

# WHY STUDY RETROVIRUSES?

## \* PATHOGENIC CONSEQUENCES: A HISTORY OF DISCOVERY...

EQUINE ANEMIA 1904 (EIAV)

CHICKEN LEUKOSIS 1907 (ALV)

CHICKEN SARCOMA 1911 (RSV)

MOUSE MAMMARY CARCINOMA 1930'S (MMTV)

MOUSE LEUKEMIA 1951 (MLV)

ADULT T CELL LEUKEMIA/LYMPHOMA 1980 (HTLV)

ACQUIRED IMMUNODEFICIENCY SYNDROME 1983 (HIV)

(OTHER PATHOLOGY INCLUDES ARTHRITIS, NEUROLOGICAL DISEASES, OSTEOPETROSIS, MANY CANCERS, ETC.)

## \* UNUSUAL LIFE CYCLE

TEMIN'S PROVIRUS HYPOTHESIS:

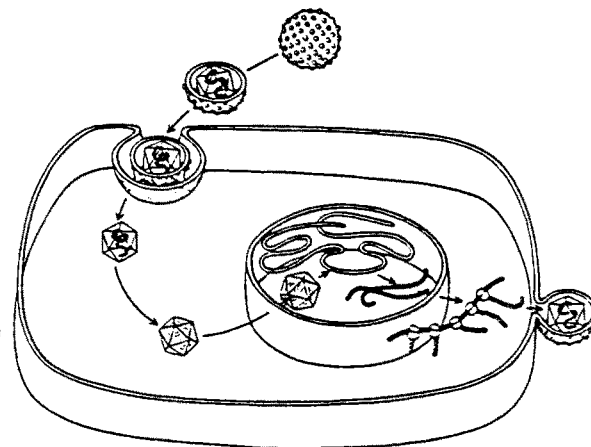
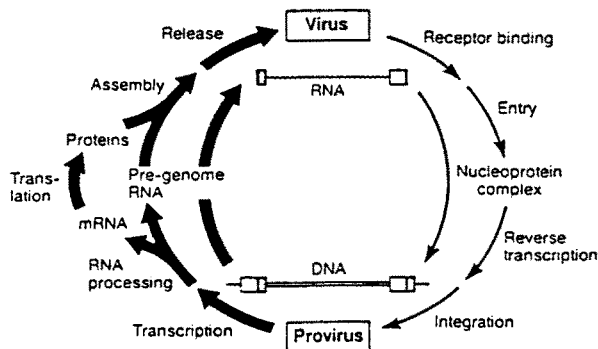
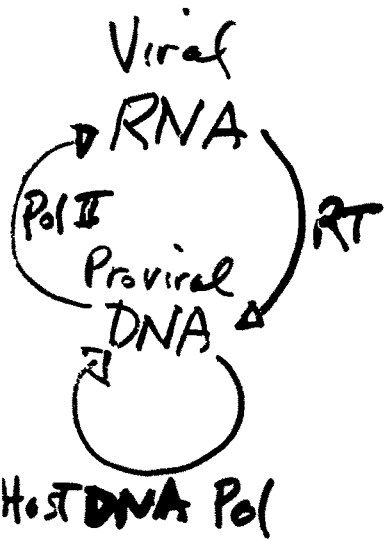
THE CLUES: INHIBITORS OF DNA SYNTHESIS AND OF DNA

-DIRECTED RNA SYNTHESIS BLOCK INFECTION OF AN RNA VIRUS

THE EVIDENCE: REVERSE TRANSCRIPTASE IN VIRIONS

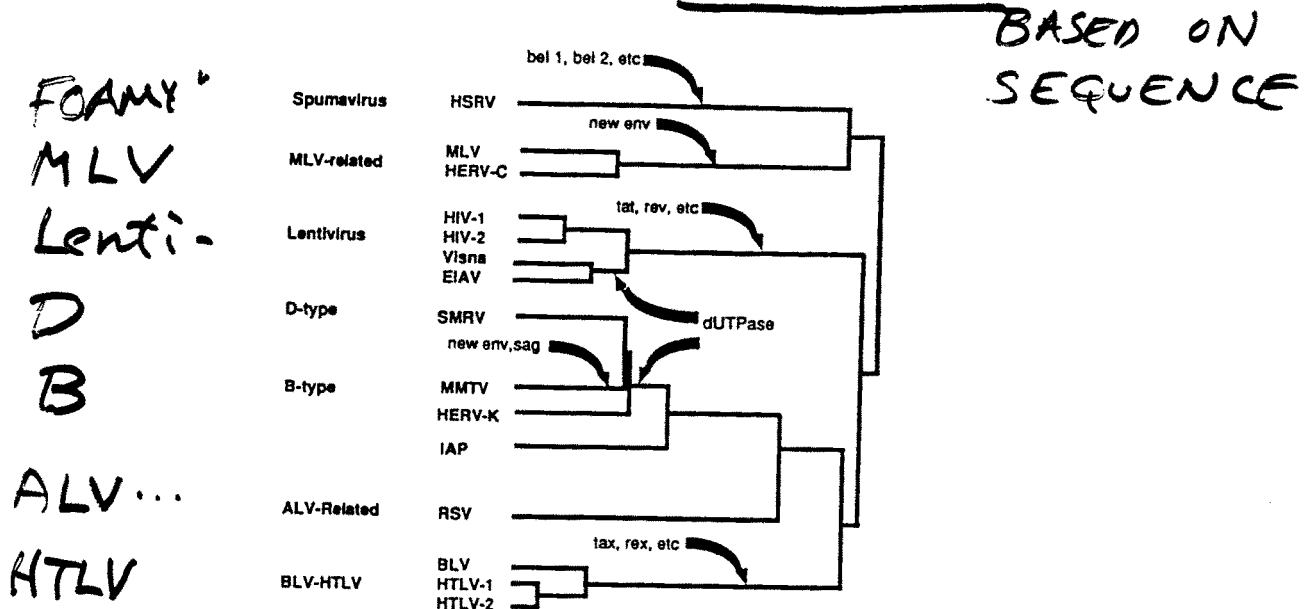
TS MUTANTS OF POL GENE

DNA IN INFECTED CELLS (HYBRIDIZATION, TRANSFECTION, CLONING)



# WHY STUDY RETROVIRUSES?

\* WIDE DISTRIBUTION IN NATURE...THE SEVEN CLASSES



(OTHER MEANS OF CLASSIFICATION: PATHOLOGY, PARTICLE TYPE, HOST)

TRANSMISSION: BLOOD, MILK (MMTV, HTLV), CONGENITAL INFECTION, VIA GERM LINE (ENDOGENOUS PROVIRUSES)

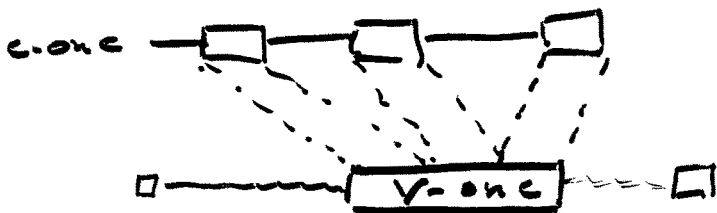
\* EVOLUTIONARY IMPLICATIONS OF ENDOGENOUS PROVIRUSES,

OTHER RETRO-ELEMENTS (NEXT LECTURE)

