

Transfected plasmid DNA is incorporated into the nucleus via nuclear envelope reformation at telophase

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1 **Transfected plasmid DNA is incorporated into the nucleus via nuclear**
2 **envelope reformation at telophase**

3

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25 CLEM, iCLEM

26

27 **Abstract**

28 DNA transfection is an important technology in the life sciences, wherein nuclear entry
29 of DNA is necessary to achieve expression of the exogenous DNA. Non-viral vectors
30 and their transfection reagents are useful as safe tools for transfection. However, they
31 have no effects for transfection of non-proliferating cells, the reason for which remains
32 unknown. This study aimed to elucidate the mechanism by which transfected DNA
33 enters the nucleus for gene expression. To monitor the intracellular behavior of
34 transfected DNA, we introduced a plasmid bearing *lacO* repeats and RFP-coding
35 sequences into cells expressing GFP-LacI and observed plasmid behavior and RFP
36 expression. RFP expression appeared only after mitosis. Electron microscopy showed
37 that plasmids were wrapped with nuclear envelope (NE)-like membranes or associated
38 with chromosomes at telophase. Depletion of BAF, which is involved in NE
39 reformation, delayed plasmid RFP expression. These results suggest that transfected
40 DNA is incorporated into the nucleus via NE reformation at telophase.

41

42

43 **Introduction**

44 Transfection to introduce DNA plasmids into cells is an essential technology in the life
45 sciences and medical fields, for example, in gene therapy. Non-viral vectors and
46 transfection reagents, including cationic lipids or non-liposomal lipids, have been
47 developed as safe tools for introducing exogenous DNA into cells. DNA transfected
48 with these reagents is incorporated into the endosome via endocytosis and exposed to
49 the cytosol upon endosome rupture¹⁻⁵, where it is detected by DNA sensor molecules<sup>6-
50 10</sup>. However, it remains unclear how the transfected DNA in the cytosol enters the
51 nucleus. It is empirically known that DNA transfection, which leads to gene expression
52 from foreign DNA, is effective in proliferating cells but not in non-proliferating cells.
53 Therefore, it has been speculated that mitotic events, such as the nuclear envelope (NE)
54 breakdown, are important for nuclear entry of the transfected DNA. However, this has
55 not been experimentally proven yet.

56 Several attempts have been made to visualize the behavior of transfected DNA
57 and the cellular responses to it^{8,11,12}. These include methods using fluorescent dye-
58 conjugated DNA¹¹, the *lacO*/GFP-LacI visualizing system¹¹, fluorescent nanodiamond
59 particles coated with DNA¹², and beads conjugated with DNA (DNA-beads)⁸. In the
60 DNA-bead method, it has been shown that DNA-beads are internalized into cells via
61 endocytosis^{8,13}, entrapped in endosomes, and exposed to the cytosol when the endosome
62 ruptures⁸. Endosome rupture stimulates autophagosome formation in the region of the
63 transfected DNA in a p62-dependent manner¹⁴, leading to its degradation. However,
64 DNA on the beads simultaneously induces NE formation around the DNA-bead in a
65 barrier-to-autointegration factor (BAF)-dependent manner⁸. Once the NE forms around
66 the DNA beads, autophagy is suppressed⁸.

67 BAF is a DNA binding protein¹⁵⁻¹⁸ involved in NE assembly¹⁹⁻²¹. It binds to
68 foreign DNA, such as transfected DNA and viral DNA, in the cytosol^{8,15,22-23}, where it
69 assembles the NE-like membrane containing LEM domain proteins, such as emerin,
70 around the transfected DNA⁸. This BAF-dependent NE assembly is thought to occur
71 competitively with the assembly of the autophagy membrane^{8,24}. However, the
72 biological significance of NE assembly on transfected DNA present in the cytosol is
73 unclear.

74 The NE is a double-membrane structure consisting of the inner and outer nuclear
75 membranes. The outer membrane is connected to the ER and contains proteins shared
76 with the ER, whereas the inner membrane contains transmembrane proteins specific to
77 the NE. Among the inner nuclear membrane proteins, LEM domain NE proteins,
78 including lamina-associated polypeptide 2 (Lap2), emerin, MAN1, and Lem2, are
79 highly conserved among metazoans^{25,26}. LEM domain proteins bind BAF with the LEM
80 domain^{19,27-29}. Beneath the inner nuclear membrane, there is a nuclear lamina consisting
81 of lamins^{30,31}. There are fenestrated structures, called nuclear pores, spanning the
82 double-membrane structure of the NE^{32,33}. The nuclear pore complex (NPC) is formed
83 at the pores. Nucleocytoplasmic transport between the nucleus and cytoplasm is carried
84 out through the NPC.

85 The NE is disassembled at the beginning of mitosis (prometaphase) and reformed
86 around chromosomes at the end of mitosis (telophase) in higher eukaryotic cells,
87 including human cells. During NE reformation at telophase, emerin and other LEM
88 domain proteins assemble at the “core region” of the chromosome mass in a BAF-
89 dependent manner, whereas lamin B receptor (LBR) and NPC components assemble in
90 the “non-core region” in a BAF-independent manner^{19,21,34-36}. The ESCRT-III protein

91 complex functions to seal gaps in the nuclear membranes during NE reformation at
92 telophase³⁷.

93 This study aimed to understand when and how transfected exogenous DNA enters
94 the nucleus. For this purpose, we transfected cells with DNA plasmids designed to
95 visualize the plasmids and monitor their expression. Here, we show that transfected
96 DNA is incorporated into the nucleus when the NE is being reformed during telophase.
97 Our findings will contribute to understanding how foreign DNA behaves within cells,
98 providing useful information for generating improved transfection reagents and
99 techniques.

100

101 **Results**

102 **Protein expression from exogenous DNA occurs only after mitosis**

103 To visualize transfected DNA and its expression, we generated the following
104 experimental system. We constructed a non-viral DNA plasmid (pLacO-pEF1 α -mRFP)
105 carrying a *lacO* repeat sequence (256 repeats, about 10 kbp) and a sequence expressing
106 monomeric red fluorescent protein (hereafter RFP) driven by the EF1 α promoter (Fig.
107 1a). We introduced it into human HeLa cells stably expressing EGFP-LacI (hereafter
108 GFP-LacI) using a non-liposomal lipid transfection reagent. Thus, when the DNA
109 plasmid enters the cytosol after endosome rupture, it is visualized by binding to GFP-
110 LacI, forming fluorescent puncta that are easily detected by fluorescence microscopy
111 (Fig. 1a; arrows, Fig. 1b). Plasmid that has not entered the cytosol cannot bind GFP-
112 LacI, and therefore will not be fluorescent (arrowheads, Fig. 1b).

113 We first assayed efficacy of the GFP-LacI signal to detect transfected
114 exogenous DNA in the cytosol. After removal of transfection reagents, cells were fixed

115 and subjected to indirect immunofluorescence staining using anti-BAF. We examined
116 the overlap between the GFP-LacI signal and BAF, because it is known to bind
117 exogenous DNA immediately after it is released into the cytosol by endosome
118 rupture^{8,14,24}. We found that the punctate cytoplasmic GFP-LacI signal, overlapped with
119 that of BAF (Fig. 1b), indicating that our system enabled the visualization of cytosolic
120 plasmid DNA. We also used the DNA-specific fluorescent dye, 4',6-diamidino-2-
121 phenylindole (DAPI), to confirm that the puncta contained DNA. Some of the DAPI
122 puncta did not colocalize with the GFP-LacI signal (arrowheads, upper panels, Fig. 1b),
123 suggesting that some plasmids were retained in endosomes. The GFP-LacI-positive
124 regions of DNA puncta colocalized with the LEM domain NE protein emerlin (lower
125 panels, Fig. 1b), suggesting that they were surrounded by an NE-like membrane
126 immediately after release into the cytosol. This result is consistent with the report that
127 DNA-bead complexes become wrapped with NE-like membranes within 5-10 min after
128 endosome rupture⁸.

129 We then measured the timing of gene expression from transfected DNA by the
130 appearance of RFP fluorescence (Fig. 1c, d). Time-lapse analysis was carried out
131 immediately after transfection reagents were replaced with culture medium; images
132 were taken every 10 min for 18 h using DeltaVision or LSM880 microscopes. We
133 found that RFP signals appeared only in cells that had proceeded through mitosis (total
134 number of cells tested, n=594) (Fig. 1c, d; Supplementary movies 1, 2), suggesting that
135 gene expression from the transfected plasmid does not occur during interphase before
136 mitosis. RFP expression was detected 2-5 h after the onset of mitotic chromosome
137 segregation (median 150 min, n=594) (Fig. 1d). Since the G1 phase lasts for 6-8 h in the
138 HeLa cell line³⁸, our data suggest that gene expression occurs during the G1 phase in

139 most cells (Fig. 1c, d). In these cells, the RFP signal appeared to be comparable in the
140 two daughter cells. However, in a small fraction of the cells, RFP fluorescence appeared
141 later, after 500 min (6 h 20 min). In these cells, the RFP signal often was confined to
142 only one of the daughter cells. The reason for this bias remains unknown.

