



# VIROLOGY LIVE

WITH VINCENT RACANIELLO

## Reverse transcription and integration

Session 9

Virology Live

Fall 2021

*"One can't believe impossible things," said Alice.  
"I dare say you haven't had much practice," said  
the Queen. "Why, sometimes I've believed as many  
as six impossible things before breakfast."  
--LEWIS CARROLL, Alice in Wonderland*

# Tumor virus history

- 1908 - Discovery of chicken leukemia virus, Bang & Ellerman
- 1911 - Discovery of Rous sarcoma virus, Peyton Rous (Nobel Prize 55 years later)
- Called tumor viruses
- Found to have RNA genomes



Vilhelm Ellerman



Oluf Bang



Peyton Rous

## Howard Temin's insight

- RNA tumor viruses caused permanent changes in cells (transformation)
- Viral DNA was integrated into host genome
- Became permanent part of host DNA
- Provirus hypothesis



## David Baltimore's insight

- (+) RNA viruses: No RdRp in particle
- (-) RNA viruses: RdRp in particle
- An enzyme to copy (+) RNA to DNA must be in virus particle





# Baltimore and Temin independently discovered RT in RNA tumor virus particles (Nobel Prize, 1975)

## RNA-dependent DNA Polymerase in Virions of Rous Sarcoma Virus

INFECTION of sensitive cells by RNA sarcoma viruses requires the synthesis of new DNA different from that synthesized in the *S*-phase of the cell cycle (refs. 1, 2 and unpublished results of D. Boettiger and H. M. T.); production of RNA tumour viruses is sensitive to actinomycin D<sup>3,4</sup>; and cells transformed by RNA tumour viruses have new DNA which hybridizes with viral RNA<sup>5,6</sup>. These are the basic observations essential to the **DNA provirus** hypothesis—replication of RNA tumour viruses takes place through a DNA intermediate, not



## RNA-dependent DNA Polymerase in Virions of RNA Tumour Viruses

DNA seems to have a critical role in the multiplication and transforming ability of RNA tumour viruses<sup>1</sup>. Infection and transformation by these viruses can be prevented by inhibitors of DNA synthesis added during the first 8–12 h after exposure of cells to the virus<sup>1–4</sup>. The necessary DNA synthesis seems to involve the production of DNA which is genetically specific for the infecting virus<sup>5,6</sup>, although hybridization studies intended to demonstrate virus-specific DNA have been inconclusive<sup>1</sup>. Also, the formation of virions by the RNA tumour viruses is sensitive to actinomycin D and therefore seems to involve DNA-dependent RNA synthesis<sup>1–4,7</sup>. One model which explains these data postulates the transfer of the information of the infecting RNA to a DNA copy which then serves as template for the synthesis of viral RNA<sup>1,3,7</sup>. This model requires a unique enzyme, an RNA-dependent DNA polymerase.



Listen to TWiV #100 (Baltimore) for more insight

# Reverse transcriptase

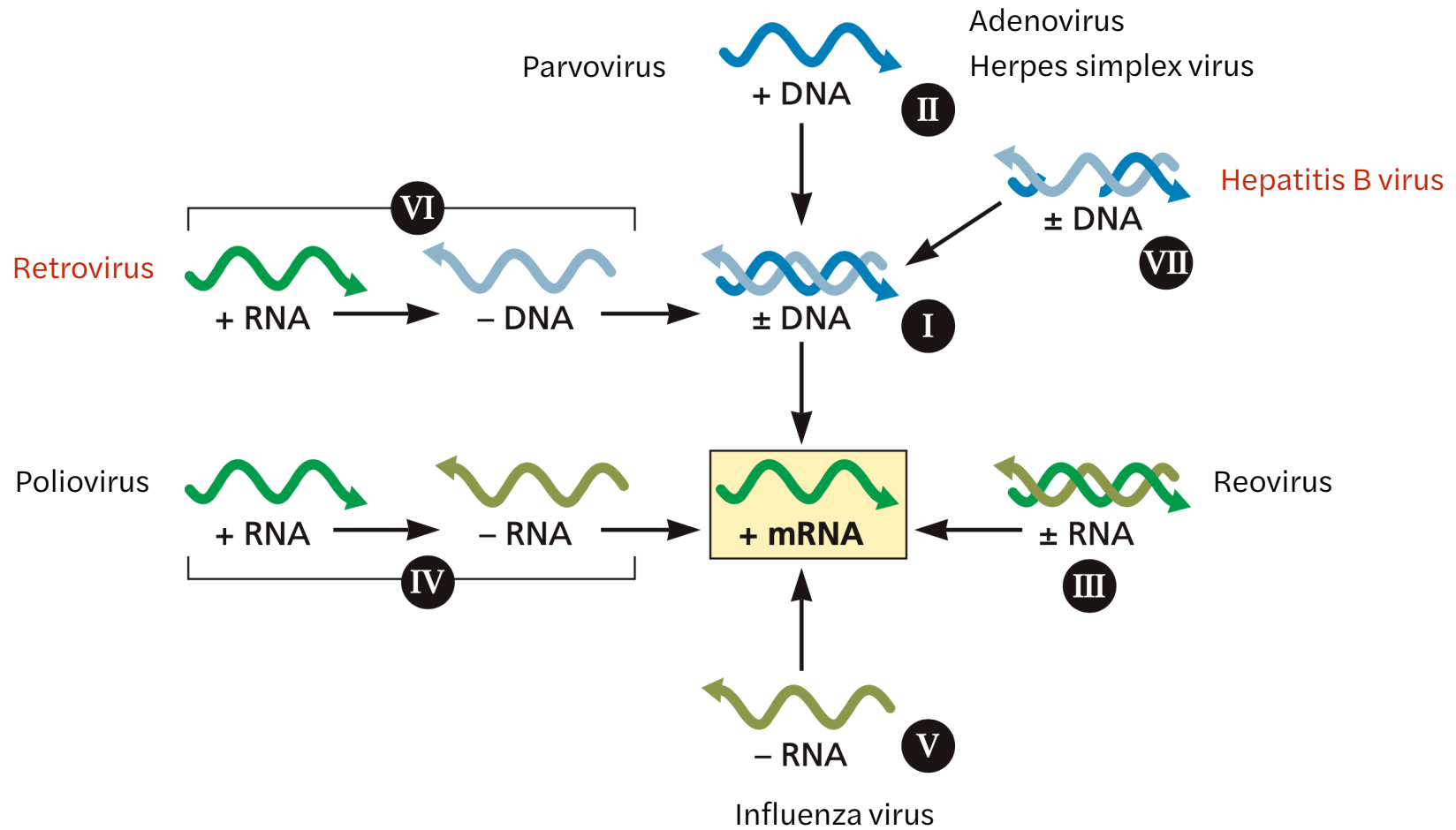
- Retroviruses got their name because of their ability to reverse the flow of genetic information

DNA => RNA => protein

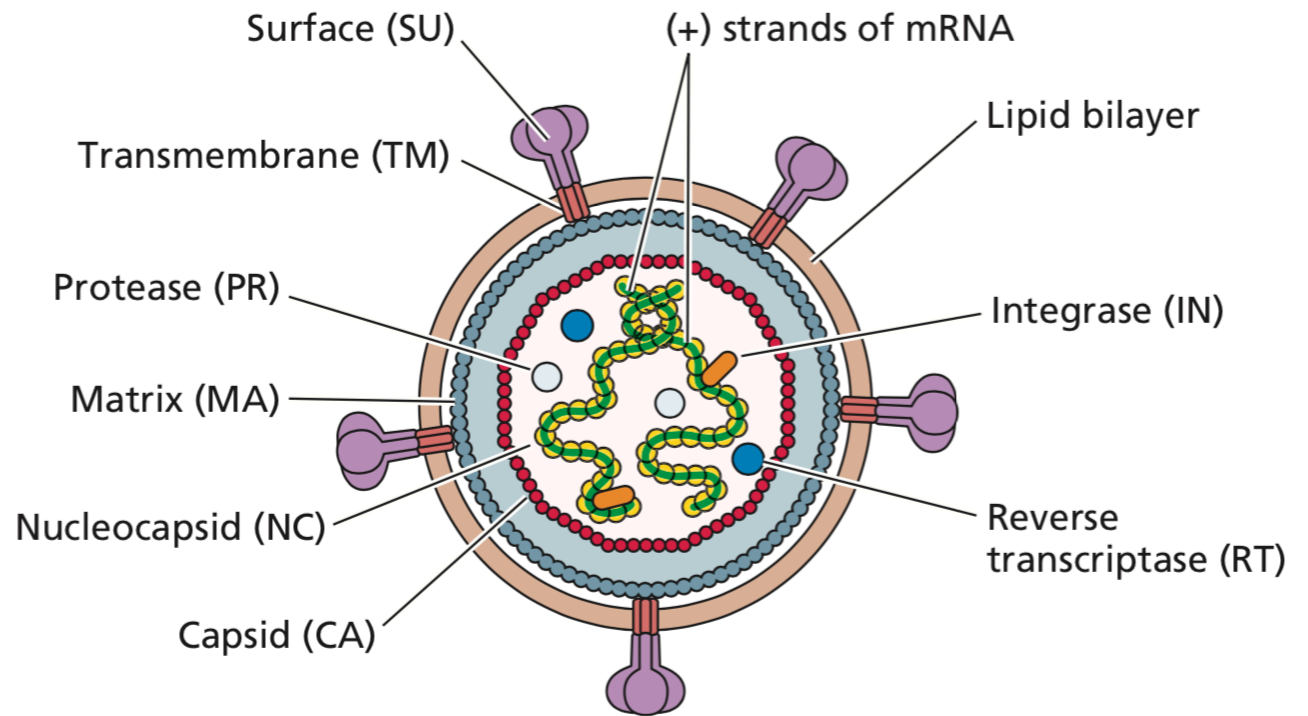
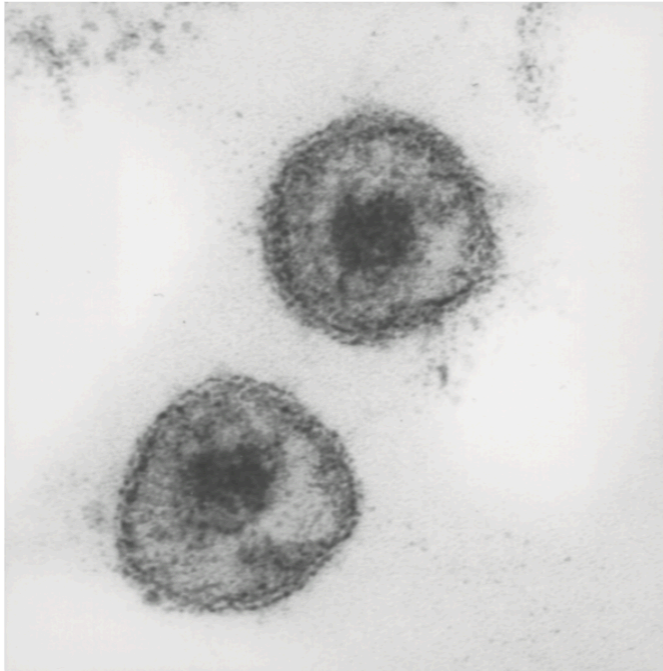
- RT discovery revolutionized molecular biology (e.g. SARS-CoV-2 diagnosis may utilize RT-PCR)