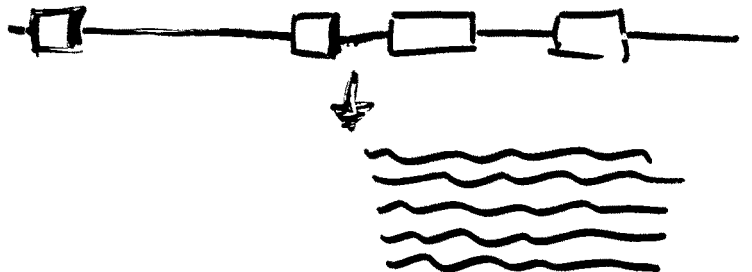
\* NOVEL STRATEGIES FOR EXPRESSION (TAT, REV, FRAMESHIFTING)

\* ONCOGENIC MECHANISMS:

TRANSDUCTION



INSERTIONAL ACTIVATION

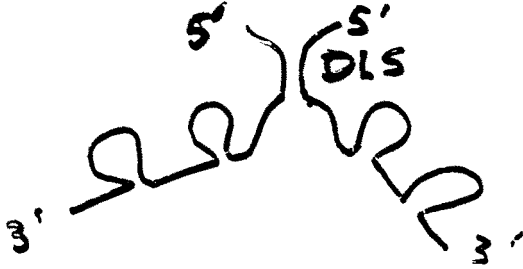


\* UTILITY AS GENETIC VECTORS



# RETROVIRAL GENOMES....

HOMODIMERS, DIMER LINKAGE, ASSOCIATED tRNA



NON-RANDOM tRNA,  
MAINLY PRIMER

IMPLICATIONS OF "PSEUDODIPLOIDY": HETEROZYGOSIS, RECOMBINATION

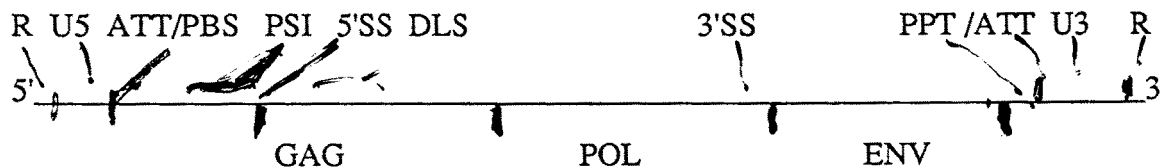
2 RNA subunits → 1 PROVIRUS

## COMMON ELEMENTS

CIS ACTIVE: SITES FOR PRIMING RT AND FOR INTEGRATION (ATT SITES);

PROMOTERS, ENHANCERS, AND RNA PROCESSING SIGNALS  
(SPLICING AND POLYADENYLATION);

TRANSLATION, DIMERIZATION, PACKAGING SIGNALS.



TRANS ACTIVE: GAG----> MATRIX, CAPSID, NUCLEOCAPSID

POL----> PROTEASE, RT, INTEGRASE

ENV-----> SURFACE, TRANSMEMBRANE GP

## IDIOSYNCRATIC ELEMENTS:

TRANSCRIPTIONAL REGULATORS: TAX, TAT, BEL GENES AND TARGETS

REGULATORS OF RNA METABOLISM: REX AND REV AND TARGETS

OTHER LENTIVIRUS GENES: NEF, VIF, VPU, VPR, VPX

SUPERANTIGEN (SAG) GENE OF MMTV; dUTP-ASES OF LENTIVIRUSES

TRANSUCED GENES (MOSTLY ONCOGENES)

# HIV-1

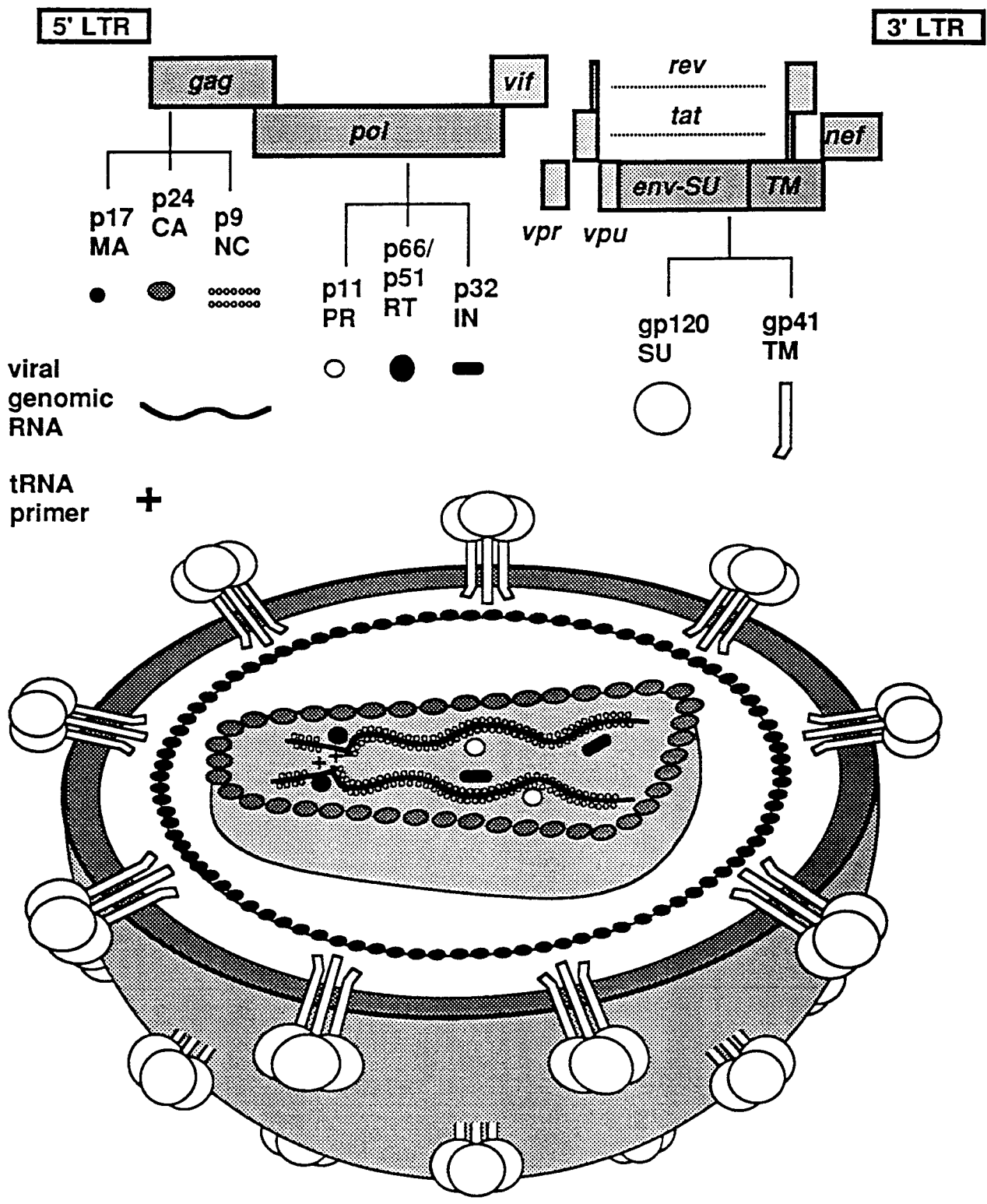


FIGURE 3. Virion morphology.

