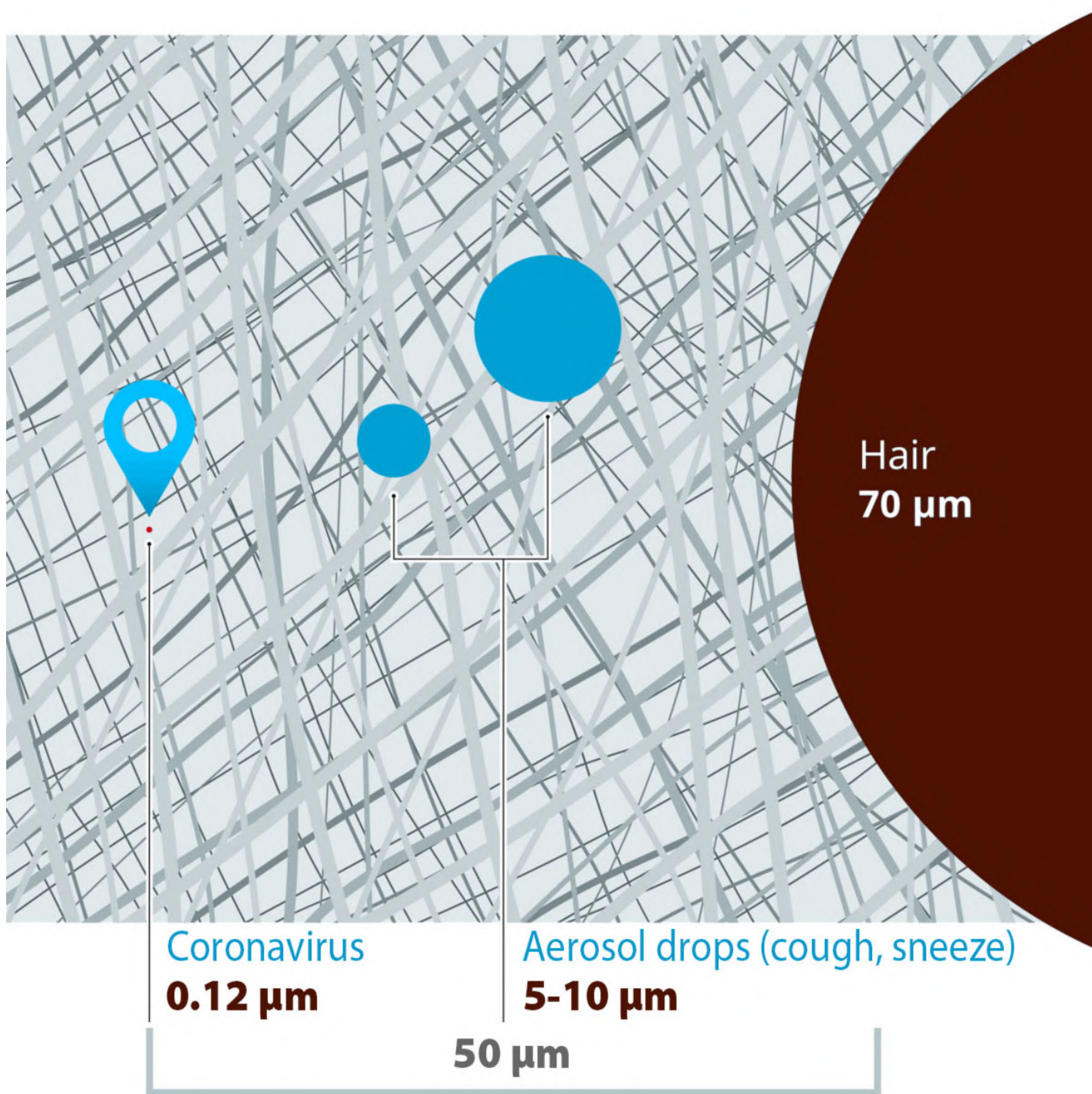
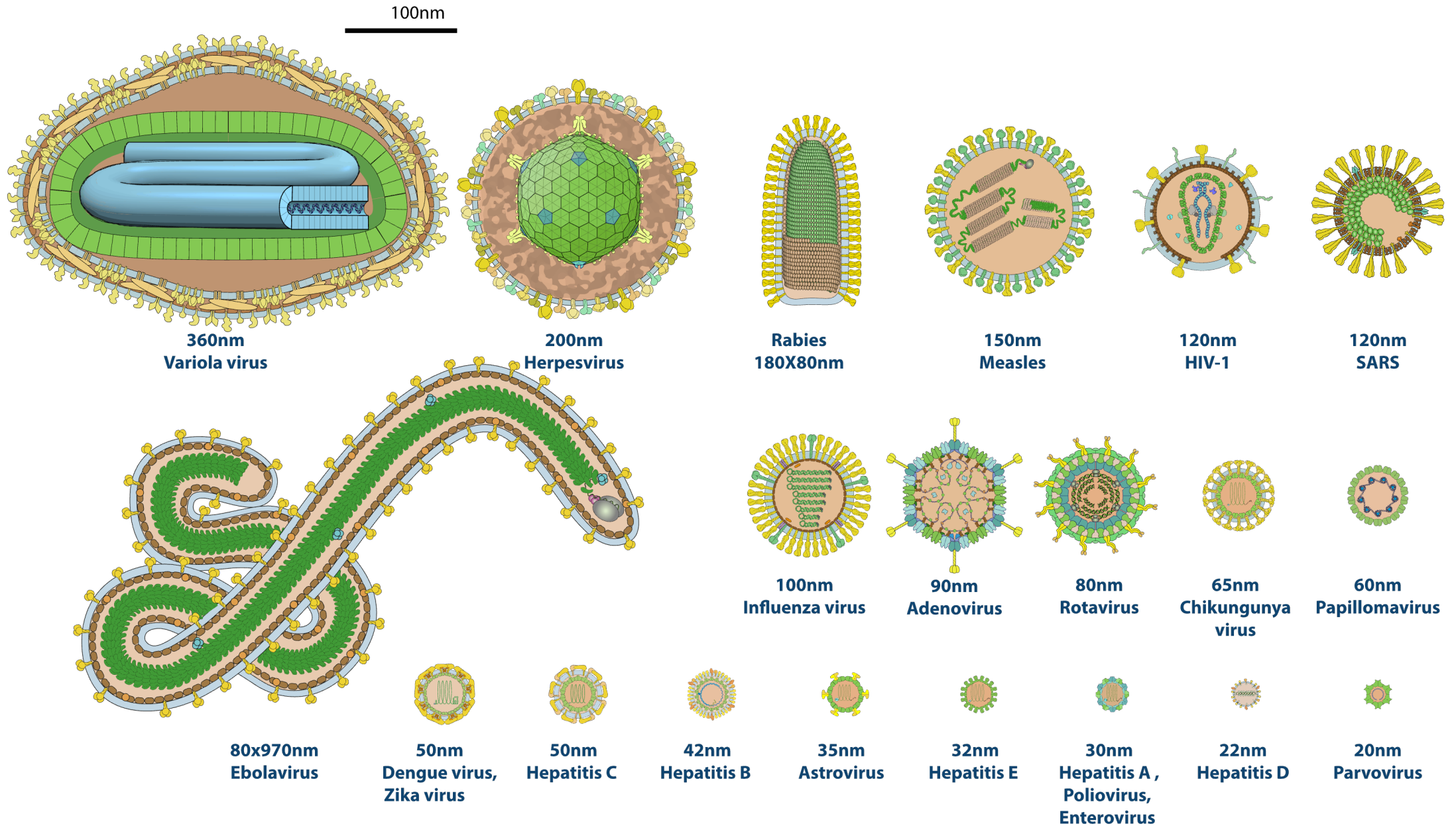


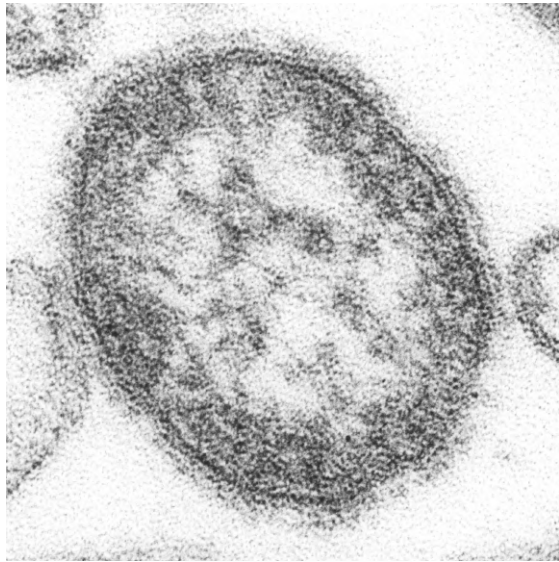
A2.3

Viruses

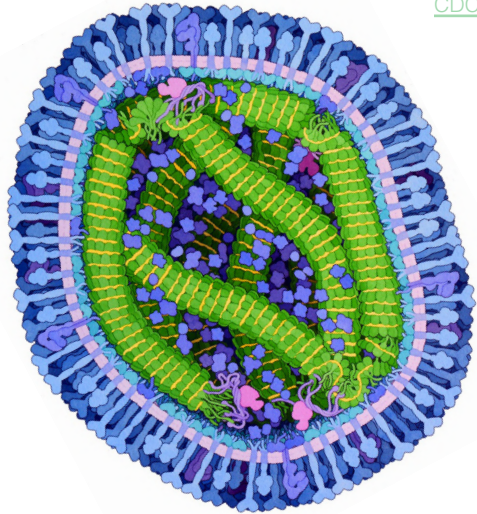
How can viruses exist with so few genes?
In what ways do viruses vary?