# Viruses with RT

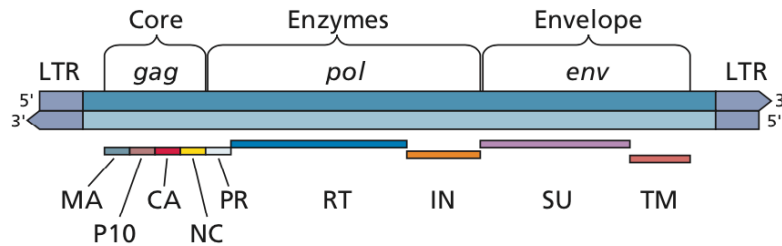


# Rous sarcoma virus, a retrovirus



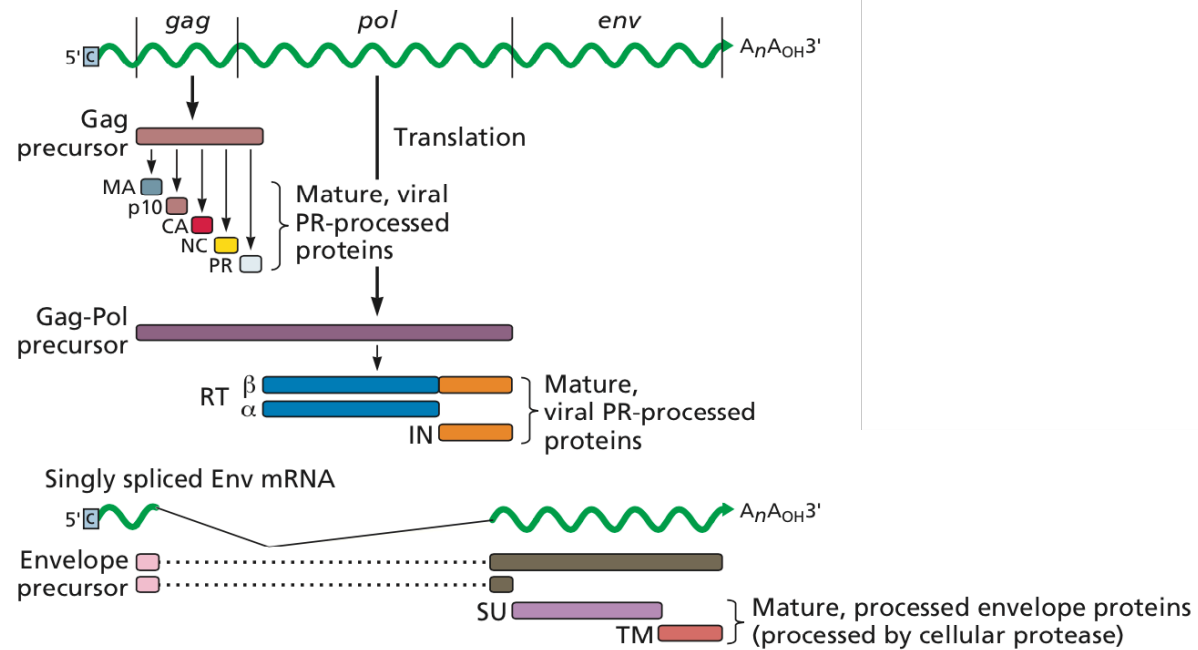
## Simple genome (ALV)

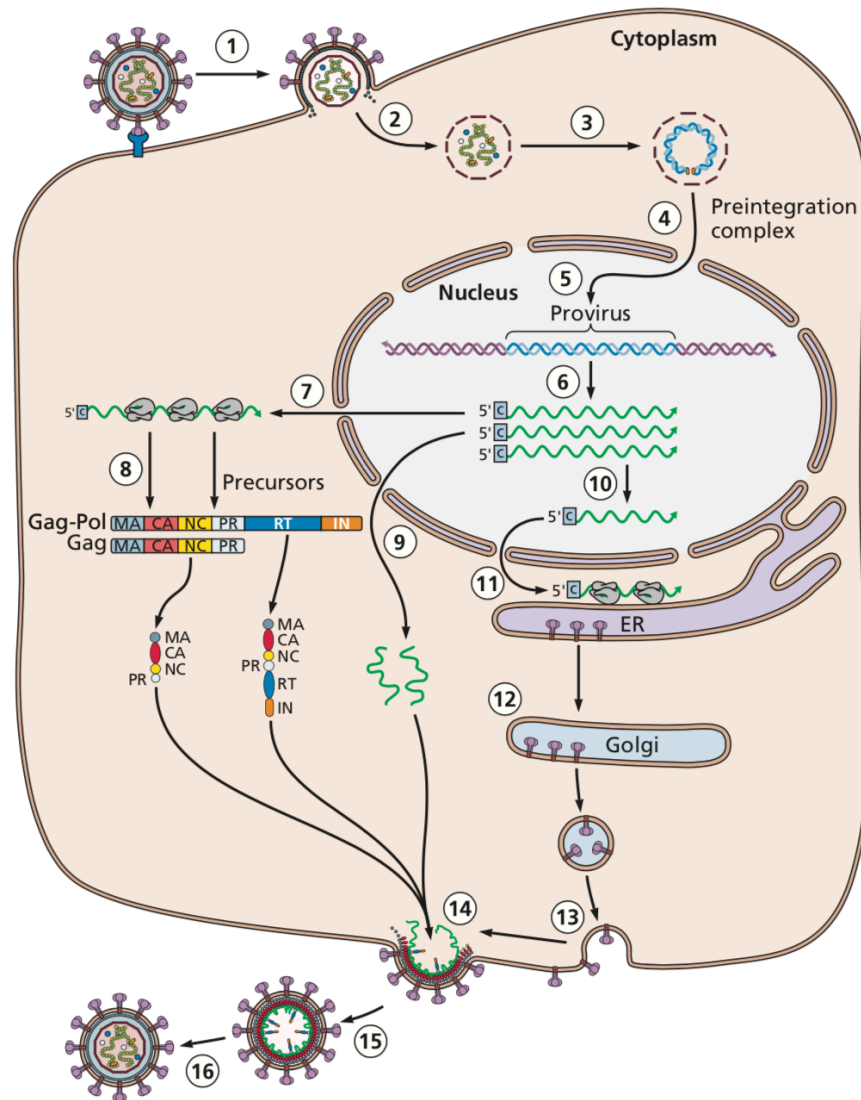
Proviral DNA



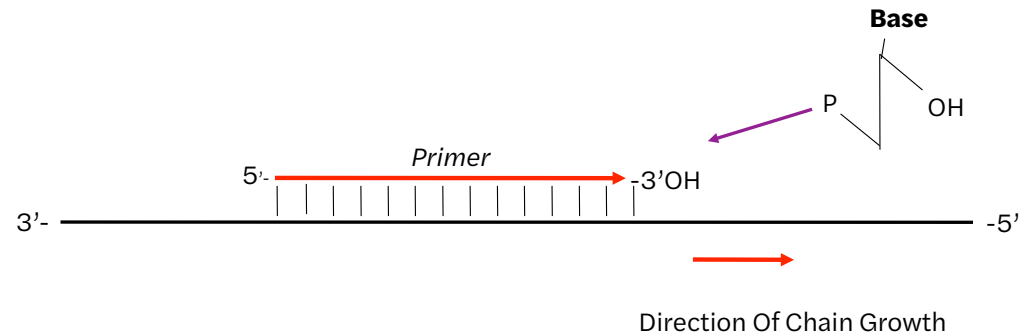
## Genome expression

Genomic RNA, Gag-Pol mRNA, pre-mRNA





# Reverse transcriptase

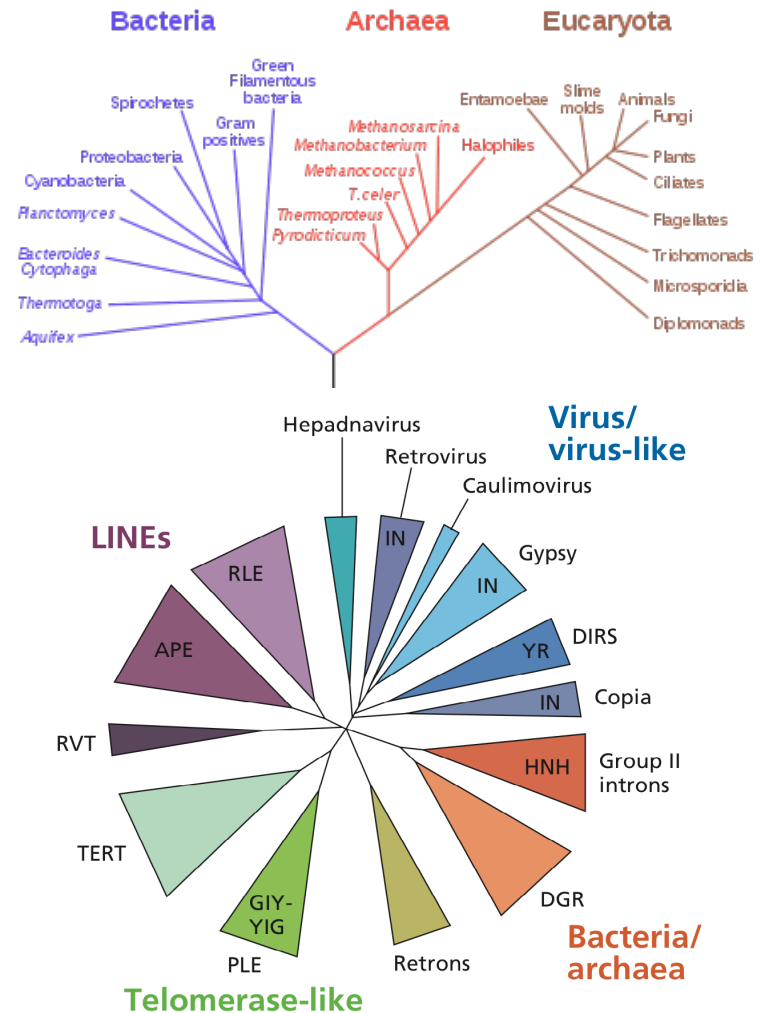


- Primer can be DNA or RNA
- Template can be RNA or DNA
- Only dNTPs, not rNTPs, are incorporated

# RT

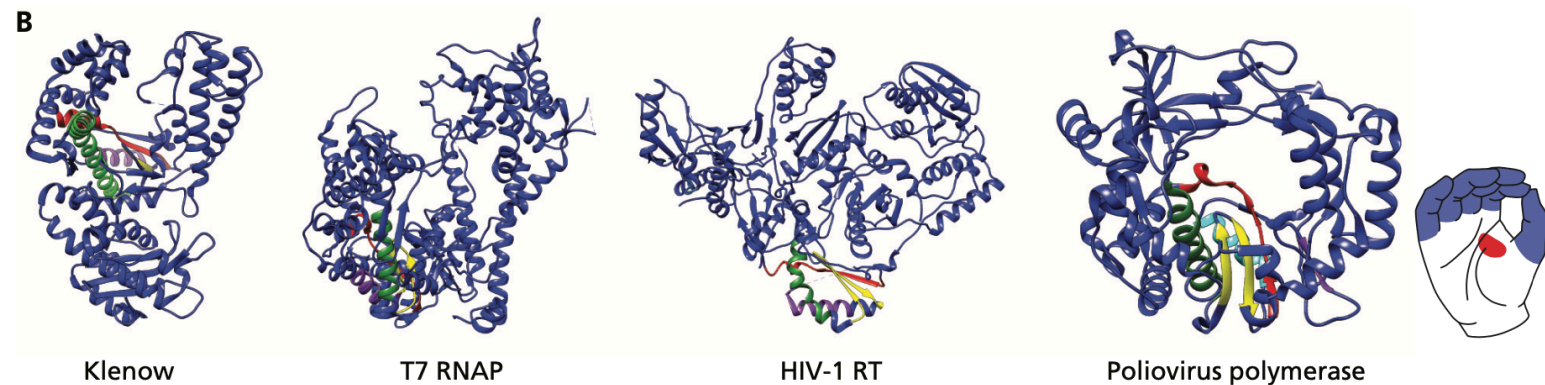
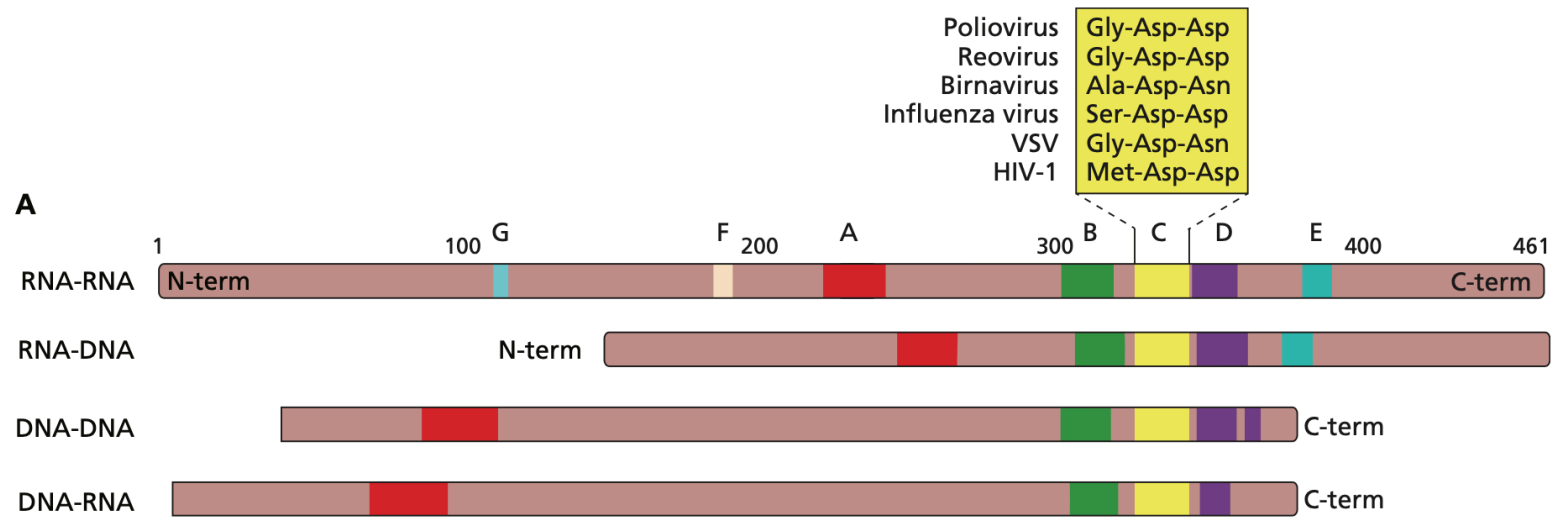
- Bacteria, Archaea, eukaryotes have RT activity
- Therefore RT evolved before the separation of Archaea, bacteria, and eukaryotes
- RT might be the bridge between early RNA world and modern DNA world
- RT also in HBV, *Caulimoviridae*