COURTESY OF PAUL LUCIW

# DETERMINANTS OF VIRUS ENTRY AND HOST RANGE

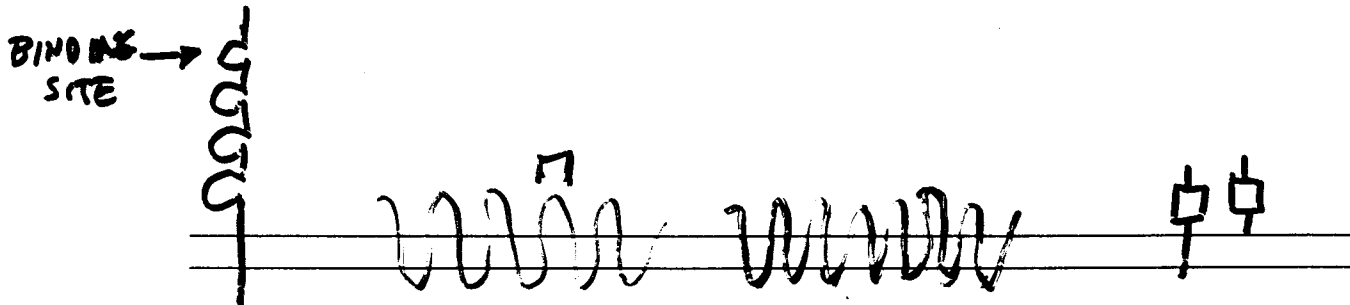
ENTRY MEDIATED BY VIRAL SURFACE ENV PROTEIN AND VARIETY OF CELL SURFACE RECEPTORS WITH WIDE RANGE OF NORMAL FUNCTIONS...

INFECTION BLOCKED BY: ABSENCE OF RECEPTOR OR CO-FACTOR

HOMOLOGOUS INTERFERENCE, DOWN REG.

ANTIBODY AGAINST SU-ENV OR RECEPTOR

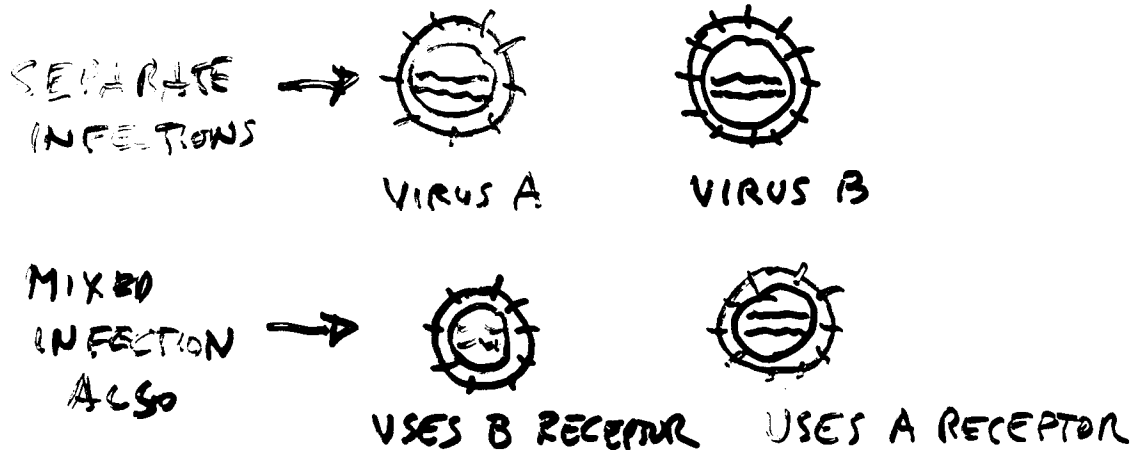
SOLUBLE RECEPTOR



VIRUS :	HIV	GALV/FLV-B	MLV-ECO	RSV-A
RECEPTOR :	CD4 (REQUIRES COFACTOR)	?TRANSPORTER	BASIC AA PERMEASE	LDL-R RELATED
pH :	INDEP.	?	DEP.	INDEP.

SURFACE INTERACTIONS ARE MAJOR DETERMINANT OF VIRAL HOST RANGE AND TISSUE TROPISM

RESTRICTION CAN BE OVERCOME BY FORCED FUSION, PROVIRAL TRANSFECTION, AND PSEUDOTYPE FORMATION

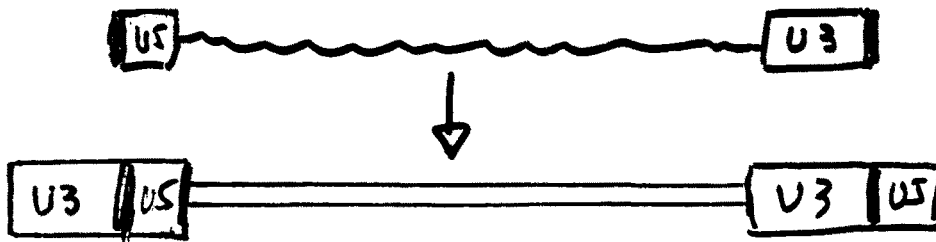


## EARLY EVENTS: THE NUCLEOPROTEIN COMPLEX

A PRODUCT OF ENTRY AND UNCOATING,  
COMPOSITION AND STRUCTURE UNCERTAIN



VEHICLE FOR REVERSE TRANSCRIPTION IN CYTOPLASM, TO MAKE  
FULL-LENGTH LINEAR DUPLEX WITH LONG TERMINAL REPEATS



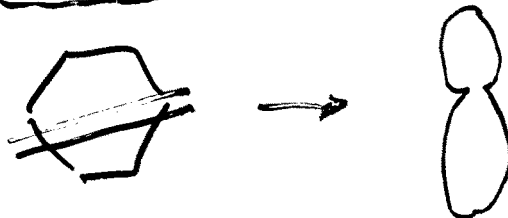
### MIGRATION TO THE NUCLEUS:

\* REQUIREMENT FOR NUCLEAR MEMBRANE BREAKDOWN IN MITOSIS (MLV)  
VS. DIRECT ENTRY (HIV)

\* PRESUMPTIVE DEAD ENDS: CIRCLES WITH ONE OR TWO LTRS



### INTEGRATION MACHINE FOR INSERTING DNA INTO CHROMOSOMES



HOST RESTRICTION VIA NPC: SOME FV-1 ALLELES RESTRICT  
GROWTH OF SOME STRAINS OF MLV (VIA CAPSID DETERMINANT),  
OPERATING AFTER ENTRY AND BEFORE INTEGRATION....

# PRINCIPLES OF RETROVIRAL REVERSE TRANSCRIPTION

## \* PROPERTIES OF ENZYME...RNASE H AND DNA POLYMERASE

HOLOENZYME (INTRINSIC TEMPLATE:PRIMER, DISRUPTED VIRIONS)

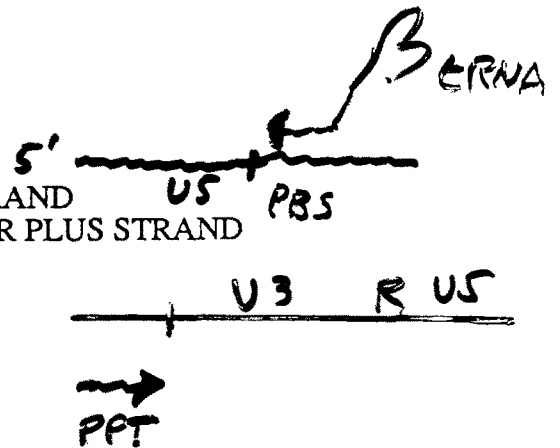
ISOLATED ENZYME (FROM VIRIONS, MADE IN BACTERIA)