143 We also investigated gene expression from transfected DNA in U2OS cells, a
144 human osteosarcoma-derived cell line. We found that gene expression did not occur in
145 pre-mitotic cells, similar to HeLa cells. The timing of gene expression in U2OS cells
146 was 2-5 h (median 150 min) after the onset of chromosome segregation (n = 107)
147 (Supplementary Fig. 1). This median value was the same as that obtained for HeLa
148 cells. These results indicate that gene expression from exogenous DNA transfected
149 using a non-viral transfection reagent occurs only after mitosis in these cell lines. Our
150 findings provide clear evidence for the first time that non-viral transfection is effective
151 only for proliferating cells.

152

153 **Dynamic behavior of exogenous DNA during mitosis**

154 Protein expression from plasmid DNA did not occur in some cells, even in the presence
155 of GFP-LacI-positive clusters. To understand the difference in the behavior of plasmid
156 DNA between cells with and without gene expression, we characterized the dynamic
157 behavior of GFP-LacI clusters during mitosis. Time-lapse analysis of GFP-LacI in
158 living mitotic cells revealed that in most, GFP-LacI clusters were disjointed at
159 prometaphase with dispersion into smaller puncta during mitosis (Fig. 2a;
160 Supplementary movies 1, 3). In contrast, in some cells, GFP-LacI clusters remained
161 intact during mitosis. Thus, we investigated whether this difference in mitotic behavior
162 was related to gene expression. Most cells with GFP-LacI puncta that dispersed during

163 mitosis expressed RFP during the subsequent interphase (62 out of 68 cells observed)
164 (Supplementary Fig. 2a), whereas the cells without dispersion did not (6 out of 68 cells)
165 (Supplementary Fig. 2b), suggesting that dispersion of plasmids may be important for
166 gene expression in the subsequent interphase.

167 We next investigated whether the presence or absence of NE was involved in this
168 GFP-LacI dispersion. Indirect immunofluorescence staining showed that GFP-LacI
169 puncta colocalized with emerin in interphase cells (lower panels, Fig. 1b). In
170 prometaphase cells, in which NE breakdown is ongoing, some of the GFP-LacI puncta
171 colocalized with emerin (arrowheads, upper panels in Fig. 2b), but others did not (lower
172 panels in Fig. 2b). Notably, small puncta (arrows, Fig. 2b') were found near the larger
173 puncta with no emerin signals (lower panels in Fig. 2b; Supplementary Fig. 2c),
174 suggesting that dispersion occurred upon NE breakdown. In metaphase cells, GFP-LacI
175 puncta did not colocalize with emerin (Fig. 2c). Many small puncta of GFP-LacI of the
176 same size were often found in metaphase cells (white puncta, Fig. 2d; Supplementary
177 movie 4). These results suggest that the dispersion of GFP-LacI puncta during mitosis is
178 caused by the breakdown of the NE-like membrane surrounding them. Fig. 2e
179 summarizes the behavior of the GFP-LacI puncta during mitosis.

180

181 **Plasmid DNA is incorporated into the nucleus via NE reformation at telophase**

182 As gene expression occurs only after mitosis (Fig. 1; Supplementary Fig. 1), we
183 hypothesized that plasmid DNA was incorporated into the nucleus via NE reformation
184 at the end of mitosis. In fact, small puncta of GFP-LacI, probably containing a single
185 copy or a few copies of plasmid DNA, were frequently observed near the telophase
186 chromosomes (arrows, Fig. 3a). To visualize the structural details of these GFP-LacI

187 puncta, we performed correlative light and electron microscopy (CLEM) on telophase
188 cells undergoing NE reformation (Figs. 3b-f; Supplementary Fig. 3). We selected
189 telophase cells approximately 10 min after the onset of chromosome segregation for
190 CLEM analysis, in which most regions of the chromosomal mass were expected to be
191 surrounded by NE, although some regions may not yet be²¹. At the “core region” of the
192 chromosome mass, the NE containing LEM domain proteins is assembled in a BAF-
193 dependent manner, whereas in the “non-core region”, the NE containing LBR and NPC
194 is assembled in a BAF-independent manner (right drawing in Fig. 3a)^{19,21,36}.

195 CLEM showed that puncta near the chromosomes at telophase were surrounded
196 by a double-membrane structure similar to the NE or endoplasmic reticulum (ER)
197 (labeled “dot 1” in Figs. 3b, c). This NE-like membrane was fused with the NE on the
198 chromosome (Fig. 3c), suggesting that the transfected DNA is incorporated into the
199 nucleus via membrane fusion. Multiple additional structures were found: dot 2 in Fig.
200 3d was inside the nucleus (Fig. 3e), while dot 3 was outside and sandwiched between
201 two sets of ER or NE precursor membranes (Fig. 3e). Dot 4 in Fig. 3d was attached to
202 the chromosomes (“Type 4” in Fig. 3f; Supplementary Fig. 3l-o). Relatively larger
203 puncta surrounded by ER or NE precursor membranes were also found in the cytoplasm
204 (“Type 1” in Fig. 3f; Supplementary Fig. 3a-d). We classified these puncta into five
205 types on the basis of their morphologies obtained by CLEM, as indicated in Fig. 3f (also
206 Supplementary Fig. 3): Type 1 (cytoplasmic circle) is a relatively large circular
207 structure covered with a double membrane, which is often found around the GFP-LacI
208 puncta present in the cytoplasm. Type 2 (sandwich) is an electron-dense structure
209 squeezed between two double membranes, suggestive of ER or NE precursor
210 membranes (Fig. 3e; Supplementary Fig. 3e-g). Type 3 (membrane fusion) is a structure

211 in which a relatively large circular structure covered with a double membrane is fused
212 to the NE (Fig. 3c; Supplementary Fig. 3h-k). Type 4 (attached to chromosome) is a
213 structure at the edge of the chromosomal region where the NE is not present
214 (Supplementary Fig. 3l-o). Type 5 (inside the nucleus) is a structure inside the nucleus
215 (between chromosomes) (Supplementary Fig. 3p-s). Besides these types, we frequently
216 observed large GFP-LacI puncta in the cytoplasm. These puncta were found in
217 degradative organelles, including lysosomes and autophagosomes.

218 To confirm the CLEM results, immuno-CLEM analysis was performed using an
219 anti-GFP antibody conjugated with Alexa and nanogold (Fig. 4). GFP-LacI puncta co-
220 stained with Alexa, locating near the telophase chromosome mass, were examined (Fig.
221 4a, b; arrows indicate dots 1-5). Dots 1 and 2 in Fig. 4 were located inside a relatively
222 large circle surrounded by a double membrane structure similar to NE or ER (Fig. 4c),
223 so were classified as type 1. Dot 3 was located on the edge or inside of the
224 chromosomal region (Fig. 4d) and classified as type 4 or 5 (compare with a typical
225 example of Types 4 and 5 in Supplementary Fig. 4). Dots 4 and 5 were located inside
226 the nucleus (Fig. 4e) and were classified as type 5 (compare with a typical example of
227 “Type 5” in Supplementary Fig. 4). These results suggest that at least part of the
228 transfected DNA is attached to the chromosome or located in the nucleus. Therefore,
229 our results suggest that plasmid DNA attaches telophase chromosomes and is
230 subsequently incorporated into the nucleus via NE reformation at the end of mitosis.
231 Fig. 4f shows the typical localization of GFP-LacI puncta at telophase.

232

233 **BAF-dependent NE reformation at telophase facilitates nuclear entry**

234 To understand the molecular mechanism underlying nuclear entry of exogenous DNA in
235 telophase, we depleted NE proteins and examined the effect on gene expression from
236 the transfected DNA. BAF, emerlin and Lem2 were selected as target proteins for
237 depletion because all are known to be involved in NE reformation during telophase^{21,39-}
238 ⁴¹. We first employed the auxin-inducible degron (AID)-induced degradation system to
239 specifically deplete a target protein at the time of interest; for example, only during
240 mitosis^{42,43}. For this purpose, we integrated a sequence of mClover3-mAID into each of
241 the three target genes (*BAFNI*, *EMD*, and *LEMD2*) using CRISPR/Cas9 editing⁴⁴. We
242 produced *LEMD2* mutant cell lines (HeLa/mClover3-mAID-Lem2) expressing an
243 mClover3-mAID-Lem2 protein as the sole Lem2 protein (Supplementary Fig. 5), but
244 were not successful for BAF and emerlin. Western blotting (WB) of lysates of cells
245 treated with indole-3-acetic acid (IAA) for 60 min showed that Lem2 was almost lost
246 (Supplementary Fig. 5b), consistent with the time-lapse microscopy data
247 (Supplementary Fig. 5c; Supplementary Movie 5).

248 We examined the effect of Lem2 depletion on exogenous gene expression from
249 the transfected plasmid using time-lapse microscopy (Fig. 5a). The fluorescence of the
250 mClover3-mAID-Lem2 protein was almost completely lost in telophase (“IAA, 0:10” in
251 Fig. 5a) by the addition of IAA in metaphase (“IAA, -1:10” in Fig. 5a), suggesting that
252 depletion of Lem2 worked as expected. Time-lapse analysis of individual cells showed
253 that the timing of gene expression in Lem2-depleted cells was unchanged from that of
254 control cells (Fig. 5b), suggesting that Lem2 does not influence the expression of
255 exogenous DNA at telophase.