## Phylogenetic Tree of Life

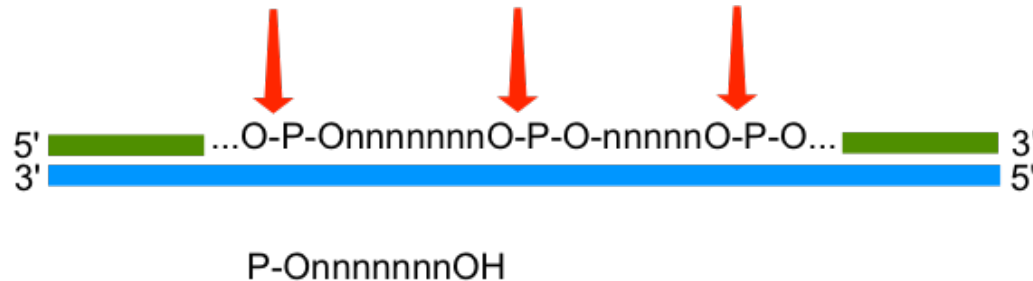




# Sequence relationships among polymerases

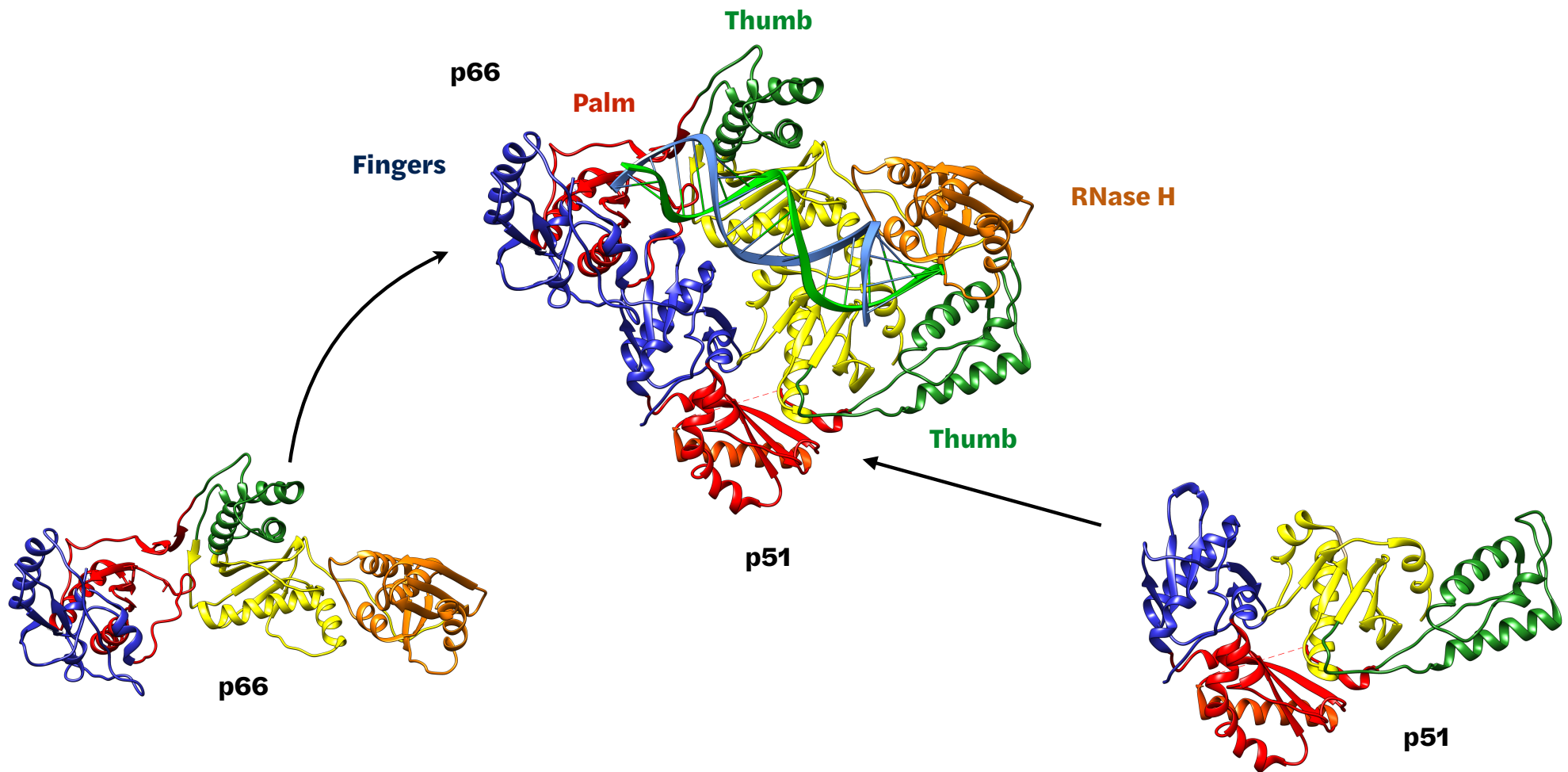


## RNAse H: A second activity of RT

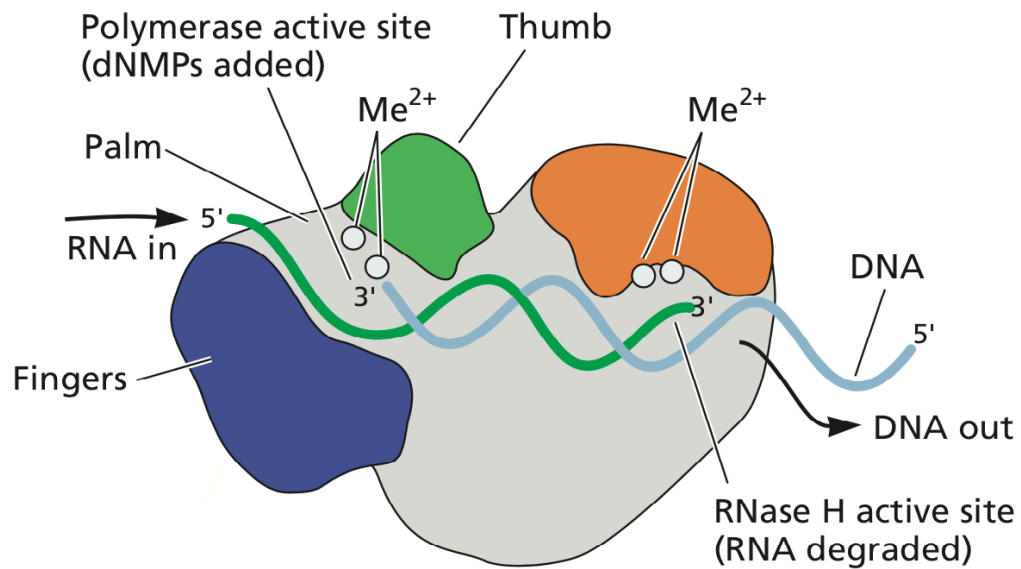


- Cleaves RNA only when in duplex form
- RNA can be in RNA:RNA or RNA:DNA duplexes
- Makes endonucleolytic cleavages
- Produces short oligonucleotides with 5'-phosphate, 3'-OH

# HIV-1 Reverse transcriptase



# HIV-1 Reverse transcriptase



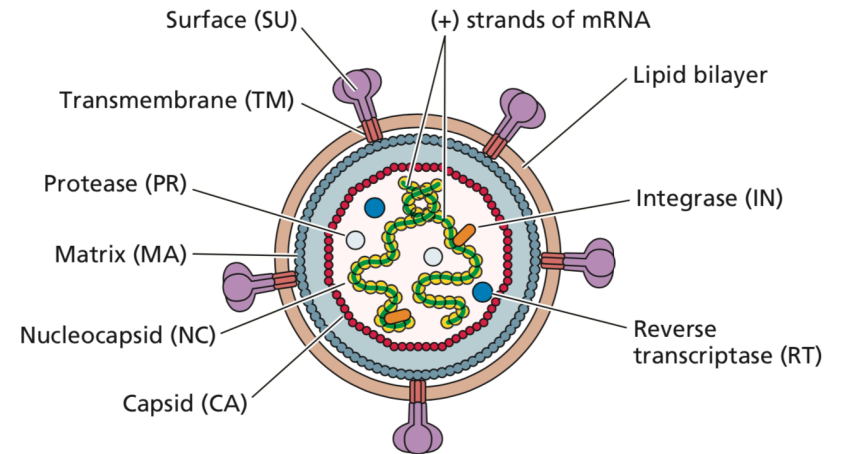
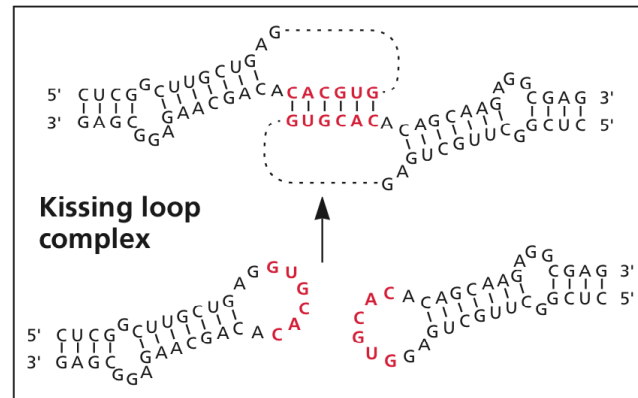
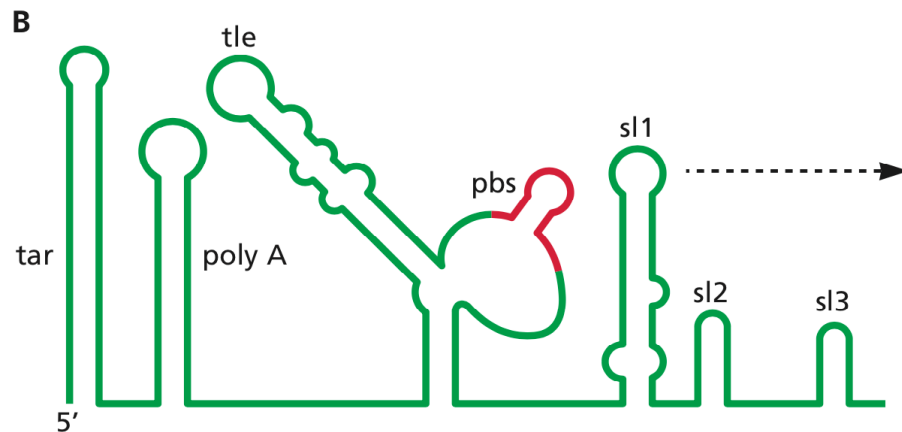
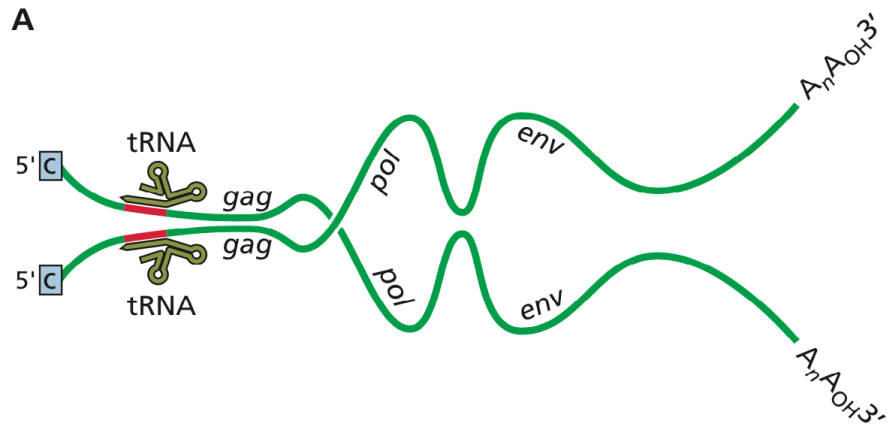
DNA synthesis is slow (4 h per 9 kb genome) and error prone (1 misincorporation per  $10^4$  to  $10^6$  nt)

**Go to:**

**[b.socrative.com/login/student](https://b.socrative.com/login/student)  
room number: virus**

Reverse transcriptase has revolutionized molecular biology. Which statement about the enzyme is not correct?