## \* HETERODIMERS (HIV, RSV) VS. MONOMERS (MLV)

	RT	RNASE H	IN
RSV	_____	_____	_____
HIV	_____	_____	_____
MLV	_____	_____	_____

## \* TEMPLATES AND PRIMERS

NATURAL PRIMERS: tRNA FOR FIRST (MINUS) STRAND  
POLYPURINE OLIGO-RNA FOR PLUS STRAND



EXOGENOUS TEMPLATE-PRIMERS: mRNA-OLIGO dT  
GAPPED DNA  
POLY rA: dT OR POLY rC: dG

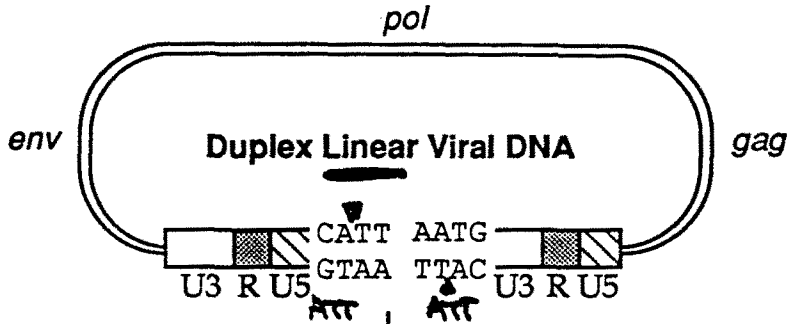
## \* JUMPS, METASTABILITY, ERRORS

## \* INHIBITORS: NUCLEOSIDE ANALOGS, OTHERS

AZT  
ddI  
ddC

Nevirapine  
Pyridinone  
TIBO  
etc.

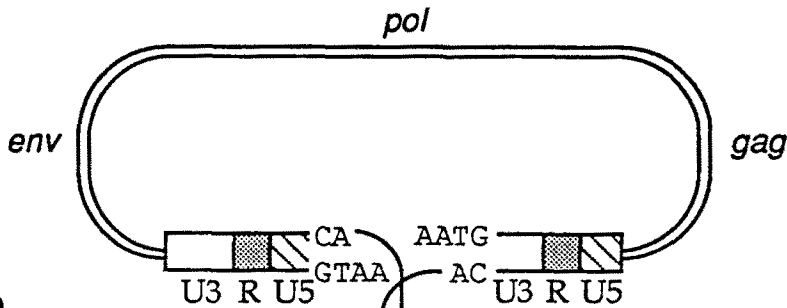
# CURRENT VIEW OF INTEGRATION



**Step 1**

Nicking of linear viral DNA by IN and removal of 3' dinucleotides

TRANSESTERIFICATION (OR ALCOHOL) DONE



**Step 2a**

Staggered cleavage of target cell DNA by IN

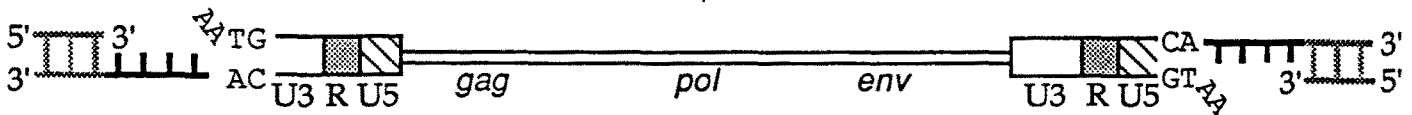
COORDINATED NO ATP REQ.



**Step 2b**

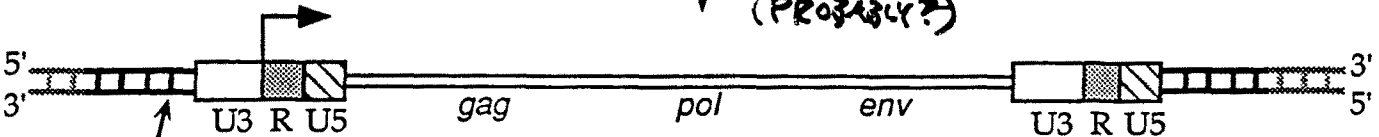
Joining (i.e., strand transfer) by IN

TRANSESTERIFICATION 3' OH AS DONOR



**Step 3**

Cell enzymes (PROBABLY?)  
 -removal of mismatched dinucleotides  
 -fill-in of gaps  
 -ligation



Short direct repeats in cellular DNA

Integrated Provirus



# PRINCIPLES OF RETROVIRAL INTEGRATION

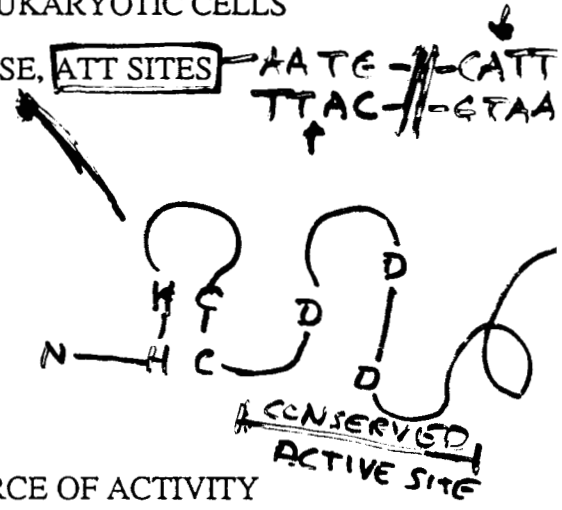
THE BEST UNDERSTOOD RECOMBINATION EVENT IN EUKARYOTIC CELLS

MACHINERY: NUCLEOPROTEIN COMPLEX, INTEGRASE, ATT SITES

PRECISION AT VIRAL ENDS, MANY SITES IN TARGET

NO REQUIREMENT FOR ENERGY SOURCE

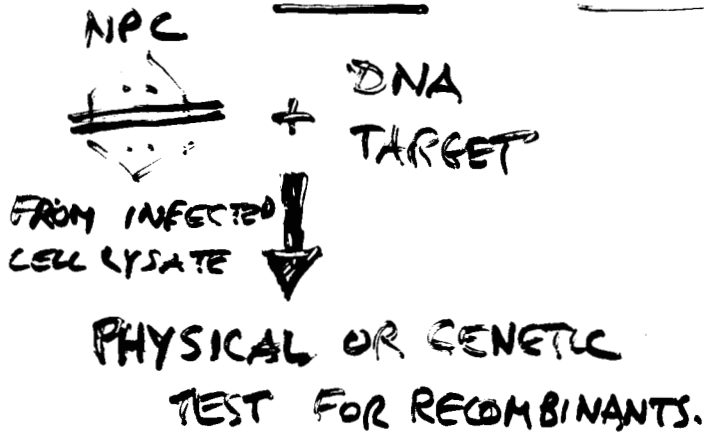
ESSENTIAL FOR REPLICATION AND PERSISTENCE



