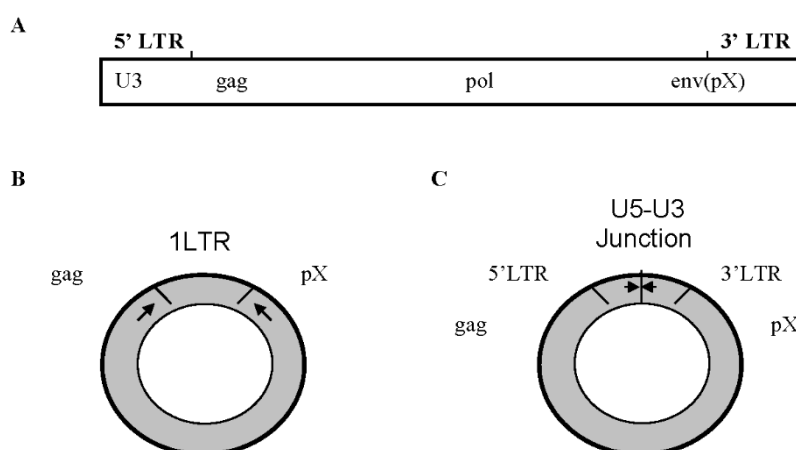


# Supplementary Materials: Long Terminal Repeat Circular DNA as Markers of Active Viral Replication of Human T Lymphotropic Virus-1 *in Vivo*

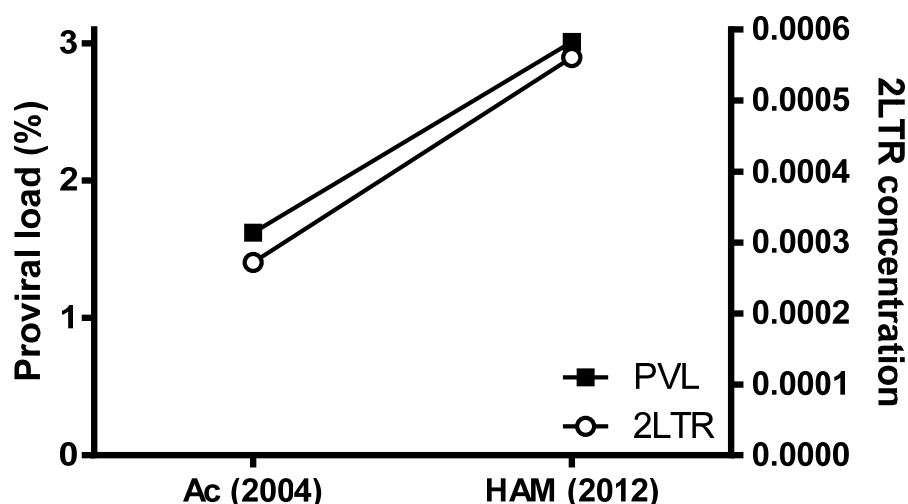
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**Table S1.** Demographics of the patients with HTLV-1 from whom DNA samples for LTR quantification were taken.

Demographics	Frequency (%)	Range
Female	33 (70)	
Afro-Caribbean	37 (80)	
African	4 (8%)	
Caucasian	4 (8 %)	
Persian	1 (2%)	
South American	1 (2%)	
Age	Years	
Average age at sampling	56.1	19–80



**Figure S1.** Structures of extrachromosomal DNA in HTLV infected cells and primer design rational. Linear viral DNA (A) is synthesized in the cytoplasm after cell infection and can circularise to form 1- (B) or 2-LTR DNA circles (C). 1LTR DNA circles are formed through the deletion of either LTR; 2LTR DNA circles are formed at the U3-U5 junction where the LTRs from each end of the HTLV-1 DNA connect. Specific primer pairs (arrows) were designed to selectively amplify circularised DNA (sequences are shown in Table 1). LTR: long terminal repeats; gag, pol, env(pX): gene regions coding for viral proteins.



**Figure S2.** PVL and 2LTR DNA circle levels increase in an incident patient progressing from asymptomatic HTLV-1 carrier to being diagnosed with HAM. The PVL (■) and 2LTR levels (○) were quantified in PBMCs from an incident asymptomatic patient in 2004 and from the same patient after disease progression and HAM diagnosis in 2012. Incident patient demographics: male; 20 years of age at diagnosis; black ethnicity (skin colour, self-definition). His risk factors were: breast-feeding from an HTLV positive mother and his brother was also HTLV positive. Blood transfusion denied. His date of diagnosis was on his enrolment into the cohort: 10 October 2004 when he was asymptomatic. The patient was seen again on 9 May 2009 and he was still asymptomatic but Rossolimo's sign was present. Onset of symptoms: 9 September 2011 (HAM/TSP).



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