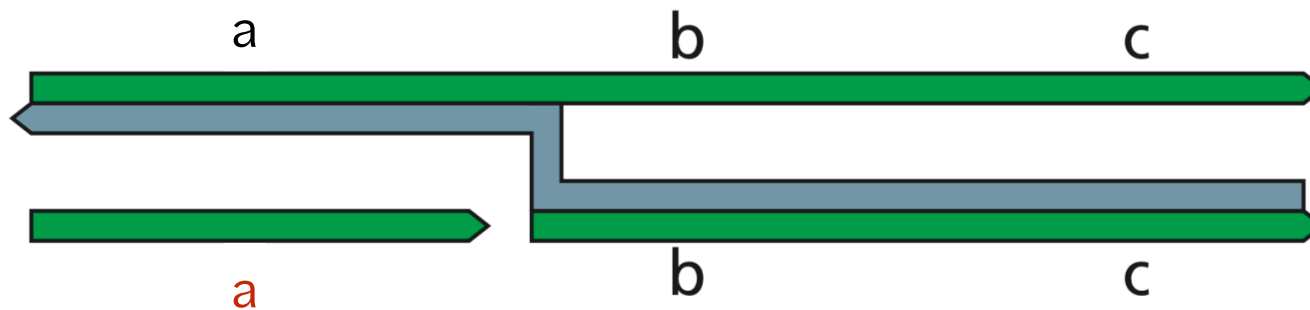
- A. RT is unique to retroviruses
- B. RT is packaged in the retrovirus particle
- C. The RT protein also has RNase H activity
- D. The name of the enzyme comes from its ability to reverse the flow of genetic information
- E. Might have bridged the ancient RNA world and the DNA world



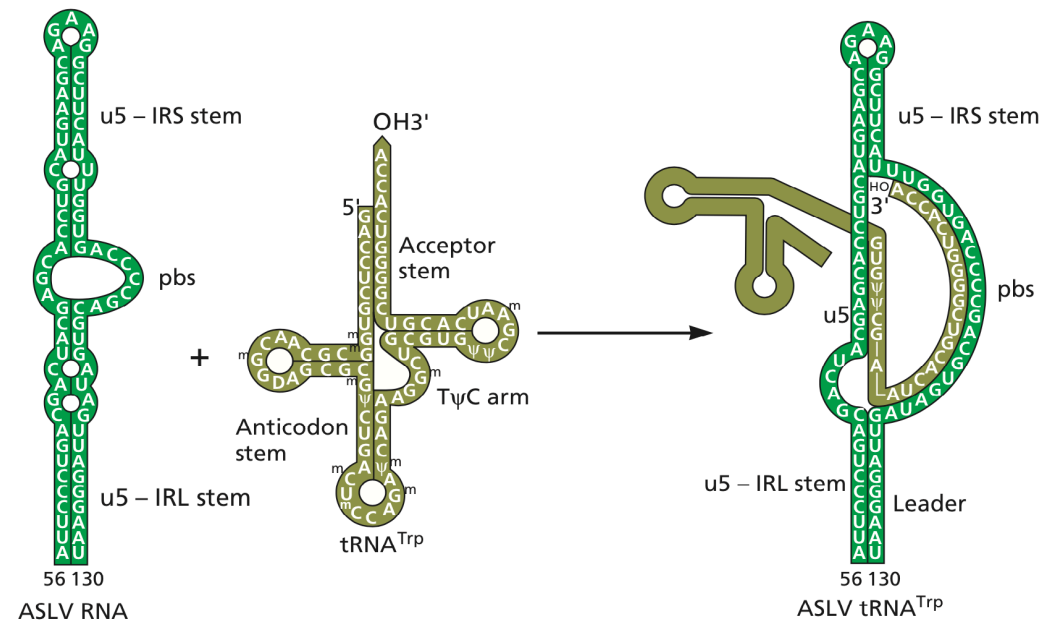
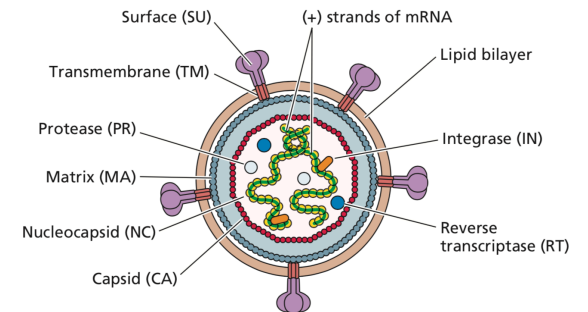
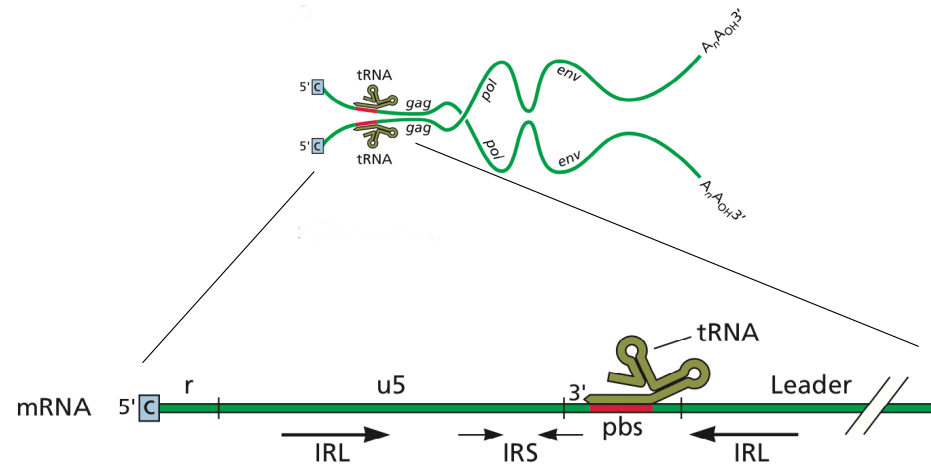
RNA coated with NC protein  
50-100 molecules RT per virus particle

## RNA dimer

- Explains why retroviruses are relatively resistant to UV and ionizing radiation
- Two copies of all genes
- Recombination (copy-choice) during reverse transcription builds one functional genome

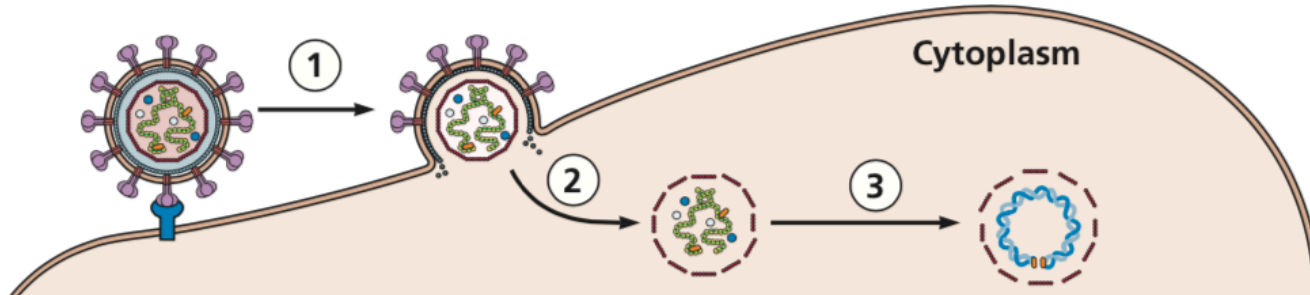


# Primer tRNA binding to retroviral genome



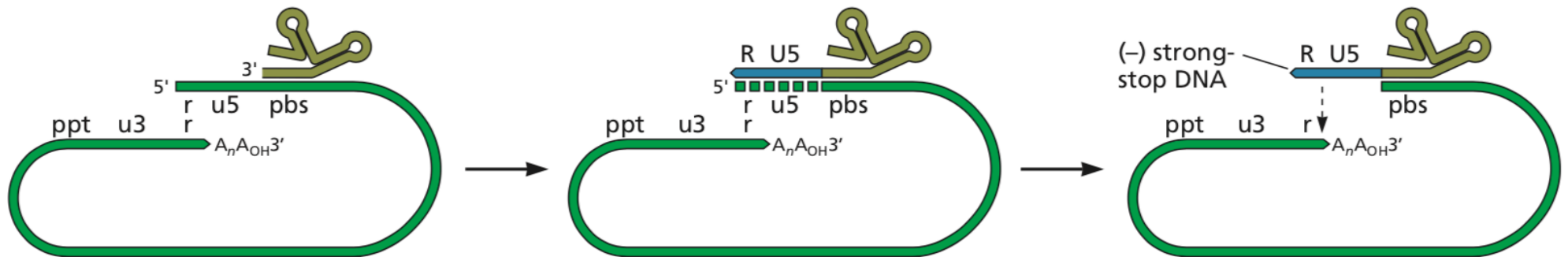


# DNA synthesis: cytoplasmic

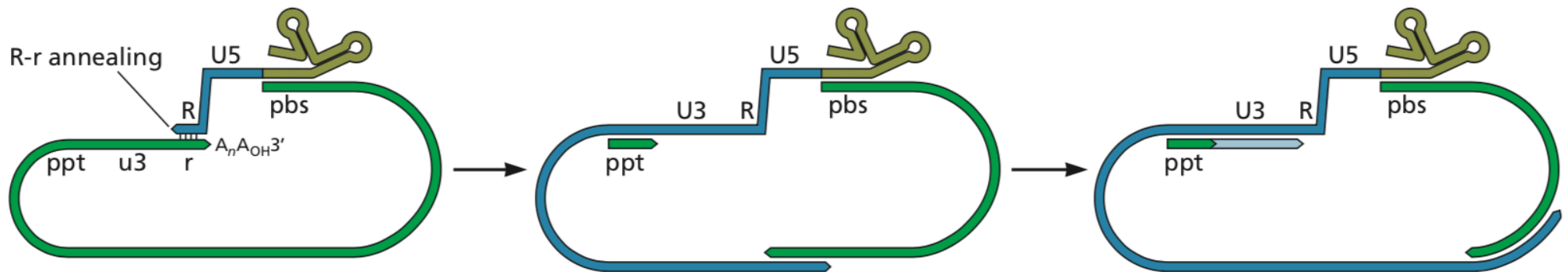


## Initiation of (–) strand DNA synthesis

The 5' end of the viral RNA genome is degraded by the RNase H activity of RT as the (–) strand DNA is synthesized.