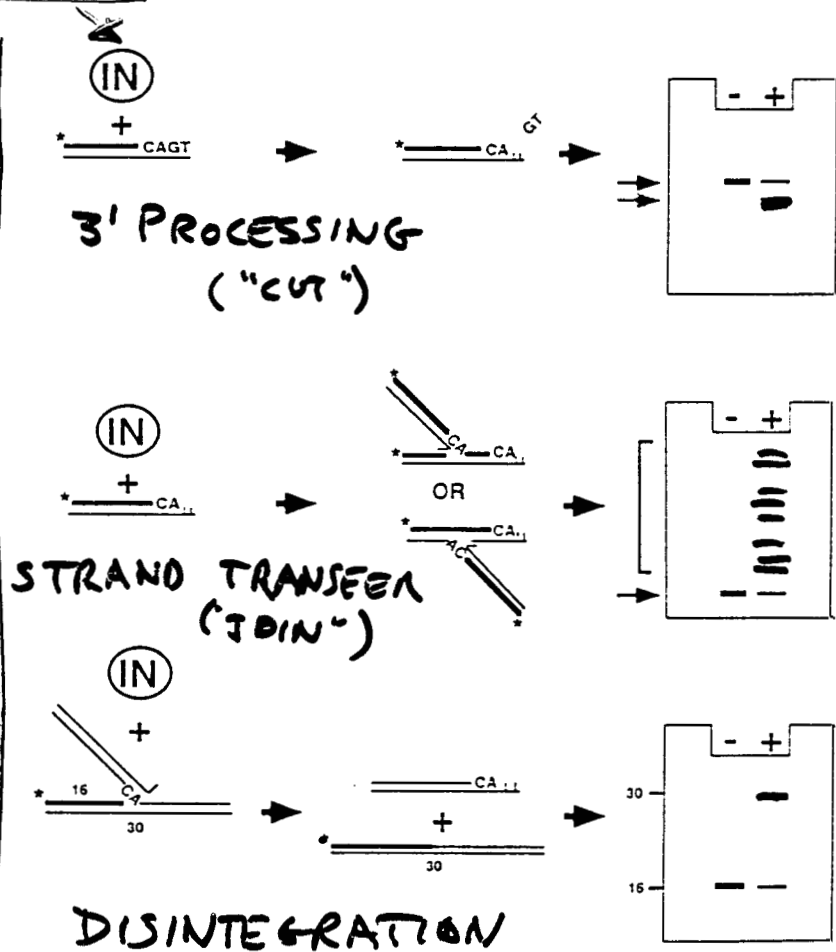
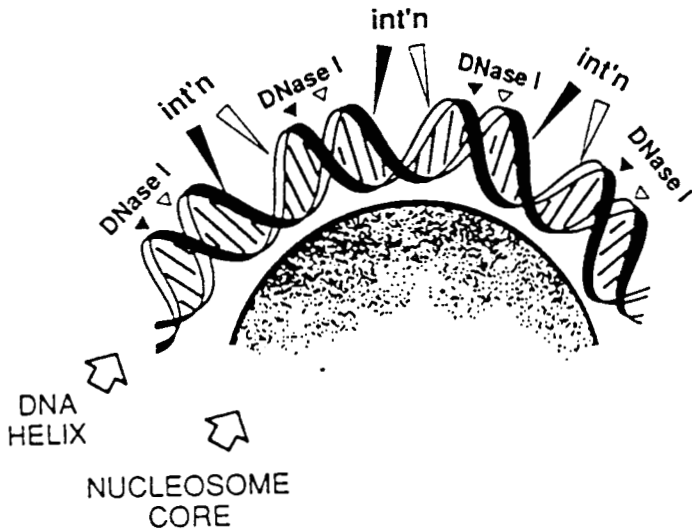
## ASSAYS:

INFECTED CELLS...MAPPING, CLONING, PCR

IN VITRO...NPC VS. PURIFIED INTEGRASE AS SOURCE OF ACTIVITY



## ROLE OF NUCLEOSOMES



# CONTROL OF PROVIRAL TRANSCRIPTION

## 1) HOST TRANSCRIPTION FACTORS INTERACT WITH TARGETS IN U3

- \* DETERMINES LEVEL OF EXPRESSION
- \* INFLUENCES TISSUE TROPISM AND PATHOGENIC TARGET

MLV: ERYTHRO VS. T CELL LEUKEMIA

MMTV: MAMMARY VS. T CELL TUMORS

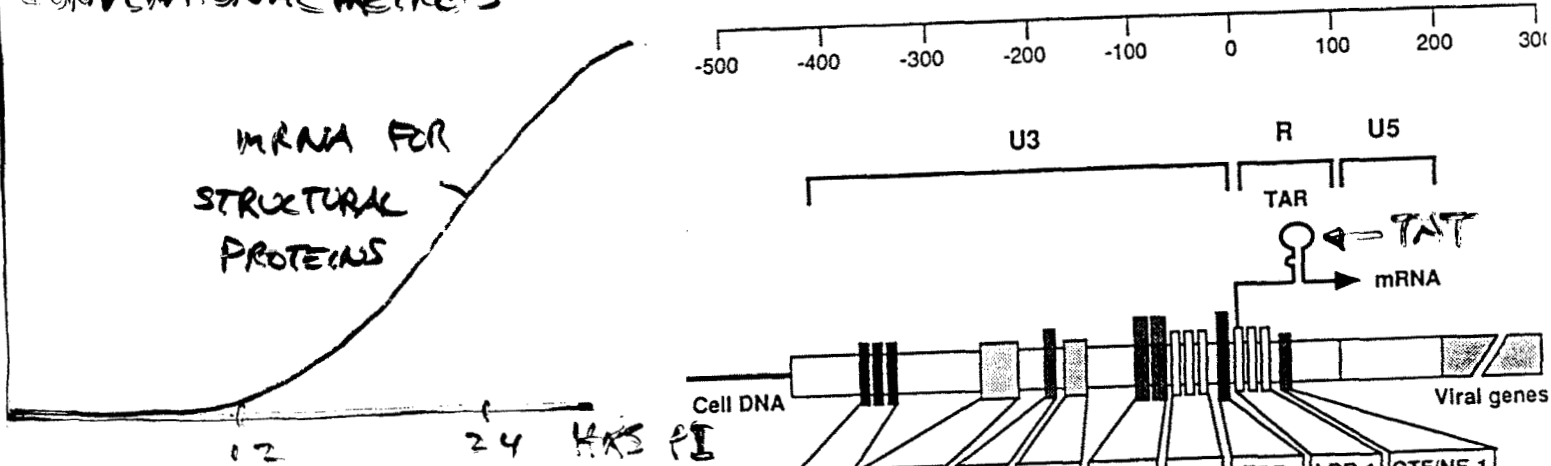
## 2) VIRAL FACTORS AS TRANSCRIPTIONAL CO-ACTIVATORS

- \* TAX OF HTLV/BLV
- \* BEL OF FOAMY VIRUSES

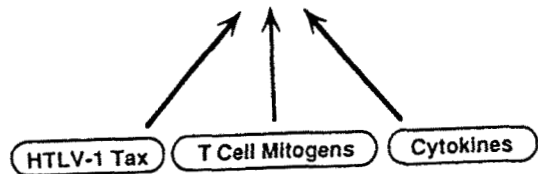
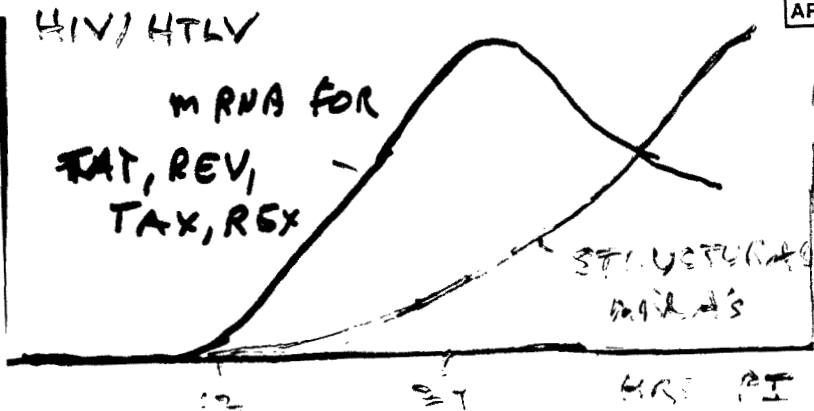
## 3) VIRAL FACTORS AS REGULATORS VIA VIRAL RNA TARGET

- \* TAT/TAR OF HIV & SIV
- \* ABSOLUTE REQUIREMENT FOR VIRUS PRODUCTION
- \* ROLE IN TEMPORAL CONTROL OF LATE EVENTS (AND LATENCY?)

