Supplementary Figure S2. Allele frequencies of polymorphic HML-2 5’ LTR transcription factor binding sites within each superpopulation. Allele frequencies were determined for the proviruses shown from 2,504 individuals from the 1000 Genomes Project and broken down by superpopulation. EAS = East Asian; AMR = Ad Mixed American; AFR = African; EUR = European; SAS = South Asian.