CDC



PDB

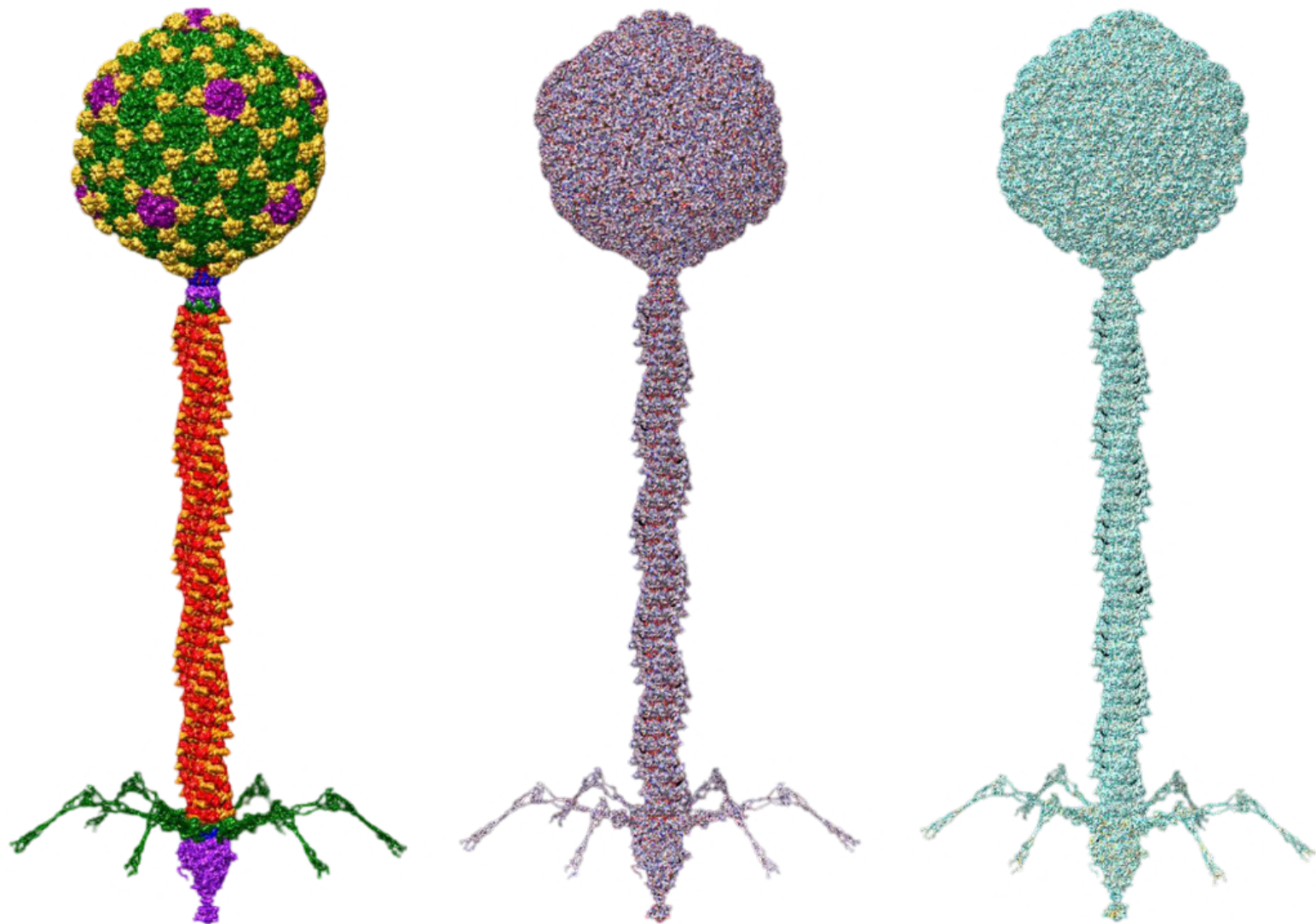
Measles virus (*Morbillivirus hominis*), is a single-stranded, negative-sense, enveloped, non-segmented RNA virus within the family of *Paramyxoviridae*.

Virus Taxa Infecting Vertebrates

DNA	dsDNA		
	 Adenoviridae Asfarviridae	 Alloherpesviridae Herpesviridae	 Iridoviridae Papillomaviridae Polyomaviridae Poxviridae
	ssDNA		dsDNA (RT)
	 Anelloviridae Circoviridae Parvoviridae		 Hepadnaviridae
RNA	ssRNA (-) and (+/-)		
	 Arenaviridae Deltavirus Bunyaviridae Orthomyxoviridae	 Mononegavirales Filoviridae Bornaviridae Paramyxoviridae Rhabdoviridae	
	ssRNA (+)		
	 Arteriviridae Coronaviridae Coronavirinae Torovirinae	 Nidovirales Astroviridae Caliciviridae Flaviviridae	 Hepeviridae Picornaviridae Nodaviridae Togaviridae
	dsRNA		ssRNA (RT)
	 Bimaviridae Picobimaviridae	 Reoviridae Sedoreovirinae Spinareovirinae	 Metaviridae Retroviridae

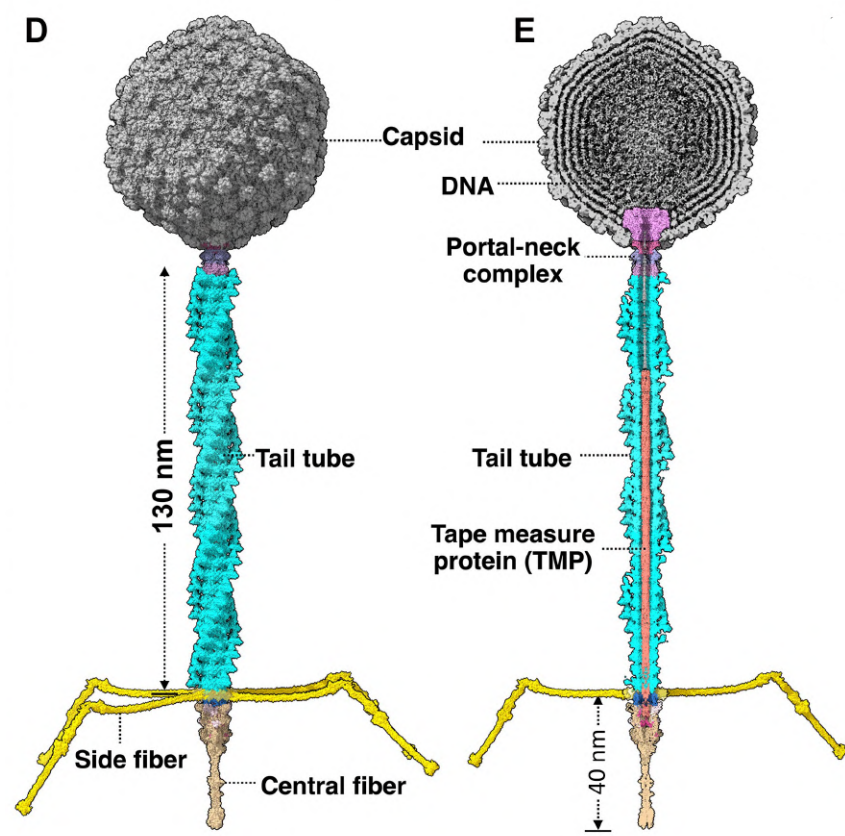
100 nm

bacteriophage lambda



[Padilla-Sanchez 2026](#)

