

Retroviruses

TRACO

September 26, 2016

Frank Maldarelli



HIV Dynamics and Replication Program

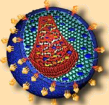
NCI-Frederick

NATIONAL
CANCER
INSTITUTE



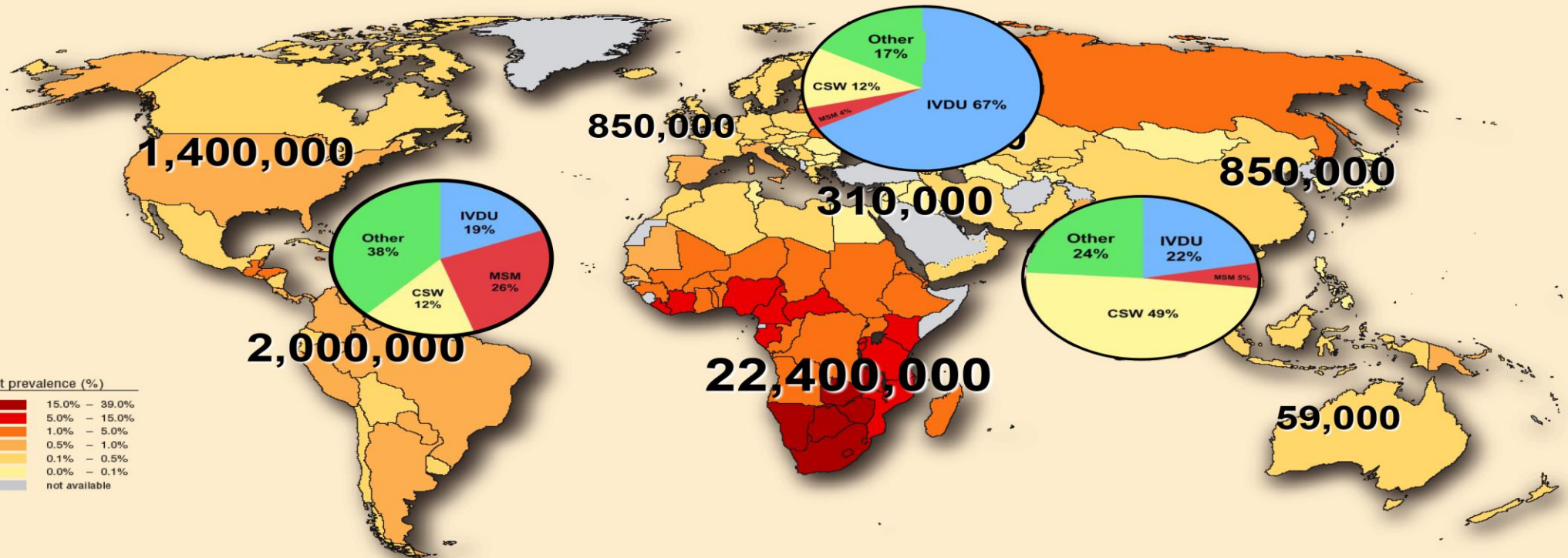
Retroviruses

- **Introduction**
- **Molecular Biology/Replication**
- **Retroviruses in Human Populations**
- **Emergence/Spread**
- **Lessons**



HIV-1 Pandemic: Risk

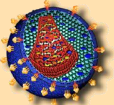
HIV-1 PANDEMIC: RISK



Total 35.3 million (range 32.2-38.8)

Retroviruses

•Molecular Biology/Replication



HIV Dynamics and Replication Program

NCI-Frederick

Retroviruses

Retroviruses

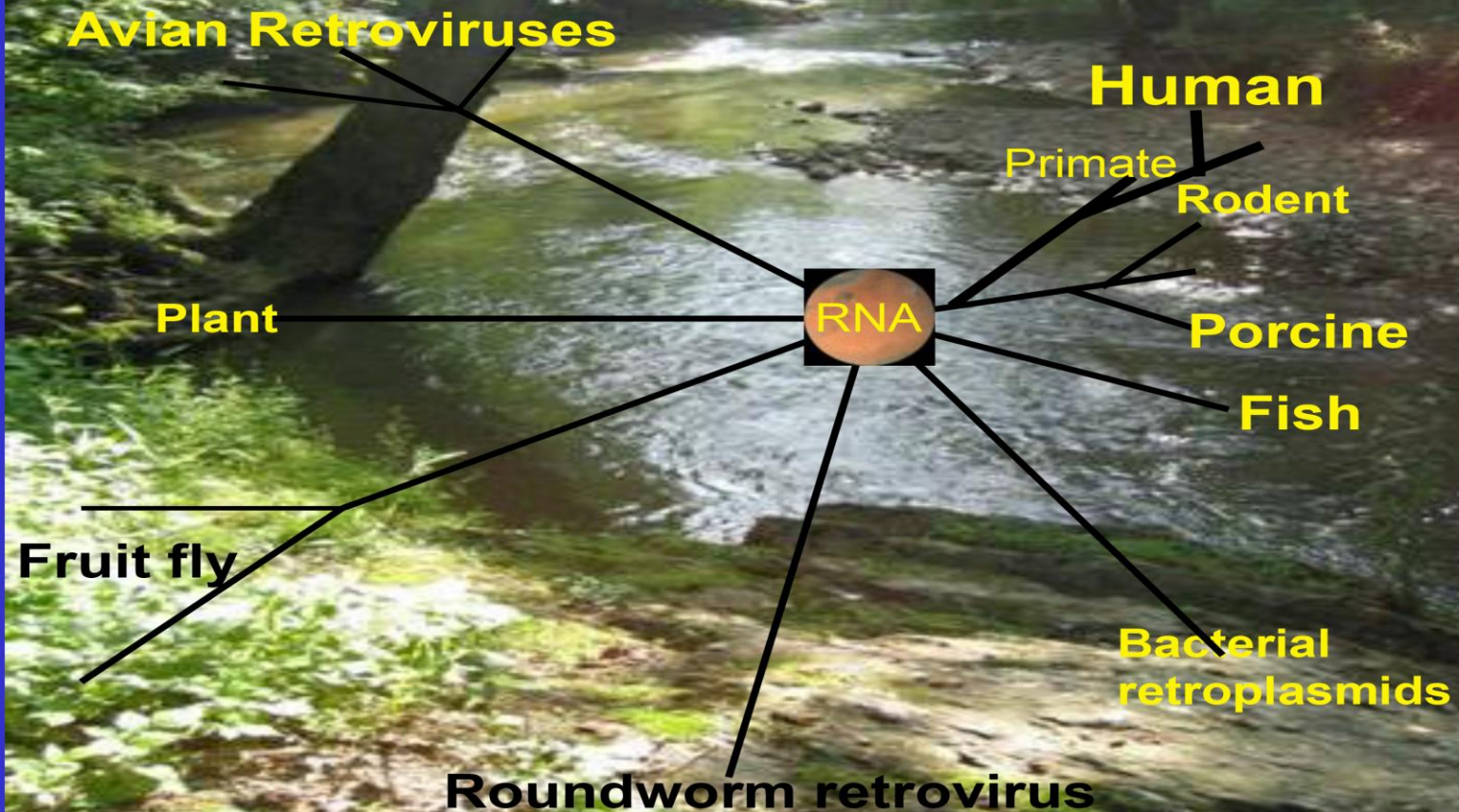
A group of RNA viruses that replicate via a DNA intermediate using Reverse Transcriptase.

A different paradigm for replication

Transition from RNA World?

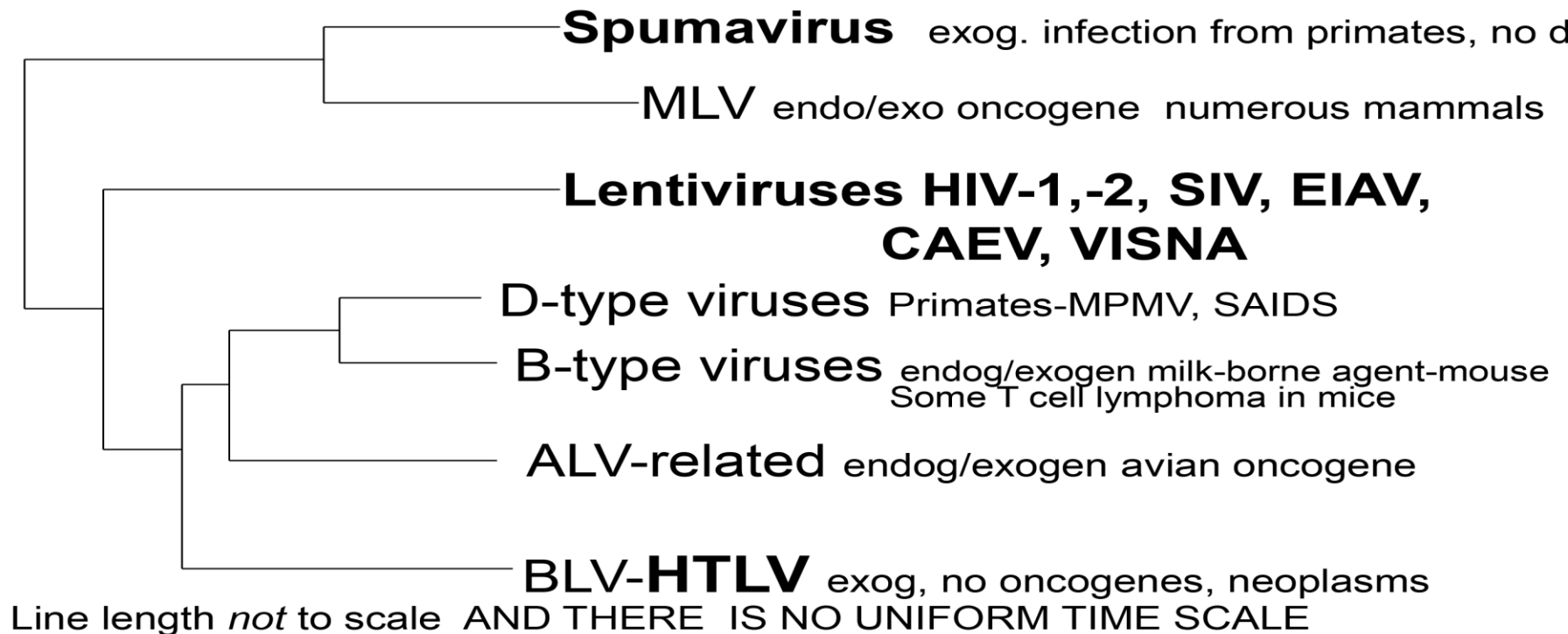
Retroelements

Reverse Transcriptase and Retroelements are all around you



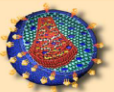
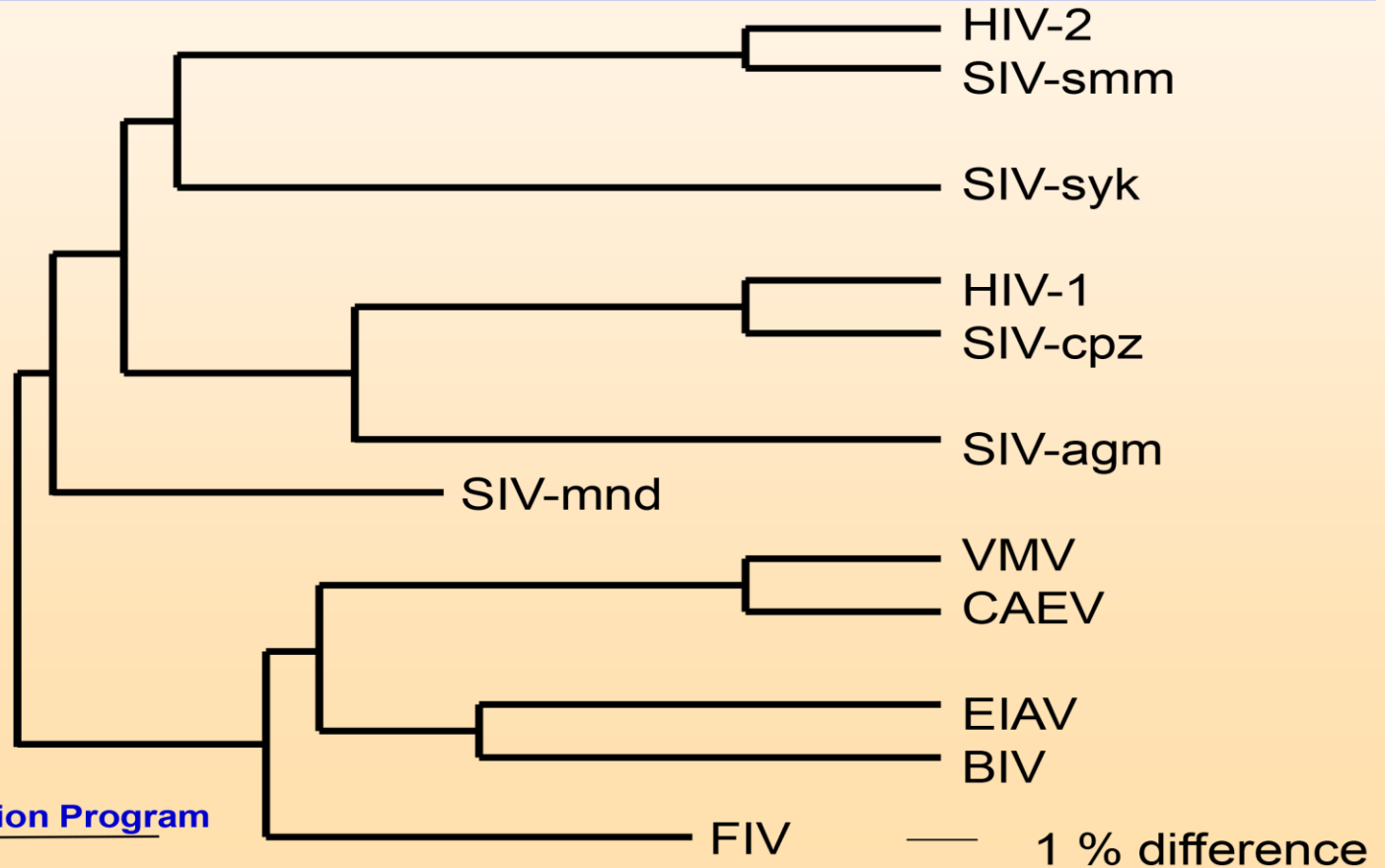
Retrovirus classification

