

Supplemental Materials

Molecular Biology of the Cell

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Fig. S1. GFP-53BP1 do not occupy nucleolus regions and YFP-NLS also segregates in small 3 μm constrictions

(A) U2OS cells were co-transfected with GFP-53BP1 and mApple-Fibrillarin, which indicates the presence of nucleoli. The representative image clearly show the exclusion of GFP-53BP1 and Hoechst stained DNA from the nucleoli.

(B) Representative images showing the segregation of YFP-NLS-MS2 inside the 3 μm pore.

Fig. S2. Acetylated histone H3 follows DNA behavior in constrictions.

(A,B) Following DNA behavior, the level of acetylated histone H3 (acetyl-H3) is enriched within the 3 μm pore constrictions, hence the intensity ratio is above one (≥ 15 cells; $n = 2$ expts). However, the enrichment of acetyl-H3 is not as high as the DNA, probing the need to test acetyl-H3 antibody specificity. (C) Indeed, inhibition of histone deacetylase by 300 nM Trichostatin A for 12 hours resulted in an increase in acetyl-H3 level, as shown by the antibody immunostaining (≥ 120 cells per group, $*p < 0.05$)

Fig. S3. Mobile protein segregation is an unavoidable physical effect of nuclear constriction.

Well over 90% of aspirated cells exhibit segregation; even those that do not appear to be within statistical uncertainty of doing so (≥ 4 cells per group; $n \geq 3$ expts).

Fig. S4 Nuclear FokI levels are higher with addition of tamoxifen and seem to accumulate over time. Nuclear FokI levels increase with the addition of tamoxifen to the 1 μM Shield1 treated cells. The amount of FokI in the nuclei is higher at 24 hours, when compared to the 6 hours treated cells (≥ 100 cells per group, $n \geq 2$ expts, $*p < 0.05$, $**p < 0.01$).

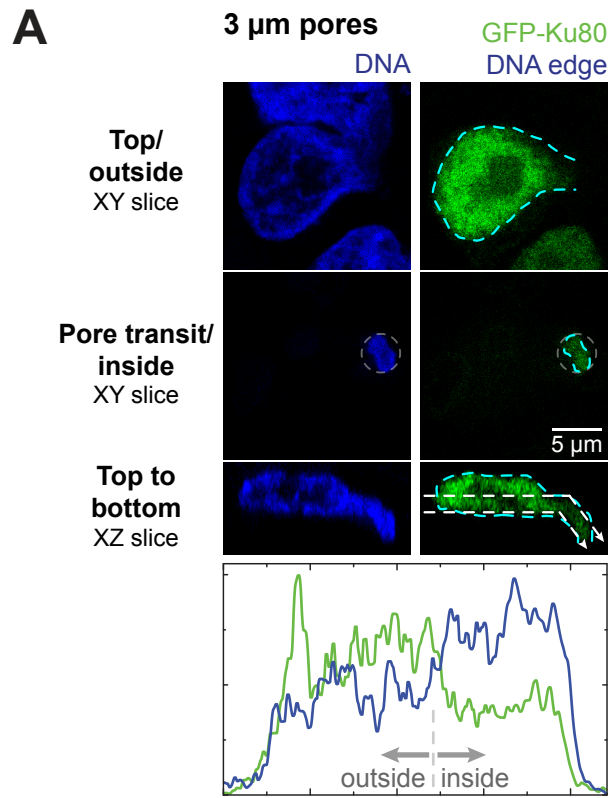
Fig. S5 H2B-GFP at the FokI foci are not depleted over time.

(A) Representative images of the observation of H2B-GFP and mCherry-FokI levels at the FokI focus region, over a period of 20 minutes. Intensity profiles across the FokI foci and the surrounding H2B-GFP show the depletion of FokI at the focus, but not H2B-GFP.

(B,C) Intensity profiles of two different nuclei, also show the depletion of FokI, but not H2B-GFP.