### CONVENTIONAL RETROVIRUS



### HIV/HTLV



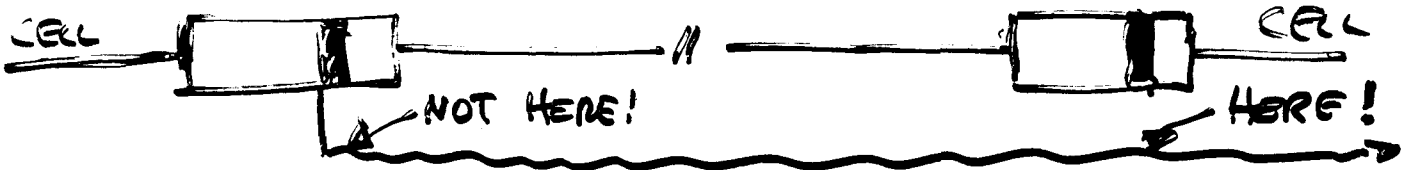
SOME FACTORS AFFECTING HIV-1 RNA SYNTHESIS



# PROCESSING OF RETROVIRAL RNA

CAPPING 5' ENDS

POLYADENYLATION: RESPONDING TO SIGNALS FROM 3' (NOT 5') LTR



SPLICING AND TRANSPORT TO CYTOPLASM:

- MAINTAINING AN APPROPRIATE RATIO OF RNA'S  
GENOME (GAG mRNA) vs ENV mRNA vs OTHER mRNA  
 (N.B.: PRE-mRNA IS GENOME AND ALSO mRNA)

- FOR SIMPLE GENOMES: CIS SIGNALS SUFFICE

- FOR COMPLEX GENOMES:

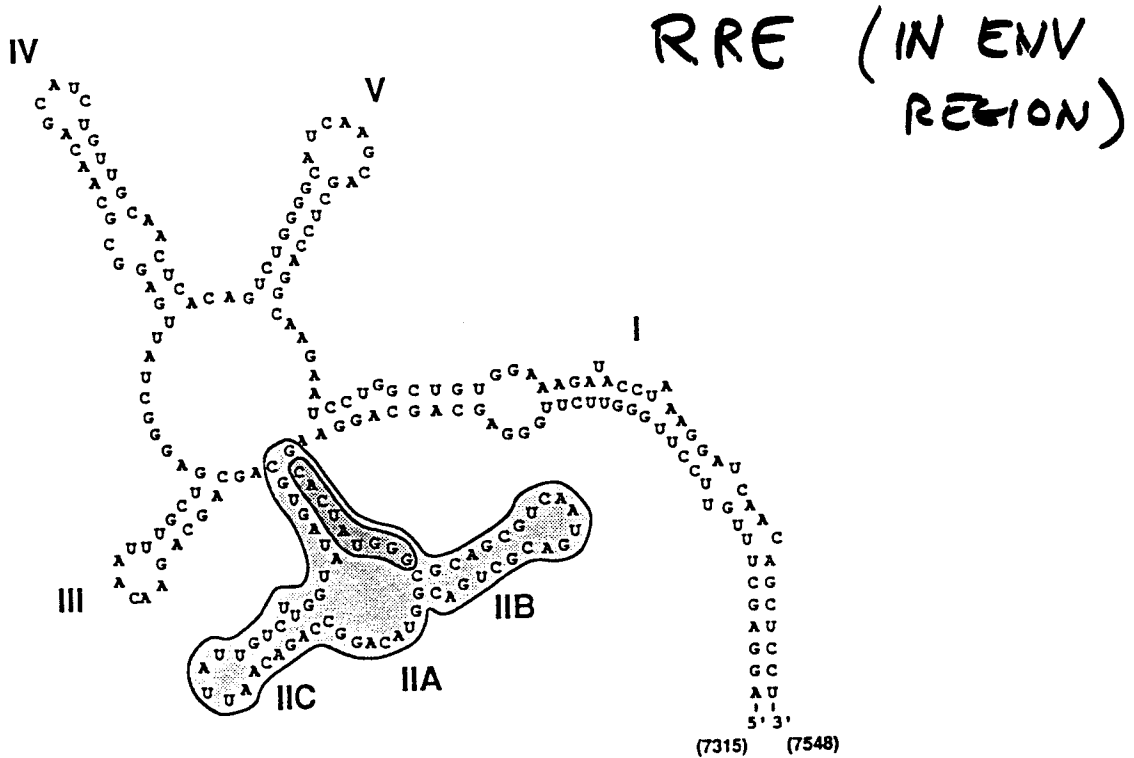
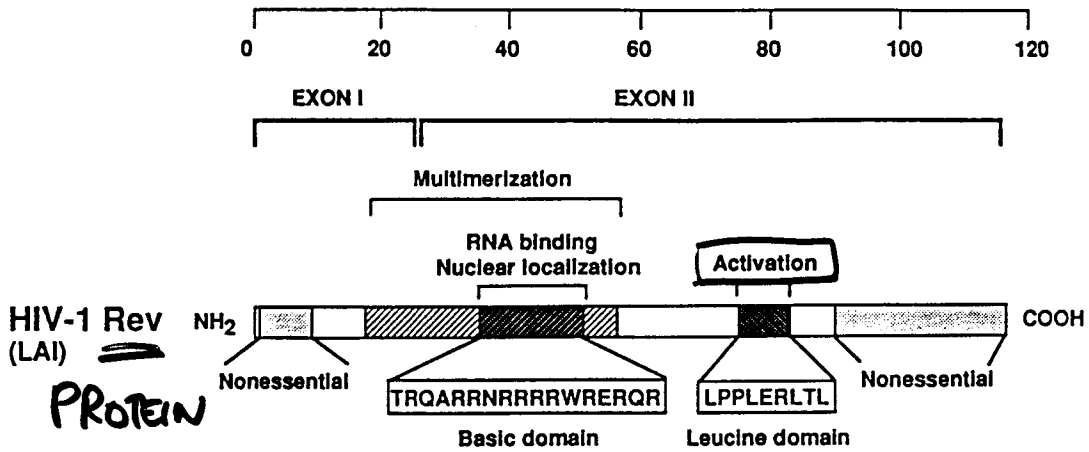
APPEARANCE OF INCOMPLETELY SPLICED mRNA IN

CYTOPLASM GOVERNED BY REX OR REV VIA RXRE OR RRE

## CYTOPLASMIC RNA NORTHERNS

	MLV, ALV, etc.	HIV REV <sup>+</sup>	HIV REV <sup>-</sup>
GENOMIC gag mRNA	—	—	—
ENV mRNA (spliced)	—	—	—
TAT, REV, etc.	—	—	—