Retroviruses Classification by RT Sequence into Seven Families



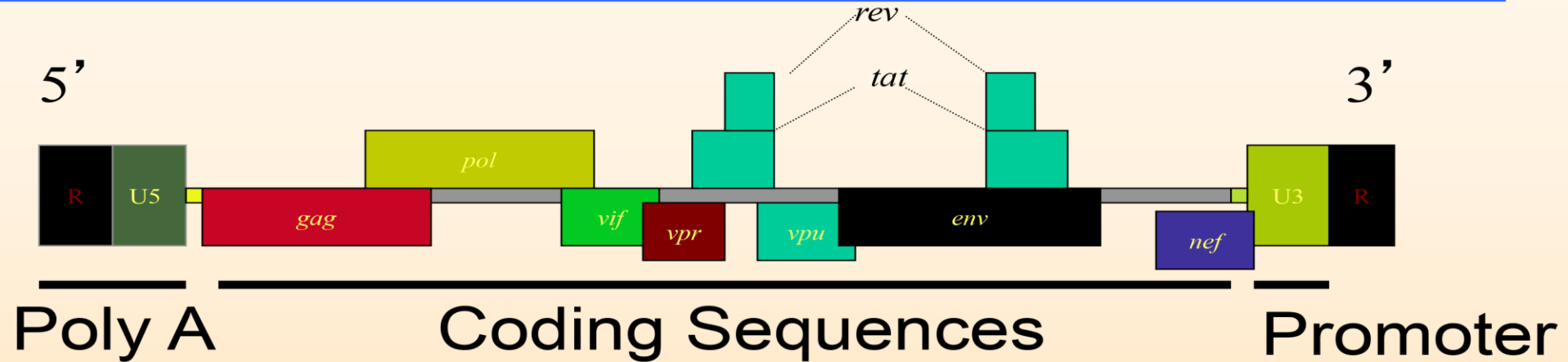
Lentivirus Relationships

Lentivirus Relationships

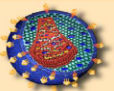


Retrovirus conventions

Retroviruses Conventions



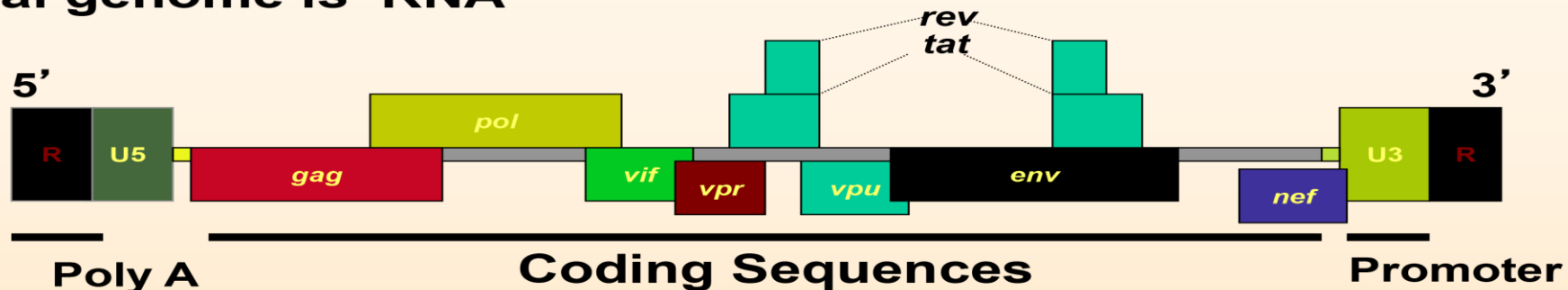
Names of genes in lower case *italics*, e.g., *pol*, *env*
Protein gene products are capitalized, e.g., Reverse Transcriptase, Gp120



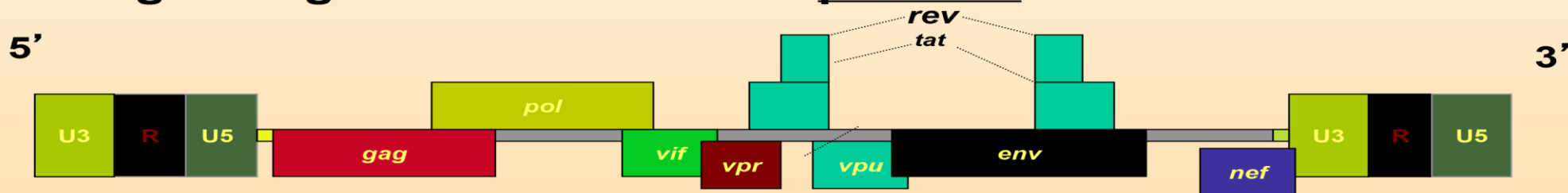
Retrovirus

Retroviruses Conventions

The viral genome is RNA

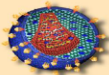


The integrated genome is called the provirus



Names of genes in lower case *italics*, e.g., *pol*, *env*

Protein gene products are capitalized, e.g., Reverse Transcriptase, Gp120



Retroviruses

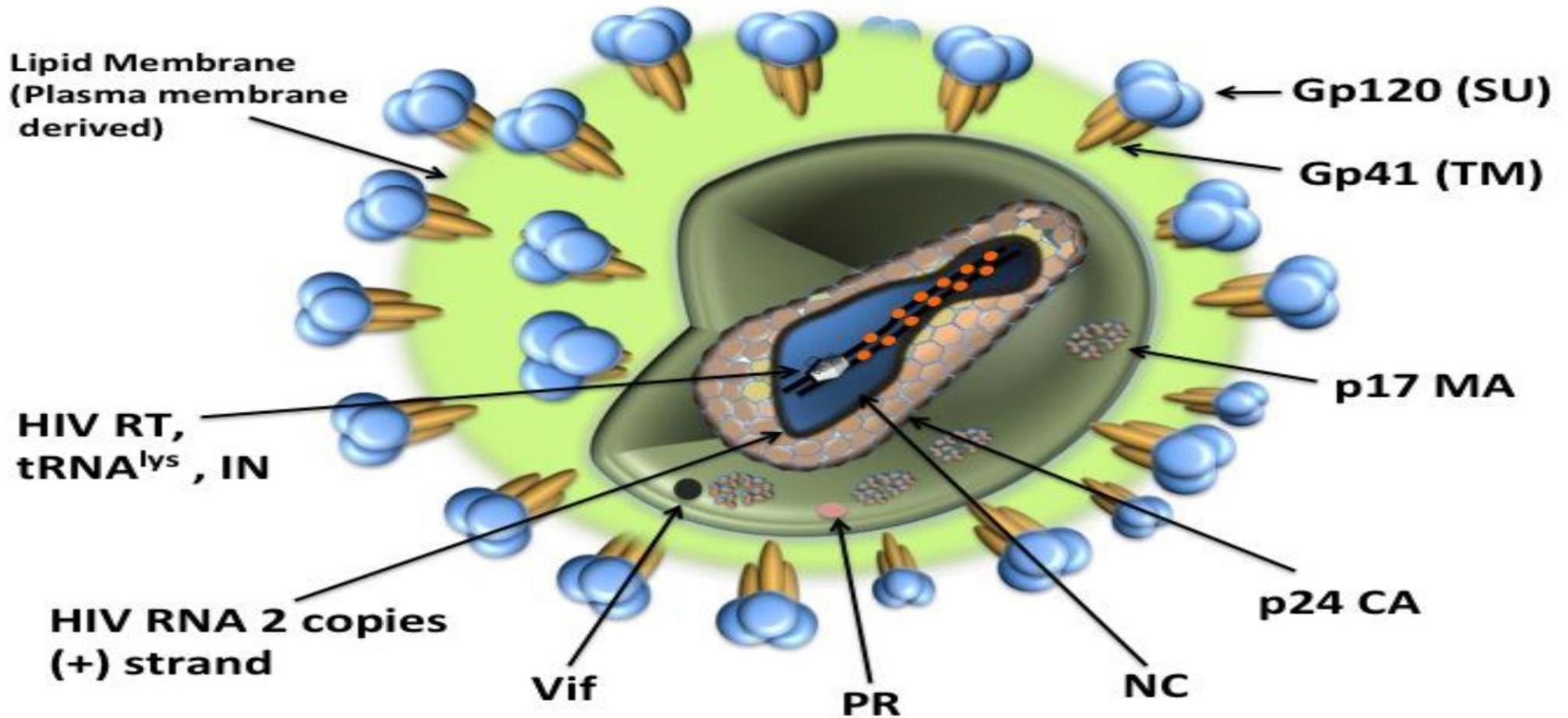
Retroviruses Glossary

- *gag: group antigen*
- *pol: polymerase*
- *env: envelope*
- *tat: Transactivator*
- *rev: Regulator of Expression of Virion proteins*
- U3: unique sequence in 3' region
- U5: Unique sequence in 5' region
- R: Repeat sequence
- PBS Primer binding site for initiation of RT
- Ppt: polypurine tract primer for RT
- TAR: Tat activating sequence
- RRE: Rev responsive element
- Provirus: copy of retrovirus that is integrated into host genome



HIV virion

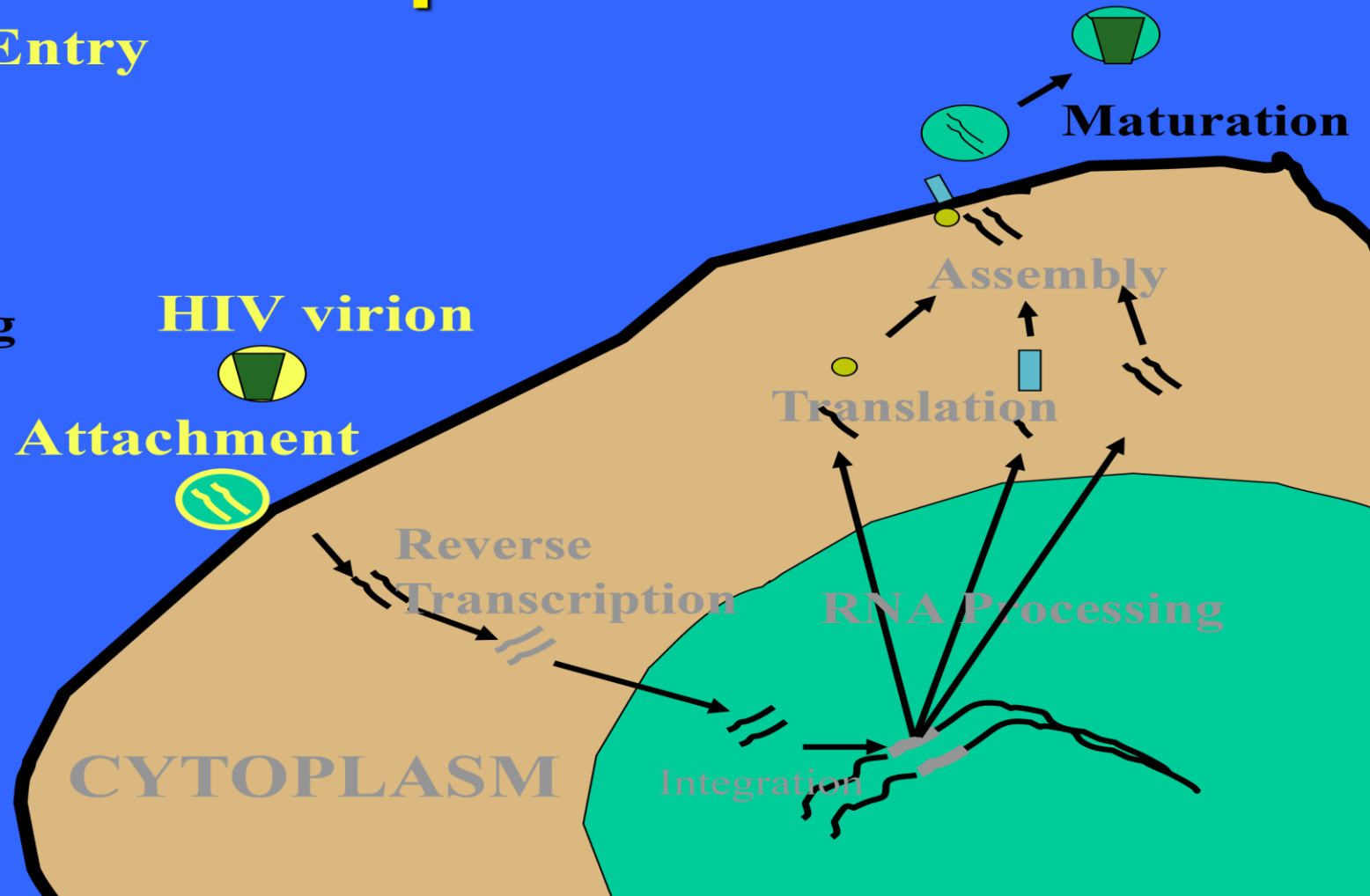
HIV Virion



HIV replication

HIV Replication

- **Attachment/Entry**
- **Reverse Transcription**
- **Integration**
- **Transcription**
- **RNA Processing**
- **Translation**
- **Assembly**
- **Maturation**



HIV attachment and Entry

HIV Attachment and Entry

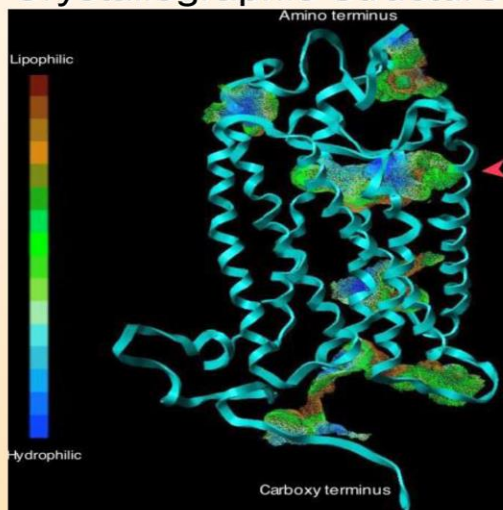
- **Virus Factors**
 - Attachment: Env glycoprotein gp120
 - Entry: Env glycoprotein gp41
- **Host Cell Factors**
 - Receptor
 - CD4
 - Co-receptor (major)
 - CXCR4
 - CCR5