Figure S1

GFP-Ku80 also segregates in small 3 μm constrictions



Nuclear GFP-53BP1 proteins are excluded from the nucleoli

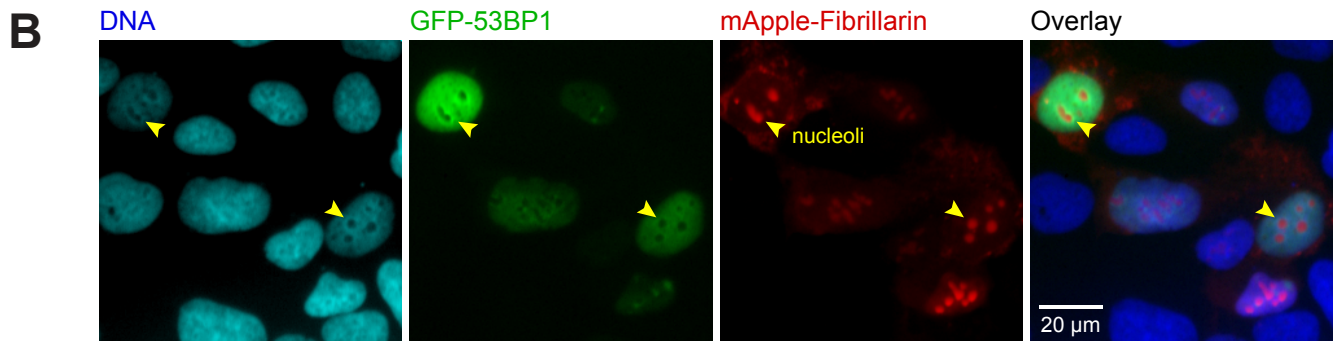


Figure S2

Acetylated histone H3 follows DNA and other chromatin-bound proteins behaviour in constrictions

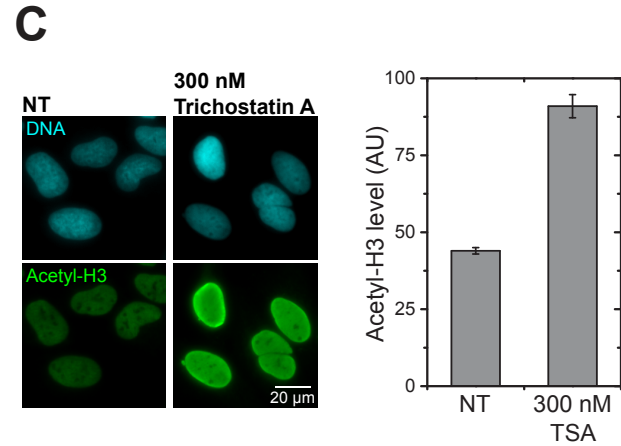
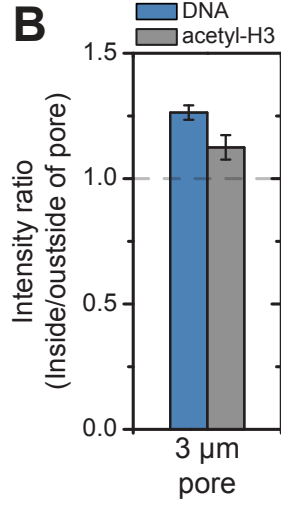
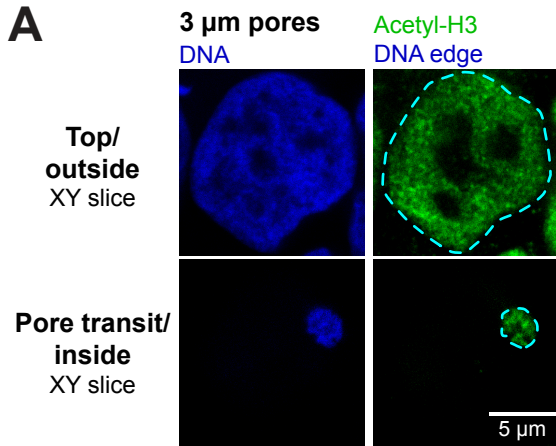


