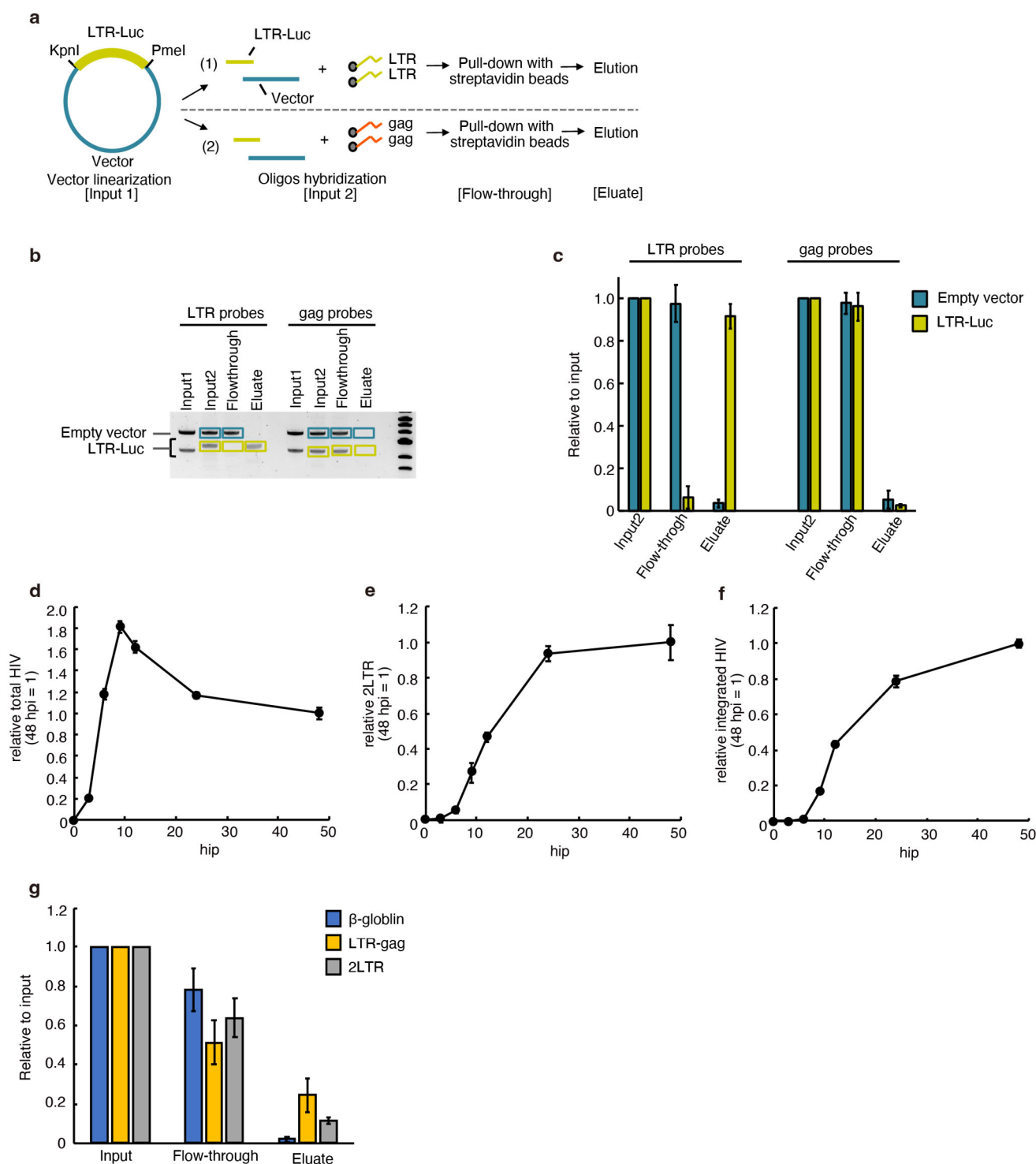




Extended Data Fig 1. Optimization of PICh method to the isolation of unintegrated HIV-1 DNA in vitro and in infected cells.