256 Next, we used small interfering RNA (siRNA) treatment to deplete the target
257 proteins. HeLa cells were treated with siRNA against *BAFNI* (iBAF, Fig. 5c-e), with

258 siRNA against luciferase used as a control (iLuc, Fig. 5c-e). WB showed that the
259 amount of BAF was reduced to 8% under these conditions (Fig. 5d). Then, cells
260 depleted of BAF were transfected with the plasmid and gene expression was assessed in
261 individual cells by fluorescence microscopy (Fig. 5c). The timing of gene expression in
262 cells treated with *BAFNI* siRNA (median 180 min, n=136) was delayed relative to that
263 of the control cells (median 140 min, n=129) (Fig. 5e). To confirm that BAF was truly
264 depleted in the cells assayed by microscopy, a similar experiment was carried out using
265 cells expressing GFP-BAF, which produced similar results (Supplementary Fig. 6a, b).
266 Because BAF is known to be required for NE reformation at telophase²¹, these data
267 suggest that NE reformation during telophase is an important step for the entry of
268 exogenous DNA into the nucleus. We also tested the effect of emerin siRNA depletion,
269 but it has no effect on the timing of gene expression (Supplementary Fig. 6c-e).

270

271 **NE rupture can cause pre-mitotic gene expression from exogenous DNA**

272 Our data led us to hypothesize that intact NE in interphase cells blocks the entry of
273 foreign DNA into the nucleus. To test this hypothesis, we physically ruptured the NE,
274 then examined whether gene expression occurred from exogenous DNA without
275 telophase NE reformation. To do this, cells were transfected with plasmids.
276 Immediately after removal of the transfection reagent, the transfected cells were
277 microinjected with FITC-dextran to rupture the NE. FITC-dextran was used as a marker
278 for the NE rupture (Fig. 6a). Time-lapse microscopy showed that some of the NE-
279 ruptured cells expressed RFP from plasmid DNA, even in interphase cells that had
280 never undergone mitosis (Fig. 6b). Similarly, premitotic gene expression from
281 transfected DNA was observed in cells depleted of NE proteins. When emerin was

282 depleted with siRNA treatment, premitotic gene expression from the transfected DNA
283 was observed (5 of 131 cells tested; Fig. 6c, d). No premitotic gene expression was
284 observed in the control siRNA treatment (129 cells tested; Supplementary Fig. 6e).
285 These results demonstrate that transient NE rupture caused by physical punctation or
286 depletion of the NE proteins allowed nuclear entry of the transfected DNA, resulting in
287 gene expression from foreign DNA, even in premitotic cells.

288

289 **Discussion**

290 Multiple cellular barriers to the delivery of DNA plasmids to the nucleus are known to
291 exist, including intracellular uptake, endosome acidification, autophagy, immune
292 sensing pathways, and nuclear entry^{45,46}. This study focused on how transfected DNA
293 enters the nucleus. We found that gene expression occurs only after telophase, with
294 plasmids associated with telophase chromosomes enwrapped by the NE. Therefore, we
295 concluded that NE reformation at telophase is important for the nuclear entry of the
296 transfected DNA. This conclusion indicates that NE acts as a physical barrier to the
297 delivery of foreign DNA into the nucleus, and that NPC-mediated nuclear transport of
298 foreign DNA does not occur in interphase cells. If premitotic gene expression happened
299 to occur in interphase cells, transient NE rupture might have been induced in these cells.
300 This hypothesis is consistent with our data showing that premitotic gene expression
301 occurs in cells in which the NE has been ruptured by a microinjection needle (Fig. 6a,
302 b), and in cells in which emerin and BAF were depleted (Fig. 6c, d). Similar transient
303 NE ruptures have been reported in some diseases, including laminopathy and
304 cancers^{47,48}. In addition, BAF has been reported to play a role to repair NE rupture⁴⁹,
305 consistent with our finding.

306 It has been reported that most plasmids form clusters in the cytosol after
307 endosomal escape, regardless of transfection method; moreover, the clusters are
308 relatively immobile during interphase¹¹. The clusters remaining at metaphase are
309 asymmetrically partitioned with a bias to the younger centrosomes in dividing cells¹¹.
310 We also found similar plasmid clusters in the cytosol. However, most of them were
311 dispersed at the beginning of mitosis to smaller puncta. The small puncta were scattered
312 throughout the cell and distributed to daughter cells (Fig. 2a; Supplementary movies 1
313 and 3), whereas the large puncta were retained in one cell as reported previously¹¹.
314 Because gene expression from foreign DNA occurred only in cells with small puncta,
315 the small puncta that are incorporated into the reassembling nucleus at telophase are
316 associated with efficient gene expression from foreign DNA in the G1 phase. The
317 biological significance of plasmid clusters in mitosis remains unclear.

318 Transfected DNA in the cytosol activates DNA sensor signaling pathways^{6,7,50}.
319 Binding of DNA sensors to the cytosolic DNA is hypothesized to be a defense response
320 against infection. Many DNA sensor proteins have been identified, including cyclic
321 GMP-AMP synthase (cGAS)⁵¹ and their common signaling adaptor stimulator of
322 interferon (IFN) gene (STING)⁵², DNA-dependent activator of interferon regulatory
323 factors/Z-DNA binding protein 1 (DAI/ZBP1)⁵³, interferon activated gene 204 (p204)⁵⁴,
324 DEAD box helicase 60 (DDX60)⁵⁵, and BAF^{8,24}. DAI/ZBP1 and p204 have been
325 reported to bind cytosolic plasmid DNA within 15 min, whereas DDX60 does not⁵⁶.
326 BAF binds the cytosolic DNA within seconds⁸ and plays a role in assembling the NE-
327 like membrane around the plasmid DNA within approximately 10 min⁸. This BAF-
328 dependent NE assembly protects foreign DNA from autophagy^{8,24}. Although BAF is
329 expected to function as a defense against viral infection^{23,57,58}, it unexpectedly assists in

330 the integration of the human immunodeficiency virus (HIV) genome into the host
331 genome^{15,16}. In addition, a recent report showed that BAF protects against the cGAS-
332 STING response to prevent innate immune activation^{59,60}. Collectively, these data
333 suggest that BAF can suppress the innate immune response and act to safely carry the
334 cytosolic DNA to the next cell cycle stages by wrapping it in the NE-like membrane.

335 In summary, our data suggest that NE acts as a barrier for foreign DNA to enter
336 the nucleus, and that foreign DNA is incorporated into the nucleus when the NE is
337 being reassembled in telophase. Our findings provide important insights not only for
338 understanding cellular responses during DNA transfection, but also for the development
339 of transfection reagents and methods.

340

341 **Materials and Methods**

342 **Plasmids.** We constructed the plasmid carrying a *lacO* repeat sequence and a sequence
343 expressing RFP driven by the EF1 α promoter (pLacO-pEF1 α -mRFP) as follows: First,
344 to generate the mRFP-C1 vector, the DNA fragment encoding mRFP was amplified by
345 PCR from the mRFP-pRSETB vector using the primer set mRFP-FWD3-Nhe (5'-
346 GAGCTAGCATGGCCTCCTCCGAGGACGTCA-3') and mRFP-REV2-Bgl (5'-
347 TAAGATCTGGCGCCGGTGGAGTGGCGGCC-3'), and the PCR product was
348 inserted into the pEGFP-C1 vector (Clontech, Takara Bio Inc., Kusatsu, Japan) at the
349 *NheI* and *BglIII* sites. The resulting plasmid is called the mRFP-C1 vector. Next, to
350 generate the pEF1 α -mRFP plasmid, the DNA fragment encoding the EF1 α -promoter
351 region was amplified by PCR from the pBOS-H2B-GFP vector (Stratagene, La Jolla,
352 CA) using the primers EF1 α -prom.FWD-Ase (5'-
353 GGGATTAATCGTGAGGCTCCGGTGCCCGTCAGTGG-3') and EF1 α -prom.REV-

354 Nhe (5'-GGGCTAGCCTCACGACACCTGAAATGGAAGAAAAAACTTTG-3'),
355 and the PCR product was inserted into the mRFP-C1 vector at the *AseI* and *NheI* sites.
356 The resulting plasmid is called the pEF1 α -mRFP plasmid. Then, the pEF1 α -mRFP
357 plasmid was digested with *SalI*, then circularized with a ligation kit (Ligation High
358 Ver.2, Toyobo Co., Osaka, Japan) after filling the gaps with *E. coli* DNA polII Klenow
359 fragment (Takara Bio Inc., Kusatsu, Japan) to generate the pEF1 α -mRFP-C1- Δ *SalI*
360 vector. The pEF1 α -mRFP-C1- Δ *SalI* vector was inserted with the sequence of
361 TAGTCGAC to generate a new *SalI* site in front of the EF1 α promoter region to
362 generate the *SalI*-pEF1 α -mRFP-C1 vector. Finally, to generate pLacO-pEF1 α -mRFP,
363 the DNA fragment containing a *LacO*-repeat sequence was obtained by digestion of the
364 pSV-dhfr DNA plasmid, then inserted into the *SalI*-pEF1 α -mRFP-C1 vector at its *SalI*
365 site using Ligation High Ver.2 (Toyobo Co., Osaka, Japan).