### First template exchange

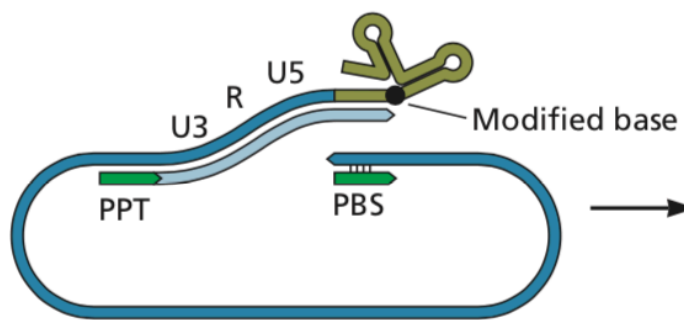


ppt = polypurine tract

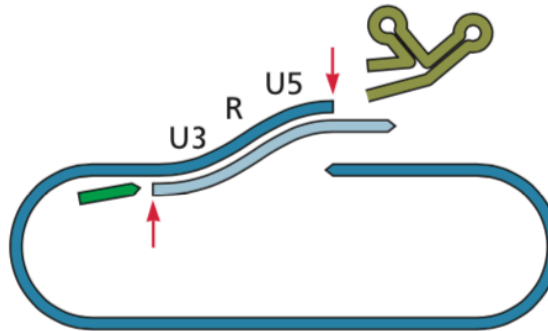
## (+) strand DNA synthesis

The pbs sequence is copied twice:

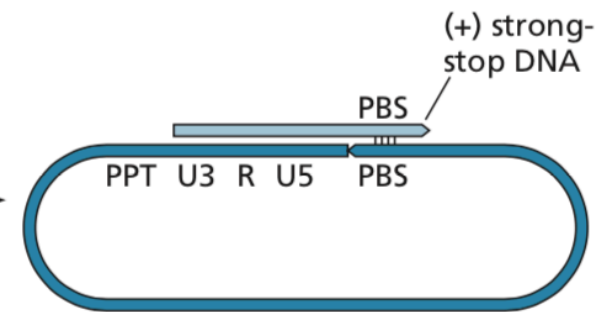
- once from the RNA genome
- once from the tRNA primer



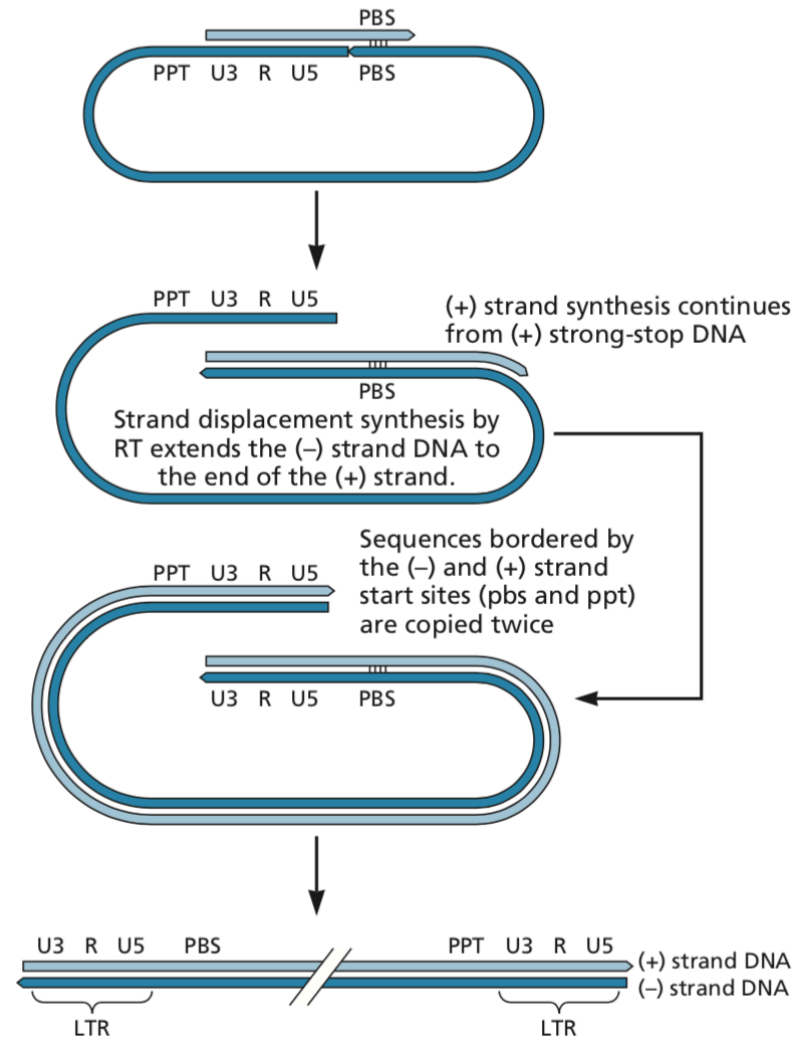
RNase H endonuclease activity of RT removes both primer RNAs

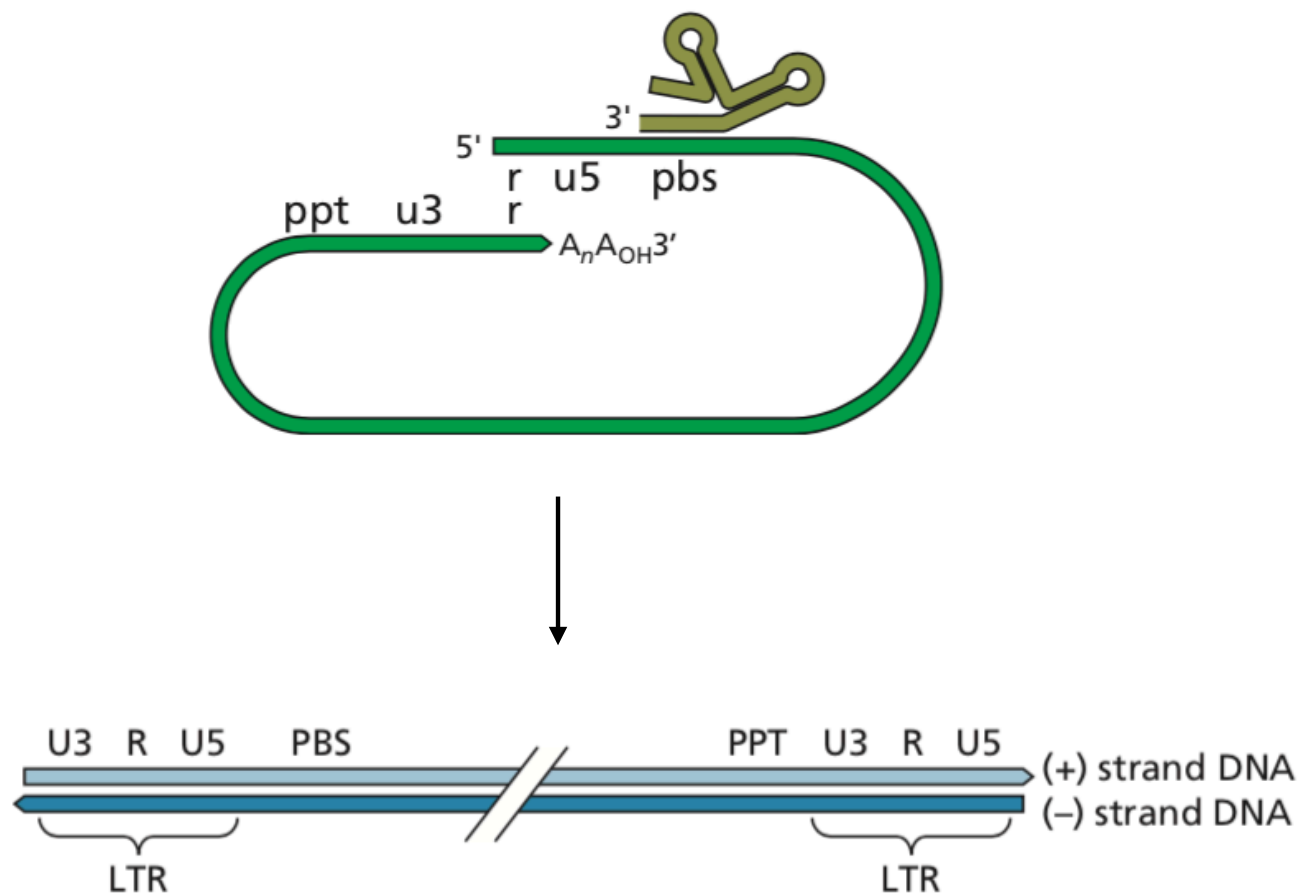


DNA ends are juxtaposed by annealing at complementary PBS sequences



**Second template exchange is facilitated by annealing of PBS sequences**





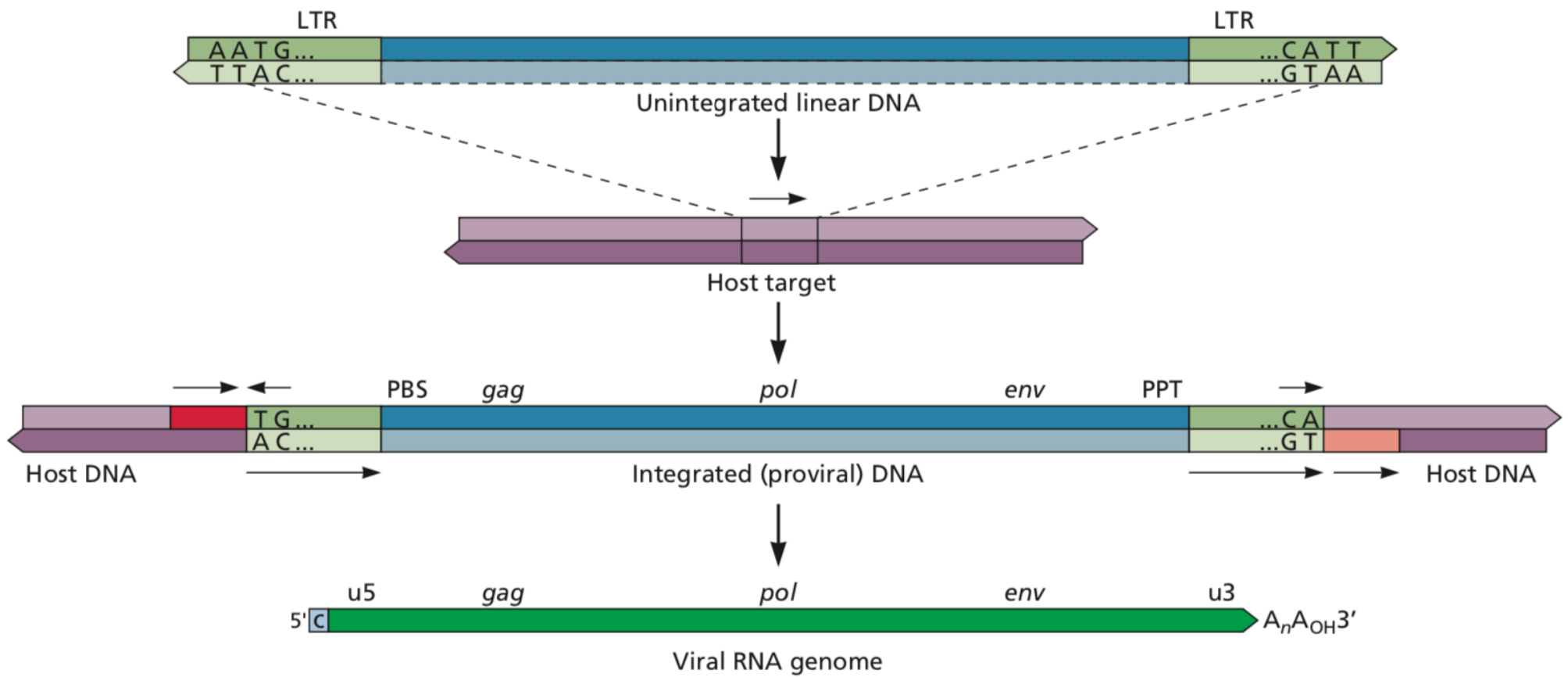
Animation of reverse transcription: <https://youtu.be/RYwVnzYf4V8>