bacteriophage lambda

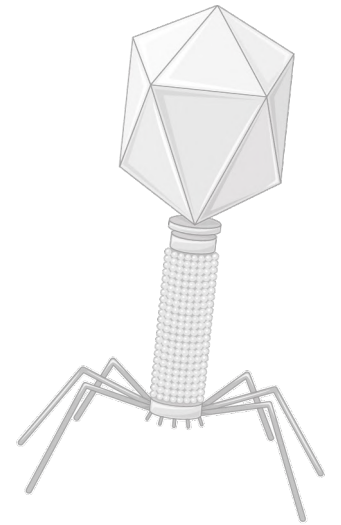
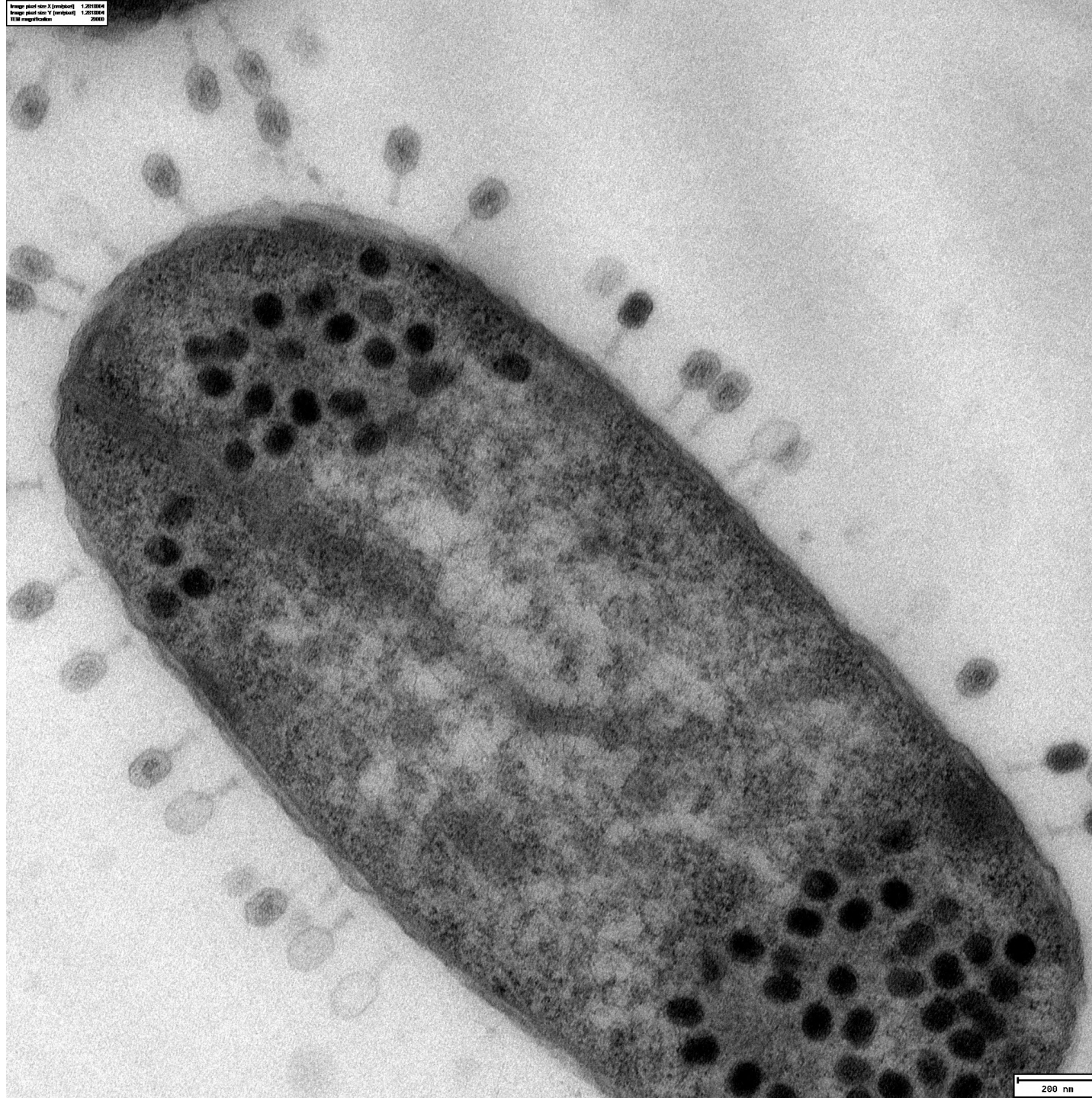


[Structural basis of bacteriophage \$\lambda\$ -lambda infection initiation Sci. Adv. 11.eadw7914\(2025\). DOI:10.1126/sciadv.adw7914CC-BY-NC](#)

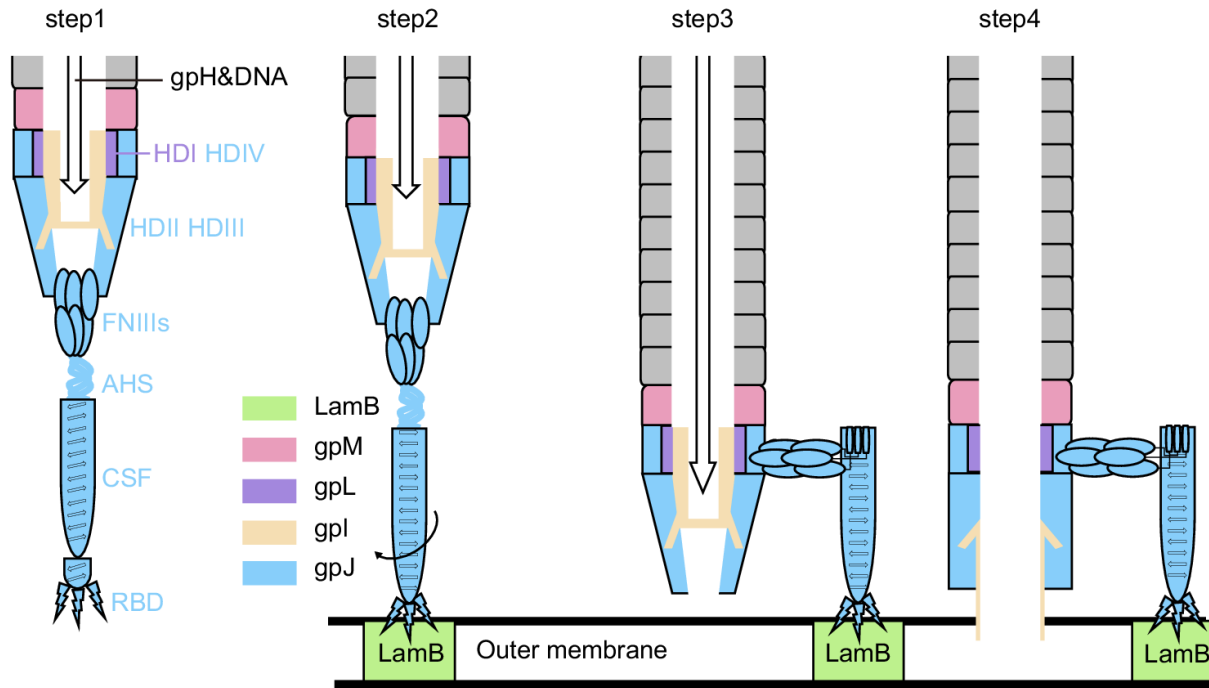
Some
bacteriophages have
contractile tails.

[animation](#)

Lambda does not
have a contractile
tail.



