

Description of Additional Supplementary Files

Supplementary Data 1: Overview of the statistical tests performed in this study (Figures 3C, 3E, and 4B).

Supplementary Data 2: Masterfile of the single-sorted p24+ cells for STIP-Seq analyses performed on 6 participants (MRC01, MRC04, MRC08, MRC15, MRC21, MRC25). For each sorted cell, the table contains information about the stimulation conditions, the phenotype (TCM/TTM/TEM/TTd), the integration site, the list of all near full length PCRs that have been tested, and the proviral genome classification.