366 To construct the plasmid expressing EGFP-fused LacI (GFP-LacI), EGFP was
367 amplified by PCR from the pEGFP-C1 vector (Clontech, Takara Bio Inc., Kusatsu,
368 Japan) using the primers 5'-
369 TGTGACCGGGCGCCTACTATGGTGAGCAAGGGCGAGGAGCT-3' and 5'-
370 CGAATTCGCTAGCTCTAGACTTGTACAGCTCGTCCATGC-3', and the PCR
371 product was inserted into the PB-EF1 α -MCS-IRES-Neo vector (PB533A-2, System
372 Biosciences, Mountain View, CA, USA) using *XbaI* sites to generate the PB-EF1 α -
373 EGFP-MCS-IRES-Neo vector. Next, DNA encoding LacI was amplified from the
374 MK9-38/pYC19 vector using the primers (5'-
375 ATTCGAATTTAAATCGGATATGGTGAAACCAGTAACGTTATA-3' and 5'-
376 CTCAGCGGCCGCGGATCTTACAGCTGCATTAATGAATCGGCC-3') and the

377 PCR product was inserted into the PB-EF1 α -EGFP-MCS-IRES-Neo vector using a
378 *Bam*HI site to generate the plasmid expressing GFP-LacI.

379

380 **Cell strains and culture.** HeLa and U2OS cells were obtained from the Riken Cell
381 Bank (RCB007, Tsukuba, Japan) and the Health Protection Agency (London, UK),
382 respectively. To generate HeLa cells stably expressing GFP-LacI (HeLa/GFP-LacI),
383 they were transfected with plasmids encoding GFP-LacI using Effectene (Qiagen,
384 Hilden, Germany), according to the manufacturer's instructions. Cells were selected
385 with 800 μ g/ml Geneticin (11811-031, Life Technologies, Thermo Fisher Scientific,
386 Tokyo, Japan). Several clones were isolated from the survivors and their expression
387 levels were evaluated using WB. The selected clones were maintained in DMEM
388 containing 10% fetal bovine serum (FBS) and 200 μ g/ml Geneticin (11811-031, Life
389 Technologies, Thermo Fisher Scientific, Tokyo, Japan). We established *LEMD2* mutant
390 cell lines (HeLa/mClover3-mAID-Lem2) expressing an mClover3-mAID-Lem2 protein
391 as the sole Lem2 protein as describe below (see section "Depletion of NE proteins" in
392 Materials and Methods).

393

394 **Antibodies.** Primary mouse monoclonal antibodies included anti-GFP for indirect
395 immunofluorescence staining and WB (Cat No. 632381, Lot No. 1004037, Clontech,
396 USA), anti- α -tubulin (clone DM1A, Cat No. T9026, Lot No. 017H4838, Sigma), and
397 anti-BAF for WB (BANF1, Cat No. H00008815-M01, Lot No. 6132-S3, Abnova,
398 Taipei, Taiwan). Primary rabbit polyclonal antibodies included anti-GFP antibody for
399 iCLEM (Cat No. 600-401-215, Lot No. 24437, Rockland, Limerick, PA, USA), anti-
400 emerin (ED1)⁶¹, anti-BAF (PU38143)⁶², anti-Lem2 (LEMD2, Cat No. HPA017340, Lot

401 No. B96759, Atlas Antibodies, Bromma, Sweden), and anti- β -actin (Cat No. ab8227,
402 Lot No. GR14272-3, Abcam). Alexa594-conjugated secondary antibodies against
403 mouse IgG (Cat No. A11037, Thermo Fisher Scientific) and rabbit IgG (Cat No.
404 A11032, Thermo Fisher Scientific) were used to detect the primary antibodies.

405

406 **Transfection and live-cell imaging.** HeLa/GFP-LacI cells were seeded on 35-mm
407 glass-bottom culture dishes (MatTek, MA, USA) at a concentration of 2×10^5 per dish
408 one day before imaging. For transfection, the cells were incubated for 4 h with
409 Effectene transfection reagent (Qiagen) containing 0.5 $\mu\text{g}/\text{dish}$ pLacO-pEF1 α -mRFP.
410 During this incubation, Hoechst 33342 was added to a final concentration of 100 ng/ml
411 and incubated for 15 min to stain DNA. After transfection reagents were replaced with
412 culture medium, cells were subjected to time-lapse imaging. Images for each
413 wavelength were obtained every 10 min ($1 \mu\text{m} \times 7$ z-stacks for a single time point
414 unless otherwise specified) through an oil-immersion objective lens UApo/340 (40x,
415 NA = 1.35, Olympus, Tokyo, Japan) using the DeltaVision fluorescence microscope
416 system (GE Healthcare Japan, Tokyo, Japan) in a temperature-controlled (37°C) room;
417 this microscope system is equipped with a high-sensitive sCMOS camera (pico.edge
418 4.2, PCP AG, Kelheim, Germany). Images were also obtained every 10 min ($2 \mu\text{m} \times 5$
419 z-stacks for a single time point) through a water-immersion objective lens C-Apo40
420 (40x, NA=1.2, Carl Zeiss, Jena, Germany) on a confocal fluorescence microscope
421 system LSM880 (Carl Zeiss), maintained at 37°C using a stage warmer. Scanning
422 conditions were as follows: pinhole size 2.5 A.U.; laser power 0.5-1%; zoom 1; scan
423 speed 6; averaging 2; tiling 5×5 .

424

425 **Immunofluorescence staining.** HeLa/GFP-LacI cells were transfected with plasmid
426 using Effectene as described above. Immediately after removal of the transfection
427 reagent, cells were fixed with a fixative (3.7% formaldehyde and 0.2% glutaraldehyde
428 in PBS) and incubated with anti-BAF or anti-emerin antibodies as previously
429 described^{19,62}. Anti-rabbit IgG-conjugated Alexa-594 was used as the secondary
430 antibody. Fluorescence images were obtained using an Olympus oil-immersion
431 objective lens PLAPON60xOSC (NA = 1.40) on the DeltaVision microscope system.
432 Z-stacks of images (typically 20-30 focal planes at 0.5 μm intervals) were obtained and
433 deconvoluted using software provided with the microscope system, SoftWorx.
434

435 **Super-resolution microscopy.** HeLa/GFP-LacI cells were transfected with plasmid
436 using Effectene as described above. After removal of the transfection reagent, cells
437 were fixed with a fixative (3.7% formaldehyde and 0.2% glutaraldehyde in PBS). Then,
438 cells were observed by LSM880 Airyscan system (Carl Zeiss) equipped with Plan-
439 Apochromat 63x/1.40 M27 objective lens using immersion oil (#518F, Carl Zeiss) at
440 room temperature (RT, approximately 26°C). Pinhole size was set to 384 μm . Hoechst
441 and GFP were excited with 405 and 488 nm laser, respectively. The fluorescent signals
442 for Hoechst and GFP were collected using the following beam splitters and bandpass
443 filters: MBS-405 and BP 420-480 for Hoechst and MBS488/561 and BP 495-550 for
444 GFP. Images were acquired at 988 pixels each with 0.037 $\mu\text{m}/\text{pixel}$. A z-stack image set
445 was acquired for 133 focal planes with 0.159 μm intervals. After processing for super-
446 resolution image, brightness and contrast were changed for better visibility in Zen 2.3
447 SP1 software without changing gamma.
448

449 **Correlative light and electron microscopy (CLEM).** HeLa/GFP-LacI cells were
450 transfected with plasmid using Effectene as described above. After removal of the
451 transfection reagent, cells were fixed with 2.5% glutaraldehyde for 1 h. CLEM was
452 performed with some modifications of our previously published method²¹. Briefly, after
453 washing with PBS, the telophase cells 8-10 min after metaphase to anaphase transition
454 were selected by morphology, then subjected to fluorescence microscopy (FM) to
455 obtain 3D images (typically 40-60 focal planes at 0.2 μm intervals) using a low-
456 chromatic-aberration oil-immersion lens PLAPON60xOSC (NA = 1.40, Olympus) on a
457 DeltaVision microscope system. The images were deconvoluted to remove out-of-focus
458 images using software provided with the microscope system. Samples were post-fixed
459 with 1% OsO₄ (3002, Nisshin EM, Tokyo, Japan), stained with 2% uranyl acetate
460 (8473-1M, Merck, Darmstadt, Germany), and then embedded in Epon812 (T024,
461 TAAB, England). Cells observed by FM were identified in the block. Ultrathin sections
462 of 80 nm were prepared and further stained with 4% uranyl acetate, followed by a
463 commercial ready-to-use solution of lead citrate (18-0875-2, Sigma-Aldrich, St. Louis,
464 MO, USA). Electron microscopy (EM) images were acquired using a JEM-1400
465 electron microscope (80 kV; JEOL, Tokyo, Japan). To correlate the FM and EM
466 images, we selected the FM image corresponding to the EM image from the 3D FM
467 images by eye using criteria including nuclear shapes and areas of bead cross-sections.
468

469 **Immuno-CLEM (iCLEM).** HeLa/GFP-LacI cells were transfected for 4 h with 0.2
470 $\mu\text{g}/\text{dish}$ pLacO-pEF1 α -mRFP using Effectene as described above. After removal of
471 transfection reagents, cells were fixed with a mixture of 3.7% formaldehyde and 0.2%
472 glutaraldehyde for 15 min. After washing with PBS, cells were permeabilized with