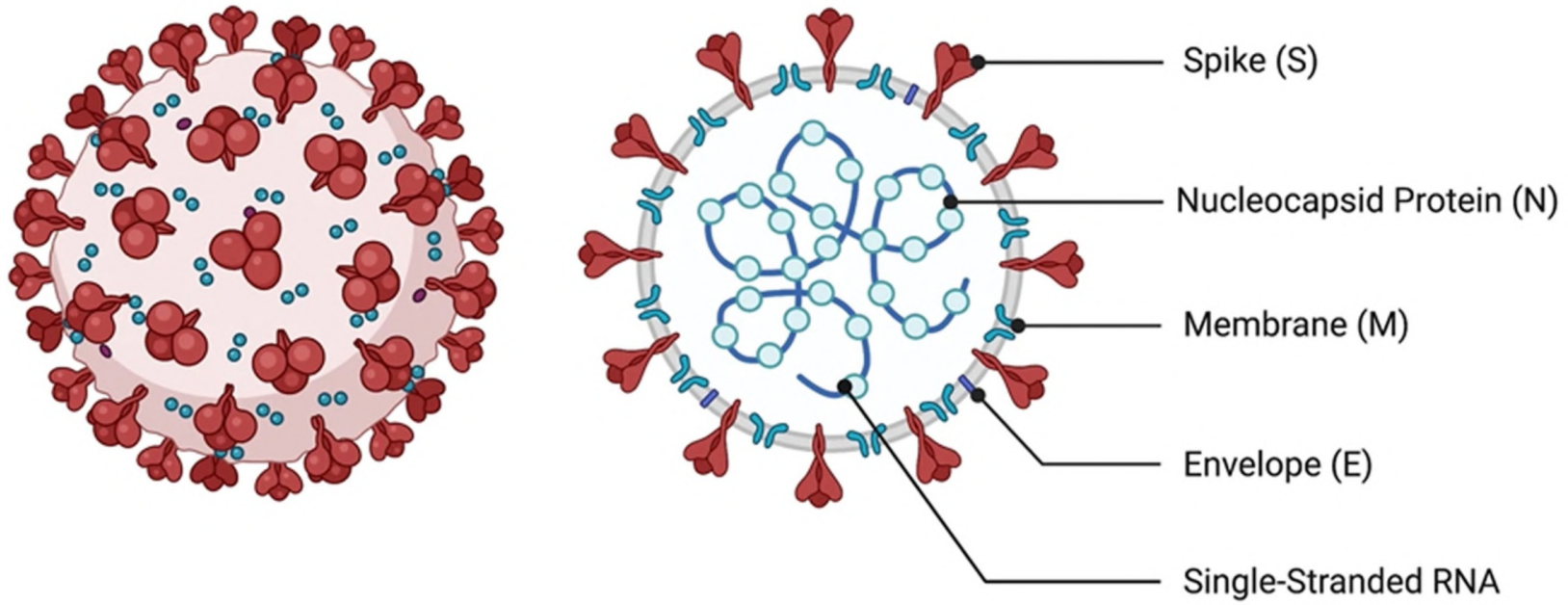
Extra detail - Tail Tube attachment of Lambda phage



Ge, X., Wang, J. Structural mechanism of bacteriophage lambda tail's interaction with the bacterial receptor. Nat Commun 15, 4185 (2024). <https://doi.org/10.1038/s41467-024-48686-3>

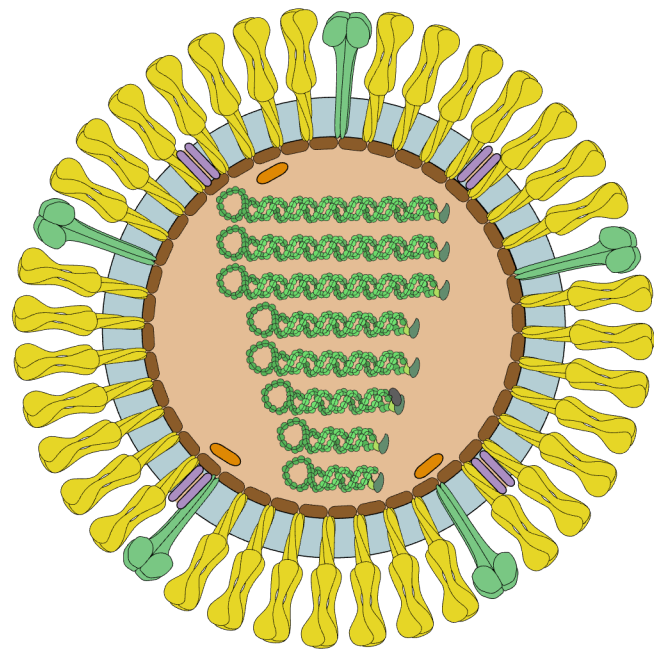
The LamB gene makes the maltoporin protein which acts as a receptor for the Lambda phage tail.

SARS CoV 2 Virus



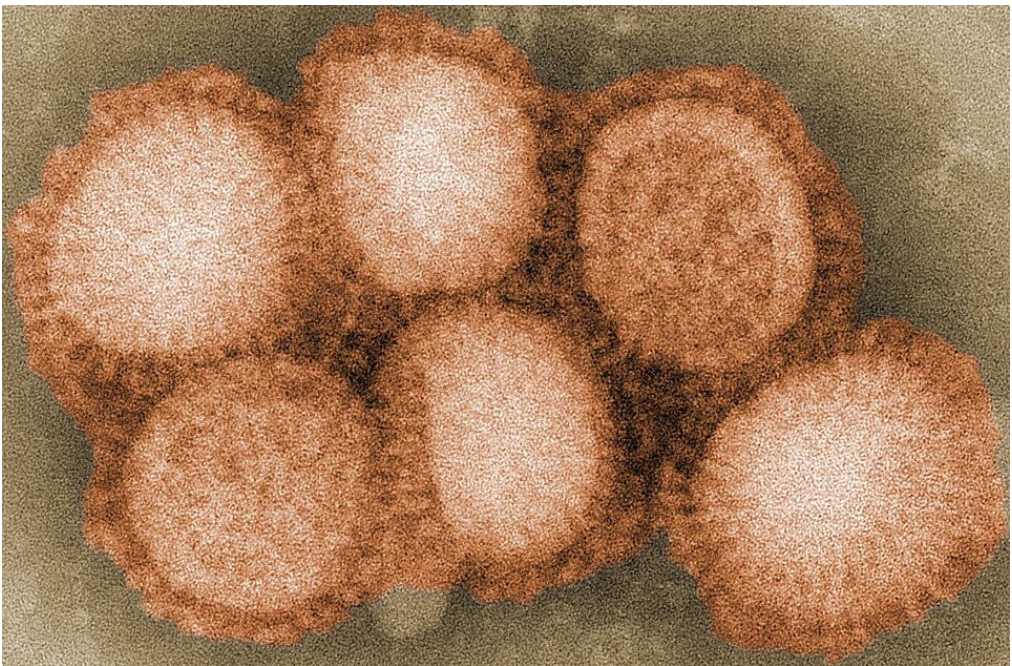
[Jamison et al 2022](#)

Influenza Virus

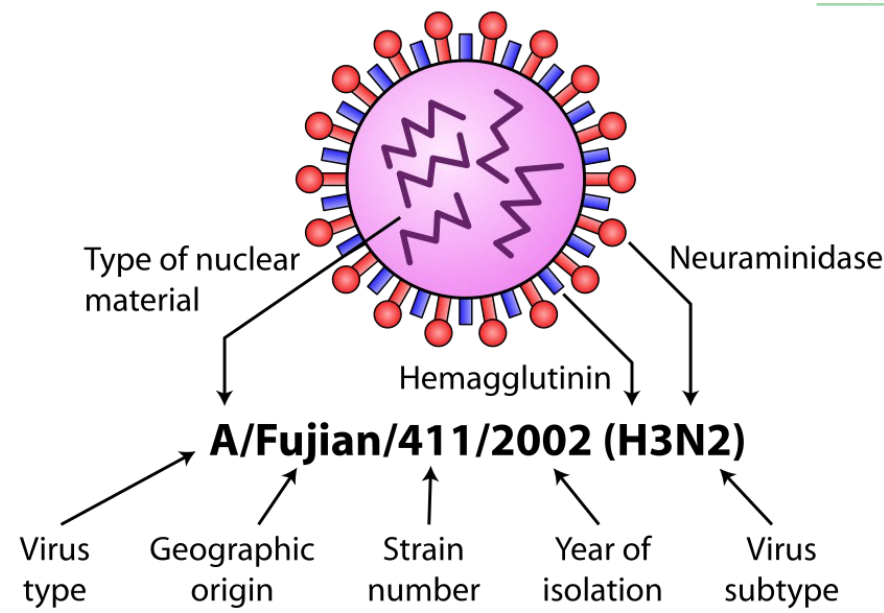


[Viralzone](#)