# REV AND RRE OF HIV-1

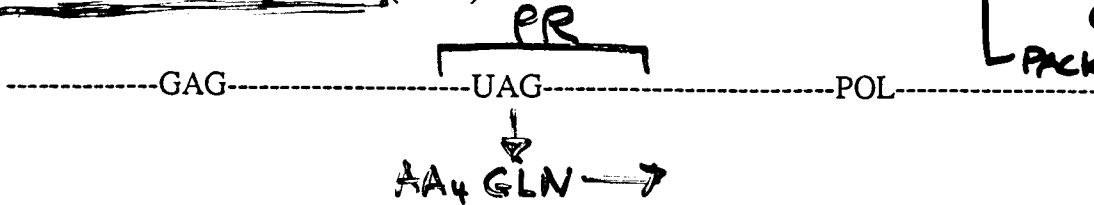


# CONTROL OF RETROVIRAL TRANSLATION

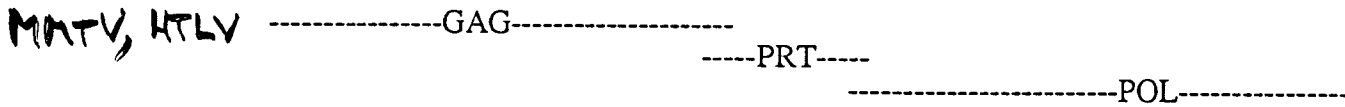
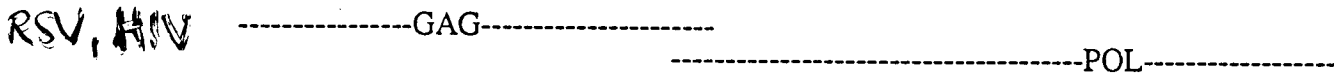
EXPRESSION OF POL GENES VIA TRANSLATIONAL READ THROUGH

AVOIDS SPLICING  
OR INTERNAL  
START  
MAINTAINS RATIO  
GAG/GAG-POL  
PACKAGING POL

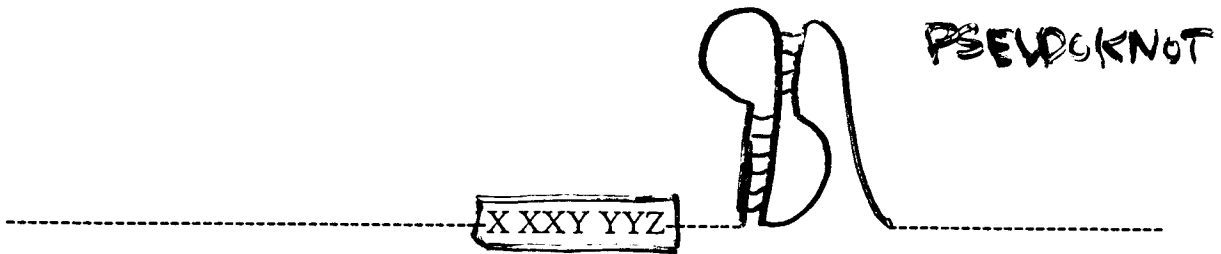
1) NONSENSE SUPPRESSION (MLV)



2) MINUS-ONE FRAMESHIFTING---ONCE (RSV, HIV) OR TWICE (MMTV, HTLV)



SIGNALS: SHIFTY SITE, RNA STRUCTURE 3' OF SITE, SPECIAL tRNA



MECHANISM: SIMULTANEOUS SLIPPAGE AFTER DECODING YYZ

FREQUENCY: 3% TO 20%

OTHER VENUES: CORONAVIRUSES, RETROTRANSPOSONS, YEAST DS RNA, BACTERIAL TRANSPOSONS, PLANT VIRUSES, E.COLI DNA X.....

NO EUKARYOTIC HOST GENES AS YET

# RULES GOVERNING RETROVIRAL ASSEMBLY

## CAPSID-CAPSID INTERACTIONS DRIVE ASSEMBLY OF CORE

GAG PRECURSOR SUFFICES

INTERACTIONS WITH CA PORTION OF GAG-POL INCLUDE POL

## MATRIX DETERMINES SITE OF ASSEMBLY

ACETYLATION/MYRISTYLATION REQUIRED TO GET TO MEMBRANE

MATRIX VARIANTS FOR ASSEMBLY AT PLASMA MEMBRANE (MOST),  
CYTOSOL (TYPE D), OR ER (MUTANTS)

NUCLEOCAPSID REGION INTERACTS WITH PSI SITE ON VIRAL RNA,  
AND FACILITATES DIMERIZATION

RT PROMOTES INCLUSION OF APPROPRIATE TRANSFER RNA (PRIMER)

ENV PROTEINS MODIFIED, OLIGOMERIZED IN ER AND GOLGI

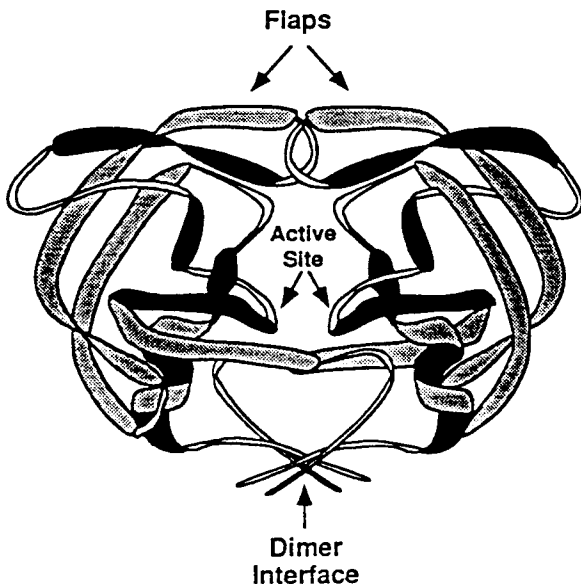
BUDDING AT SITE OF ENV PROTEIN: MOST HOST TM PROTEINS EXCLUDED



HOW DOES CORE RECOGNIZE ENV-RICH PLASMA MEMBRANE?

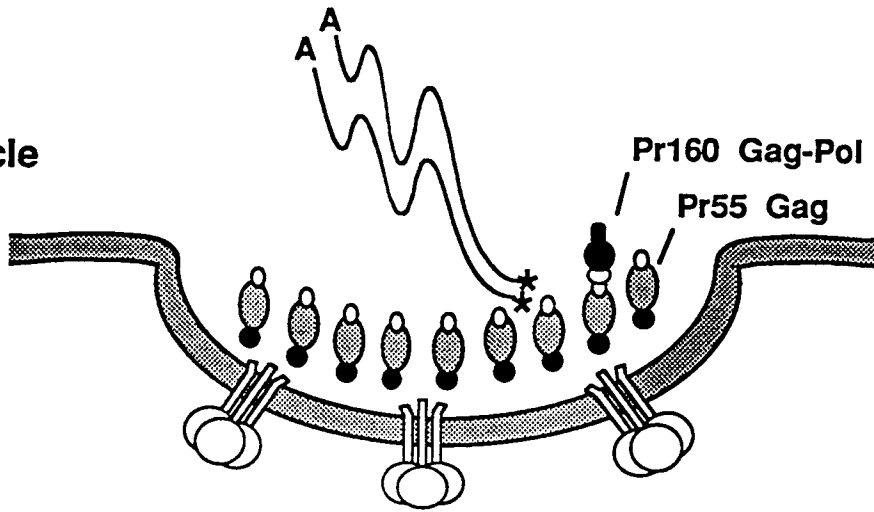
MATURATION: DIMERIZATION OF PROTEASE REQUIRED FOR ACTIVITY,  
CLEAVES GAG AND GAG-POL AT SPECIFIC SITES