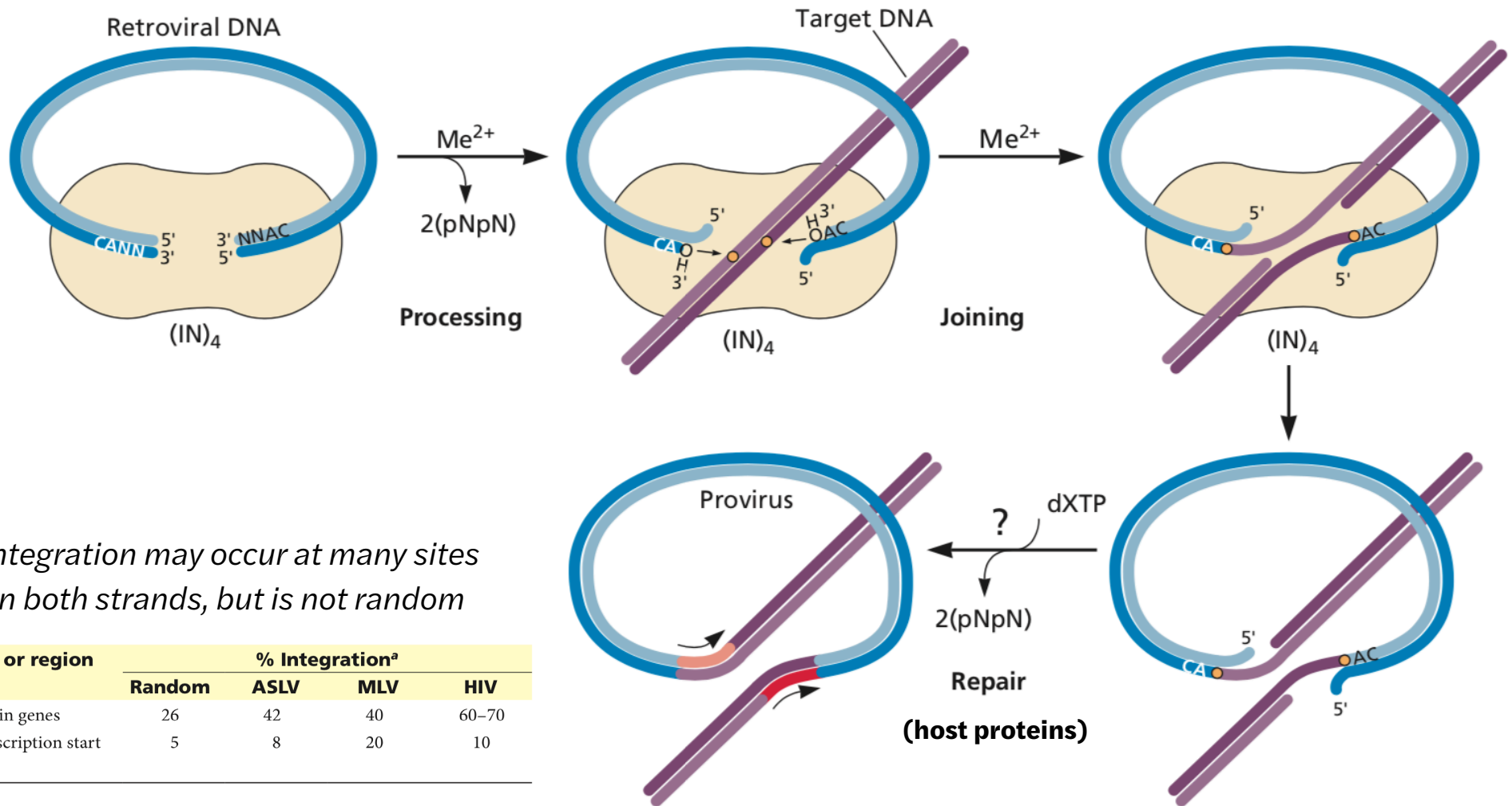
**Go to:**

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room number: virus**

Which of the following steps occur during reverse transcription of retroviral genomic RNA?

- A. Priming of (-) DNA synthesis by tRNA
- B. Two template exchanges
- C. Degradation of the viral RNA by RNase H
- D. Generation of two LTRs
- E. All of the above



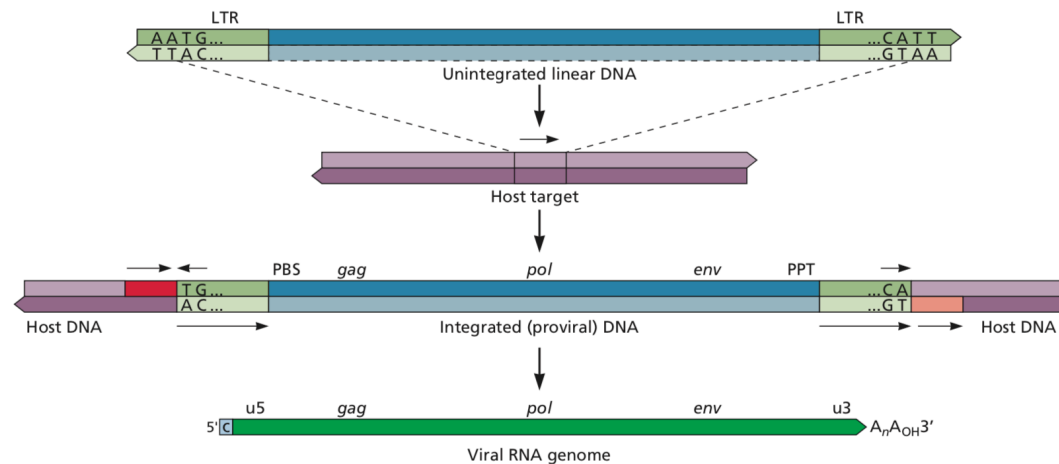


*Integration may occur at many sites on both strands, but is not random*

Site or region	% Integration <sup>a</sup>			
	Random	ASLV	MLV	HIV
Within genes	26	42	40	60–70
Transcription start sites	5	8	20	10

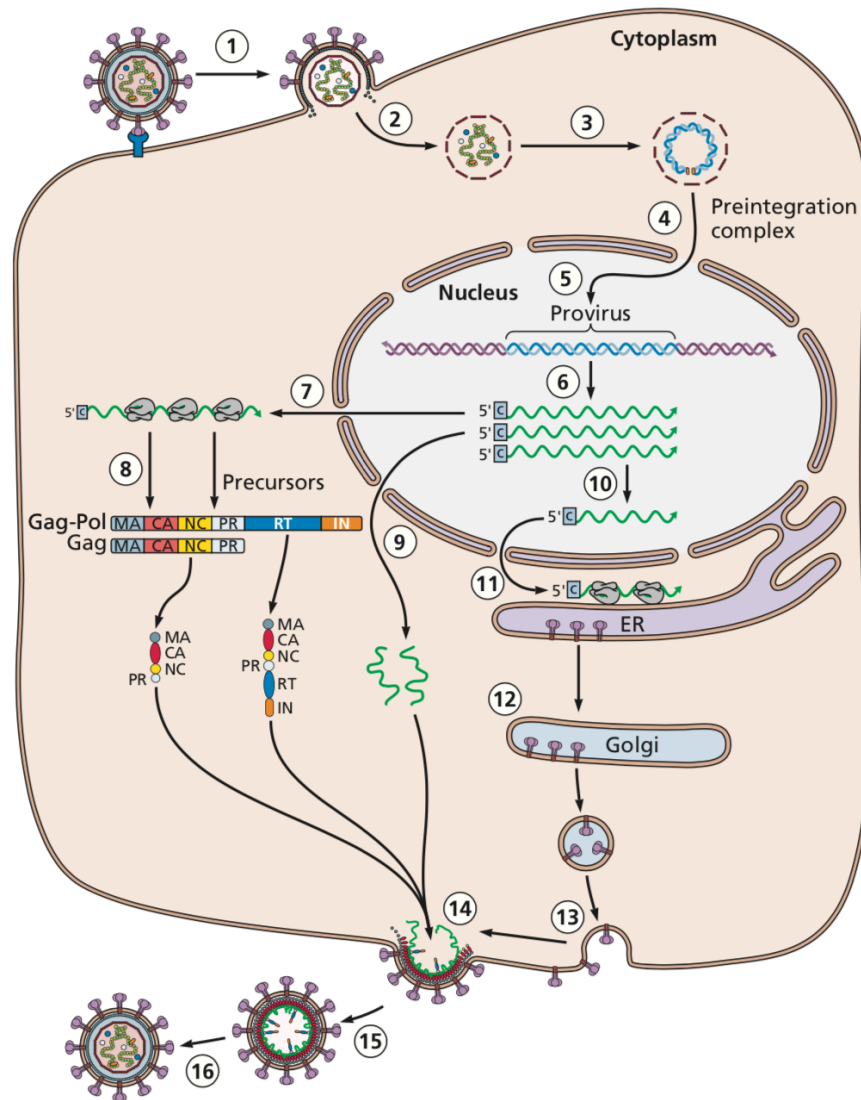




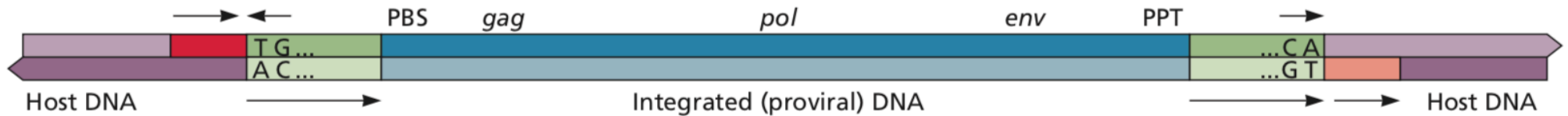


- One DNA produced from two RNAs by RT
- Strong promoter (the LTR) built during RT
- Proviral DNA directs the host transcription machinery to synthesize many copies of viral mRNA
- Viral mRNA is translated into viral proteins OR encapsidated into virus particles

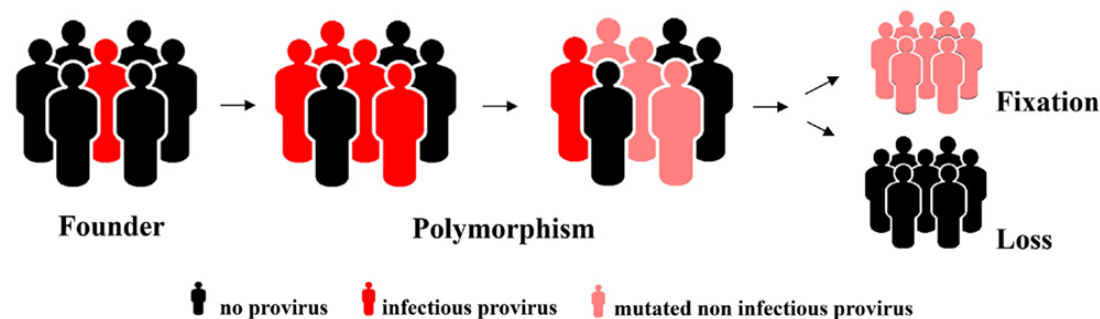
There is *no viral DNA replication* and *no viral RNA replication*



# Provirus is a permanent part of host genome

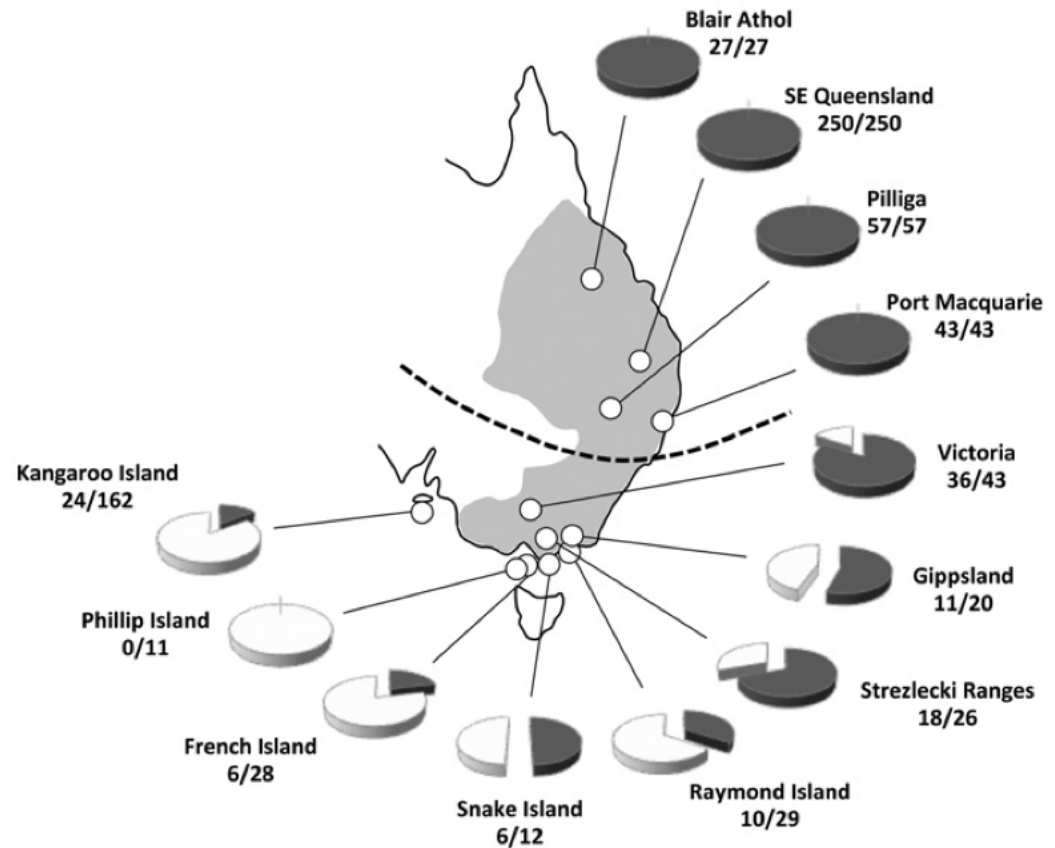


- No mechanism for precise excision of integrated provirus
- Only way out of genome is transcription by host RNA pol II
- Cell genomes are littered with ancient and modern retroelements - via endogenization