Figure S3

Scatter plot of intensity ratio data shows >90% of (DNA measurement ≥ 1.0) and (mobile factors ≤ 1.0)

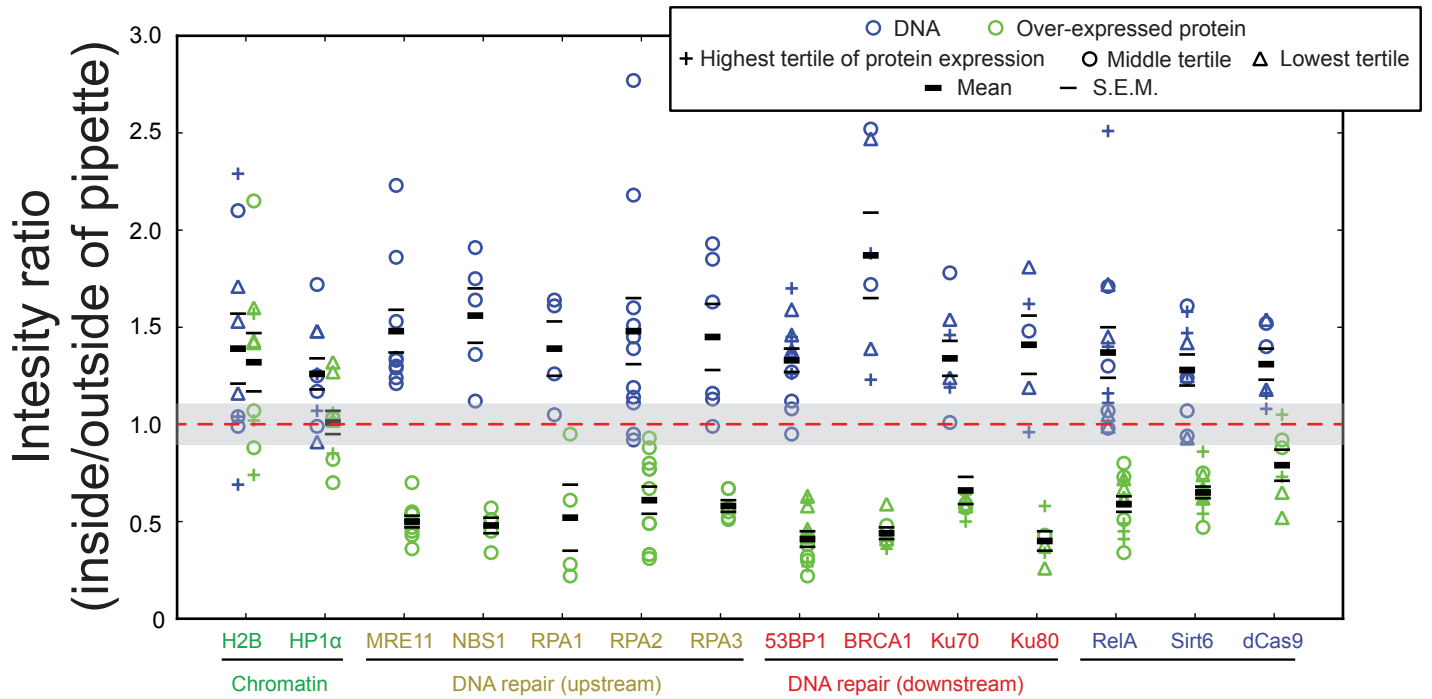


Figure S4

Nuclear FokI levels are higher with addition of Tamoxifen and seem to accumulate over time

