## Human Immunodeficiency Virus Type 1 nucleotide sequence unanimity alignments, Complete Pol Region. Highlighted regions indicate the nucleotide sequence encoding for the Protease, Reverse Transcriptase & Integrase enzymes.

	Ribsomal slip regionRibsomal slip 2085
нхв2	 TTT TTT AGG GAA GAT CTG GCC TTC CTA CAA GGG AAG GCC AGG GAA TTT TCT TCA GAG CAG ACC AGA GCC AAC AGC
MXHIV00086	*** ***
MXHIV00327 MXHIV00395	*** ***
MXHIV00401	*** *
MXHIV00422	*** ***
HXB2 MXHIV00086 MXHIV00395 MXHIV00401 MXHIV00422 HXB2 MXHIV0086 MXHIV00327 MXHIV00395 MXHIV00422	=
MXHIV00422	G A G G C   <mark>A AG</mark>   <mark> GAA</mark>
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	Protease (p15)
	2379
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	2557       ATT 500       ATT 500
	Protease (p15)
	Xerminal domain 2454 2528
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	GGA CAT AAA GCT ATA GGT ACA GTA TTA GTA GGA CCT ACA CCT GTC AAC ATA ATT       GGA AGA AAT CTG TTG ACT CAG         A G A A

	Protease (p15)> < Reverse transcriptase (p66, p51)   p66 domain and p51 domain start	
	Terminal domain> <	2603
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	$\begin{bmatrix} 1 \\ 1 \\ 1 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 1 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 1 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 \\ 2 \\ 2 \\ 2 \end{bmatrix}$ $\begin{bmatrix} 1 $	AT GGC
	Reverse transcriptase (p66, p51) Finger subdomain 2604	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	 CCA AAA GTT AAA CAA TGG CCA TTG ACA GAA GAA AAA ATA AAA GCA TTA GTA GAA ATT TGT ACA GAG ATG GAA 	A
	Reverse transcriptase (p66, p51) Reverse transcriptase (p66, p51)	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	GAA       GGG       AAA       ATT       GGG       CCT       GAA       AAT       CCA       TAC       AAT       ACT       CCA       GTA       TTT       GCC       ATA       AAG       AAA       GAC	
	> Reverse transcriptase (p66, p51)Palm subdomain >  <palm subdomain<="" td=""><td></td></palm>	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	ACT       AAA       TGG       AGA       ATA       GAA       GAT       GAT       TC       AGA       GAA       CTT       AAA       AAA       AAA       AAA       TTA       GAA       GAT       TC       AGA       AAA       AAA       AAA       TTC       TC       AGA       GAA       CTT       AAA       AAA       AAA       AAA       TTC       TC       AGA       GAA       CTT       AAA       AAA       AAA       AAA       TTC       TTC       AGA       AAA       AAA </td <td>G</td>	G
	Reverse transcriptase (p66, p51) Palm subdomain 1	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	GTT   C
	Reverse transcriptase (p66, p51)	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	I         CCC TTA GAT GAA GAC TTC AGG AAG TAT ACT GCA TTT ACC ATA CCT AGT ATA AAC AAT GAG ACA CCA GGG A         AG	
	Finger subdomain	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	TAT       CAG       TAT       CAG       GTG       CTT       CCA       CAG       GGA       TGG       AAA       GGA       TCA       CCA       AAT       TTC       CAA       AGC       ATG       ACA       AAA       ATC	

HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	ATA GTG CTG CCA GAA AAA GAC AGC TGG ACT GTC AAT GAC ATA CAG AAG TTA GTG       GGG AAA TTG AAT TGG GCA AGT
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	

	Reverse transcriptase (p66, p51)
	Connection subdomain
HXB2 MXHIV00086	GAG CCA TTT AAA AAT CTG AAA ACA GGA AAA TAT GCA AGA ATG AGG GGT GCC CAC ACT AAT GAT GTA AAA CAA TTA 
MXHIV000327 MXHIV00395	AG
MXHIV00401 MXHIV00422	A
	Reverse transcriptase (p66, p51)
	Connection subdomain
HXB2 MXHIV00086	ACA GAG GCA GTG CAA AAA ATA ACC ACA GAA AGC ATA GTA ATA TGG GGA AAG ACT CCT AAA TTT AAA CTG CCC ATA
MXHIV00327 MXHIV00395	A G CT A
MXHIV00401 MXHIV00422	C- G
	Reverse transcriptase (p66, p51)
	Connection subdomain
HXB2 MXHIV00086	CAA AAG GAA ACA TGG GAA ACA TGG TGG ACA GAG TAT TGG CAA GCC ACC TGG ATT CCT GAG TGG GAG TTT GTT AAT
MXHIV00327 MXHIV00395	
MXHIV00401 MXHIV00422	G G G T
	Reverse transcriptase (p66, p51)
	Connection subdomain> RNAse H ( 3804 3869  3870
HXB2 MXHIV00086	ACC CCT CCC TTA GTG AAA TTA TGG TAC CAG TTA GAG AAA GAA CCC ATA GTA GGA GCA GAA ACC TTC TAT GTA GAT
MXHIV00327 MXHIV00395	
MXHIV00401 MXHIV00422	TA  TAC
	Reverse transcriptase (p66, p51)
	RNAse H (p15)
HXB2 MXHIV00086	GGG GCA GCT AAC AGG GAG ACT AAA TTA GGA AAA GCA GGA TAT GTT ACT AAT AGA GGA AGA CAA AAA GTT GTC ACC
MXHIV00327 MXHIV00395	
LIVIT 1 0 0 0 2 2 2	
MXHIV00401	
MXHIV00401	G-C -A G-G T T-T G-C -A G-G
MXHIV00401	
MXHIV00401	G-C -A G-G T T-T G-C G-C G-C
MXHIV00401 MXHIV00422 HXB2	G-C -A G-G T T-T T-T G-C G-G G-G T T-T
MXHIV00401 MXHIV00422 HXB2 MXHIV00086 MXHIV00327	
MXHIV00401 MXHIV00422 HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401	$\begin{array}{cccccccccccccccccccccccccccccccccccc$
MXHIV00401 MXHIV00422 HXB2 MXHIV00086 MXHIV00327 MXHIV00395	$\begin{array}{cccccccccccccccccccccccccccccccccccc$
MXHIV00401 MXHIV00422 HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
MXHIV00401 MXHIV00422 HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
MXHIV00401 MXHIV00422 HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422 HXB2	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
MXHIV00401 MXHIV00422 HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	$\begin{array}{cccccccccccccccccccccccccccccccccccc$