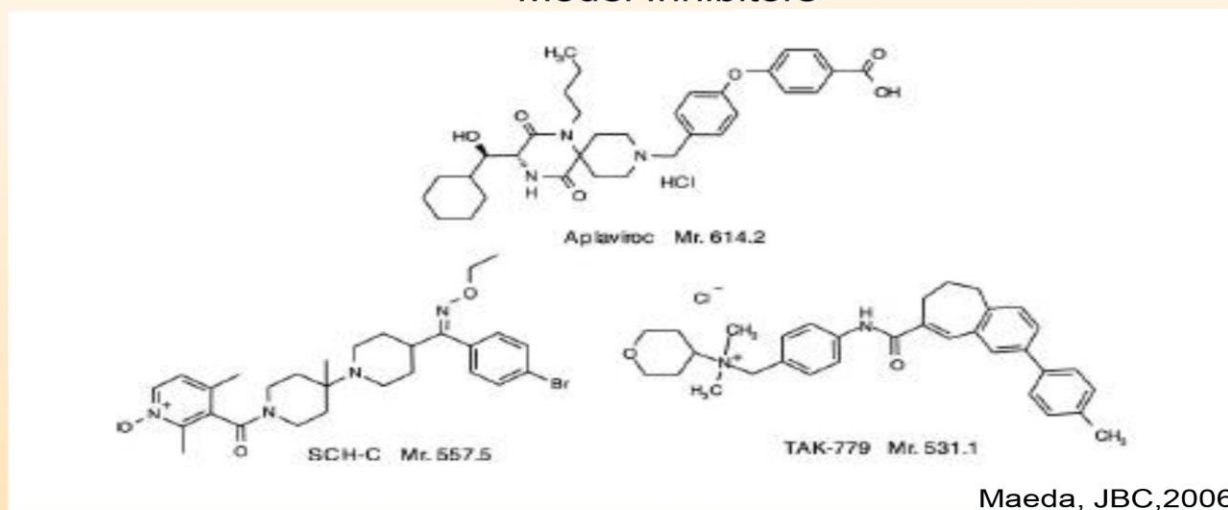
HIV coreceptor blockade

HIV Coreceptor Blockade

Crystallographic Structure



Model Inhibitors

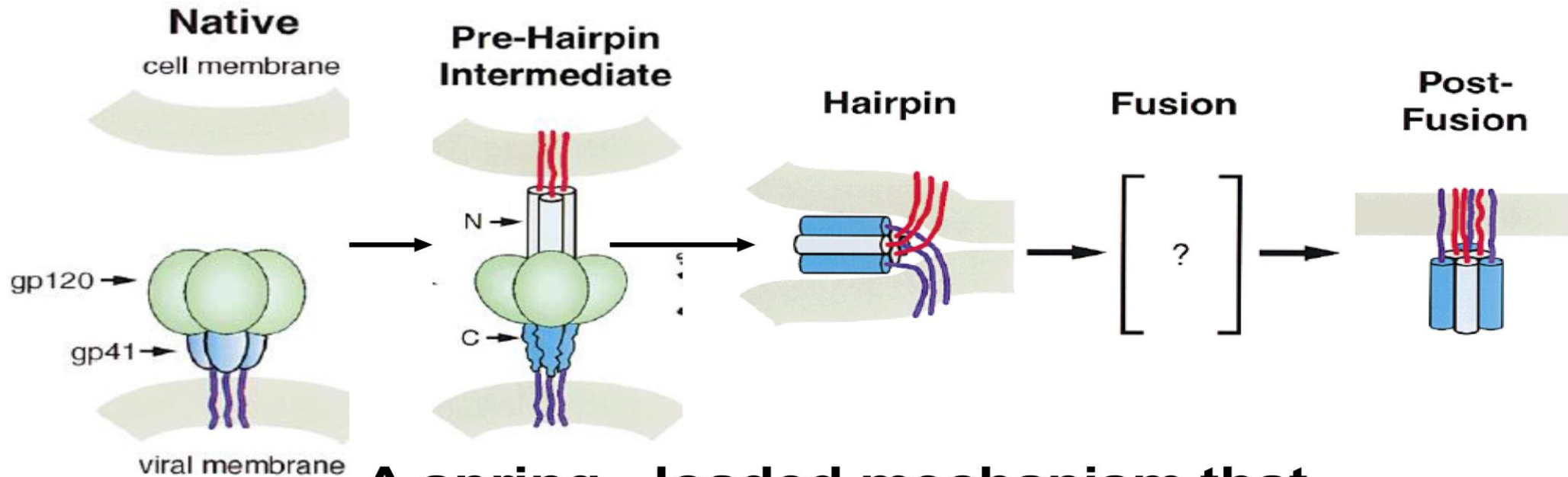


- Multiple binding domains predicted
 - Binding disrupts structure generally
 - Does not require blocking CCR5-gp120 interaction
 - Potential for simultaneous inhibition
- Resistance emerges by reducing affinity for drug



HIV Fusion-Gp41

HIV Fusion-Gp41



A spring - loaded mechanism that drives the membranes together to overcome a high energy barrier to

HIV gp41

HIV gp41

Coiled coil in a hydrophobic cavity

Spring-loaded mechanism

T-20

D amino acid interacting (PS Kim)

8582 JI ET AL.

J. VIROL.

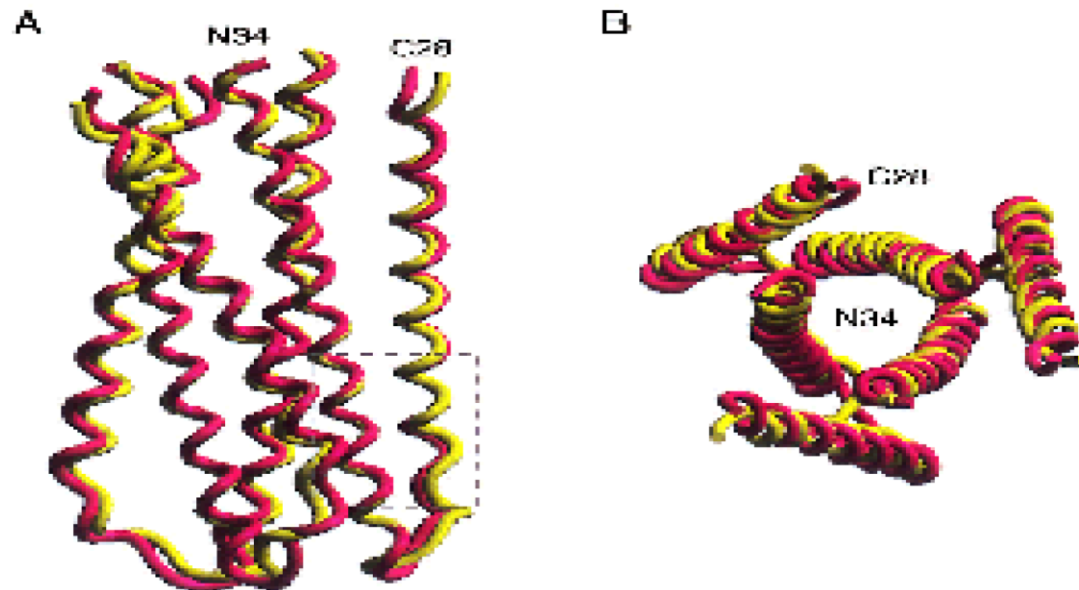


FIG. 4. Overall views of the mutant L568A and W571R cores. (A) Side view. The amino termini of N34 and the carboxyl termini of C28 are at the top. Helices in L568A (yellow) and W571R (pink) were used for the superposition. The bottom of the central N34-coiled-coil surface contains three symmetry-related hydrophobic cavities (one is outlined by the box). (B) View from the top, looking down the threefold axis of the trimer. The same color coding as in panel A is used. Figures were

Uncoating

HIV Replication

• Attachment/Entry

y

• Reverse
Transcription

• Integration

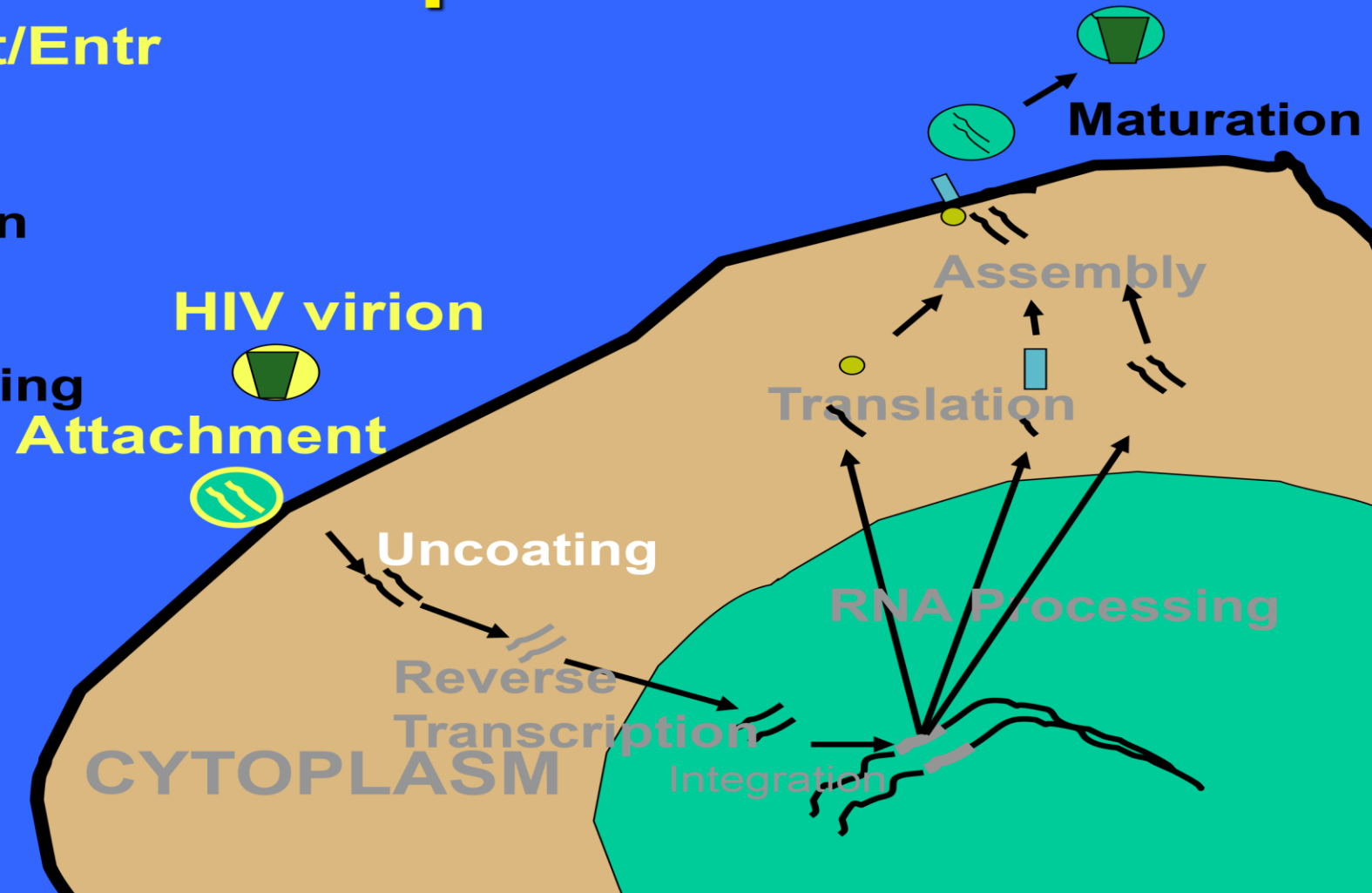
• Transcription

• RNA Processing

• Translation

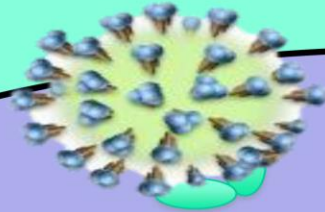
• Assembly

• Maturation

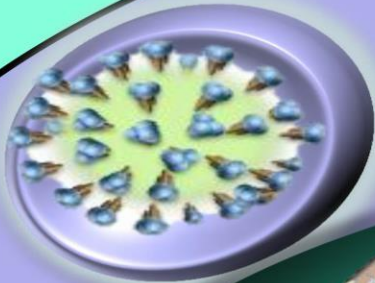


Uncoating

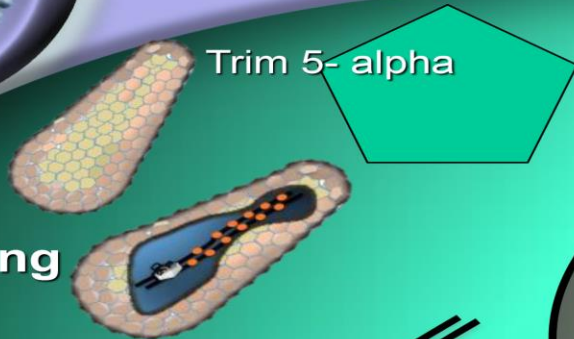
Attachment



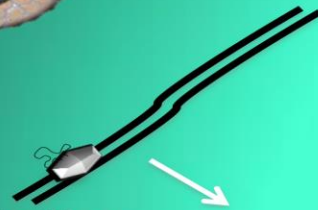
Fusion



Uncoating



Reverse transcription



HIV DNA

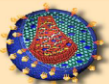
Integration into Host Genome



Post-Entry Events

HIV Post – Entry Events

- **Uncoating is a fundamental step in virus replication**
 - **Restricts replication**
 - **Source of host range restriction**
- **Requires interactions between viral and cellular factors**
- **Virus**
 - **Gag**
- **Cell**
 - **Trim 5 – alpha**



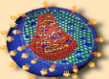
Post entry events

HIV Post – Entry Events

Host Trim5 Alpha

VIRUS

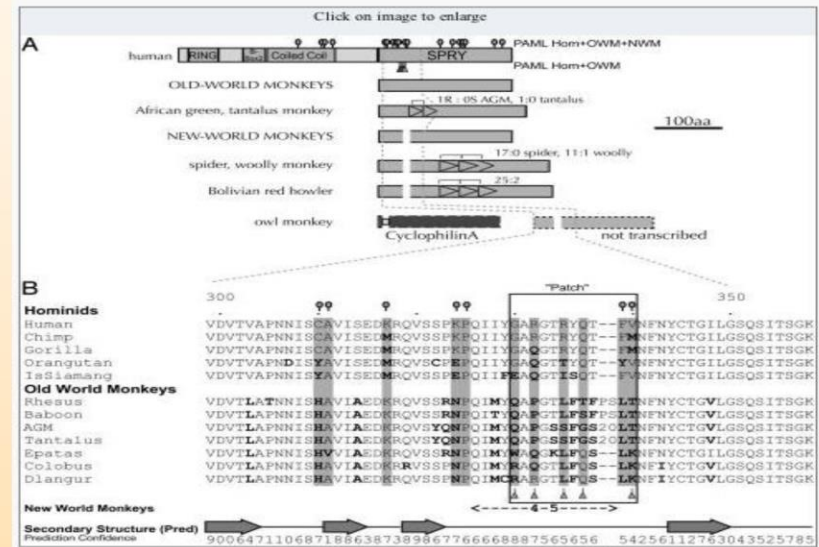
	Human	Chimp	Monkey
HIV	Infection	Infection	NO INFECTION
SIV Chimp	INFECTION	Infection	Poor infection
SIV Monkey	INFECTION	Poor infection	Infection



