Supplementary Figure 2  DNA methylation in the 5’ LTR promoter in HTLV-1-infected clones

(a) Region amplified by PCR for bisulfite sequencing. The three Tax-responsive elements are indicated in red. (b) Schematic representation of DNA methylation. Each bar represents a PCR fragment that was cloned and sequenced. Open circles indicate unmethylated cytosine, and closed circles methylated cytosine. The numbers correspond to the positions shown in Figure 5a. Clone A does not have CpG at position 6, so this position is omitted. (c) DNA methylation in the 5’ LTR promoter in mutated HTLV-1-infected clones that lack CTCF binding in the provirus.