

Supplementary Data

SUPPLEMENTARY TABLE S1. PRIMERS USED FOR POLYMERASE CHAIN REACTION
AMPLIFICATION OF HIV-1 NEAR FULL-LENGTH GENOMES

<i>Name</i>	<i>Hxb2 position</i>	<i>Region</i>	<i>Sequence (5'-3')</i>
JA153	683–707	p17	CTCTCGACGCAGGACTCGGCTTGCT
P17-894F	894–913	p17	ATGGGCAAGCAGGGAGCTGG
P17-913R	913–894	p17	CCAGTCCCTGCTTGCCCAT
P24-1300F	1300–1324	p24	ATACCCATGTTT(A/T)CAGCATTATCAG
JA155 (R)	1324–1300	p24	CTGATAATGCTGAAAACATGGGTAT
P24-1818F	1818–1838	p24	AGAAGAAATGATGACAGCATG
P24-1838R	1838–1818	p24	CATGCTGTCATCATTTCTTCT
IBPR1.1	2008–2030	gag NC	AAAAGGGCTGTTGGAAATGTGG
IBPR2.2	2733–2712	RT	GCAAATACTGGAGT(A/G)TT(G/A)TATG
IBPR3.1	2119–2140	gag P1-P6	AGGCCAGGGAATTT(T/C)C(T/C)TCAGA
IB2621PR4	2591–2621	PR	AATGCTTTTTATTTT(C/T)TCTTCTGTCAATGGC
IBRT1	2487–2505	RT	CCTACACCTGTCAACATAA
IBRT2	3649–3630	RT	TGTTTTACATCATTAGTGTG
IBRT3	2542–2560	RT	TAAATTTTCCAATTAGTCC
IBRT4	3579–3560	RT	TAAATTTGATATGTCCATG
RT3482FI	3482–3509	RT	AGAACCAGTACATGGRGTTATATTATGA
IB3626RT2	3626–3593	RT	TCCGTTAA(C/T)TGT(C/T)TTACATCATT AGTGTG(A/G)GCA
RT3864F	3864–3881	RT	CAACAAATCA(A/G)AAGACTG
P15-3881R	3881–3864	p15	CAGTCTT(T/C)TGATTTGTG
P15-4176F	4176–4194	p15	GGAGGAAATGAACAAGTAG
P15-4194R	4194–4176	p15	CTACTTGTTCATTTCCCTCC
P31-4658F	4658–4675	p31	CAATCCCCAAAGTCAAGG
P31-4675R	4675–4658	p31	CCTTGACTTTGGGGATTG
Vif5041F	5041–5059	vif	ATGGAAAACAGATGGCAGG
Vif5059R	5059–5041	vif	CCTGCCATCTGTTTTCCAT
Vif5461F	5461–5476	vif	AAGGTAGGATC(T/C)(T/C)TAC
Vif5476R	5476–5461	vif	GTA(A/G)(A/G)GATCCTACCTT
PBENV1	5968–5986	REV	CTATGGCAGGAAGAAGCGG
REV5986R	5986–5968	REV	CCGCTTCTTCCTGCCATAG
ENV6223RI	6223–6203	REV	CCACTGTCTTCTGCTCTTTC
PBENV2	6203–6223	REV	GAAAGAGCAGAAGAYAGTGGC
ENV8637FI	8637–8657	gp41	CAGGAACTAAAGAATAGTGC
PBENV4	8797A8817	NEF	TTTTGACCACTTGCCCHCCCAT
PBENV3	9036–9016	NEF	AGTCATTGGTCTTARAGGTAC
NEF9532RI	9532–9511	3'LTR	GCGAAAAGCRGCTGCTTATAT