*I had a little bird,
its name was Enza.
I opened the window,
and in-flu-enza.*



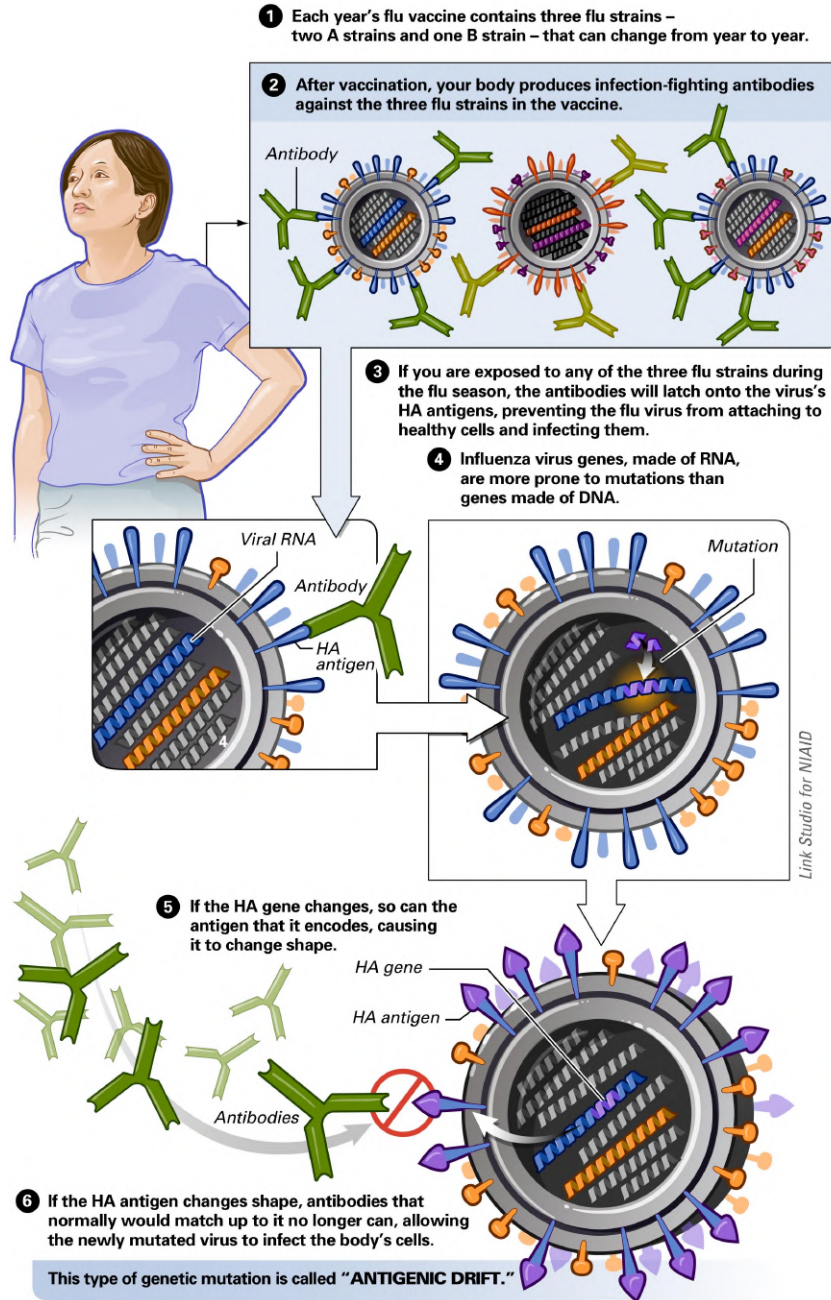
[Wikimedia](#)



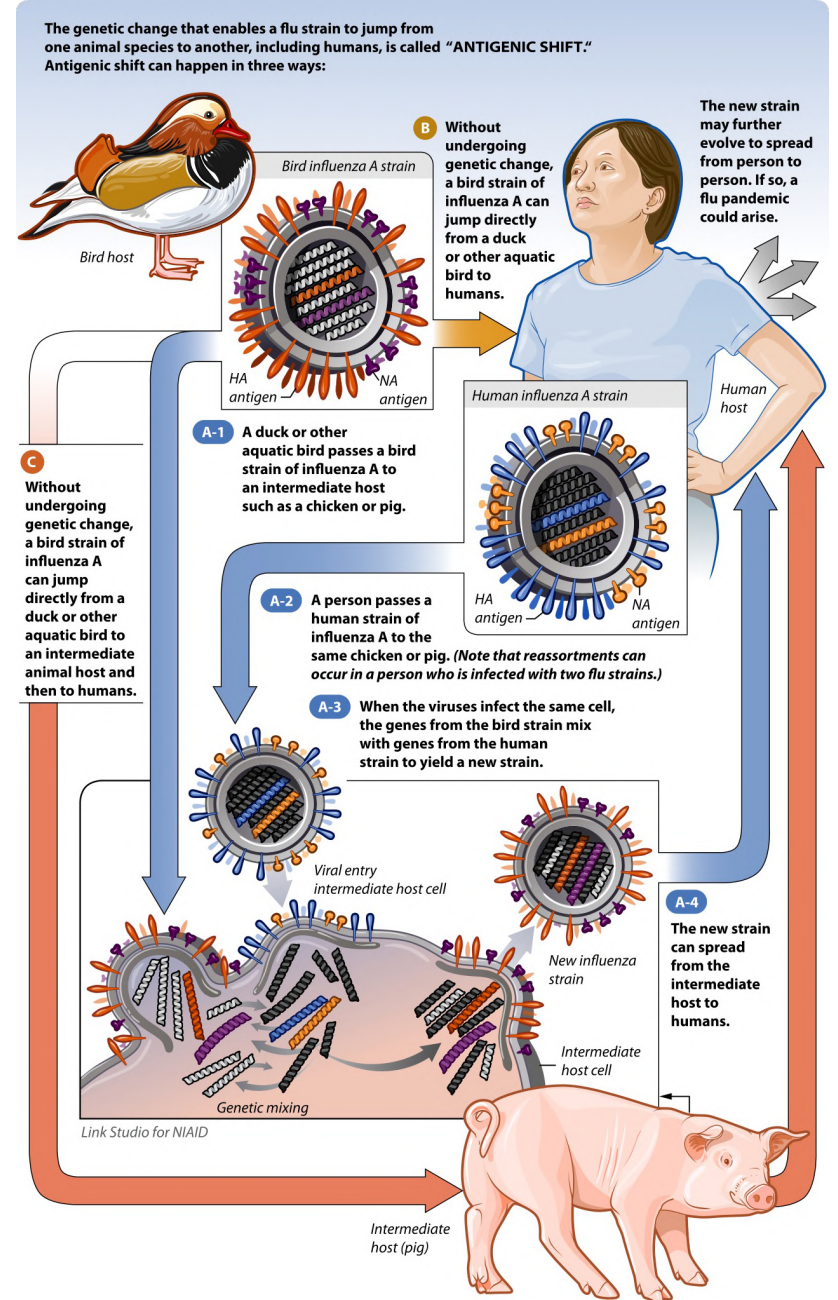
[Wikimedia](#)