473 0.01% Triton X-100 in PBS for 5 min at RT (26°C), treated twice with 0.1% sodium
474 borohydride in PBS for 15 min, and washed twice with 1% BSA in PBS for 1 h. Cells
475 were then incubated with anti-GFP antibody (1:500 dilution in 1% BSA in PBS, 600-
476 401-215, lot. No. 35459, Rockland). After extensive washing with PBS (10 min × 12
477 times), cells were incubated with a secondary Alexa594-nanogold-anti rabbit antibody
478 (1:200 dilution in 1% BSA in PBS, Nanoprobes, Yaphank, NY, USA) for 3 h at RT. A
479 z-stack of fluorescence images (typically 40–60 focal planes at 0.2-µm intervals) were
480 acquired for cells of interest using a low-chromatic-aberration PLAPON60xOSC (NA =
481 1.40) objective on a DeltaVision microscope system as described above. Then, cells
482 were fixed with 1% glutaraldehyde for 1 h and washed with PBS three times. After
483 washing twice with 100 mM Lysine-HCl/PB (pH 7.4) for 10 min, cells were subjected
484 to silver enhancement as follows: cells were washed with PBS, incubated with 50 mM
485 HEPES (pH 5.8) three times, and washed with distilled water. Cells were then treated
486 with silver-enhancing reagent at 25°C for 5 min and washed with distilled water three
487 times. Samples were post-fixed with 1% OsO₄ (3002, Nisshin EM), stained with 2%
488 uranyl acetate (8473-1M, Merck), and embedded in Epon812 (T024, TAAB) as
489 described previously²¹. Cells observed by fluorescence microscopy were identified in
490 the block. Ultrathin sections with a thickness of 80 nm were prepared and further
491 stained with 4% uranyl acetate (8473-1M, Merck), followed by a commercial ready-to-
492 use solution of lead citrate (18-0875-2, Sigma-Aldrich). EM images were acquired
493 using a JEM-1400 plus electron microscope (80 kV; JEOL, Tokyo, Japan). To correlate
494 the FM and EM images, we selected FM image corresponding to EM images from the
495 3D FM images by eye, using criteria including nuclear shape and area of bead cross-
496 section in each image.

497

498 **Depletion of NE proteins.** To deplete BAF, HeLa cells or HeLa cells stably expressing
499 GFP-BAF were treated with *BAFNI* siRNA (Qiagen) using RNAiMax (Thermo Fisher
500 Scientific) as described previously²¹. To deplete emerin, HeLa cells were treated with
501 *EMR* siRNA [AACCGTGCTCCTGGGGCTGGG, Qiagen]⁶³ using RNAiMax.
502 Luciferase siRNA (Qiagen) was used as the negative control.

503 The AID-induced protein degradation system was used to deplete Lem2 as
504 described previously⁴³. The plasmid pAID1.2-EF1a-NmClover3-mAID, was developed
505 from pAID1.2-EF1a-NGFP-mAID (Addgene plasmid #140607). The cDNA sequence
506 of Lem2 was amplified by PCR and cloned into the pAID1.2-EF1a-NmClover3-mAID
507 vector at the *EcoRV* site using the In-Fusion HD Cloning Kit (Takara Bio Inc.). The
508 pX330 plasmid targeting the human *LEMD2* gene was constructed as described
509 previously⁴⁴. The sequence 5'-GACTTACTCACCAGCTTGGATGG-3' at the intron-
510 exon boundary of the 3' end of exon 3 of the human *LEMD2* gene was selected as a
511 target for CRISPR/Cas9-mediated gene editing. These plasmids were cotransfected with
512 pAID EF1a linearized in pX330 (plasmid #140610, Addgene) into HeLa cells. After
513 selection in 8 µg/ml blasticidin S (029-18701, Wako, Osaka, Japan), several clones
514 were picked; their expression levels of endogenous and mClover3-mAID-tagged Lem2
515 were evaluated by WB using anti-Lem2 and anti-GFP antibodies, respectively. We used
516 an anti-GFP antibody to detect mClover3-mAID-Lem2 because the anti-GFP antibody
517 could cross-react with mClover3. AID-mediated protein degradation was induced by
518 adding final 0.5 mM indole-3-acetic acid (IAA).

519

520 **Western Blotting (WB).** WB was performed as previously described⁶². Briefly, cells
521 were collected and lysed in ice-cold isotonic buffer (20 mM Tris-HCl, pH 7.5, 2 mM
522 MgCl₂, 150 mM NaCl, 0.5% NP-40, Protease Inhibitor Cocktail) (final 1×10^7
523 cell/ml). Equal amounts of 2x SDS-PAGE sample buffer (without DTT) were added
524 to the lysates. After sonication, the lysates were boiled for 5 min. DTT (freshly
525 prepared from powder) was added to the samples at a final concentration of 20 mM.
526 Ten micrograms of total protein from the cell extract were loaded into each lane of an
527 SDS 10–20% gradient polyacrylamide gel or 15% polyacrylamide gel. After
528 electrophoresis, proteins were transferred to PVDF membranes using a semi-dry
529 blotting system (Atto Corp., Tokyo, Japan) and stained using the ECL western blotting
530 detection system (GE Healthcare).

531

532 **NE rupture.** HeLa/GFP-LacI cells were transfected using the transfection reagent
533 Effectene (Qiagen) with 0.3 µg/dish of pLacO-pEF1α-mRFP for 3 h. During this
534 incubation, Hoechst 33342 was added to a final concentration of 100 ng/ml and
535 incubated for 15 min to stain DNA. After removal of the transfection mixture, the
536 culture medium was replaced with fresh DMEM containing 10% FBS and then
537 subjected to fluorescence imaging. Cells in the glass-bottom culture dish were placed on
538 a stage of the DeltaVision fluorescence microscope in a temperature-controlled room
539 (37°C). Cells of interest, i.e., the cells appearing to be undergoing G1 phase, were
540 selected based on the morphology observed via bright-field microscopy; their NEs were
541 disrupted using the tip of a microinjection needle (Femtotips, Eppendorf, No.
542 5242952008 Eppendorf, Hamburg, Germany) attached to a microinjector (microinjector
543 5242 and micromanipulator 5171, Eppendorf). To confirm NE rupture, cells were

544 concomitantly microinjected with 0.5 $\mu\text{g}/\mu\text{l}$ FITC-dextran (MW, 10 kDa) as an injection
545 marker. Fluorescence images were obtained at 10-15 min intervals for 4 h.

546

547 **Data availability**

548 The authors declare that the data supporting the findings of this study are available
549 within the paper and its supplementary information files. The source data underlying
550 Fig. 1d, 5b, 5e, Supplementary Fig. 1, 6b, and 6e are provided as a source data file.

551 Further data are available from the corresponding authors upon request.

552

553 **References**

- 554 1. Conner, S. D., & Schmid, S. L. Regulated portals of entry into the cell. *Nature* 422,
555 37-44. doi: 10.1038/nature01451 (2003).
- 556 2. Hoekstra, D., Rejman, J., Wasungu, L., Shi., F, & Zuhorn, I. Gene delivery by
557 cationic lipids: in and out of an endosome. *Biochem. Soc. Trans.* **35**, 68-71. doi:
558 10.1042/BST0350068 (2007).
- 559 3. Zylberberg, C., Gaskil, K., Pasley, S., Matosevic, S. Engineering liposomal
560 nanoparticles for targeted gene therapy. *Gene Ther.* **24**, 441-452. doi:
561 10.1038/gt.2017.41 (2017).
- 562 4. Brock DJ, Kondow-McConaghy HM, Hager EC, Pellois JP. Endosomal Escape and
563 Cytosolic Penetration of Macromolecules Mediated by Synthetic Delivery Agents.
564 *Bioconjugate Chem.* **30**, 293-304. doi: 10.1021/acs.bioconjchem.8b00799 (2019).
- 565 5. Ita K. Polyplexes for gene and nucleic acid delivery: Progress and bottlenecks. *Eur.*
566 *J. Pharm. Sci.* **150**, 105358. doi: 10.1016/j.ejps.2020.105358 (2020).
- 567 6. Takaoka, A., & Taniguchi, T. Cytosolic DNA recognition for triggering innate
568 immune responses. *Adv. Drug Deliv. Rev.* **60**, 847-57.
569 doi:10.1016/j.addr.2007.12.002 (2008).
- 570 7. Yanai, H., Savitsky, D., Tamura, T., & Taniguchi, T. Regulation of the cytosolic
571 DNA-sensing system in innate immunity: a current view. *Curr. Opin. Immunol.* **21**,
572 17-22. doi: 10.1016/j.coi.2009.01.005 (2009).
- 573 8. Kobayashi, S. et al. BAF is a cytosolic DNA sensor that leads to exogenous DNA
574 avoiding autophagy. *Proc. Natl. Acad. Sci. USA* **112**, 7027-7032. doi:
575 10.1073/pnas.1501235112 (2015).

