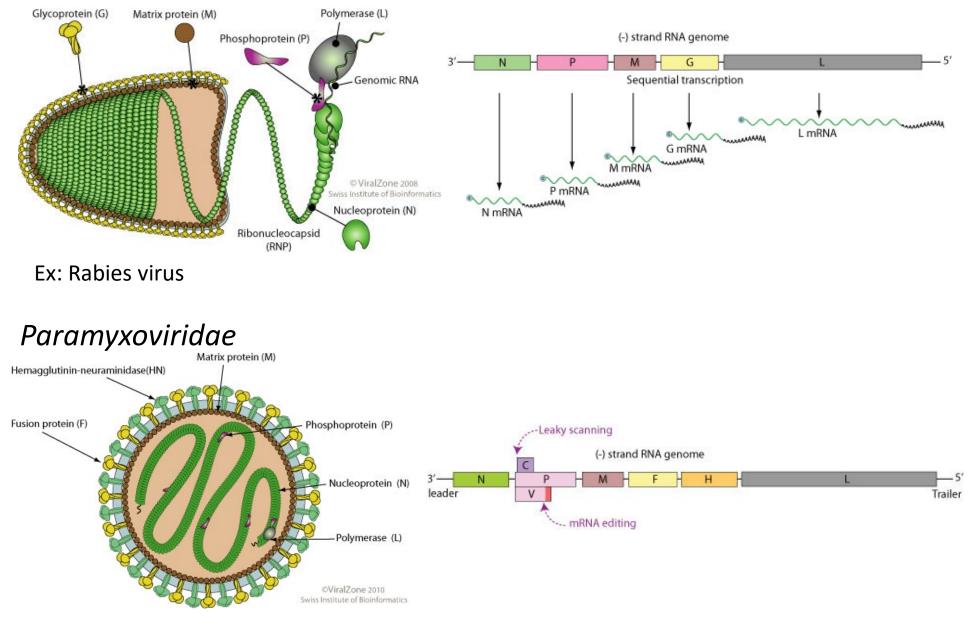
Non segmented genome Filoviridae, Paramyxoviridae, Rha

Filoviridae, Paramyxoviridae, Rhabdoviridae



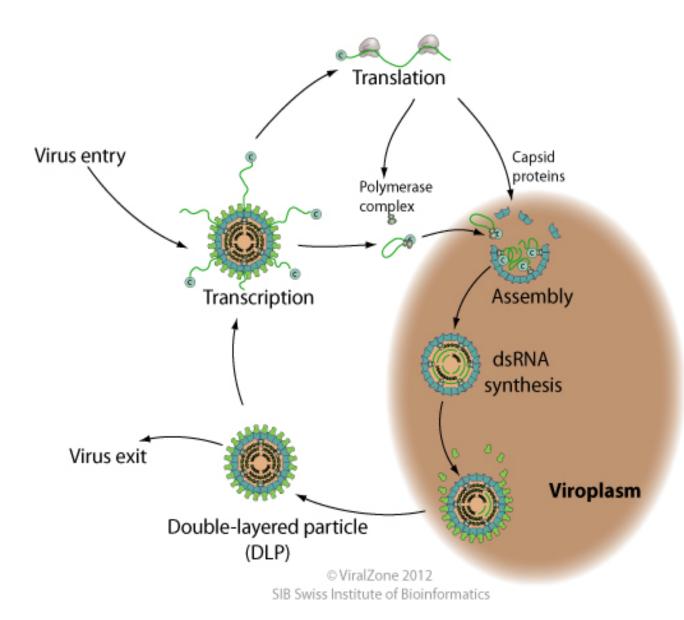
The place of the gene in the genome influences the quantity of transcript formed

Rhabdoviridae



Ex: Measles Virus, Mumps Virus

Reoviridae multiplication



innate immunitydsRNA sensors

Transcription begins in a partially decapsidated particle

Synthesis of the –RNA strand inside the capsid

Neither dsRNA nor -RNA in the cytoplasm

Replication of viruses involving a reverse transcriptase

- Reverse transcriptase (RT) : production of DNA from +RNA
- inverse of cellular transcription
- Retrovirus: genome
 - +RNA \longrightarrow DNA