Influenza Virus

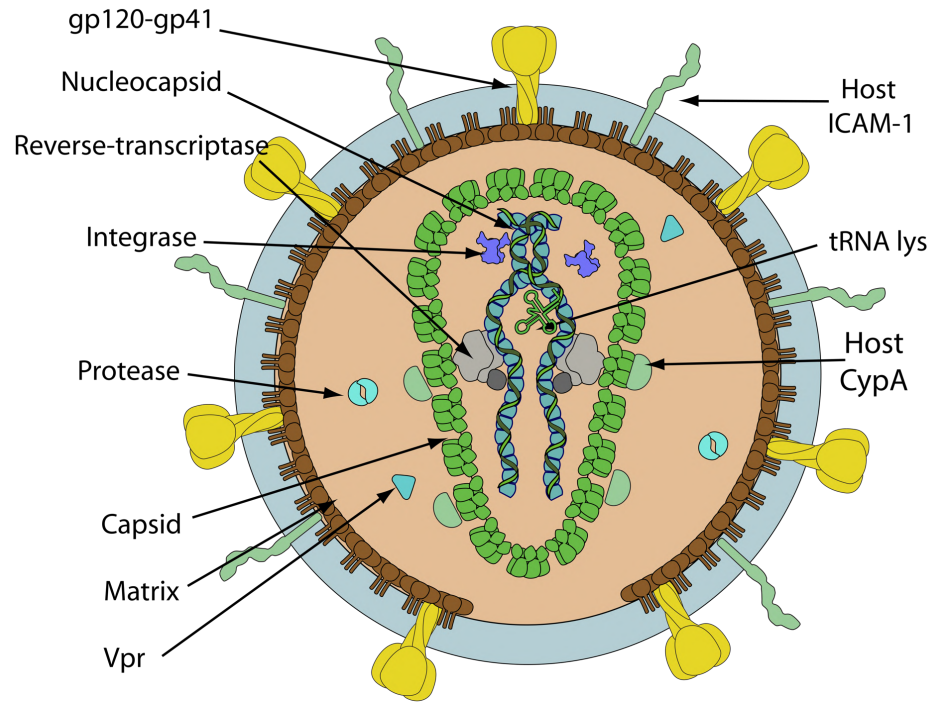
antigenic drift



antigenic shift



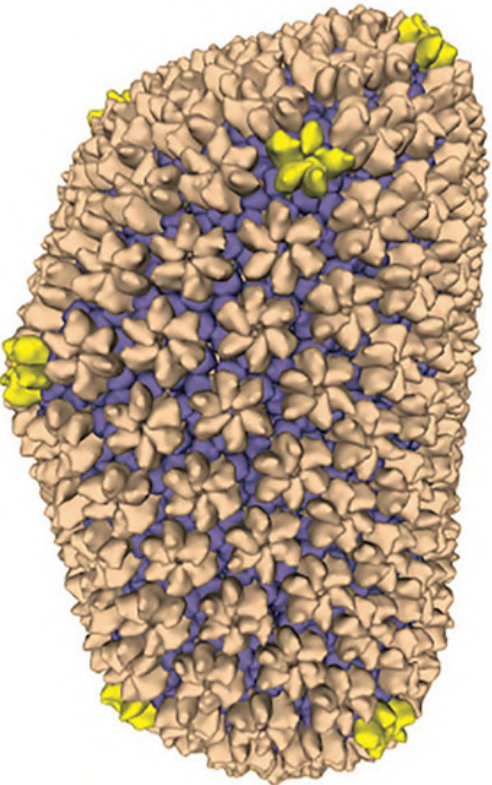
HIV Structure



MATURE

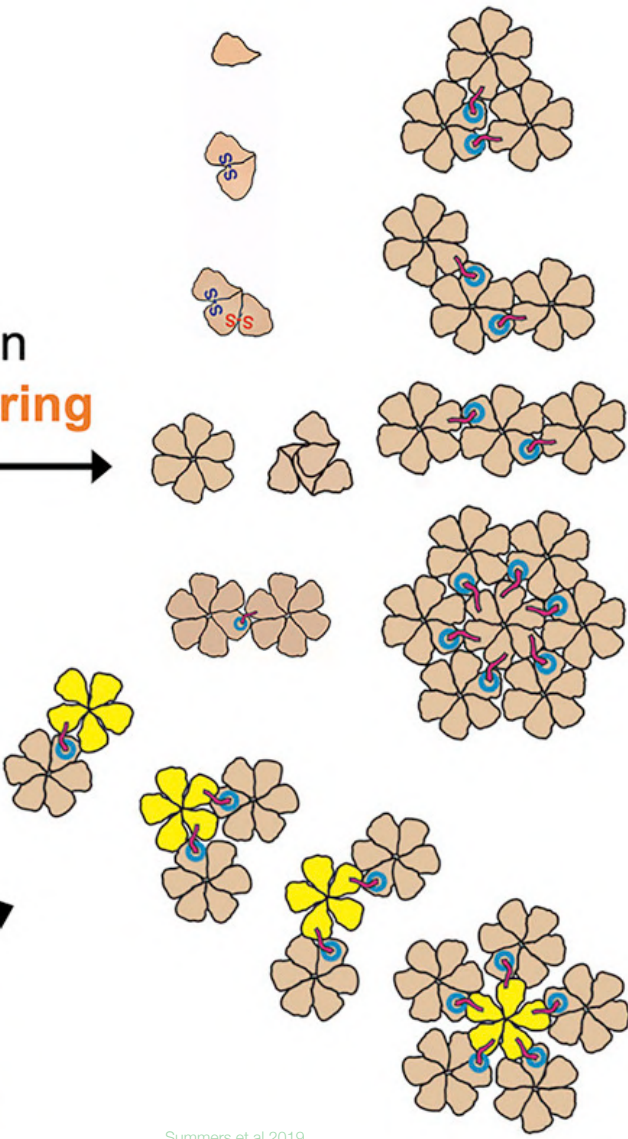
[Viralzone SIB Swiss Institute of Bioinformatics](#)