- 576 9. Chen, Q., Sun, L., & Chen, Z. J. Regulation and function of the cGAS-
577 STING pathway of cytosolic DNA sensing. *Nat. Immunol.* **17**, 1142-9. doi:
578 10.1038/ni.3558 (2016).
- 579 10. Hopfner, K. P., & Hornung, V. Molecular mechanisms and cellular functions
580 of cGAS-STING signalling. *Nat. Rev. Mol. Cell Biol.* **21**, 501-521. doi:
581 10.1038/s41580-020-0244-x (2020).
- 582 11. Wang, X., Le, N., Denoth-Lippuner, A., Barral, Y., & Kroschewski, R. Asymmetric
583 partitioning of transfected DNA during mammalian cell division. *Proc. Natl. Acad.*
584 *Sci. USA* **113**, 7177-82. doi: 10.1073/pnas.1606091113 (2016).
- 585 12. Petrakova, V., Benson, V., Buncek, M., Fiserova, A., Ledvina, M., Stursa, J.,
586 Cigler, P., & Nesladek, M. I Imaging of transfection and intracellular release of
587 intact, non-labeled DNA using fluorescent nanodiamonds. *Nanoscale* **8**, 12002-12.
588 doi: 10.1039/c6nr00610h.PMID: 27240633 (2016).
- 589 13. Kobayashi, S., Kojidani, T., Osakada, H., Yamamoto, A., Yoshimori, T., Hiraoka,
590 Y., & Haraguchi, T. Artificial induction of autophagy around polystyrene beads in
591 nonphagocytic cells. *Autophagy* **6**, 36-45 (2010).
- 592 14. Tsuchiya, M., Ogawa, H., Koujin, K., Kobayashi, S., Mori, C., Hiraoka, Y., &
593 Haraguchi, T. Depletion of autophagy receptor p62/SQSTM1 enhances the
594 efficiency of gene delivery in mammalian cells. *FEBS Lett.* **590**, 2671-80. doi:
595 10.1002/1873-3468.12262 (2016)
- 596 15. Lee, M. S., & Craigie, R. A previously unidentified host protein protects retroviral
597 DNA from autointegration. *Proc. Natl. Acad. Sci. USA* **95**, 1528-1533.
598 doi:10.1073/pnas.95.4.1528 (1998).

- 599 16. Chen, H. & Engelman, A. The barrier-to-autointegration protein is a host factor for
600 HIV type 1 integration. *Proc. Natl. Acad. Sci. USA* **95**, 15270-15274.
601 doi:10.1073/pnas.95.26.15270 (1998).
- 602 17. Umland, T. C., Wei, S.-Q., Craigie, R. & Davies, D. R. Structural basis of DNA
603 bridging by barrier-to-autointegration factor. *Biochemistry* **39**, 9130-9138.
604 doi:10.1021/bi000572w (2000).
- 605 18. Zheng, R., Ghirlando, R., Lee, M. S., Mizuuchi, K., Krause, M. & Craigie, R.
606 Barrier-to-autointegration factor (BAF) bridges DNA in a discrete, higher order
607 nucleoprotein complex. *Proc. Natl. Acad. Sci. USA* **97**, 8997-9002.
608 doi:10.1073/pnas.150240197 (2000).
- 609 19. Haraguchi, T., Koujin, T., Segura-Totten, M., Lee, K. K., Matsuoka, Y., Yoneda,
610 Y., Wilson, K. L., & Hiraoka, Y. BAF is required for emerin assembly into the
611 reforming nuclear envelope. *J. Cell Sci.* **114**, 4575-4585 (2001).
- 612 20. Gorjánác, M., Klerkx, E. P., Galy, V., Santarella, R., López-Iglesias, C., Askjaer,
613 P. & Mattaj, I. W. *Caenorhabditis elegans* BAF-1 and its kinase VRK-1 participate
614 directly in post-mitotic nuclear envelope assembly. *EMBO J.* **26**, 132-143.
615 doi:10.1038/sj.emboj.7601470 (2007).
- 616 21. Haraguchi, T., Kojidani, T., Koujin, T., Shimi, T., Osakada, H., Mori, C.,
617 Yamamoto, A., & Hiraoka, Y. Live cell imaging and electron microscopy revealed
618 dynamic processes of BAF-directing nuclear envelope assembly. *J. Cell Sci.* **121**,
619 2540-2554. doi: 10.1242/jcs.033597 (2008)
- 620 22. Wiebe, M. S., & Traktman, P. Poxviral B1 kinase overcomes barrier to
621 autointegration factor, a host defense against virus replication. *Cell Host Microbe* **1**,
622 187-197. doi:10.1016/j.chom.2007.03.007 (2007).

- 623 23. Ibrahim, N, Wicklund, A, Jamin, A, Wiebe, M. S. Barrier to autointegration factor
624 (BAF) inhibits vaccinia virus intermediate transcription in the absence of the viral
625 B1 kinase. *Virology* **444**, 363-73. doi: 10.1016/j.virol.2013.07.002 (2013).
- 626 24. Kobayashi, S., & Haraguchi, T. A novel pathway to detect and cope with
627 exogenous dsDNA. *Commun. Integr. Biol.* **8**, e1065361,
628 doi:10.1080/19420889.2015.1065361 (2015).
- 629 25. Wagner, N., & Krohne, G. LEM-domain proteins: new insights into lamin-
630 interacting proteins. *Int. Rev. Cytol.* **261**, 1–46 (2007).
- 631 26. Brachner, A., & Foisner, R. Evolvement of LEM proteins as chromatin tethers at
632 the nuclear periphery. *Biochem. Soc. Trans.* **39**, 1735-1741.
633 doi:10.1042/BST20110724 (2011).
- 634 27. Cai, M., Huang, Y., Ghirlando, R., Wilson, K. L., Craigie, R., & Clore, G.M.
635 Solution structure of the constant region of nuclear envelope protein LAP2 reveals
636 two LEM-domain structures: one binds BAF and the other binds DNA. *EMBO J.*
637 **20**, 4399-407. doi: 10.1093/emboj/20.16.4399 (2001).
- 638 28. Lee, K. K., Haraguchi, T., Lee, R. S., Koujin, T, Hiraoka, Y., & Wilson, K. L.
639 Distinct functional domains in emerin bind lamin A and DNA-bridging protein
640 BAF. *J. Cell Sci.* **114**, 4567-73 (2001).
- 641 29. Shumaker, D. K., Lee, K. K., Tanhehco, Y. C., Craigie, R., & Wilson, K. L. LAP2
642 binds to BAF.DNA complexes: requirement for the LEM domain and modulation
643 by variable regions. *EMBO J.* **20**, 1754-64. doi: 10.1093/emboj/20.7.1754 (2001).
- 644 30. Moir, R. D., & Goldman, R. D. Lamin dynamics. *Curr. Opin. Cell Biol.* **5**, 408-11.
645 doi: 10.1016/0955-0674(93)90004-a (1993).

- 646 31. Tenga, R., & Medalia, O. Structure and unique mechanical aspects
647 of nuclear lamin filaments. *Curr. Opin. Struct. Biol.* **64**, 152-159. doi:
648 10.1016/j.sbi.2020.06.017 (2020).
- 649 32. Hampoelz, B., Andres-Pons, A., Kastritis, P., & Beck, M. Structure and Assembly
650 of the Nuclear Pore Complex. *Annu. Rev. Biophys.* **48**, 515-536. doi:
651 10.1146/annurev-biophys-052118-115308 (2019).
- 652 33. Bilir, Ş., Kojidani, T., Mori, C., Osakada, H., Kobayashi, S., Koujin, T., Hiraoka,
653 Y., & Haraguchi, T. *Genes Cells* **24**, 338-353. doi: 10.1111/gtc.12677 (2019).
- 654 34. Shimi, T., Koujin, T., Segura-Totten, M., Wilson, K. L. Haraguchi, T. & Hiraoka,
655 Y. Dynamic interaction between BAF and emerin revealed by FRAP, FLIP and
656 FRET analyses in living HeLa cells. *J. Struct. Biol.* **147**, 31-41 (2004).
- 657 35. Dechat, T., Gajewski, A., Korbei, B., Gerlich, D., Daigle, N., Haraguchi, T.,
658 Furukawa, K., Ellenberg, J., & Foisner, R. LAP2alpha and BAF transiently localize
659 to telomeres and specific regions on chromatin during nuclear assembly. *J. Cell Sci.*
660 **117**, 6117-28. doi: 10.1242/jcs.01529. (2004).
- 661 36. Clever, M., Mimura, Y., Funakoshi, T., & Imamoto, N. Regulation and
662 coordination of nuclear envelope and nuclear pore complex assembly. *Nucleus* **4**,
663 105-14. doi: 10.4161/nucl.23796 (2013).
- 664 37. Olmos, Y., Hodgson, L., Mantell, J., Verkade, P., & Carlton, J.G. ESCRT-III
665 controls nuclear envelope reformation. *Nature* **522**, 236-9. doi:
666 10.1038/nature14503 (2015).
- 667 38. Haraguchi, T., Kaneda, T., & Hiraoka, Y. Dynamics of chromosomes and
668 microtubules visualized by multiple-wavelength fluorescence imaging in living

669 mammalian cells: effects of mitotic inhibitors on cell cycle progression. *Genes*
670 *Cells* **2**, 369-80. doi:10.1046/j.1365-2443.1997.1280326.x (1997).

