Additional materials

Additional file 1: Table S1. Characteristics of study participants at time of pyrosequencing.

PID ^a	Viral load		Age/ length	
	Plasma ^b	Cell- associated ^c	of infection (years)	CD4 % ^d
S 1	4.0	2.6	4.5	24
S2	4.8	4.5	1.5	30
S3	4.9	3.4	4.8	22
S4	4.9	3.3	2.1	17
S 5	5.2	5.0	4.2	22
S6	5.7	5.5	6.1	2

^a Patient identification. Subjects included in pyrosequencing study were infected perinatally by maternal transmission for more than one year, were naïve to combination antiretroviral therapy with viral RNA levels >4 log₁₀ copies per ml of plasma and genetic analysis of HIV-1 envelope by conventional clonal sequencing.

^b Log₁₀ HIV-1 RNA copies per ml of plasma by measuring HIV-1 *gag* gene (COBAS® AMPLICOR HIV-1 MONITOR Test, v1.5, Roche, Pleasanton, CA, USA).

 $^{^{\}mathrm{c}}$ Log₁₀ HIV-1 gag copies/10 $^{\mathrm{6}}$ CD4 $^{\mathrm{+}}$ T cells

^d CD4% is used to normalize for the age-related lymphocytosis that occurs in pediatric subjects.