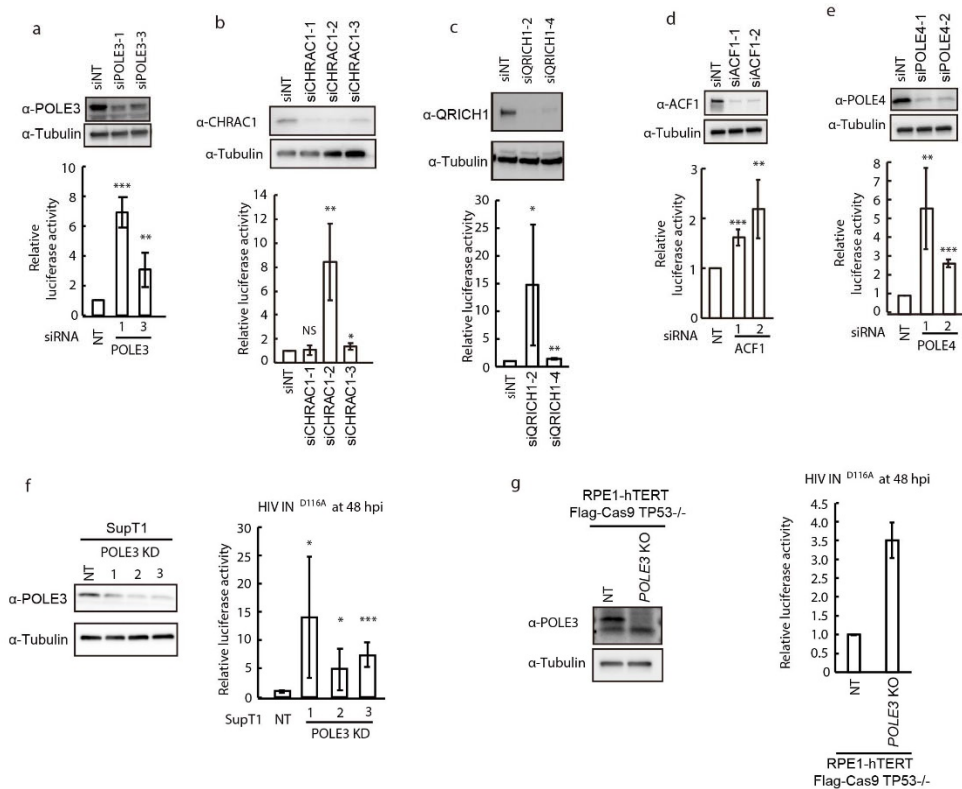
(a) Outline of the plasmid pull-down assay using LTR-luc plasmid. Linearized plasmid was mixed with PICh probes specific for the LTR (1) or Gag (2) conjugated to spacer and desthiobiotin. After hybridization, hybrids are captured using streptavidin beads and eluted with biotin.

(b) Each fraction (input, flow-through, and eluate) were analyzed by agarose gel electrophoresis and EtBr staining.

(c) Quantification of target (LTR-luc) capture represented as percent DNA relative to input 2.

(d-f) SupT1 cells were infected with a VSV-G pseudotyped HIV-Luc at a MOI of 0.8. Cells were harvested at 3, 6, 9, 12, 24 and 48 hpi. Total HIV DNA (d), 2LTR circles (e) and integrated DNA (f) were analyzed by qPCR. Data are represented as quantification relative to the value at 48 hpi. The mean values $\pm$ SD of three independent experiments are plotted.

(g) Genome capture of genomic DNA from SupT1 cells infected with HIV-Luc for 9 hrs. Genomic DNA was hybridized with probes designed for PICh, captured on streptavidin beads and eluted with biotin. Quantitative PCR analysis of input, flow through and eluate fractions using indicated primers. Results are represented as quantification relative to input. The mean values $\pm$ SD of four independent experiments are plotted.



Extended Data Fig 2. CHRAC1 and QRICH1 are not involved in unintegrated HIV-1 DNA silencing.