671 39. Asencio, C., Davidson, I. F., Santarella-Mellwig, R., Ly-Hartig, T. B., Mall, M.,
672 Wallenfang, M. R., Mattaj, I. W., & Gorjánác, M. Coordination of kinase and
673 phosphatase activities by Lem4 enables nuclear envelope reassembly
674 during mitosis. *Cell* **150**, 22-35. doi: 10.1016/j.cell.2012.04.043 (2012).

675 40. von Appen, A., LaJoie, D., Johnson, I. E., Trnka, M. J., Pick, S. M., Burlingame, A.
676 L., Ullman, K. S., & Frost, A. LEM2 phase separation promotes ESCRT-mediated
677 nuclear envelope reformation. *Nature* **582**, 115-118. doi: 10.1038/s41586-020-
678 2232-x (2020).

679 41. Dubińska-Magiera, M., Koziół K, Machowska M, Piekarowicz K, Filipczak D,
680 Rzepecki R. Emerin Is Required for Proper Nucleus Reassembly after Mitosis:
681 Implications for New Pathogenetic Mechanisms for Laminopathies Detected in
682 EDMD1 Patients. *Cells* **8**, 240. doi: 10.3390/cells8030240 (2019).

683 42. Nishimura, K., & Fukagawa T. An efficient method to generate conditional
684 knockout cell lines for essential genes by combination of auxin-inducible degra-
685 tag and CRISPR/Cas9. *Chromosome Res.* **25**, 253-260. doi: 10.1007/s10577-017-
686 9559-7 (2017).

687 43. Nishimura, K., Yamada, T., Hagihara, S., Iwasaki, R., Uchida, N., Kamura, T.,
688 Takahashi, K., Torii, K. U. & Fukagawa, T. A super-sensitive auxin-inducible
689 degra- system with an engineered auxin-TIR1 pair. *Nucleic Acids Res.* **48**, e108.
690 doi: 10.1093/nar/gkaa748 (2020).

- 691 44. Pyzocha, N. K., Ran, F. A., Hsu, P. D., & Zhang, F. RNA-guided genome editing
692 of mammalian cells. *Methods Mol. Biol.* **1114**, 269-277. doi: 10.1007/978-1-62703-
693 761-7_17 (2014).
- 694 45. Olden, B. R., Cheng, E., Cheng, Y., & Pun, S. H. Identifying key barriers in
695 cationic polymer gene delivery to human T cells. *Biomater. Sci.* **7**, 789-797. doi:
696 10.1039/c8bm01262h (2019).
- 697 46. Nguyen, J., & Szoka, F. C. Nucleic acid delivery: the missing pieces of the puzzle?
698 *Acc. Chem. Res.* **45**, 1153-62. doi: 10.1021/ar3000162 (2012).
- 699 47. Houthaeve, G., Robigins, J., Braeckmans, K., & De Vos, W. H. Bypassing Border
700 Control: Nuclear Envelope Rupture in Disease. *Physiology (Bethesda)* **33**, 39-49.
701 doi: 10.1152/physiol.00029.2017 (2018).
- 702 48. Lim, S., Quinton, R. J., & Ganem, N. J. Nuclear envelope rupture drives genome
703 instability in cancer. *Mol. Biol. Cell* **27**, 3210-3213. doi: 10.1091/mbc.E16-02-0098
704 (2016).
- 705 49. Halfmann, C. T., Sears, R. M., Katiyar, A., Busselman, B. W., Aman, L. K., Zhang,
706 Q., O'Bryan, C. S., Angelini, T. E., Lele, T. P., & Roux, K. J. Repair of nuclear
707 ruptures requires barrier-to-autointegration factor. *J. Cell Biol.* **218**, 2136-2149. doi:
708 10.1083/jcb.201901116 (2019).
- 709 50. Chiu, Y. H., Macmillan, J. B., & Chen, Z. J. RNA polymerase III
710 detects cytosolic DNA and induces type I interferons through the RIG-I pathway.
711 *Cell* **138**, 576-91. doi: 10.1016/j.cell.2009.06.015 (2009)
- 712 51. Sun, L., Wu, J., Du, F., Chen, X., & Chen, Z. J. Cyclic GMP-AMP synthase is
713 a cytosolic DNA sensor that activates the type I interferon pathway. *Science* **339**,
714 786-91. doi: 10.1126/science.1232458 (2013).

- 715 52. Burdette, D. L., Monroe, K. M., Sotelo-Troha, K., Iwig, J. S., Eckert, B., Hyodo,
716 M., Hayakawa, Y., & Vance, R. E. STING is a direct innate immune sensor of
717 cyclic di-GMP. *Nature* **478**, 515-8. doi: 10.1038/nature10429 (2011).
- 718 53. Takaoka, A., Wang, Z., Choi, M. K., Yanai, H., Negishi, H., Ban, T., Lu, Y.,
719 Miyagishi, M., Kodama, T., Honda, K., Ohba, Y., & Taniguchi, T. DAI (DLM-
720 1/ZBP1) is a cytosolic DNA sensor and an activator of innate immune response.
721 *Nature* **448**, 501-5. doi: 10.1038/nature06013. (2007).
- 722 54. Liu, C., Wang, H., Zhao, Z., Yu, S., Lu, Y.B., Meyer, J., Chatterjee, G.,
723 Deschamps, S., Roe, B. A. & Lengyel, P. MyoD-dependent induction during
724 myoblast differentiation of p204, a protein also inducible by interferon. *Mol. Cell*
725 *Biol.* **20**, 7024–7036 (2000).
- 726 55. Miyashita, M., Oshiumi, H., Matsumoto, M., & Seya, T. DDX60, a DEXD/H box
727 helicase, is a novel antiviral factor promoting RIG-I-like receptor-mediated
728 signaling. *Mol. Cell Biol.* **31**, 3802–3819. doi: 10.1128/MCB.01368-10 (2011).
- 729 56. Semenova, N., Bosnjak, M., Markelc, B., Znidar, K., Cemazar, M., & Heller, L.
730 Multiple cytosolic DNA sensors bind plasmid DNA after transfection. *Nucleic*
731 *Acids Res.* **47**, 10235-10246. doi: 10.1093/nar/gkz768 (2019).
- 732 57. Sears, R. M., & Roux, K. J. Diverse cellular functions of barrier-to-
733 autointegration factor and its roles in disease. *J. Cell Sci.* **133**, jcs246546. doi:
734 10.1242/jcs.246546 (2020).
- 735 58. Wiebe, M.S., & Jamin, A. The Barrier to Autointegration Factor: Interlocking
736 Antiviral Defense with Genome Maintenance. *J. Virol.* **90**, 3806-3809. doi:
737 10.1128/JVI.00178-16 (2016).

- 738 59. Ma, H., Qian, W., Bambouskova, M., Collins, P. L., Porter, S. I., Byrum, A. K.,
739 Zhang, R., Artyomov, M., Oltz, E. M., Mosammaparast, N., Miner, J. J., Diamond,
740 M. S. Barrier-to-Autointegration Factor 1 Protects against a Basal cGAS-STING
741 Response. *mBio* **11**, e00136-20. doi: 10.1128/mBio.00136-20 (2020).
- 742 60. Guey, B., Wischnewski, M., Decout, A., Makasheva, K., Kaynak, M., Sakar, M. S.,
743 Fierz, B., & Ablasser, A. BAF restricts cGAS on nuclear DNA to prevent innate
744 immune activation. *Science* **369**, 823-828. doi: 10.1126/science.aaw6421 (2020).
- 745 61. Yorifuji, H., Tadano Y, Tsuchiya Y, Ogawa M, Goto K, Umetani A, Asaka Y,
746 Arahata K. Emerin, deficiency of which causes Emery-Dreifuss muscular
747 dystrophy, is localized at the inner nuclear membrane. *Neurogenetics* **1**, 135-40.
748 doi: 10.1007/s100480050020 (1997).
- 749 62. Haraguchi, T., Koujin, T., Hayakawa, T., Kaneda, T., Tsutsumi, C., Imamoto, N.,
750 Akazawa, C., Sukegawa, J., Yoneda, Y., & Hiraoka, Y. Live fluorescence imaging
751 reveals early recruitment of emerin, LBR, RanBP2, and Nup153 to reforming
752 functional nuclear envelopes. *J. Cell Sci.* **113**, 779-794 (2000).
- 753 63. Harborth, J., Elbashir, S. M., Bechert, K., Tuschl, T., & Weber, K. Identification of
754 essential genes in cultured mammalian cells using small interfering RNAs. *J Cell*
755 *Sci.* **114**, 4557-65. (2001).

756

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765

766 **Author contributions**

767 TH, and YHiraoka conceived and designed the experiments. TK, TS, ŞB, HO, KN,
768 YHirano, CM, and SK performed experiments. TH, HA, YO, YC, TF, SS, and YHirano
769 analyzed the data. All the authors contributed reagents/materials/analysis tools. TH,
770 YHirano, HA and YHiraoka wrote the manuscript with the input from all the authors.