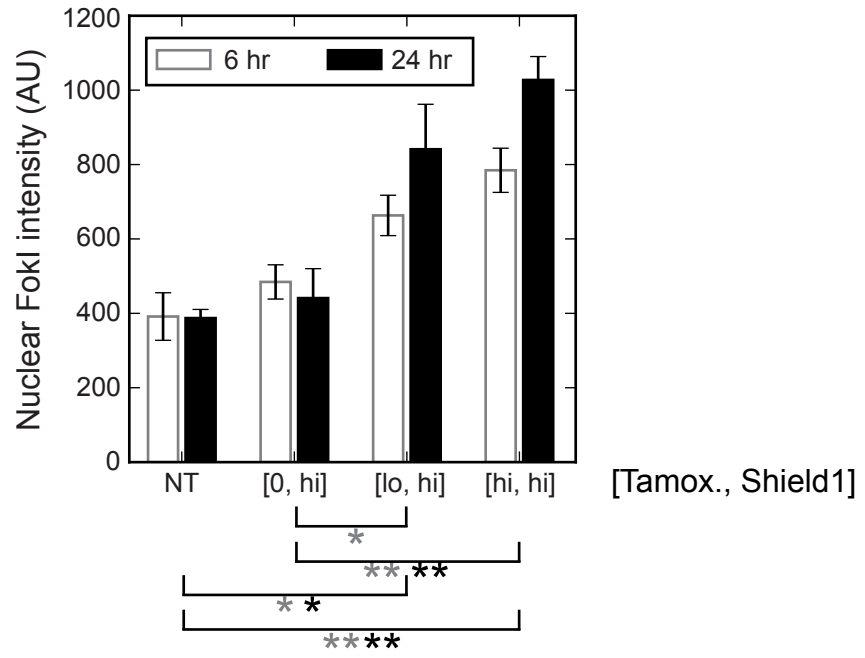
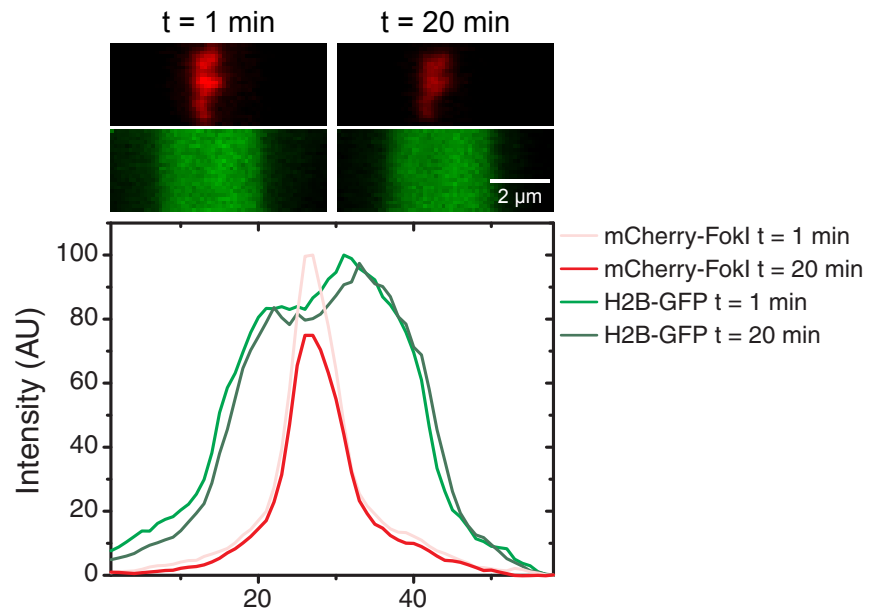
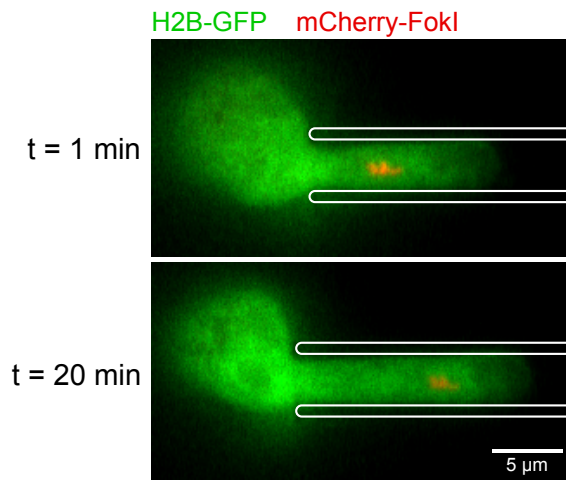