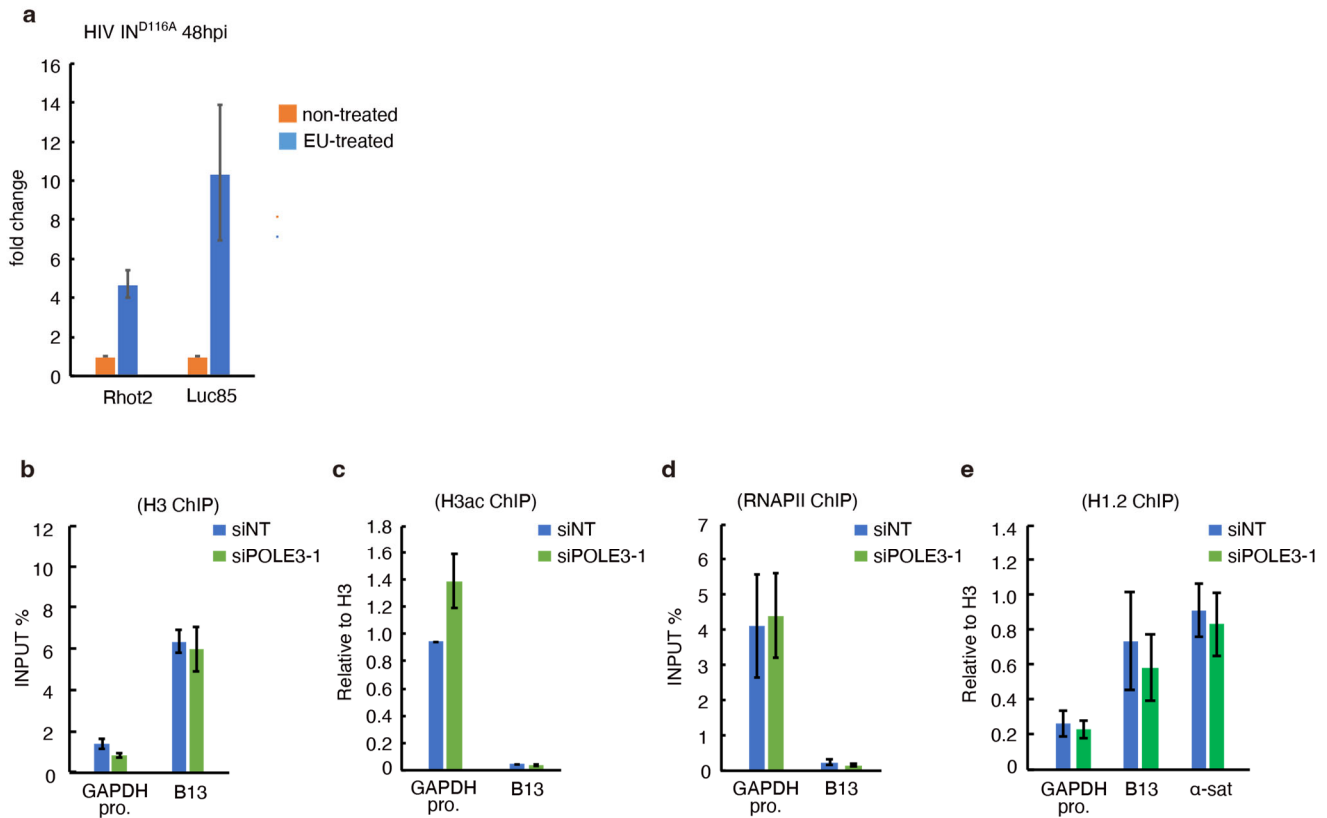
(a-e) Expression of POLE3 (a), CHRAC1 (b), QRICH1 (c), ACF1 (d) and POLE4 (e) proteins in cells treated with indicated siRNA were analyzed by immunoblotting with indicated antibodies. Luciferase assay in KD cells were performed at 48 hpi. Results are represented as luciferase expression relative to NT siRNA, and the mean $\pm$ SD values of 2 independent experiments in triplicate are plotted.

(f) Impact of knockout of POLE3 in SupT1 cells on uniHIV-1 DNA silencing. Expression of POLE3 proteins in POLE3 KO cells were analyzed by immunoblotting with indicated antibodies. Luciferase assays in POLE3 KO cells were performed at 48 hpi. Results are represented as luciferase expression relative to NT sgRNA, and the mean $\pm$ SD values of two independent experiments in triplicate are plotted.

luciferase expression relative to non-targeting (NT) siRNA, and the mean $\pm$ SD values of 6 independent experiments are plotted.