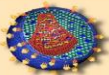
Trim 5-alpha

Positive Selection in Trim 5-alpha

- Trim 5 alpha undergoes genetic change faster than many genes
- Working hypothesis
 - human populations undergo waves of pandemics
 - Humans that survive have trim 5alpha variant that excludes infection



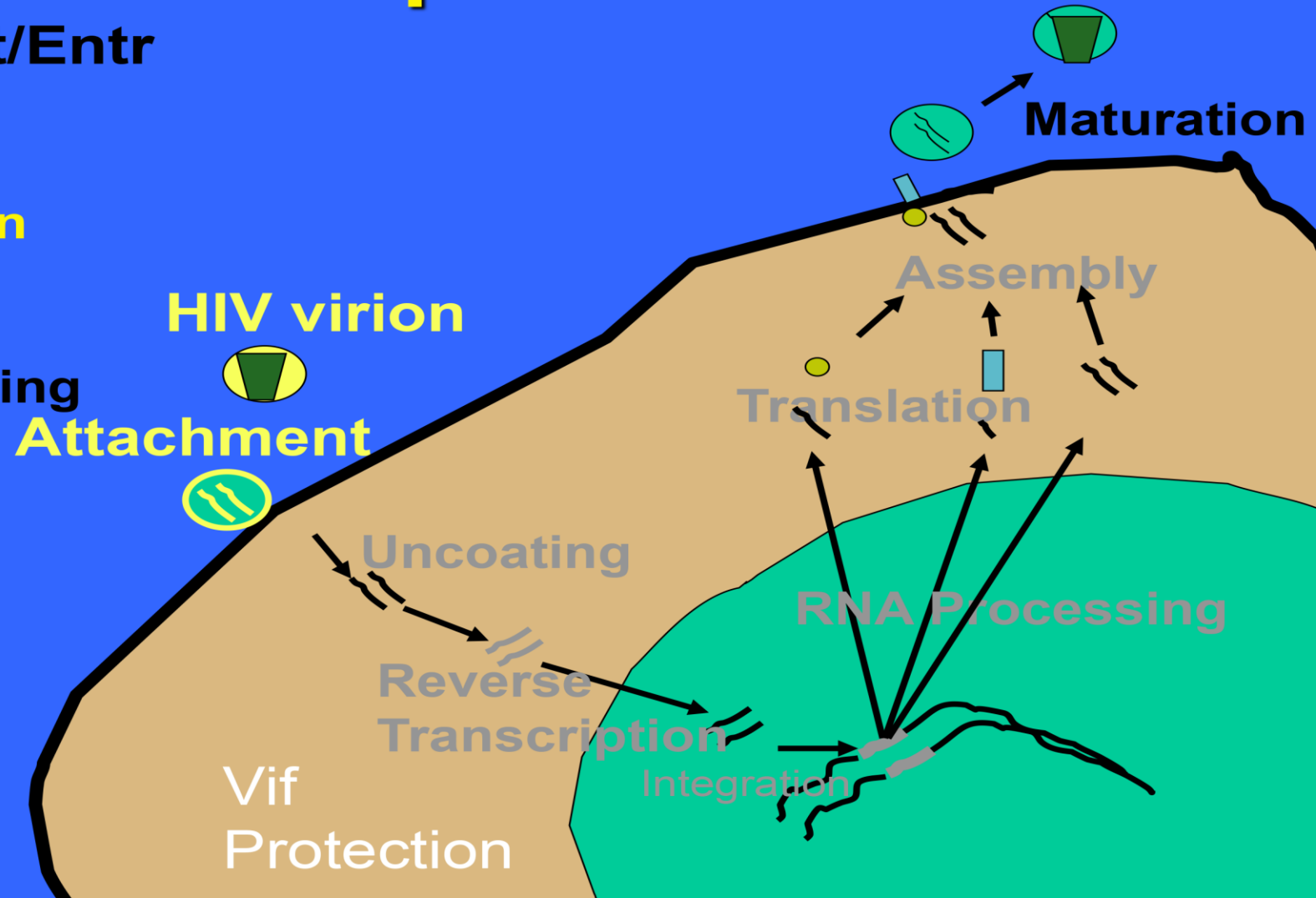
Generals are Always Fighting The Last War
 Evolution can solve this problem but it will take time



HIV replication

HIV Replication

- Attachment/Entry
- Reverse Transcription
- Integration
- Transcription
- RNA Processing
- Translation
- Assembly
- Maturation



Enzymatic Activities

Reverse Transcriptase Enzymatic Activities

- **RNA-dependent DNA Polymerase**
- **RNase H**
- **DNA-dependent DNA Polymerase**
- **Error rate on order of 1-4 / 100,000 bases synthesized**
- **Recombination occurs during reverse transcription permitting reassortment of sequences**
- **Replication rapid and error prone**

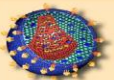
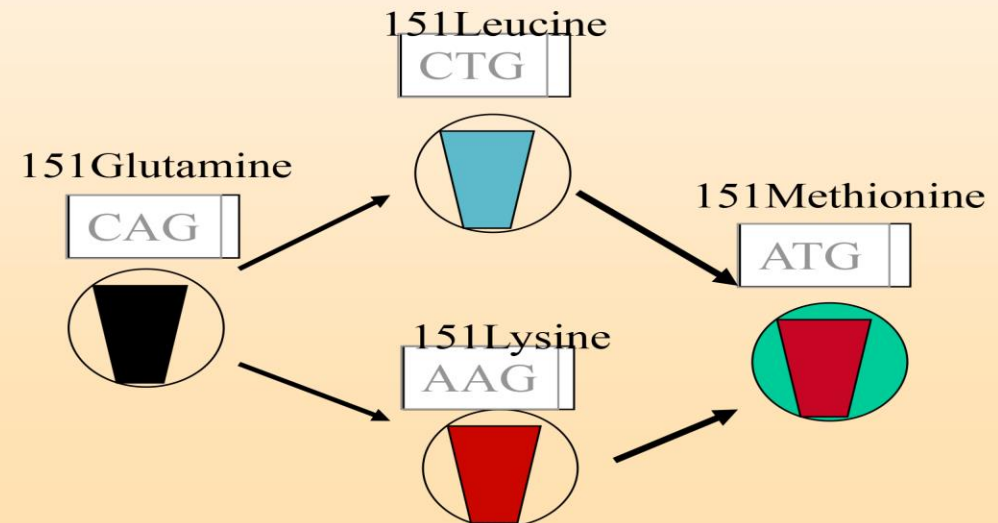
MUTANTS ARE LIKELY TO EXIST PRIOR TO THE THERAPY



Error-Prone replication

Error-Prone HIV Replication is a Pathogenic Determinant

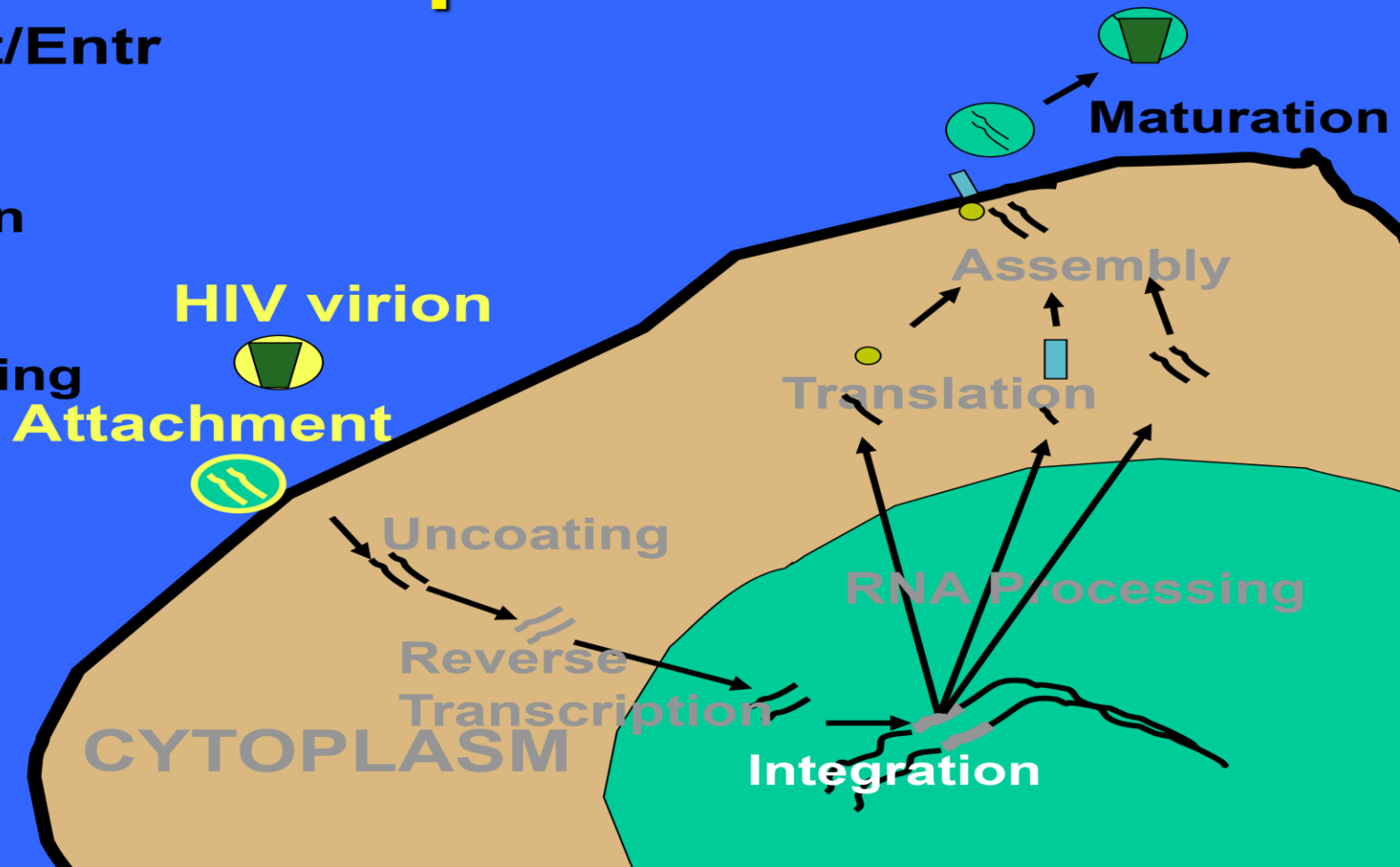
- Each round of HIV replication generates numerous mutants.
- The ability of the mutants to replicate (viral “fitness”) may vary greatly.
- The virus population can respond rapidly to a selective pressure



Integration

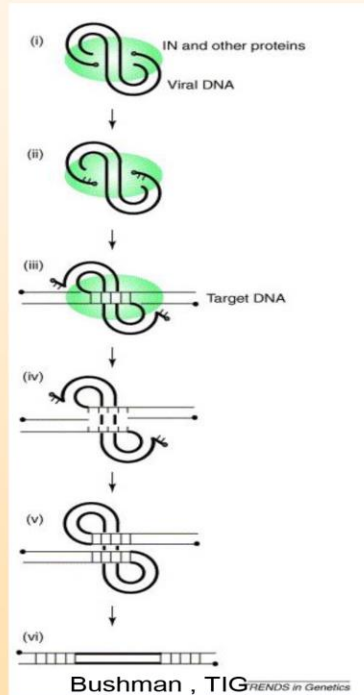
HIV Replication

- Attachment/Entry
- Reverse Transcription
- **Integration**
- Transcription
- RNA Processing
- Translation
- Assembly
- Maturation