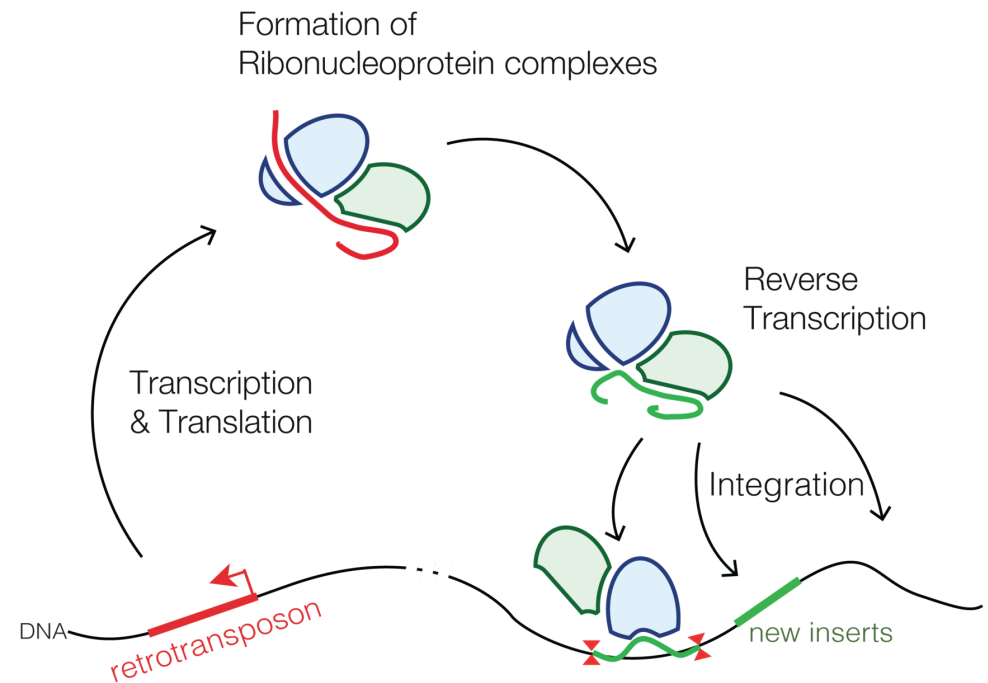
# Contemporary endogenization in Koalas

*~50,000 years ago, cross-species transmission from rodents*



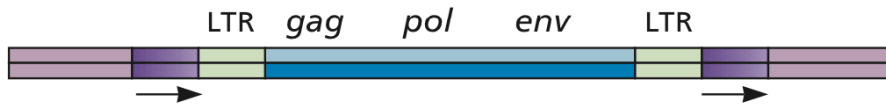
# Retroelements

- Sequences that move in the genome via RT
- Proviral DNA integrated into the germline = endogenous retroviruses, ERV
- Often replication-defective (all are defective in humans)
- ~42% of human genome comprises mobile genetic elements, including endogenous proviruses and other retroelements

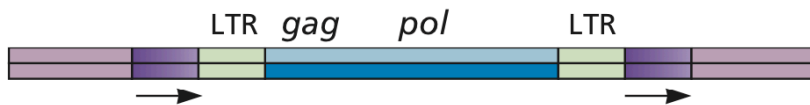


# Retroelements in the human genome

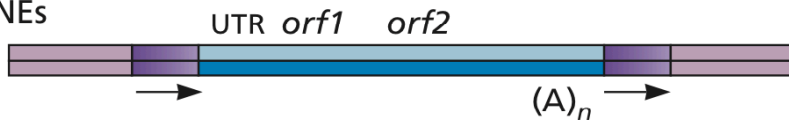
Endogenous retrovirus



Retrotransposons \*



LINEs



Processed pseudogenes



SINEs



} No RT

## Retroelements in the Human Genome 42.2%; $2.7 \times 10^6$

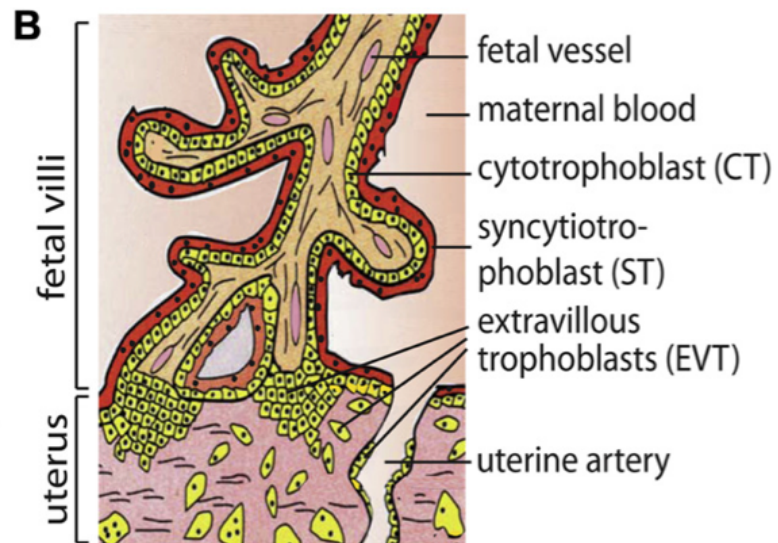
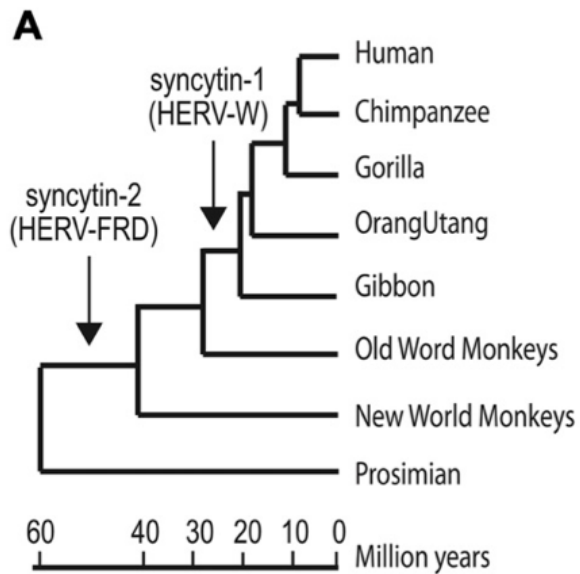
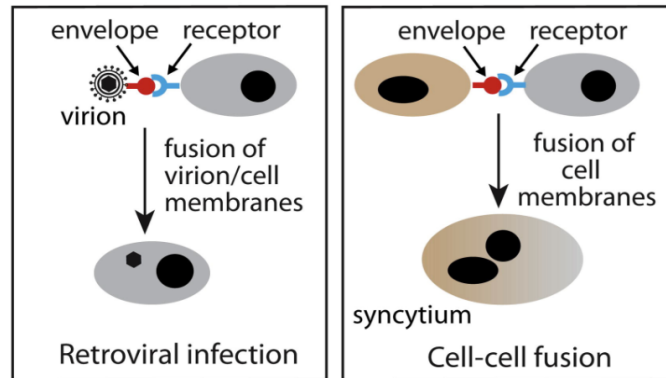
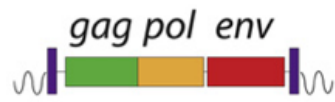
Non-LTR 33.9%; $2.4 \times 10^6$		LTR-Containing 8.3%; $0.3 \times 10^6$	
LINEs (L1)	16.9%	Endogenous Retroviral sequences (ERVs)	7.7%
(L2)	3.2		
SINEs (Alu)	10.6%	Others including Retrotransposons	0.6%
(MIR)	2.5		
Processed pseudogenes	<1.0		

<0.05% are active in human genome today

\*Likely progenitors of retroviruses



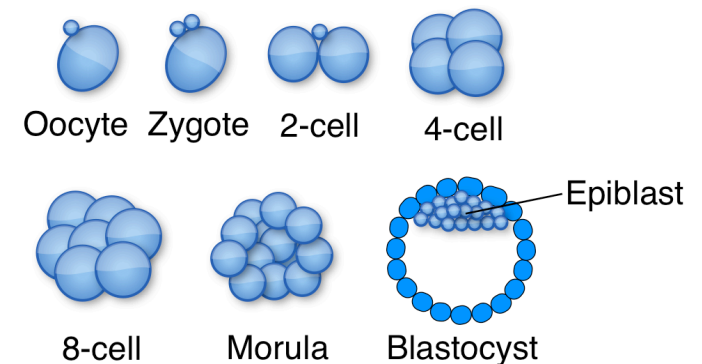
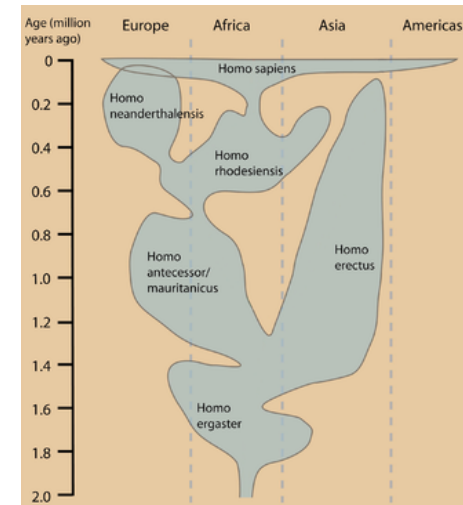
# Syncytins: Exapted retroviral env





# Retroviral influence on human embryonic development

- HERV-K, infected human ancestors ~200,000 years ago
- HERVs do not product infectious virus
- HERV-K mRNAs are produced during normal human embryogenesis
- From 8 cell stage to epiblasts
- Virus-like particles observed
- Induces an antiviral protein, IFITM1



## A retrovirus makes chicken eggshells blue



**Go to:**

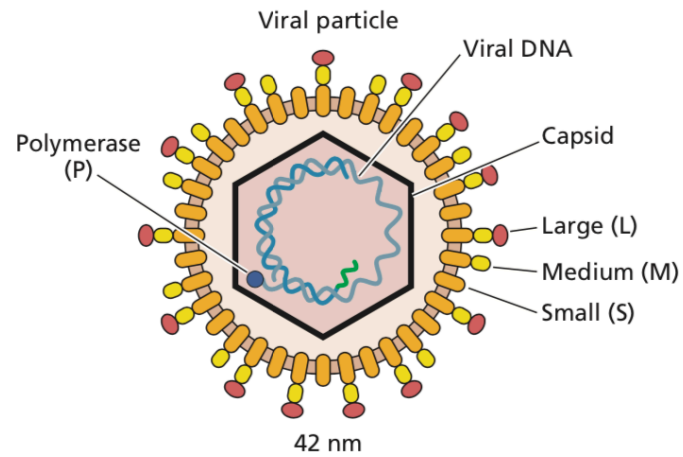
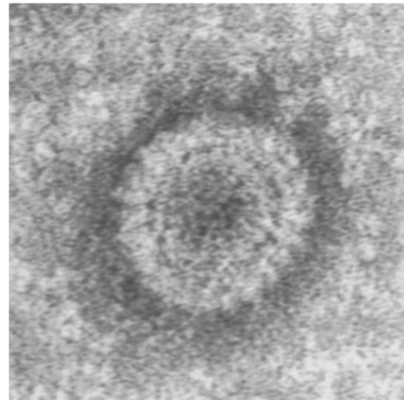
**[b.socrative.com/login/student](https://b.socrative.com/login/student)  
room number: virus**

Which of the following statements about retroelements is not correct?