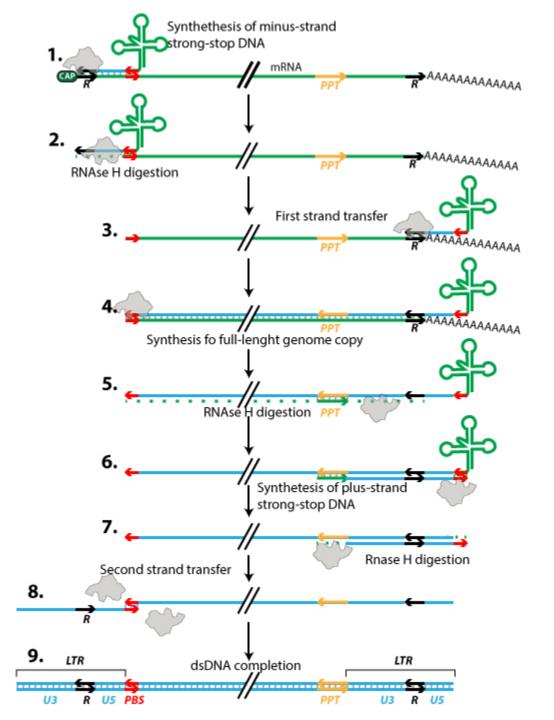
RT

 Hepatitis B virus: replication intermediate +RNA > DNA

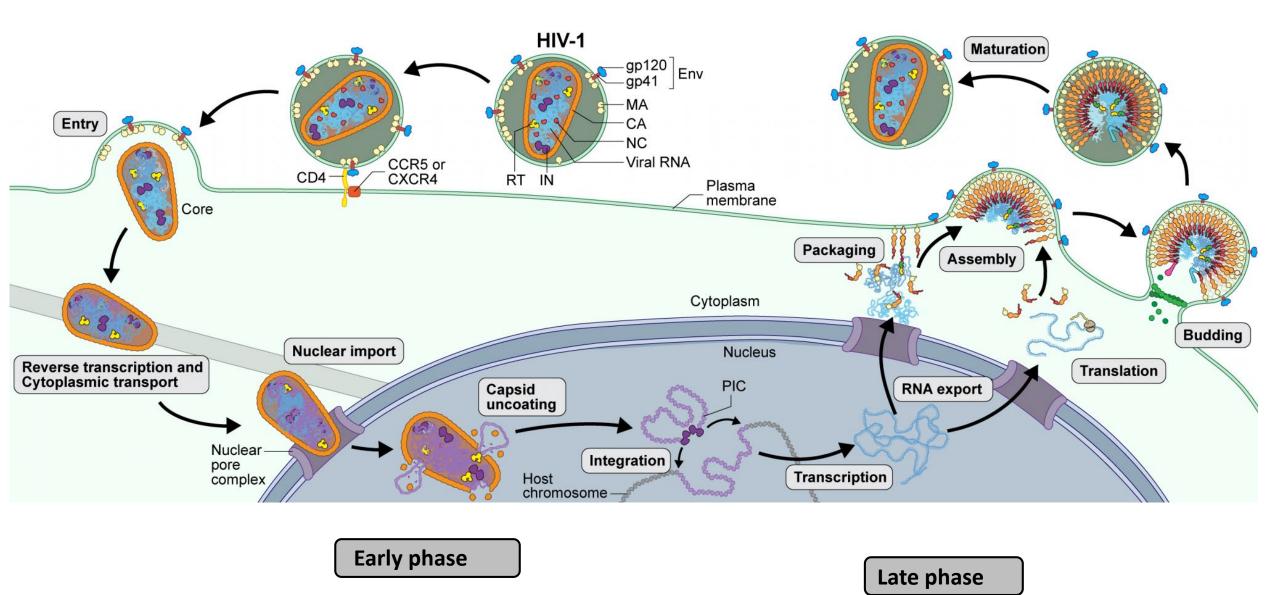
Reverse transcription RNA-DNA

- 3 activities of the RT, carried by distinct sites
- > RNA-dependent DNA polymerase
- > RNAse H
- > DNA-dependent DNA polymerase

ViralZone



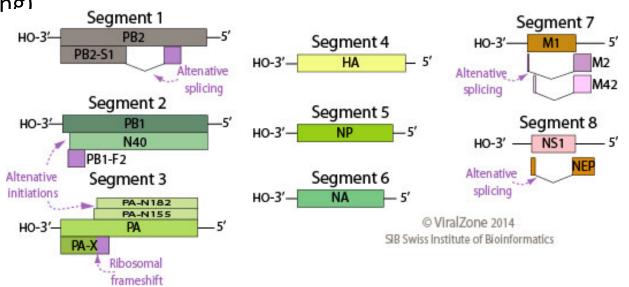
Multiplication cycle of HIV-1



scienceofhiv.org

Strategies of viral protein synthesis in RNA viruses

- Translation in eukaryotic cells
 - 1 mRNA = 1 protein
- Several strategies for RNA viruses
 - One polyprotein, cleaved to give all the necessary proteins (picornavirus, flavivirus)
 - Segmented genome 1 segment = 1 mRNA
 - Initiation of the mRNA synthesis at the level of de internal sites by RNA-dep RNA pol
 - Alternative splicing
 - Translation by itself (ribosomal frameshifting)
- Combination