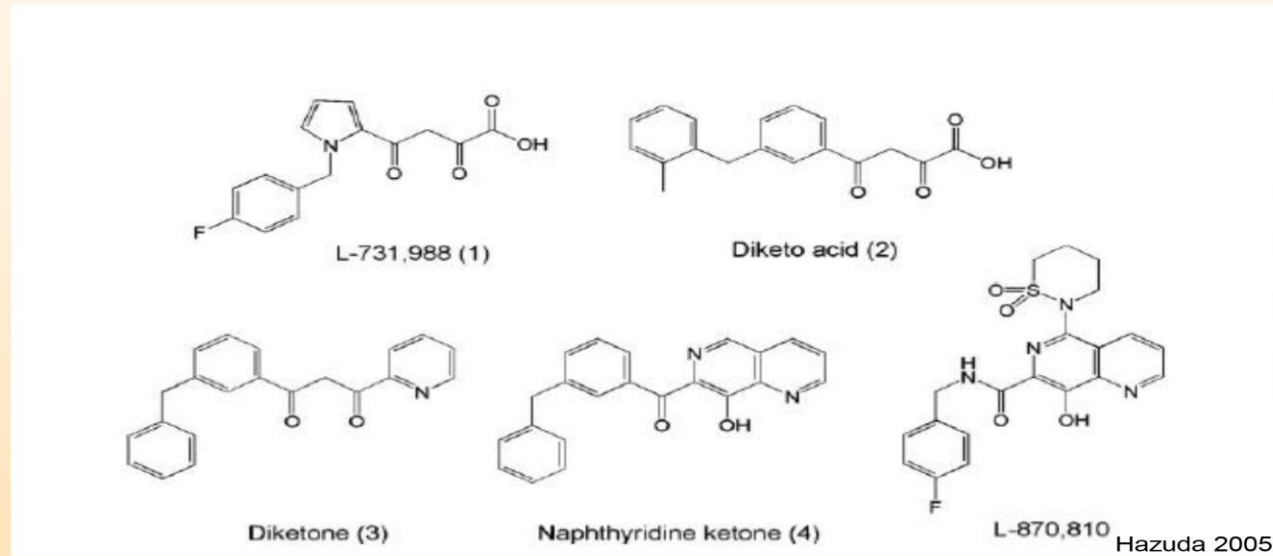


Integration

Integration



Multistep reaction



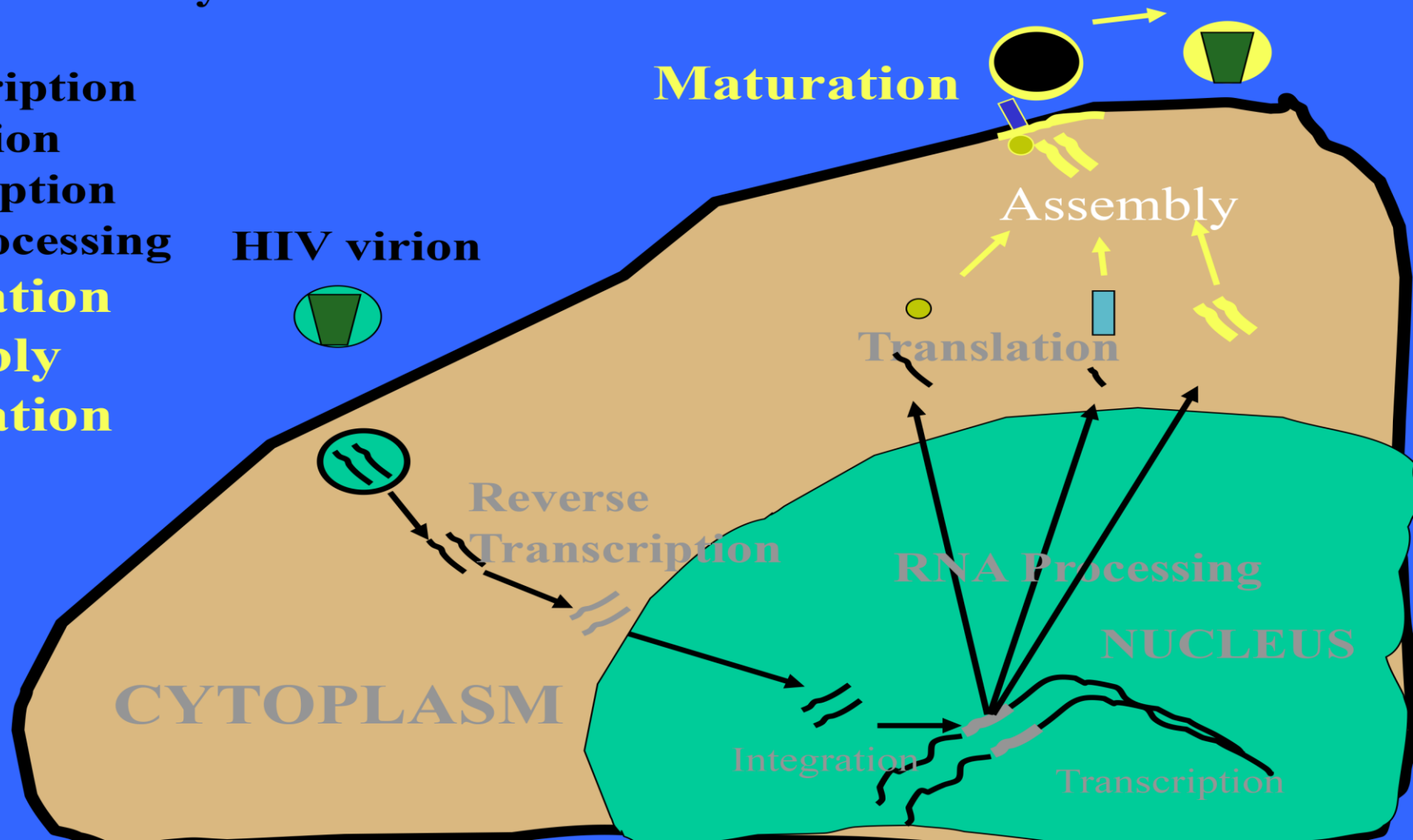
Strong inhibitors



Maturation

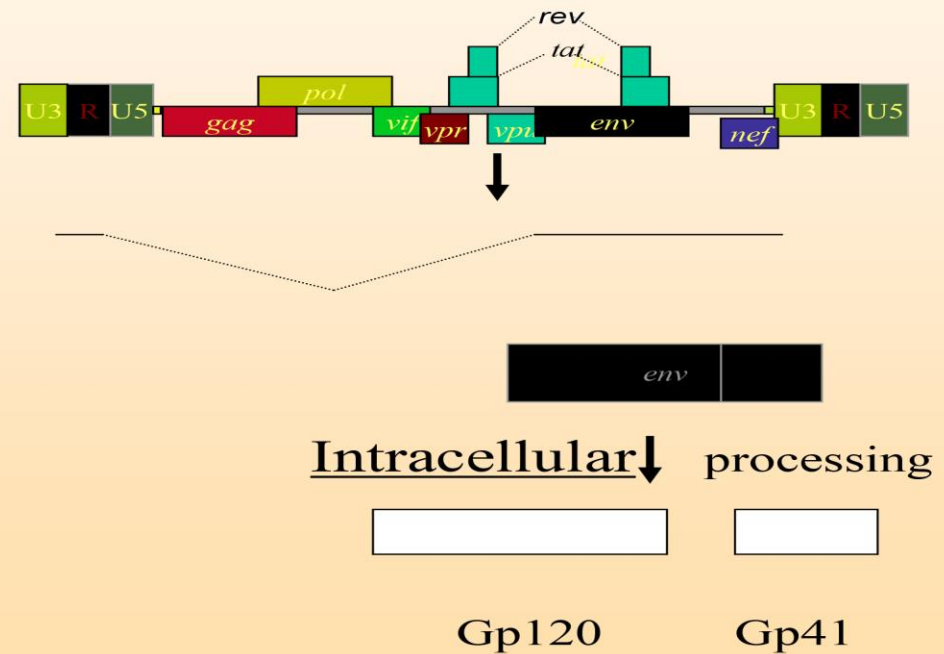
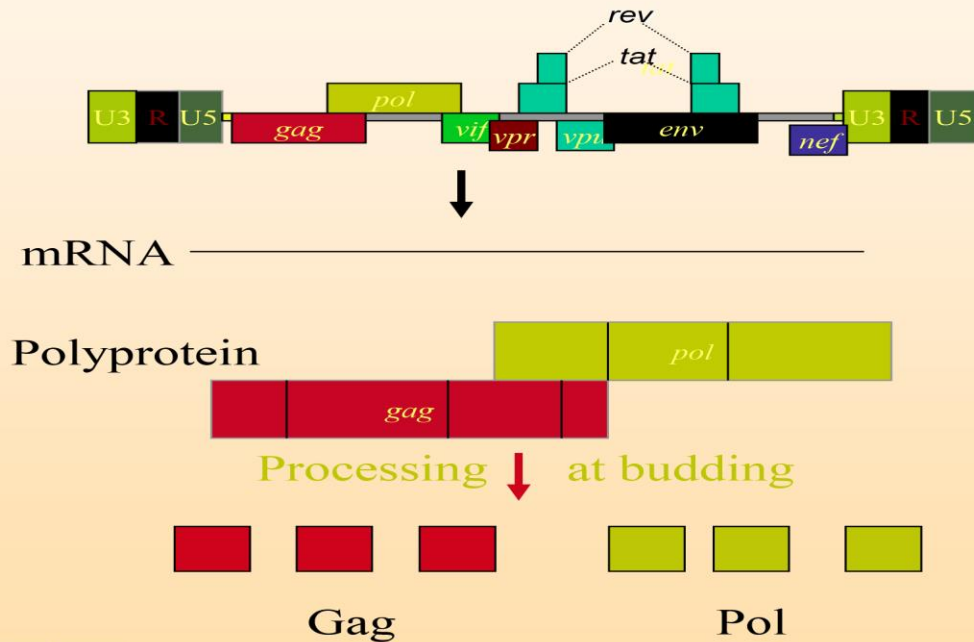
HIV Replication

- Attachment/Entry
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- Transcription
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- Maturation



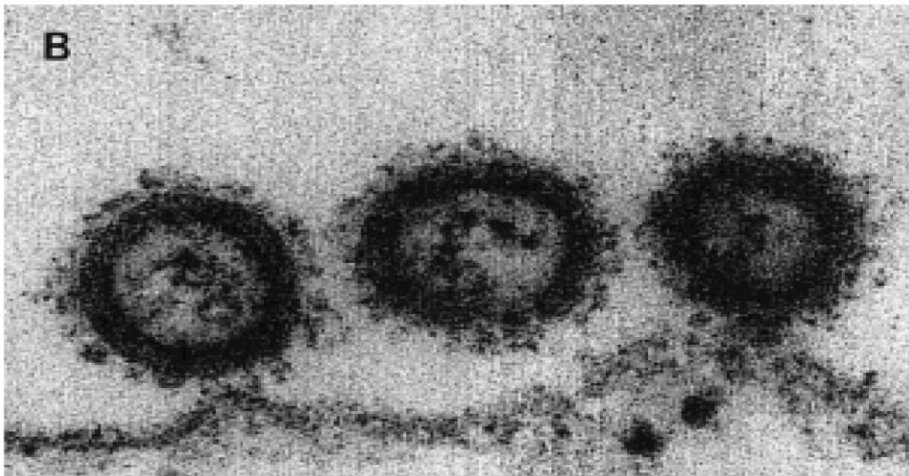
Translation

Translation of HIV *gag/pol* and *env* Paradigm: Process Polyprotein Precursors



HIV Particle Maturation

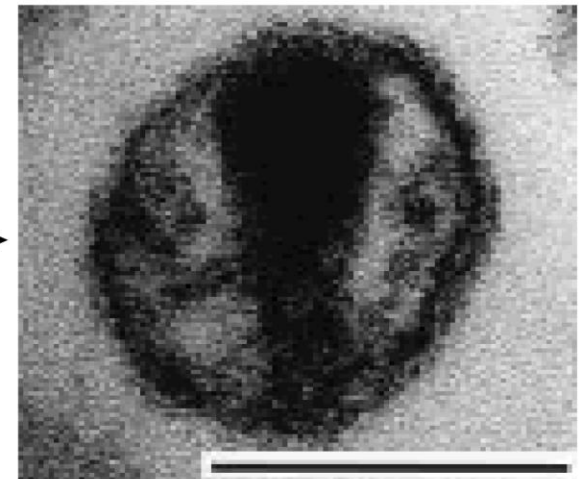
HIV Particle Maturation



Immature Particle
Noninfectious



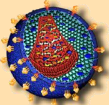
HIV
Protease



Mature Particle
Infectious

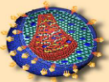
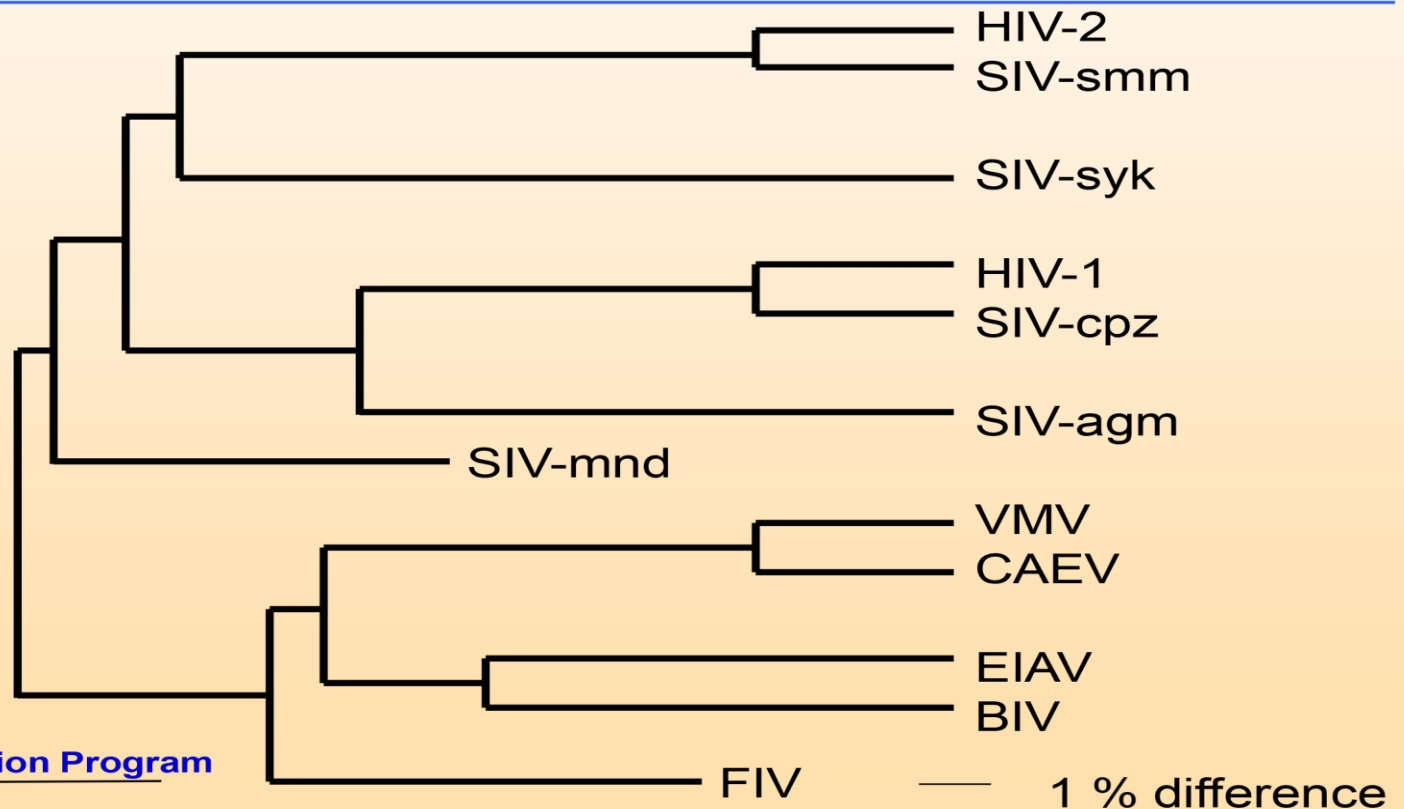
Retroviruses