	Reverse transcriptase (p66, p51)
	4104 4178
HXB2 MXHIV00086	AAT CAA ATA ATA GAG CAG TTA ATA AAA AAG GAA AAG GTC TAT CTG GCA TGG GTA CCA GCA CAC AAA GGA ATT GGA
MXHIV00327	
MXHIV00395 MXHIV00401	-G ACA
MXHIV00422	-GAAA C A
	> < Reverse transcriptase (p66, p51)> < Integrase (p31)
	N terminal domain 4179 4229   4230 4253
НХВ2	 Gga aat gaa caa gta gat aaa tta gtc agt gct gga atc agg aaa gta cta   <mark>ttt tta gat gga ata gat aag gcc</mark>
MXHIV00086 MXHIV00327	
MXHIV00395	T-C
MXHIV00401 MXHIV00422	TGC AAWYAR  YR
	Integrase (p31)
	N terminal domain
нхв2	   CAA GAT GAA CAT GAG AAA TAT CAC AGT AAT TGG AGA GCA ATG GCT AGT GAT TTT AAC CTG CCA CCT GTA GTA GCA
MXHIV00086	
MXHIV00327 MXHIV00395	ACG
MXHIV00401 MXHIV00422	ACACGG
	Integrase (p31)
	Catalitic core domain
_	
HXB2 MXHIV00086	AAA GAA ATA GTA GCC AGC TGT GAT AAA TGT CAG CTA AAA GGA GAA GCC ATG   CAT GGA CAA GTA GAC TGT AGT CCA G
MXHIV00327 MXHIV00395	
MXHIV00401 MXHIV00422	A
111111000122	
	Catalitic core domain
	4404 4478
HXB2 MXHIV00086	GGA ATA TGG CAA CTA GAT TGT ACA CAT TTA GAA GGA AAA GTT ATC CTG GTA GCA GTT CAT GTA GCC AGT GGA TAT
MXHIV00327 MXHIV00395	
MXHIV00401	
MXHIV00422	G C <u>A</u> T
	Catalitic core domain
	4479 4553
HXB2	ATA GAA GCA GAA GTT ATT CCA GCA GAA ACA GGG CAG GAA ACA GCA TAT TTT CTT TTA AAA TTA GCA GGA AGA TGG
MXHIV00086 MXHIV00327	
MXHIV00395 MXHIV00401	
MXHIV00422	A A A A A A A C C CCC
	Catalitic core domain
	4628 4628
HXB2	CCA GTA AAA ACA ATA CAT ACT GAC AAT GGC AGC AAT TTC ACC GGT GCT ACG GTT AGG GCC GCC TGT TGG TGG GCG
MXHIV00086 MXHIV00327	
MXHIV00395 MXHIV00401	
MXHIV00401 MXHIV00422	

----- Reverse transcriptase (p66, p51) -----

	Integrase (p31)
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
	Catalitic core domain
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	AAG AAA ATT ATA GGA CAG GTA AGA GAT CAG GCT GAA CAT CTT AAG ACA GCA GTA CAA ATG GCA GTA TTC ATC CAC $$
	Catalitic core domain
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	$\begin{array}{cccccccccccccccccccccccccccccccccccc$
	Integrase (p31)
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	CAA ACT AAA GAA TTA CAA AAA CAA ATT ACA AAA ATT CAA AAT TTT CGG GTT TAT TAC AGG GAC AGC AGA AAT CAA 
	N terminal domain
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	$\begin{array}{cccccccccccccccccccccccccccccccccccc$
	Integrase (p31)
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	GTA GTG CCA AGA AGA AAA GCA AAG ATC ATT AGG GAT TAT GGA AAA CAG ATG GCA GGT GAT GAT TGT GTG GCA AGT
HXB2 MXHIV00086 MXHIV00327 MXHIV00395 MXHIV00401 MXHIV00422	Integrase (p31)> 5079 5096 5114 RGA CAG GAT GAG GAT TAG *** *** *** *** *** *** AAC ATG GAA CAG CCT AG* AAC ATG GAA *** *** *** AAC ATG GA* *** **** AAC ATG GA* *** **** AAC ATG GA* *** **** AAC ATG GA* *** ****

Alignments produced from partial overlapping clones of viral RNA sequences obtained from human blood samples. Clones, contigs, alignments, annotations and maps produced by Pedro Gerardo Hernández Sánchez, Viral & Human Genomics Laboratory UASLP, México. Unanimity reformating script written by James Robinson, Anthony Nolan Research Institute, London.

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