HIV-1
Capsid



Stabilized
Capsid Assembly
Platforms

Protein
Engineering



TRIM5α
TRIMCyp

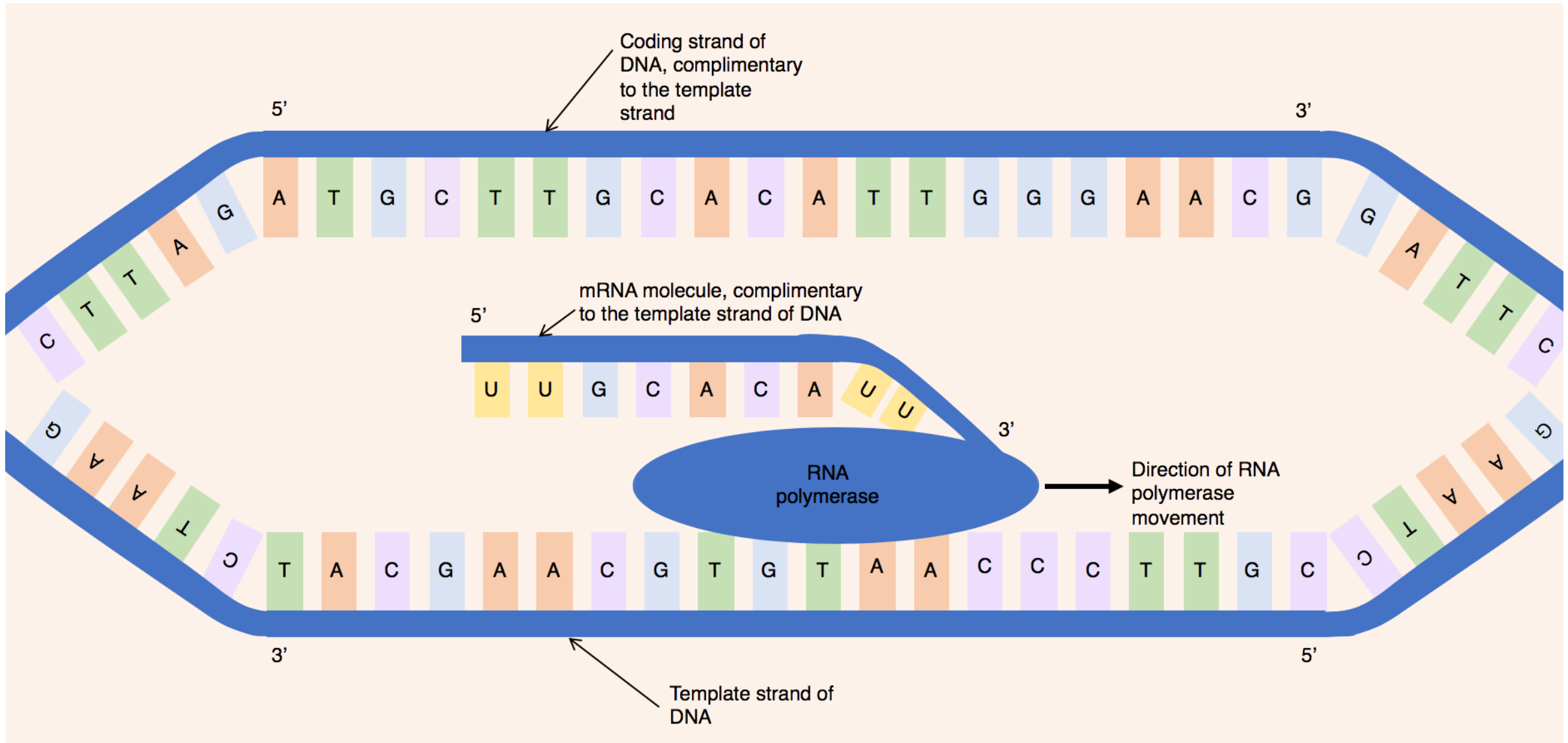
MxB

FEZ1

Host factor recognition

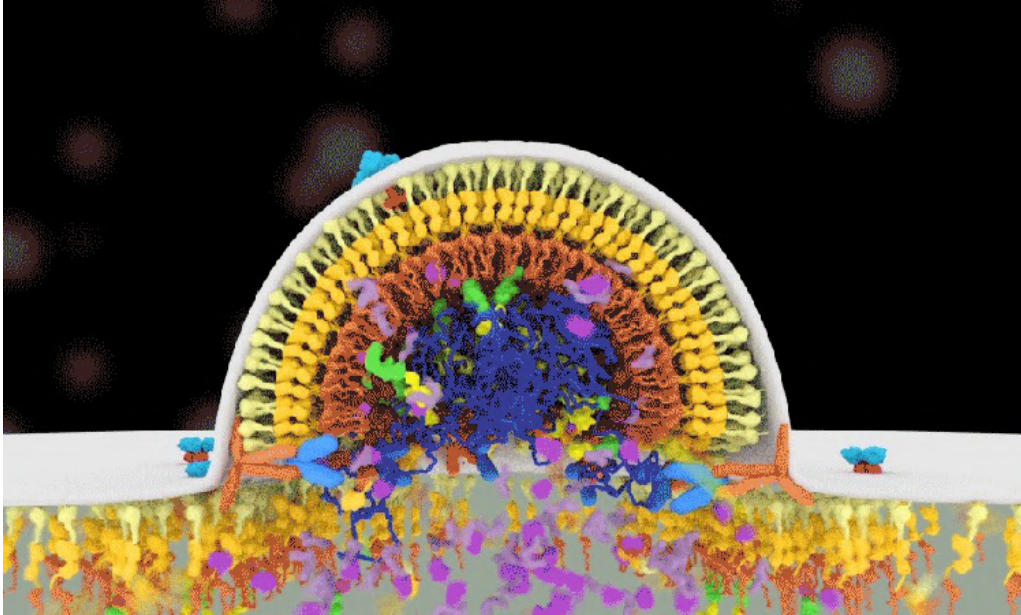
[Summers et al 2019](#)

Transcription DNA to mRNA

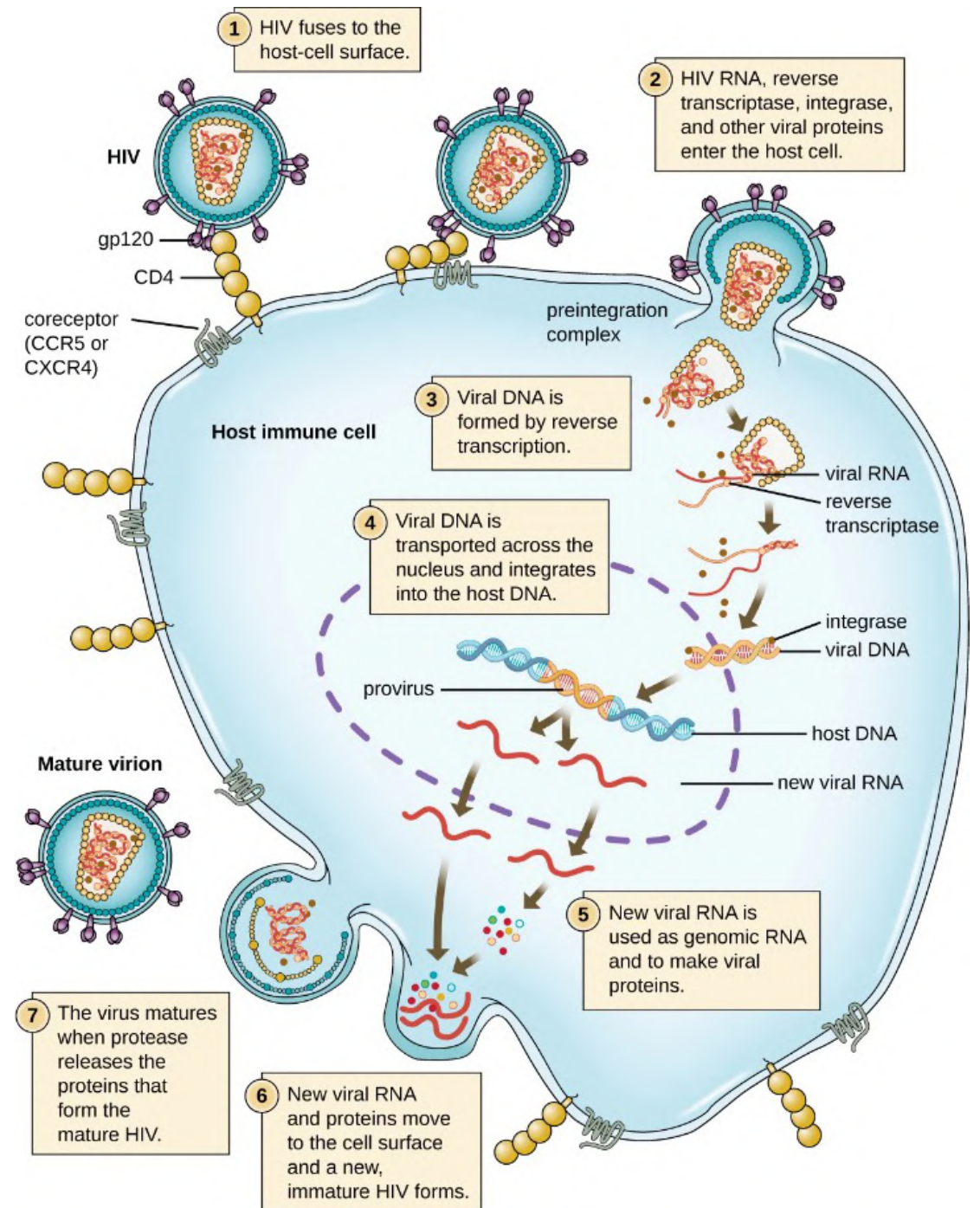


HIV Lifecycle

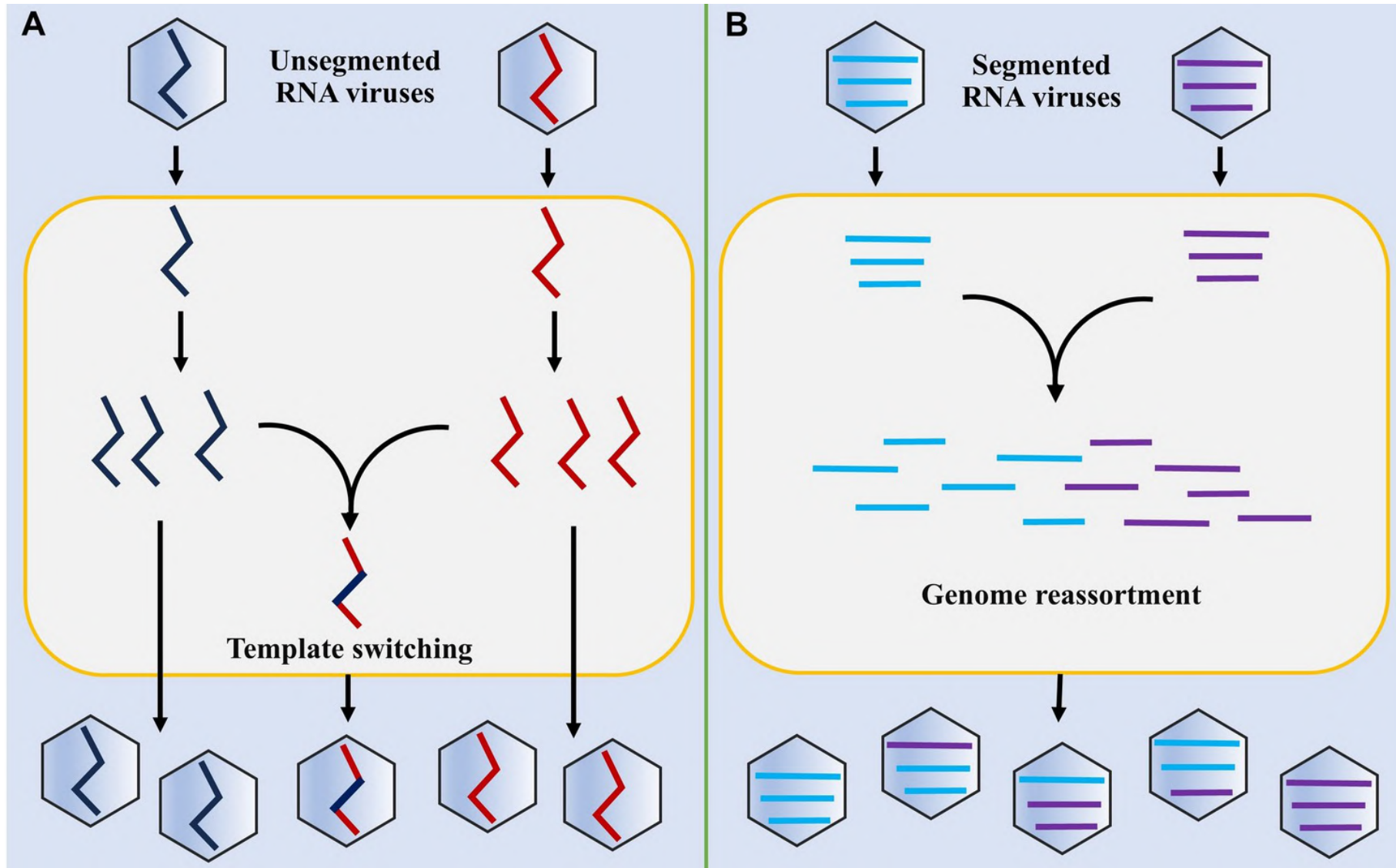
Budding of HIV



Animated and narrated by Janet Iwasa (Department of Biochemistry, University of Utah)