- **Retroviruses in Human Populations**



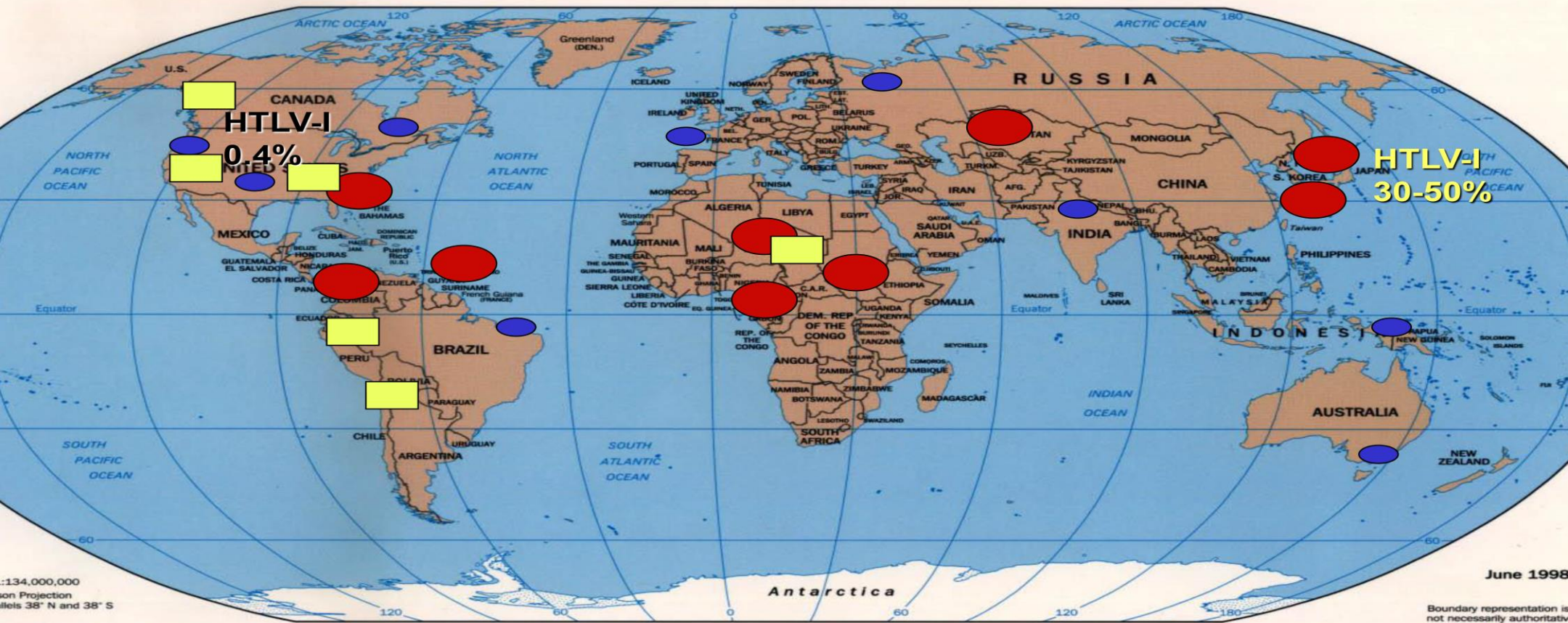
Lentivirus relationships

Lentivirus Relationships



HTLV distribution

HTLV DISTRIBUTION



Scale: 1:134,000,000
 Projection: Mercator
 Grid: 38° N and 38° S

June 1998

Boundary representation is not necessarily authoritative

HTLV-I Sporadic ● **HTLV-II Endemic (Amerindian and Pygmy tribes)**
HTLV-I High endemicity ■

HTLV-I ATL

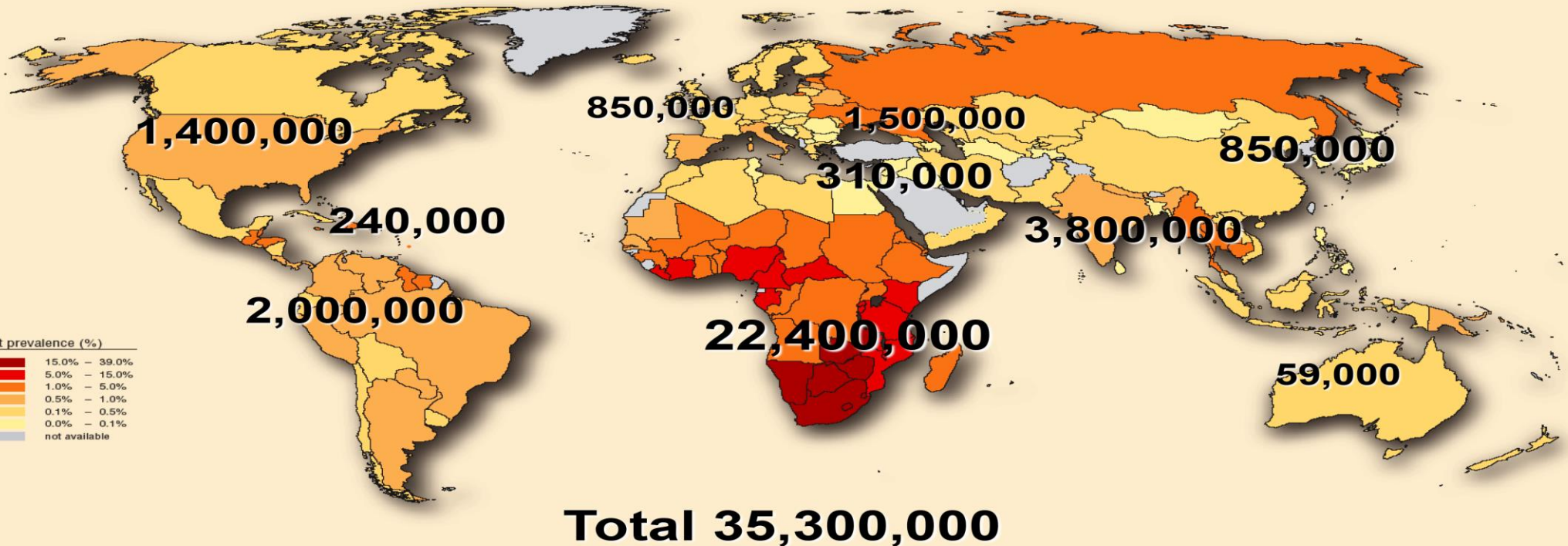
HTLV-I ATL

- **Long Latency (>30 years)**
 - **Small pediatric series in SA**
- **Epidemiology**
 - **Approximately 1% of HTLV- I infected adults**
- **Associated syndromes**
 - **Infectious**
 - **TB, MAC, Leprosy**
 - **PCP**
 - **Strongyloides**
 - **Scabies esp. Norwegian scabies**
 - **Noninfectious-hypercalcemia+lytic bone lesions**
- **Therapy-Chemotherapy, Ifn, anti-Tac**



HIV-1 pandemic risk

HIV-1 PANDEMIC: RISK



Higher Primate Origins of HIV-1

Higher Primate Origins of HIV-1

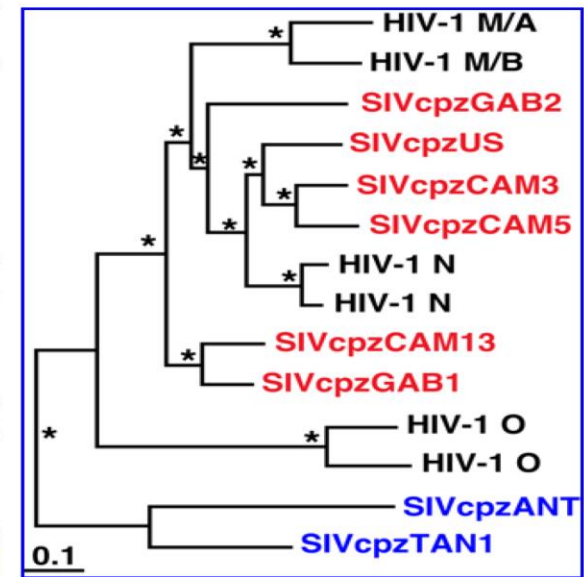
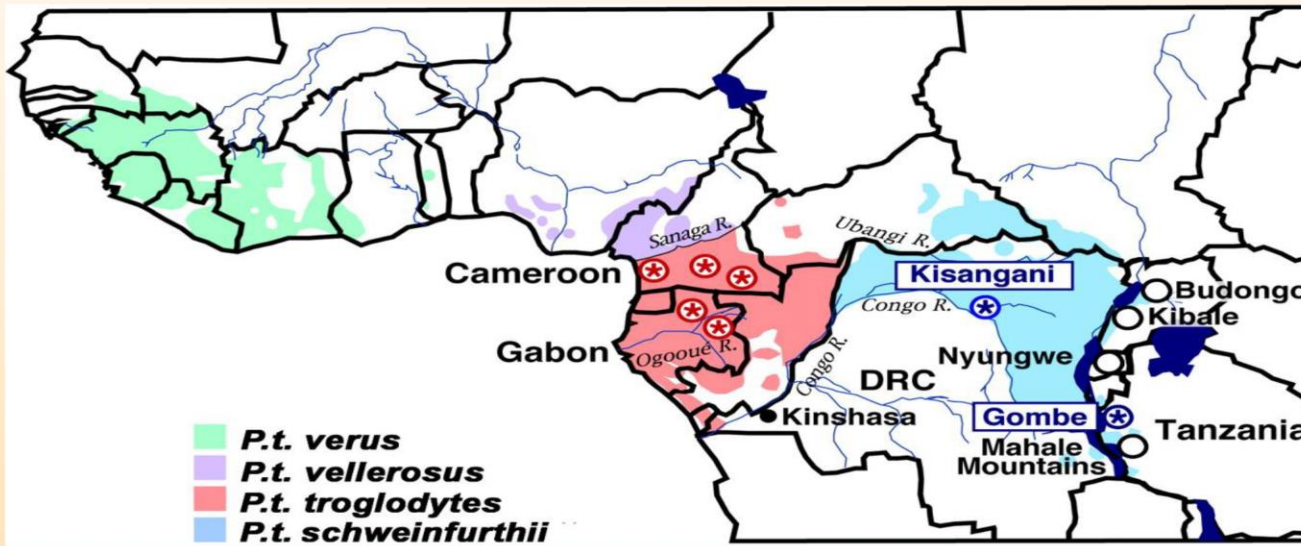
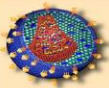
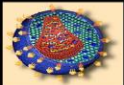
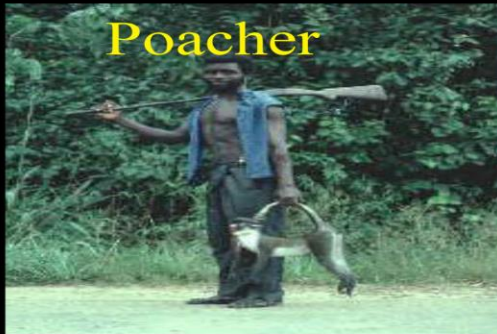


FIG. 2. Evolutionary relationships of SIVcpz and HIV-1 strains based on maximum-likelihood phylogenetic analyses of full-length envelope protein sequences (adapted from ref. 10). SIVcpz strains from *P. t. troglodytes* and *P. t. schweinfurthii* are highlighted in red and blue, respectively. Representative strains of HIV-1 groups M, N, and O were included for comparison. Asterisks indicate internal branches with estimated posterior probabilities of 95% or higher. The scale bar denotes 10% replacements per site.



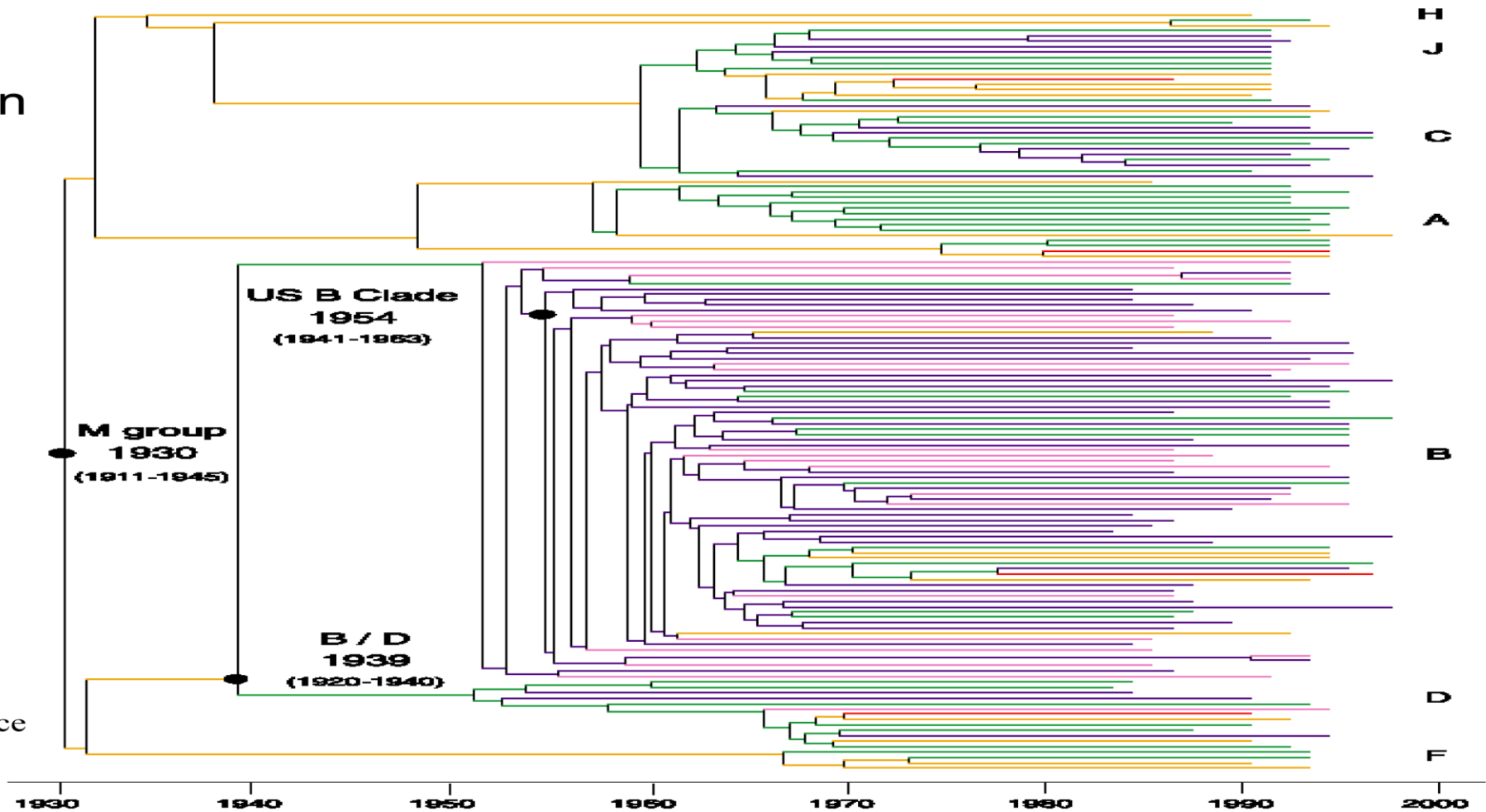
Bushmeat trade in Africa

Bushmeat Trade in Central and West Africa



HIV expansion

HIV
Expansion



Korber, Science
2000

HIV Spread

HIV Spread

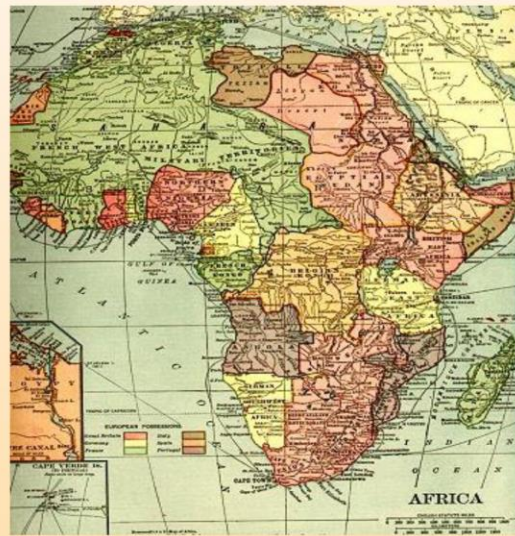
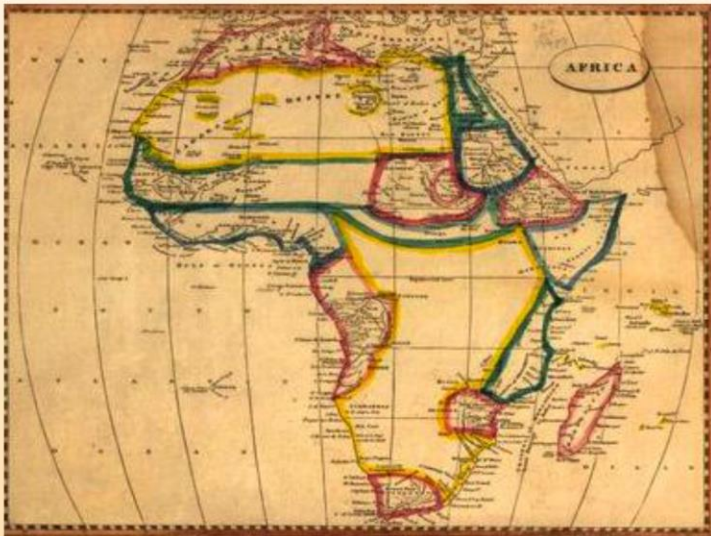
- **Biologic**
 - **Blood and body fluid**
 - **Iatrogenic**
 - **Blood transfusion**
 - **Vaccination – needles not vaccine**
 - **Mother to Child**
- **Non-Biologic**
 - **Political**
 - **Economic**
 - **Multiple Epidemics**