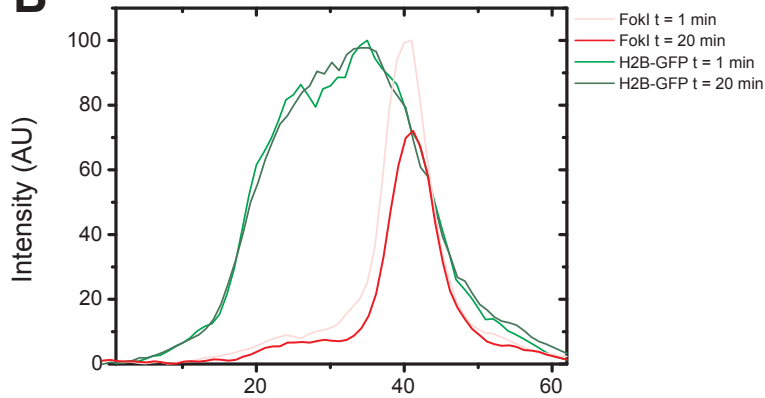
Figure S5

H2B remains in the FokI locus site during aspiration

A



B



C

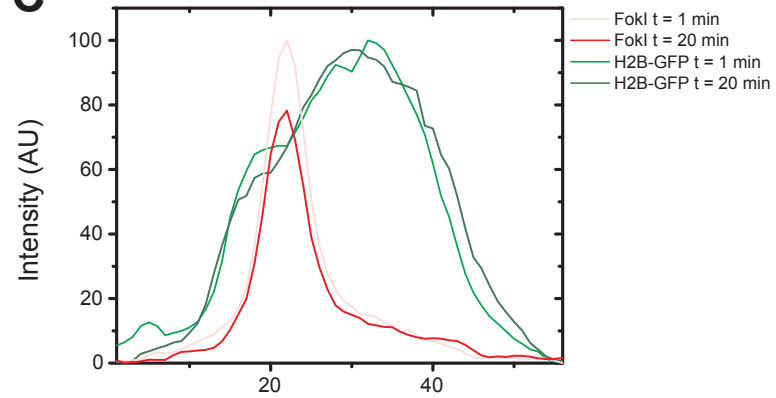


Table S1 FRAP data of non-foci bound nucleoplasmic GFP-tagged protein

GFP-tagged protein	Molecular weight ^[S1] , kDa (+GFP)	Estimated Isoelectric point, pI ^[S1] (+GFP)	Mobile fraction	recover half-time	recover time
RPA3	14 (41)	4.8 (5.2)			
GFP	27	5.4	95% ^[S2,3]	1 s	4-7 s
RPA2	29 (56)	5.4 (5.5)			
Sirt6	35 (62)	6.0 (5.7)			
YFP-MS2	42	5.6			
RelA	60 (87)	5.3 (5.4)	95% ^[S3]	1 s	7 s
RPA1	68 (95)	6.5 (5.8)	95% ^[S4]	8 s	30 s
Ku70	70 (97)	6.1 (5.8)	90% ^[S5,6]	6-7 s	20 s
Mre11	81 (108)	5.4 (5.4)	95% ^[S7]	3 s	30 s
Ku80	82 (109)	5.4 (5.4)	90% ^[S8]	1 s	3 s
NBS1	85 (112)	6.2 (5.9)	95% ^[S9,10]	0.5 s	3 s
dCas9	158 (185)	8.2 (7.8)			
MDC1	196 (223)	5.0 (5.1)	95% ^[S9,10]	2 s	7 s
BRCA1	208 (235)	5.2 (5.2)			
53BP1	214 (241)	4.5 (4.6)	100% ^[S11]	5 s	50 s

References

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