COINCIDENT WITH CORE CONDENSATION, INFECTIVITY, RT ACTIVITY



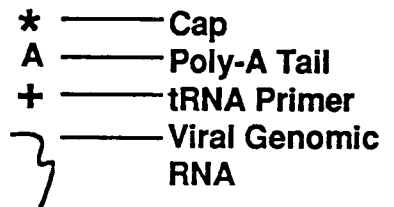
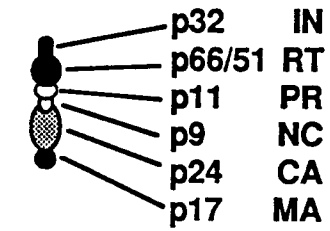
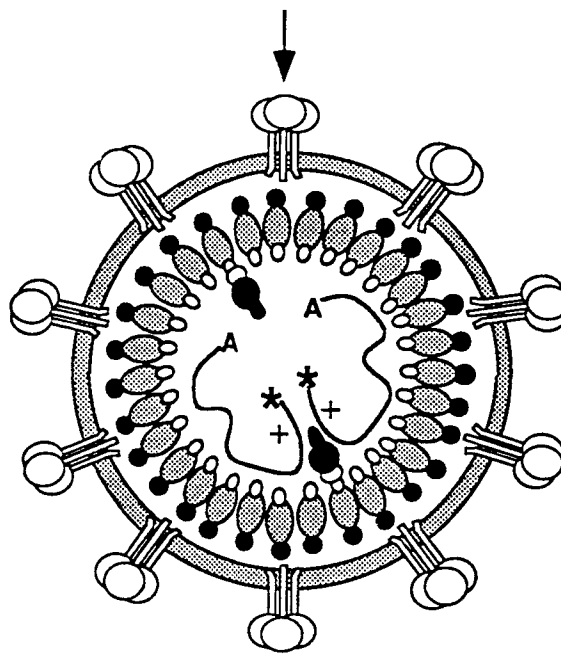
	Sites	Sequence
RSV	junction	P4 P3 P2 P1 P1' P2' P3'
<i>gag</i>	MA -	Gly - Thr - Ser - Cys - Tyr
<i>gag</i>	MA - p10	Pro - Pro - Tyr - Val - Gly
<i>gag</i>	p10 - CA	Pro - Val - Val - Ala - Met
<i>gag</i>	CA -	Ile - Ala - Ala - Ala - Met
<i>gag</i>	- NC	Gln - Pro - Leu - Ile - Met
<i>gag</i>	NC - PR	Pro - Pro - Ala - Val - Ser
<i>pol</i>	- RT	Arg - Ala - Thr - Val - Leu
<i>pol</i>	RT $\alpha$ - IN	Thr - Phe - Gln - Ala - Tyr
<i>pol</i>	IN -	Ser - Pro - Leu - Phe - Ala
HIV	-	
<i>gag</i>	MA - CA	Val - Ser - Gln - Asn - Tyr
<i>gag</i>	CA -	Lys - Ala - Arg - Val - Leu
<i>gag</i>	- NC	Thr - Ala - Thr - Ile - Met
<i>gag</i>	NC -	Arg - Pro - Gly - Asn - Phe
<i>gag</i>	- p5	Glu - Arg - Gln - Ala - Asn
<i>pol</i>	- PR	Val - Ser - Phe - Asn - Phe
<i>pol</i>	PR - RT	Cys - Thr - Leu - Asn - Phe
<i>pol</i>	RT $\beta$ - IN	Ile - Arg - Lys - Ile - Leu

**Budding  
Virus Particle**

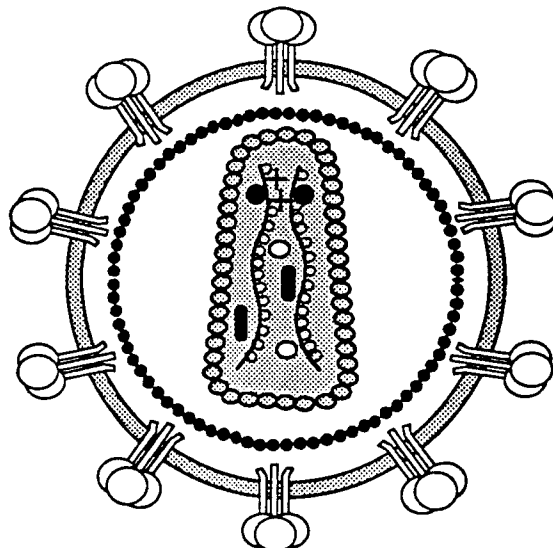


FOR VIRUS  
ASSEMBLING  
CORE AT P.M.

**Immature  
Virus Particle**



**Mature  
Virion**



. Model for virion assembly.



# RETROVIRAL GENETICS

## COMPLEMENTATION

HELPER GENOMES FOR GAG, POL, ENV COMPLEMENTATION

ENV PSEUDOTYPES (BETWEEN ANY TWO RETROVIRUSES,

RETROVIRUSES AND VSV) AFFECT HOST RANGE

EXTREME CASES:

HELPER CELLS

FOR RV VECTOR

## MUTATION

RATES SIMILAR TO OTHER RNA VIRUSES

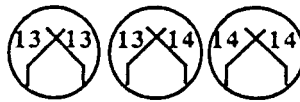
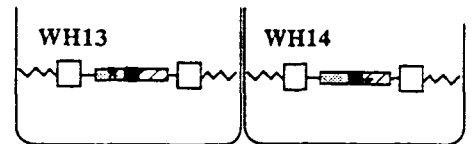
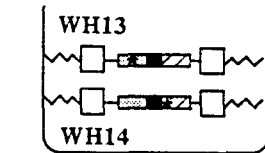
## RECOMBINATION

HIGH FREQUENCY (TOO HIGH FOR GENETIC MAPPING!)

HETEROZYGOTES ARE OBLIGATE INTERMEDIATES

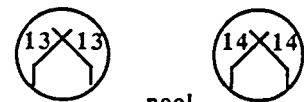
RECOMBINANTS FORMED DURING REVERSE TRANSCRIPTION

HU ←  
TEMIN  
1990



infect D17 cells

Recombinant proviruses



pool

infect D17 cells

Recombinant proviruses  
were not observed

## CONSEQUENCES OF INTEGRATION

INSERTION MUTATIONS (ACTIVATING OR INACTIVATING)

INTRODUCTION INTO GERM LINE

→ ENDOGENOUS  
PROVIRUSES

→ TRANSDUCTION OF  
HOST GENES...

## ENDOGENOUS PROVIRUSES

### LEVELS OF COMPETANCE

- \* ENCODE INFECTIOUS VIRUS (MMTV, XENOTROPIC MLV, RD114 ETC)
- \* ENCODE VIRAL PROTEINS ONLY (GAG OR ENV)
- \* NO PRODUCTS

### NUMBERS AND ORIGINS

- \* RECENT ENTRY (POST-SPECIATION)... VARIATION AMONG  
INDIVIDUALS, 0-10 COPIES (ECOTROPIC MLV, MMTV, ALV)
- \* ANCIENT SURVIVORS.... UNIFORM IN SPECIES, MAY BE 100 TO 1000  
COPIES (XENOTROPIC MLV, IAP'S, ENDOGENOUS HUMAN PV'S)

### OBSERVED GERM LINE INSERTIONS

- \* SOME VIRUS-PRODUCING MOUSE LINES
- \* INFECTION OF PRE-IMPLANTATION EMBRYOS
- \* ??INFECTION OF MALE GERM CELLS

### MUTAGENIC POTENTIAL (EXAMPLES)

- \* IN GERM LINE: HPRT, DILUTE (REVERTS BY LTR-LTR EXCISION)  
COLLAGEN ALPHA 1A (MOV 13), OTHER DEVELOPMENTALS
- \* MOVEMENT IN SOMATIC CELLS: IAP INTO PROTO-ONCOGENES

ROLE IN EVOLUTION?