HIV spread

HIV Spread

- **Modes of Transmission**
Political

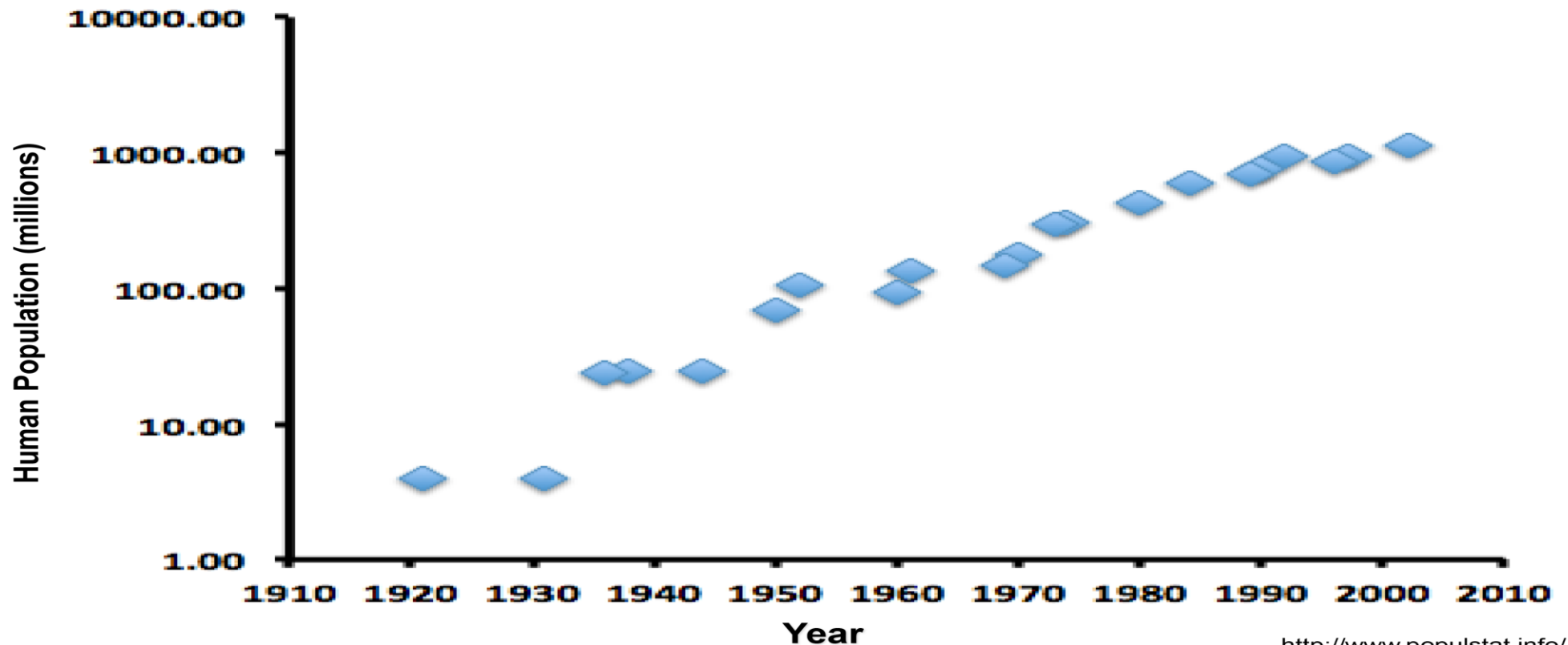


**Consequences of large political upheaval are population movement
and potential for malnutrition and immunodeficiency**



HIV and population expansion

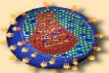
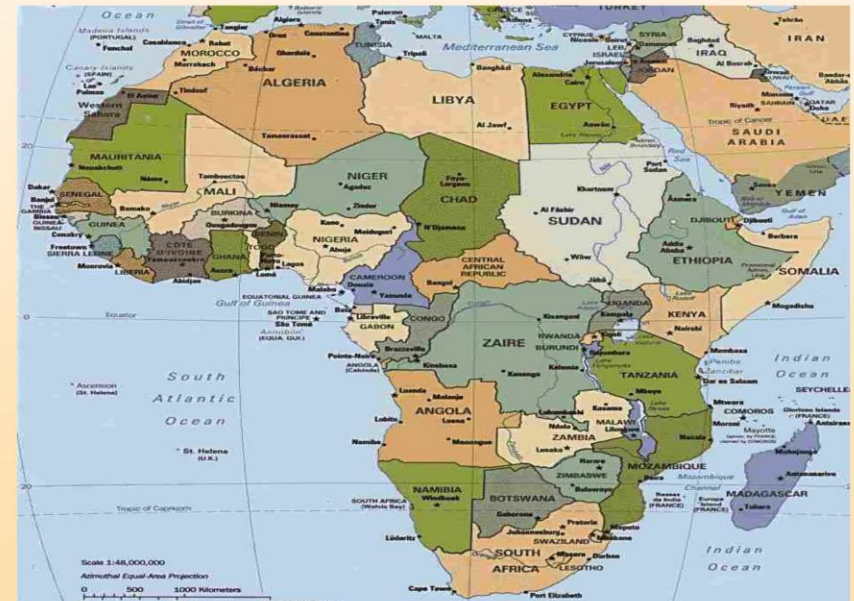
Zoonotic Transmission of HIV Coincides with Population Expansion in Africa



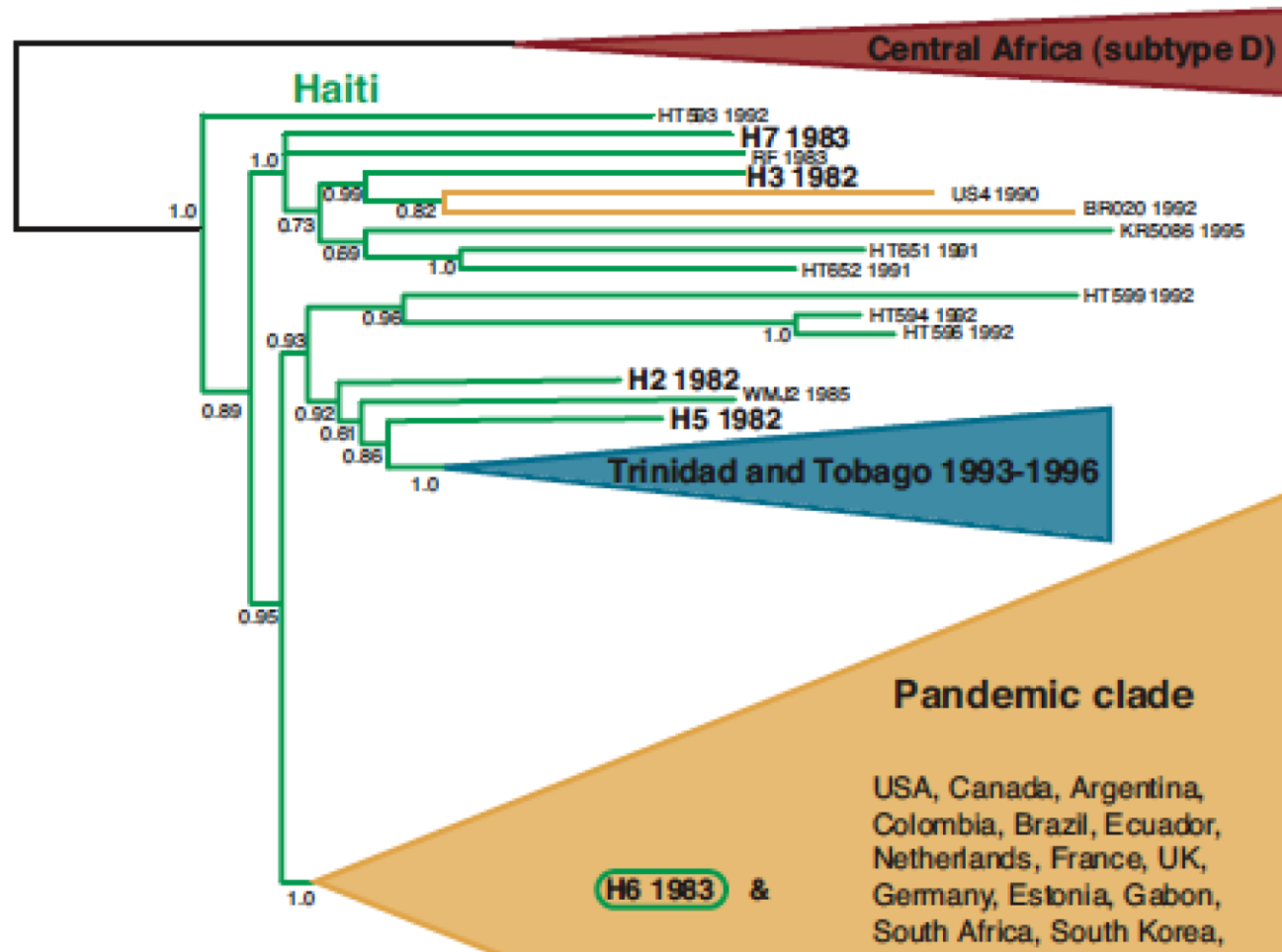
HIV spread

HIV Spread

- Modes of Transmission
Trans Africa Highway



HIV evolution



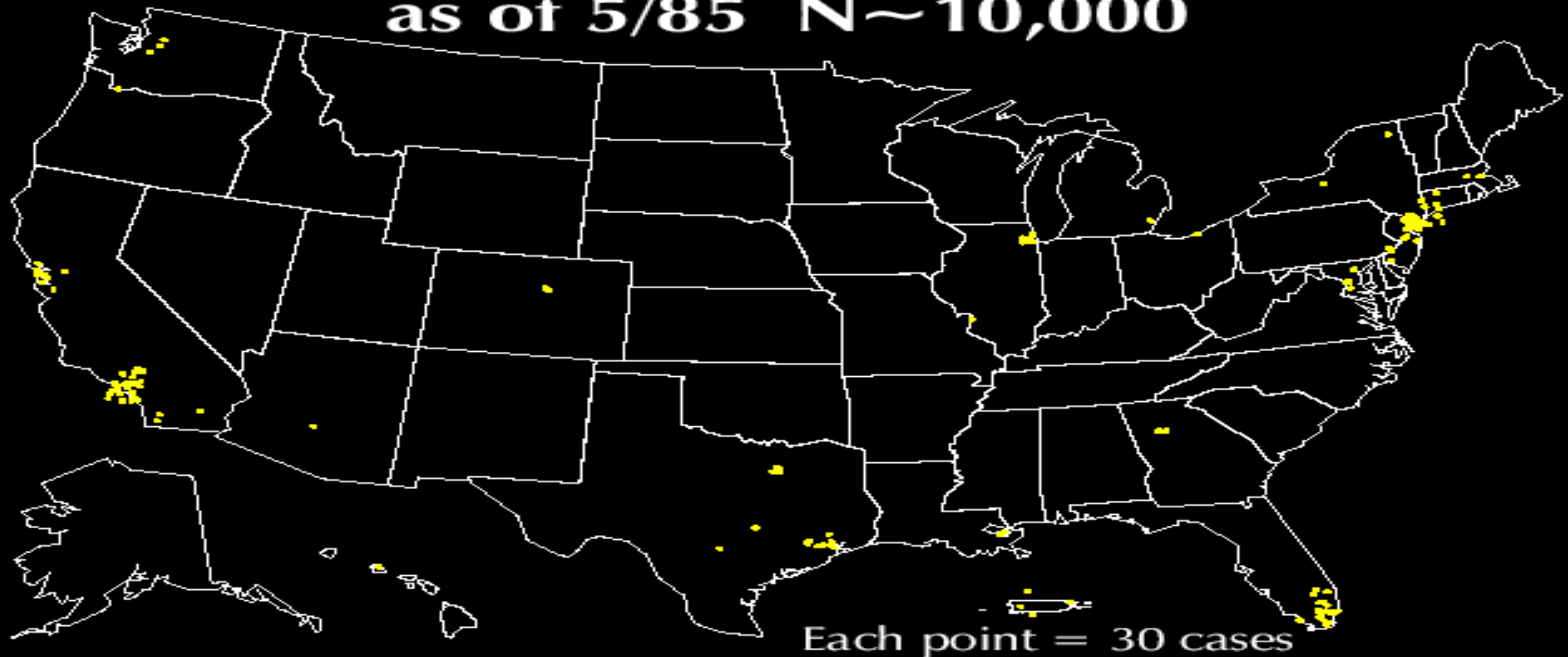
Cumulative U.S. AIDS cases as of 2/83 N = 1000