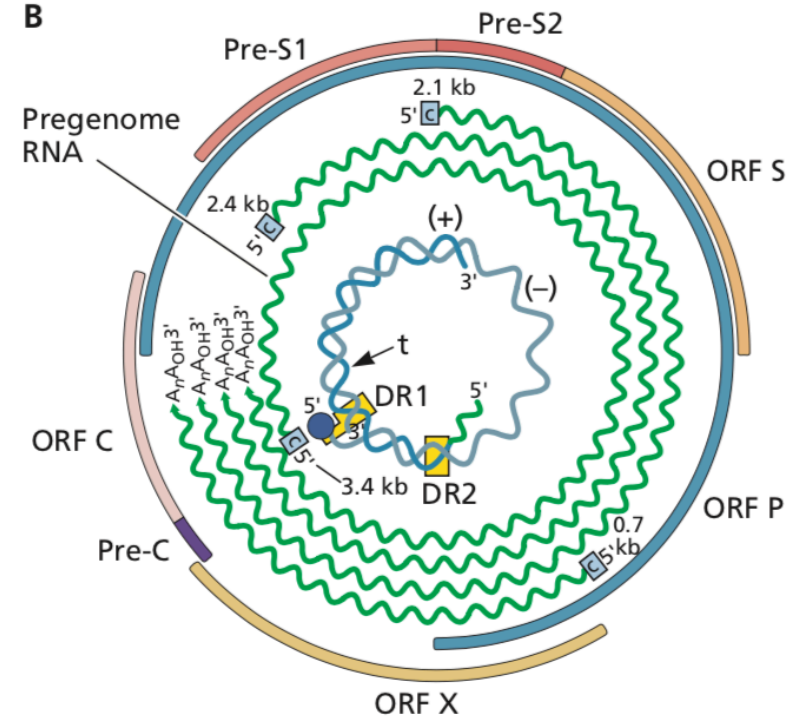
- A. There are many copies in eukaryotic genomes
- B. They are currently entering the Koala germline
- C. Those in the human genome produce infectious viruses
- D. They can be beneficial
- E. None of the above

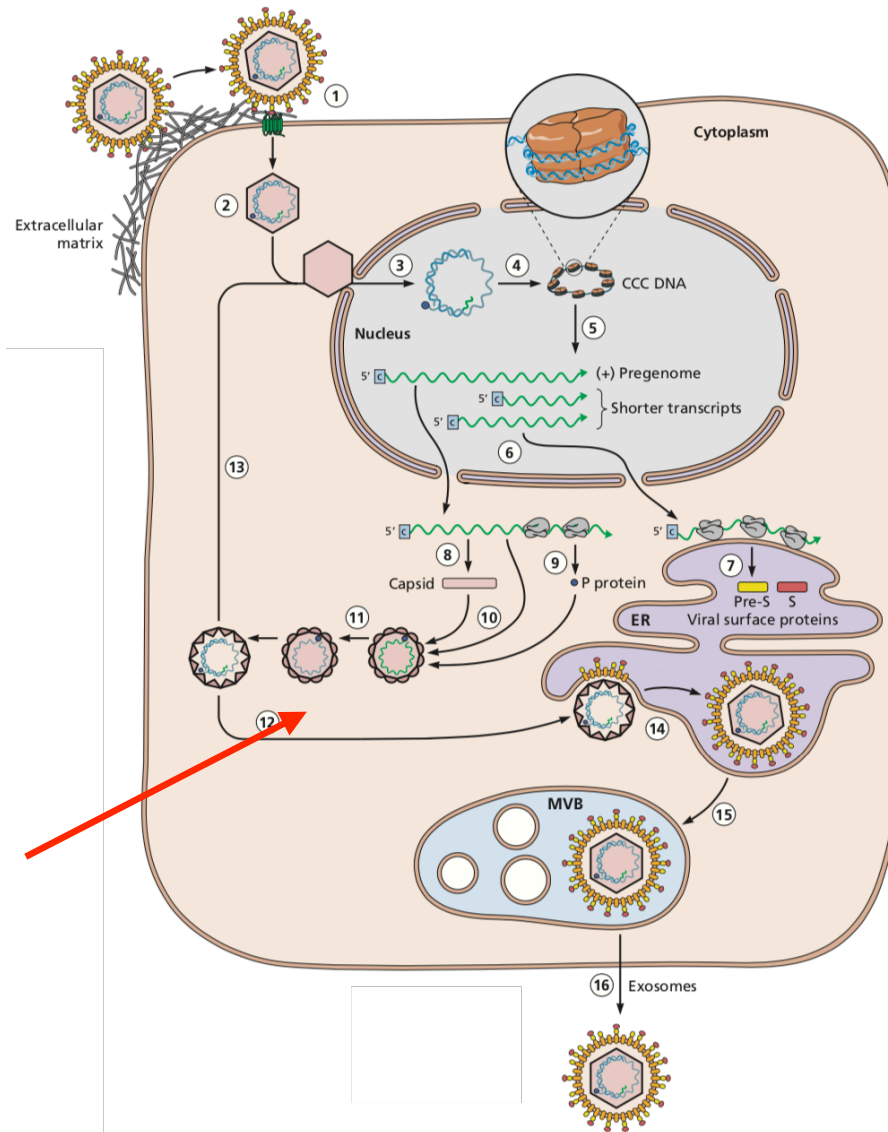
## ***Hepadnaviridae* - DNA viruses that encode RT**

**A**



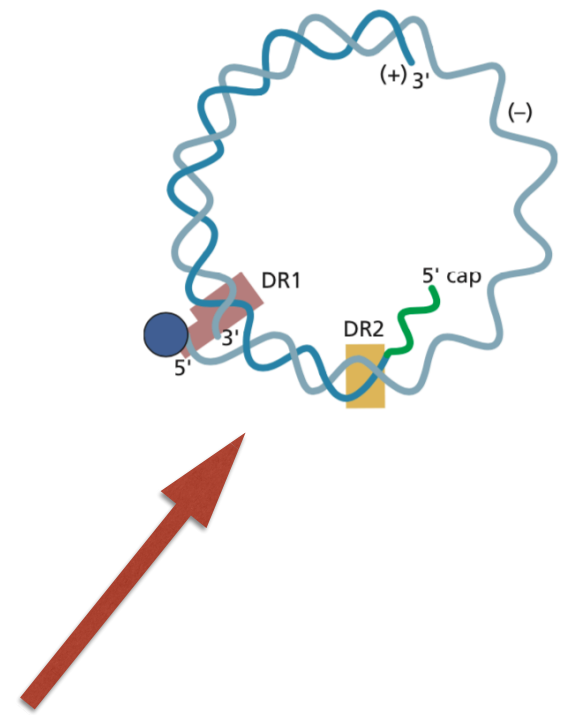
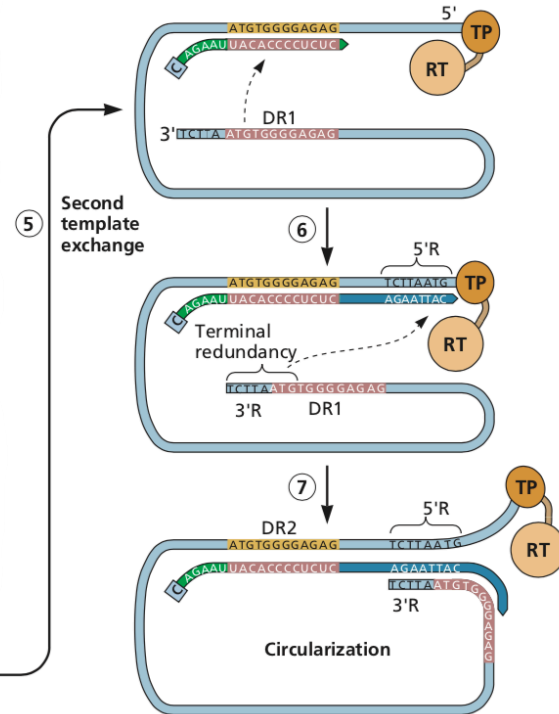
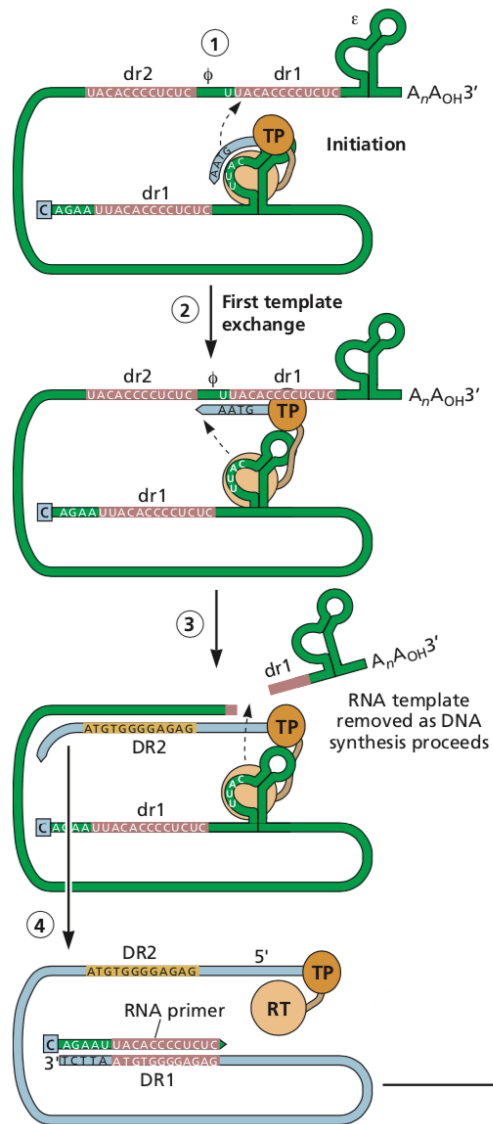
**B**

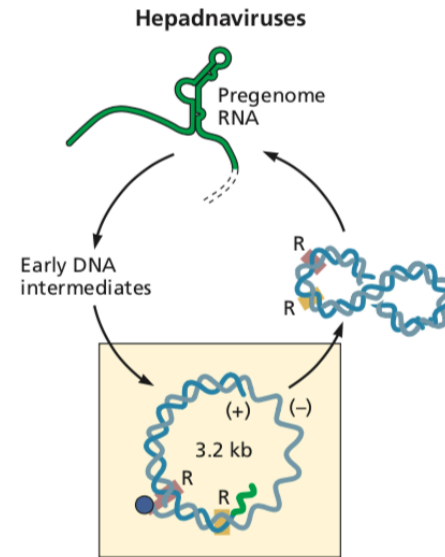
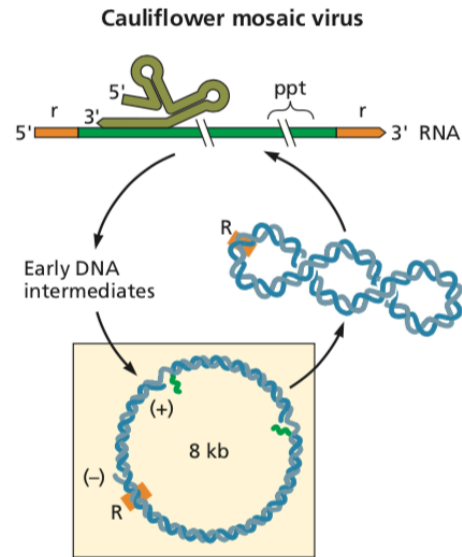




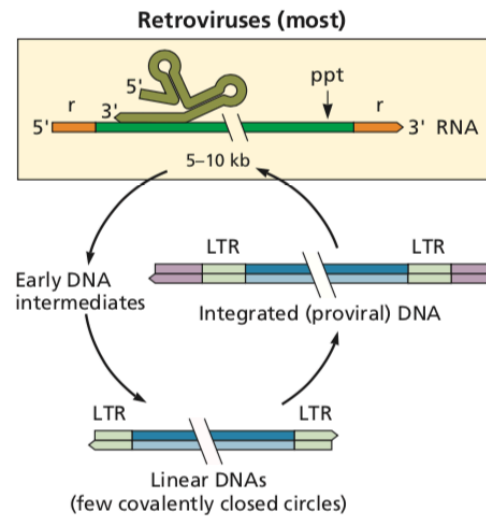
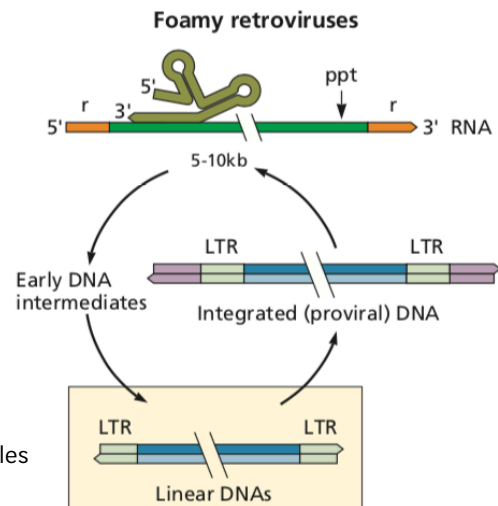
**No genome integration**

RT commences in the cytoplasm!

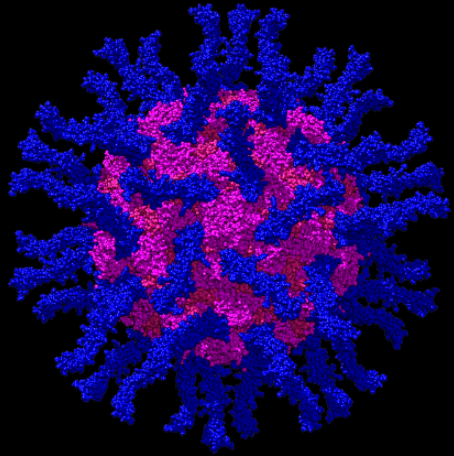




Yellow = in virus particle



DNA in 5-20% of virus particles



# **VIROLOGY LIVE**

**WITH VINCENT RACANIELLO**

**Next time: Assembly**