(e-f) Effect of POLE3 KO on uniHIV DNA silencing in RPE1-hTERT Flag-Cas9 TP53^{-/-} cells. POLE3 expression from whole cell extracts were analyzed by immunoblotting with anti-POLE3 antibody at the day of infection (e). Luciferase assays were performed at 48 hpi (f). Results are represented as luciferase expression relative to non-targeting (NT) sgRNA, and the mean $\pm$ SD values of three independent experiments in triplicate are plotted.

* $P < 0.05$, ** $P < 0.01$ independent Student's t-test.

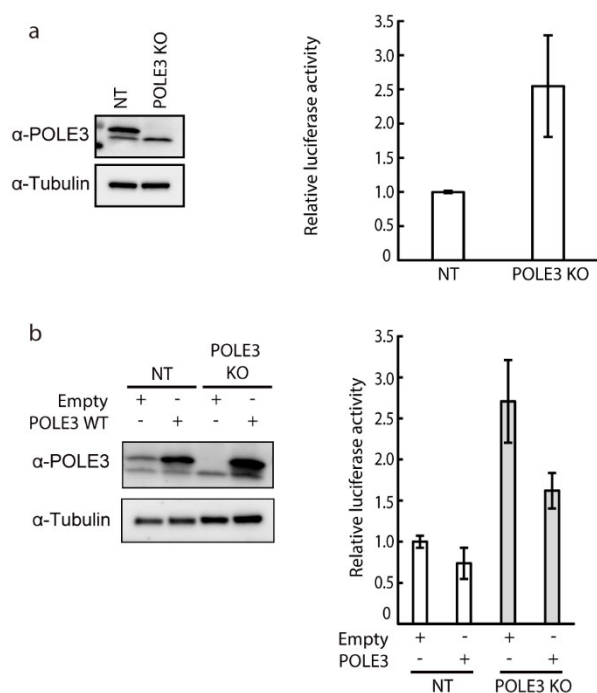


Extended Data Fig 3. Nascent HIV RNA IP and ChIP qPCR assays on human genomic region.

(a) Enrichment of EU labeled RNAs synthesized in HeLa cells infected with VSV-G pseudotyped HIVLuc IN^{D116A} at 48hpi. RNAs were extracted from HeLa cells treated with or without EU. Labeled RNAs were biotinylated, and purified with streptavidin conjugated beads. Enrichment of labeled RNAs are presented as fold change over non-labeled condition. The mean $\pm$ SD values of 2 independent experiments in triplicate are plotted.

(b-e) ChIP assay using NT (blue) and POLE3 (green) KD HeLa cells infected with VSV-G pseudotyped HIV-Luc IN^{D116A}. ChIP assays were performed at 48 hpi, using anti-H3 (b), anti-H3ac

(c), anti-RNAPII (d), anti-H1.2 (e) antibodies, corresponded to Fig 3b-e. qPCR analysis was performed using specific primers for GAPDH promotor, B13, and α -satellite. The mean $\pm$ SD values of three independent experiments are plotted. The mean $\pm$ SD values of at least 3 independent experiments are plotted.



Extended data Fig 4. POLE3 KO HeLa-P4 cells

(a) Impact of knockout of POLE3 in HeLa-P4 cells on uniHIV-1 DNA silencing. Expression of POLE3 proteins in KO cells were analyzed by immunoblotting with indicated antibodies. Luciferase assays in POLE3 KO cells were performed at 48 hpi. Results are represented as luciferase expression relative to NT sgRNA, and the mean $\pm$ SD values of at least 3 independent experiments in triplicate are plotted.

(b) Exogenous POLE3 expression in POLE3 KO cells silences uniHIV-1 DNA. POLE3 KO HeLa-P4 cells stably expressing exogenous POLE3 wt, were infected with VSVG pseudotyped HIV-Luc IN^{D116A}. Whole cell extracts were analyzed by immunoblotting with indicated antibodies. Luciferase activity was analyzed at 48 hpi. Results are represented as luciferase expression relative to NT sgRNA of 3 independent experiment in triplicate.