**Cumulative U.S. AIDS Cases
as of 2/83 N~1,000**



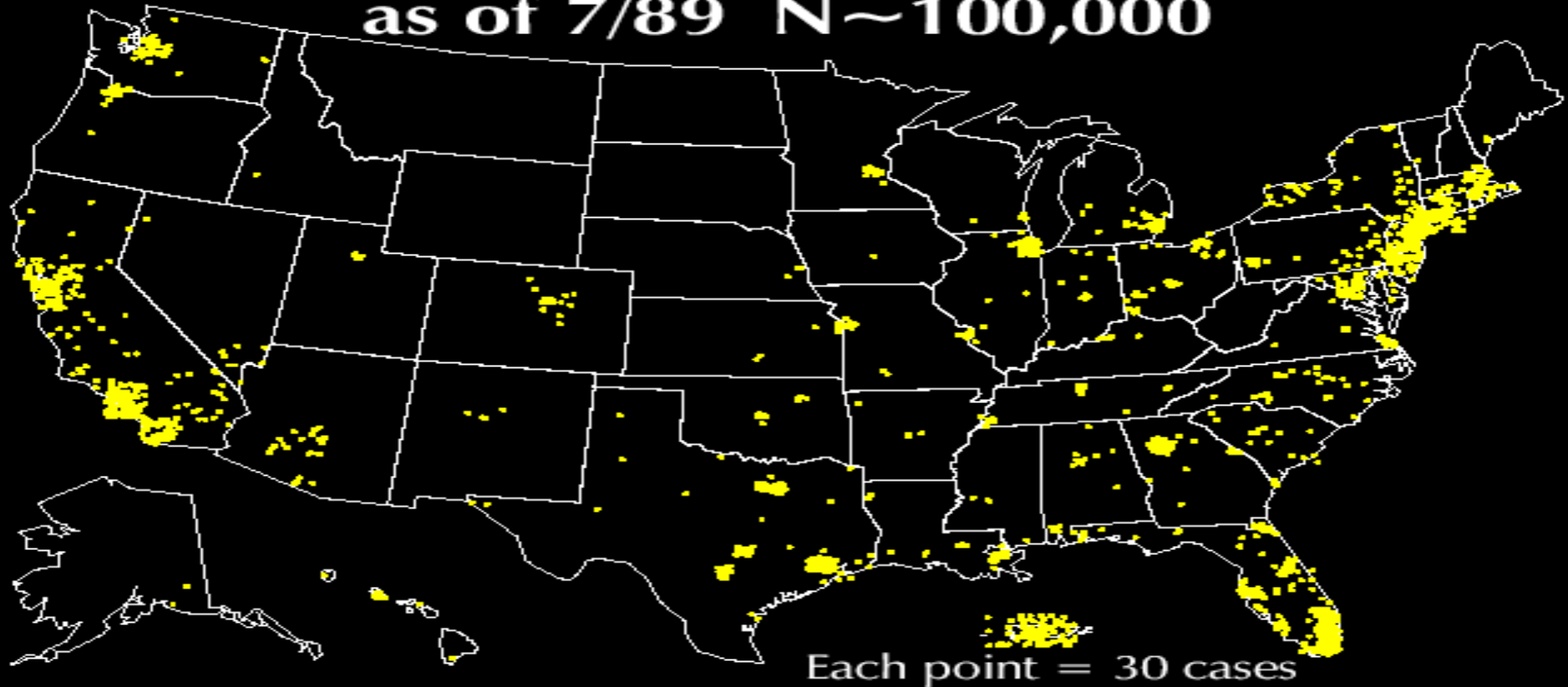
Cumulative U.S. AIDS cases as of 5/85 N = 10000

Cumulative U.S. AIDS Cases
as of 5/85 N~10,000



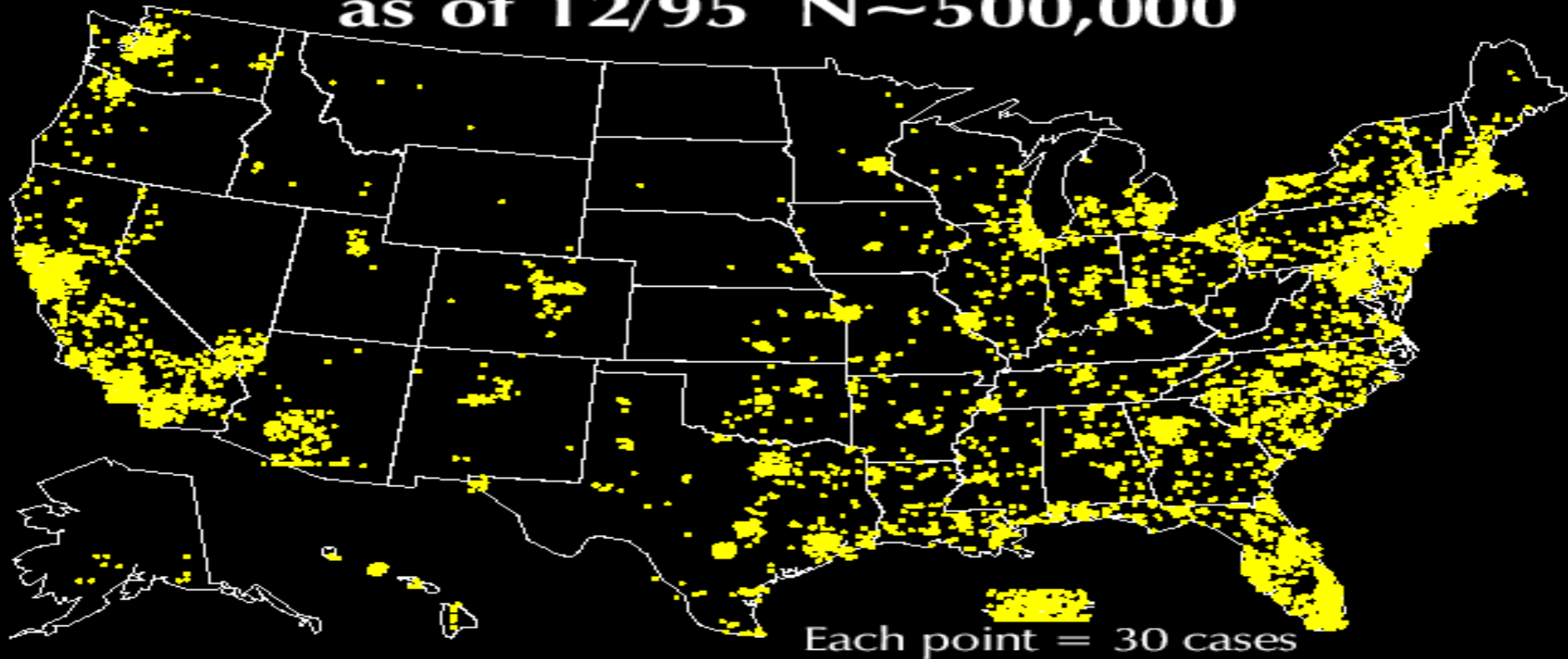
Cumulative AIDS cases as of 7/89 N = 100000

**Cumulative U.S. AIDS Cases
as of 7/89 N~100,000**



Cumulative AIDS cases as of 12/95 N = 500000

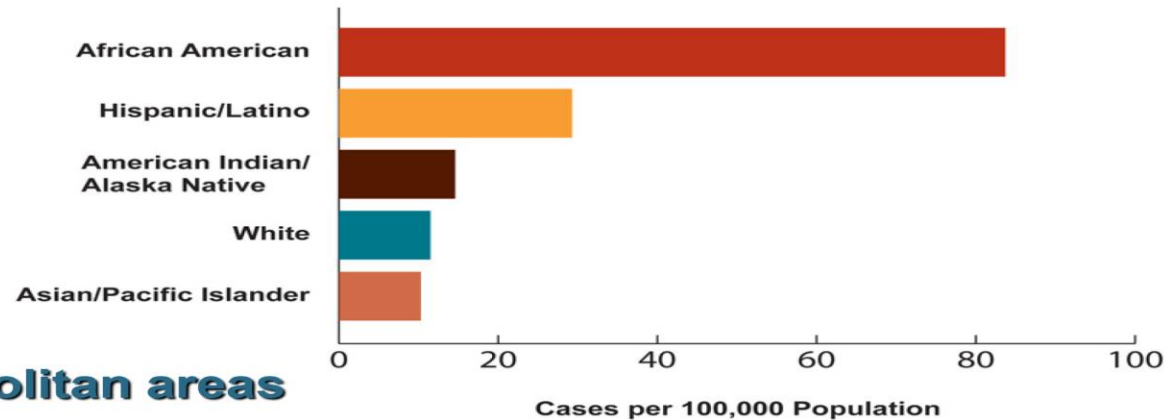
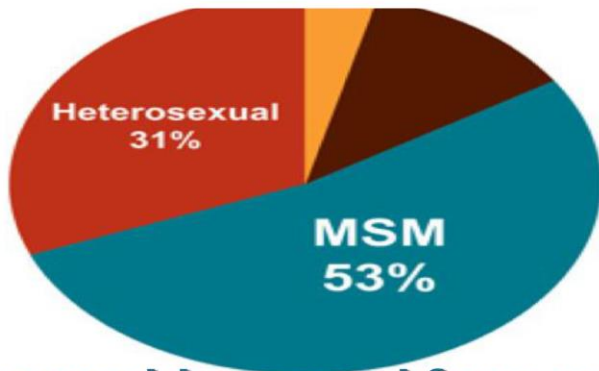
Cumulative U.S. AIDS Cases
as of 12/95 N~500,000



New cases of AIDS

New cases of HIV/AIDS—USA

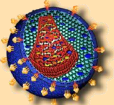
Estimates of New HIV Infections in the United States, 2006, By Transmission Category



JAMA 2008

- **Geographic spread from metropolitan areas**
 - ~12% of cases in locations with population <50,000
- **Women**
 - comprise > 25% of all AIDS cases
- **Age**
 - 11% of AIDS cases are 50+ years old
 - c.50% of persons living with HIV are >50 yo

HIV Therapy and Beyond

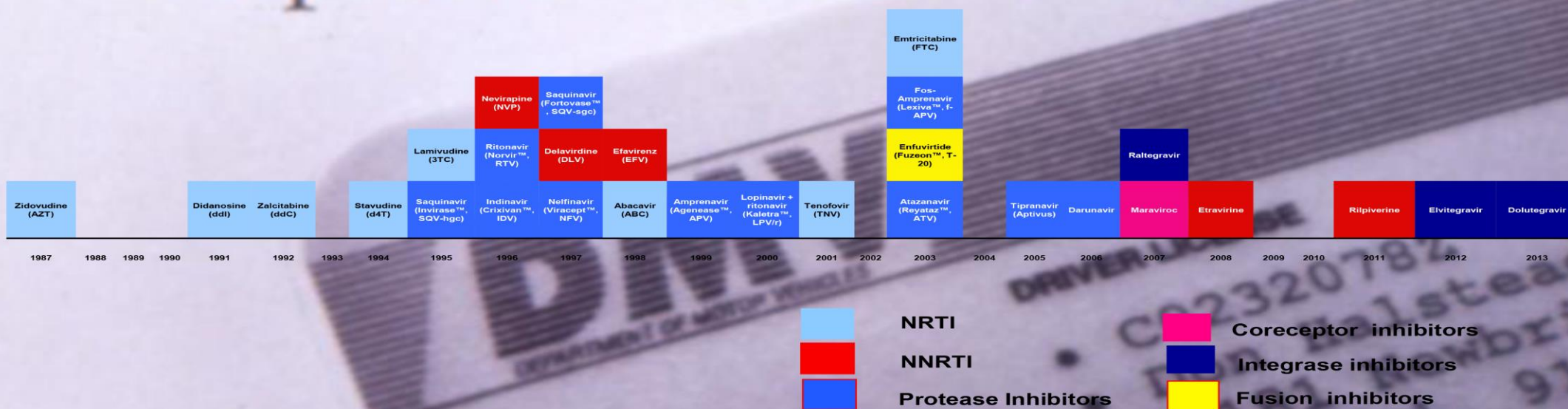


HIV Dynamics and Replication Program

NCI-Frederick

AIDS drugs

If you get the AIDS virus now, you and your license could expire at the same time.

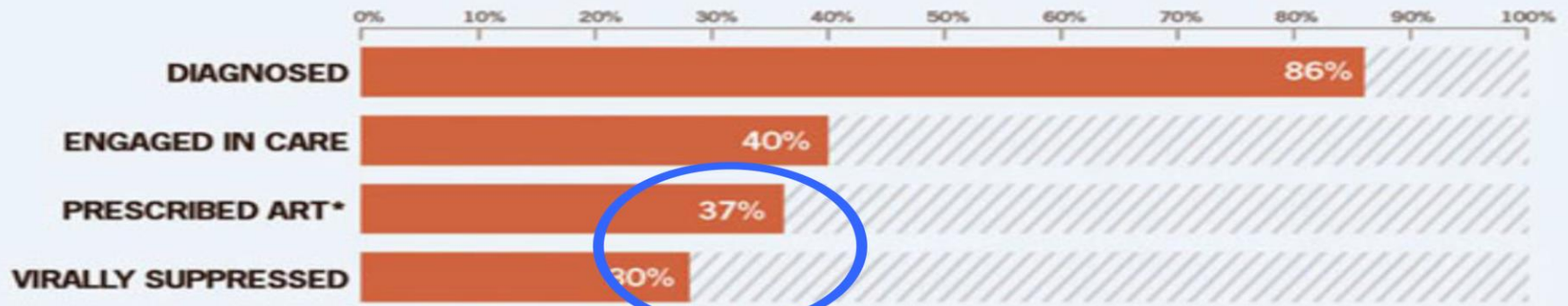


Gardner Continuum of Care

Gardner Continuum of Care

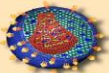
HIV Care Continuum Shows Where Improvements are Needed

In the US, 1.2 million people are living with HIV. Of those:



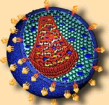
SOURCES: CDC National HIV Surveillance System and Medical Monitoring Project, 2011.

*Antiretroviral therapy



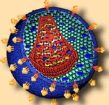
Key Advances in HIV Therapy

- PrEP
 - Adherent prophylaxis is effective
- SMART Study
 - Continuous therapy essential to avoid AIDS and other complications
- START Study
 - Earlier therapy is initiated, greater preservation of therapy



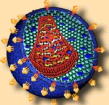
Next Advances in HIV Therapy

- Vaccines
- Cures



Retroviruses

- **Lessons**



Summary

Summary

- **Viruses are bad and should be avoided**
- **Except when they save the planet**
- **And maybe if it saves you from the next virus**
- **Epidemics are not single events**
- **Epidemics evolve**
- **Antivirals are useful**
 - **Instituted as early as possible**
 - **Adherence is essential**