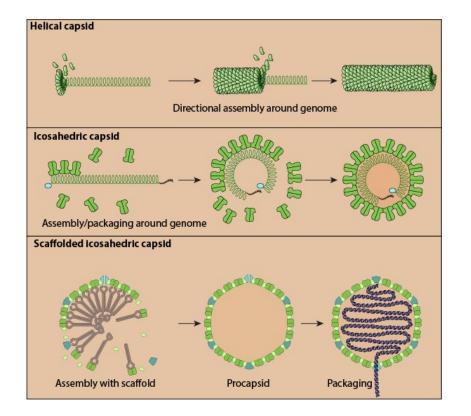
Viral cycle ends by assembly, maturation and exit of the virus

Capsid assembly occurs at different levels:

DNA viruses nucleus (except Poxvirus)

RNA viruses — cytoplasm (except influenza virus)

- Self assembly of neosynthesized proteins and nucleic acids - Procapsid
- Post translational modifications of viral proteins (cleavages, disulfide bridges, glycosylation ...)
- Oligomerization of envelope glycoproteins



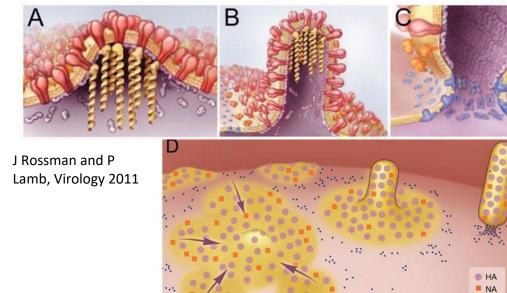
Virus release

- Naked viruses: cell lysis (ex : Adenovirus)
- Enveloped viruses:
 - Budding of the nucleocapsid through cell membranes (nuclear, ER, Golgi, plasma membranes) modified by insertion of viral glycoproteins
 - Release by budding (plasma membrane) or by exocytosis (internal membranes)

: M2

- This maturation step is absolutely required for infectivity
- Viral release possible without cell lysis, contrary to naked viruses

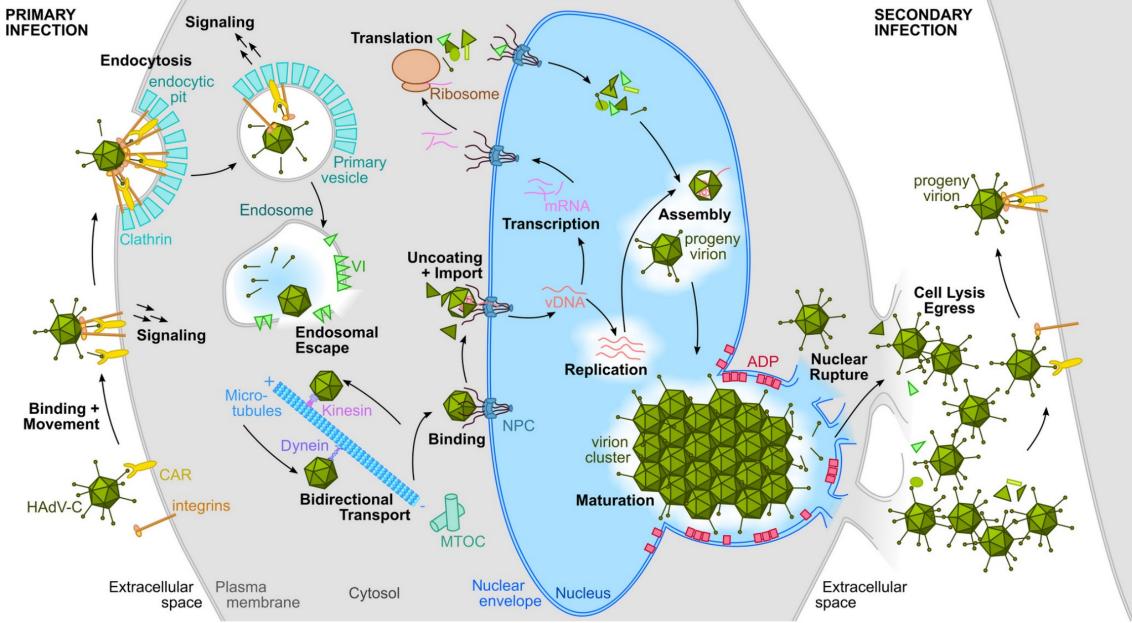
Release by budding (plasma membrane)



Asemblage et empaquetage dans l'appareil de Golgi Sortie du virus par exocytose

Release by exocytosis (Golgi apparatus)

Adenovirus exits the cell by lysis



FEBS Letters, Volume: 594, Issue: 12, Pages: 1861-1878, First published: 30 May 2020, DOI: (10.1002/1873-3468.13848)