Figure S7

A

![Graph showing average number of genes in 1MB window for integration sites and random controls. Controls include WT, G99V, N74D, N57A, P90A, HIV-1(SIVCA). The x-axis represents different mutants.]

B

![Graphs showing infected cells per field for Donors 3 and 4. The x-axis represents days post-infection (p.i.), and the y-axis represents infected cells per field. The y-axis ranges from 0 to 3000 with intervals of 500. Experiments include WT, N74D, P90A, and CA (Ba-L Env).]