Rapid Evolution in Viruses



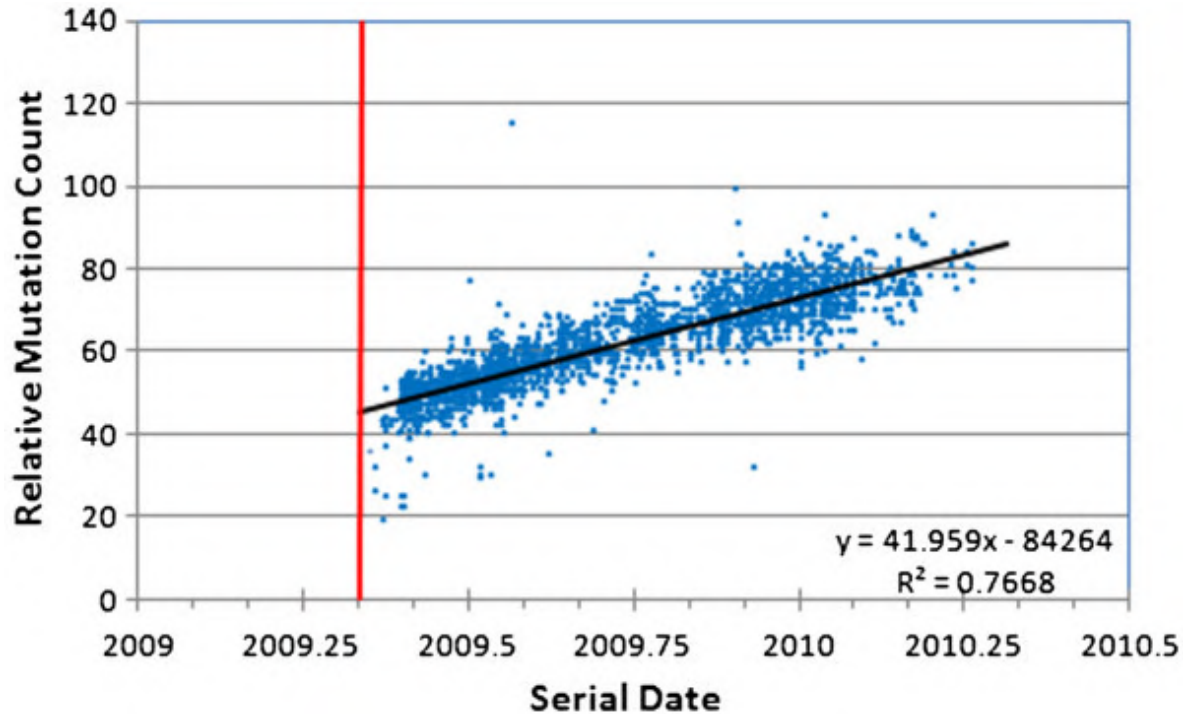
Abbasi and Alam 2025

Two viral strains of HIV in the same cell can undergo recombination of genes.

Two viral strains of influenza in the same cell can undergo reassortment of genes.

Rapid Evolution in Viruses

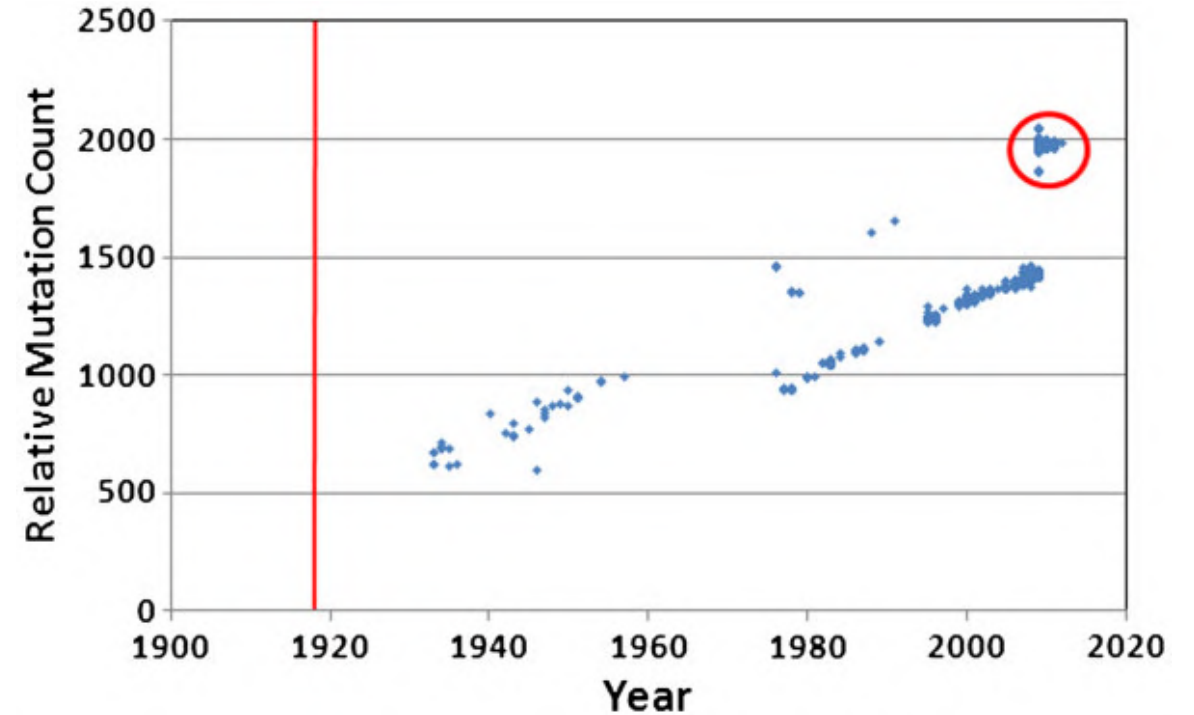
Mutation accumulation within the 2009 A/H1N1 influenza strain during the course of the 2009/2010 flu season.



The earliest sequenced genotype (California/04/2009) was used as the baseline for comparison with all subsequent genotypes (mutation counts reflect divergence from that starting sequence). The **vertical red line** indicates the date the reference sequence was collected.

[Carter and Sanford 2012](#)

Mutation accumulation in H1N1 in deeper time.



The published Brevig Mission strain from 1918 was used as the **baseline** for comparison with all available human H1N1 genomes. The 2009–2010 outbreak samples and additional samples from 2011–2012 are **circled**.

[Carter and Sanford 2012](#)

The mutations in viruses are an example of microevolution. High error rates in viruses can override natural selection and cause a virus to go extinct.

The influenza A [pandemic of 1918-20](#) after WW1 was the H1N1 subtype. It killed 2% of infected humans resulting in 50 million deaths. The human form of H1N1 went extinct in 1957 then after possibly an accidental release of stored samples in 1977 it again went extinct in 2009.

Optional extension: (not required for the IB exam)

Viruses have many beneficial ecological functions:

- o control microbial population size
- o increase or decrease microbe species diversity
- o help maintain gut health
- o facilitate horizontal gene transfer which increases phenotypic variation and adaptability
- o increase dissolved organic matter by bursting cells to recycle nutrients
- o are eaten by protists and sponges

Many medical uses of viruses await discovery including:

- o phage therapy to kill specific bacteria or
- o use of oncolytic viruses that kill specific cancerous cells
- o gene delivery of healthy DNA to specific cells

Agricultural uses of viruses await discovery including:

- o inoculating crops with mild strains or heat killed viruses can trigger defense responses that improve drought tolerance, freezing, pest resistance and protect from harmful viral strains



[James St. John](#)

A variety of *Dichanthelium lanuginosum* var. *thermale*, grows in geothermal areas of Yellowstone National Park, United States. When it is colonised by the fungus *Curvularia protuberata* and the fungus is in turn colonised by a particular virus, the grass is able to tolerate soil temperatures of up to 65 °C that would otherwise be lethal.