771

772 **Competing interests**

773 The authors declare that they have no competing interests in relation to this work.

774

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778

779

780 **Figure legends**

781 **Fig. 1. Gene expression from foreign DNA occurs after mitosis.**

782 **a.** Schematic of the experiment. Top: diagram of pLacO-pEF1 α -mRFP. This plasmid
783 carries a *lacO* repeat sequence (256 repeats, about 10 kbp) and a sequence expressing a
784 monomeric red fluorescent protein (RFP) under the EF1 α promoter. Bottom left: when
785 the transfected plasmids are exposed to the cytosol upon endosome rupture, it binds to
786 GFP-LacI and becomes fluorescent. Bottom right: when it enters the nucleus,
787 expression of RFP occurs. Therefore, RFP fluorescence is an indication of the nuclear
788 translocation of the plasmid. While using this system, gene expression was determined
789 by the appearance of RFP fluorescence.

790 **b.** HeLa/GFP-LacI cells transfected with pLacO-pEF1 α -mRFP (0.5 μ g/dish)
791 diagrammed in panel a. Immediately after removal of the transfection reagent, cells
792 were fixed and immuno-stained using anti-BAF or anti-emerin antibodies. DNA
793 (stained with DAPI, blue), GFP-LacI (green), and BAF (red) or emerin (red) in the
794 merged image. Arrows indicate DAPI signals with GFP-LacI signals. Arrowheads
795 indicate DAPI signals without GFP-LacI signals. The inset shows an image with
796 increased brightness in the boxed region. Bar, 10 μ m

797 **c.** Time-lapse images of transfected HeLa/GFP-LacI cells. DNA was stained with
798 Hoechst 33342. Immediately after removal of the transfection reagent, time-lapse
799 images for each wavelength were acquired at 10-min intervals (1 μ m \times 7 z-stacks for
800 each time-point) using DeltaVision. Maximum intensity projection images for z-stacks
801 for selected time-points are shown. The numbers represent time (h:min). Time 0
802 represents the onset of chromosome segregation. Black-and-white inverted images are

803 shown for each wavelength. Merged images, DNA (Hoechst 33342, blue), GFP-LacI
804 (green), and RFP (red). Bar, 10 μ m.

805 **d.** Timing of gene expression from transfected DNA. Timing of RFP expression was
806 determined for individual cells as shown in panel (c). Time 0 represents the onset of
807 chromosome segregation. Cell numbers tested were n=248 by DeltaVision and n=346
808 by LSM880; total number, n=594. These data are combined in this graph. M, G1, and S
809 represent the cell cycle phases estimated based on our previous report³⁸.

810

811 **Fig. 2. DNA puncta disperse during mitosis.**

812 **a.** Representative time-lapse images of HeLa/GFP-LacI cells transfected with pLacO-
813 pEF1 α -mRFP. DNA was stained with Hoechst 33342. Immediately after removal of the
814 transfection reagent, time-lapse images for each wavelength were acquired at 10-min
815 intervals (1 μ m \times 7 z-stacks for each time-point) using DeltaVision. The z-stack images
816 were deconvoluted. Maximum intensity projection images for the deconvoluted z-stacks
817 of selected time points are shown. Time 0 represents the leftmost image (680 min after
818 the start of image acquisition). Black-and-white inverted images are shown for each
819 wavelength. Merged images, DNA (red) and GFP-LacI (green). Bar, 10 μ m.

820 **b.** HeLa/GFP-LacI cells transfected with pLacO-pEF1 α -mRFP, fixed and immuno-
821 stained for emerlin as indicated in Fig. 1b. A single, representative prometaphase cell is
822 shown. The upper and lower rows show the same cell at focal planes 1.5 μ m apart.
823 Arrowheads indicate small puncta with emerlin signals. Black-and-white inverted
824 images are shown for each wavelength. Merged images, DNA (DAPI, blue), GFP-LacI
825 (green), and emerlin (red). Bars, 10 μ m.

826 **b'**. Enlarged image of the lower panel of GFP-LacI in Fig.2**b**. Arrows indicates small
827 puncta.
828 **c**. Same as panel b except that a metaphase cell is shown. Bar, 10 μ m.
829 **d**. Super-resolution images of GFP-LacI and chromosomes in a metaphase cell.
830 HeLa/GFP-LacI cells were transfected with pLacO-pEF1 α -mRFP. After transfection,
831 cells were fixed with a fixative as described in Methods. Super-resolution images were
832 obtained using LSM880 Airyscan microscope. White and magenta represent small
833 puncta of transfected DNA and chromosomes, respectively. Bar, 10 μ m.
834 **e**. Diagrams of transfected DNA in interphase, prometaphase, and metaphase cells.
835

836 **Fig. 3. CLEM images showing small puncta of transfected DNA in telophase cells.**

837 HeLa/GFP-LacI cells were transfected with pLacO-pEF1 α -mRFP, fixed and subjected
838 to indirect immunofluorescence imaging (**a**) or CLEM (**b-f**).

839 **a**. Indirect immunofluorescence images of telophase cells. After removal of the
840 transfection reagent, cells were fixed and immuno-stained for emerin. DAPI (DNA,
841 blue), GFP-LacI (green), emerin (red) are shown in the merged image. The rightmost
842 cartoon illustrates the core (red) and non-core regions (green) of the NE forming around
843 the telophase chromosomes. Arrows, positions of GFP-LacI. Bar, 10 μ m.

844 **b**. Representative CLEM of telophase cells. Green and magenta represent GFP-LacI and
845 DAPI signals, respectively. Left and right, fluorescence and electron microscopic
846 images, respectively. Middle, merged image. Bar, 5 μ m.

847 **c**. Electron micrographs of the yellow-boxed region of dot 1 in panel **b**. Arrows,
848 positions of GFP-LacI marked with green. Bar, 500 nm.

849 **d.** CLEM image. Different focal planes of the cell shown in panel **(b)**. Left and right,
850 fluorescence and electron microscopic images, respectively. Middle, merged image.
851 Arrows, positions of GFP-LacI (dots 2-4). Bar, 5 μm .
852 **e.** Higher magnification of the red-boxed region in **(d)**. Red circles represent the regions
853 of the GFP-LacI signals, marked in green. Bar, 500 nm.
854 **f.** Classification of puncta morphologies. Type 1 (cytoplasmic circle): plasmid
855 contained within a circular structure surrounded by an NE-like membrane that is present
856 in the cytoplasm. Type 2 (sandwich): plasmid is present between NE-like membranes.
857 Type 3 (membrane fusion): plasmid contained in the cytoplasmic circle fused with the
858 NE. Type 4 (attached to chromosome): plasmid attached to the telophase chromosomes.
859 Dot 4 is the same “dot 4” shown in panel **(d)**. Type 5 (inside the nucleus): plasmid
860 inside the nucleus. Middle panels, schematics of the electron microscopic images. NE is
861 shown as a double line. Red arrows indicate the positions of GFP-LacI, marked in
862 green. Electron microscope images of wider areas, including the area shown in this
863 figure, are shown in Supplementary Figure 3. Bars, 500 nm for Types 1-4 and 1 μm for
864 Type 5.

865

866 **Fig. 4. Immuno-CLEM images of telophase cells.**

867 HeLa/GFP-LacI cells were transfected with pLacO-pEF1 α -mRFP, fixed and subjected
868 to immuno-CLEM.

869 **a.** Fluorescence images. The images on the left are overviews. DNA (stained with
870 Hoechst 33342, blue), GFP-LacI (green), and Alexa-nanogold (red) are shown in the
871 merged images. Right: enlarged image of the red-boxed region. Arrows 1-5 correspond
872 to the dots analyzed in panels **b-e**.

873 **b.** Low-magnification electron micrograph of the telophase cell shown in panel (a).
874 Blue contour, telophase chromosomal region. The right image shows merged electron
875 and fluorescence micrographs. Arrows 1-5 indicate the positions analyzed; red-, blue-,
876 and yellow-boxed regions are enlarged in panels **c**, **d**, and **e**, respectively. Bar, 2 μ m.
877 **c.** The red-boxed region in panel **b**. Red lines, NE-like structures. Arrows, regions
878 where the signals are concentrated. Blue line, edge of the nucleus. Nuc, Cyt, and MT
879 label the nucleus, cytoplasm, and mitochondria, respectively. Bars, 500 nm.
880 **d.** The blue-boxed region in panel **b**. Red lines, NE-like structures. Arrows, regions
881 where the signals are concentrated. Blue line, edge of the nucleus. Nuc and Cyt label the
882 nucleus and cytoplasm, respectively. Bars, 500 nm.
883 **e.** The yellow-boxed region in panel **b**. Arrows, regions where the signals are
884 concentrated. Blue line, edge of the nucleus. Nuc and Cyt label the nucleus and
885 cytoplasm, respectively. Bars, 500 nm.
886 **f.** Diagrams of typical localizations of transfected DNA in telophase cells. Types 1-5 are
887 classified in Fig. 3f.

888

889 **Fig. 5. Depletion of BAF delays plasmid gene expression.**

890 **a.** Time-lapse images of HeLa/mClover3-mAID-Lem2 cells. HeLa/mClover3-mAID-
891 Lem2 were transfected with pEF1 α -mRFP plasmid. After removal of the transfection
892 reagent, time-lapse images for each wavelength were acquired every 10 or 15 min (2
893 μ m x 5 z-stacks for each time point) using DeltaVision. IAA (indole-3-acetic acid) or
894 ethanol as a control was added to HeLa/mClover3-mAID-Lem2 cells during live-cell
895 imaging. Maximum intensity projection images of selected time points are shown. The
896 numbers represent time (h:min). Time 0 is the onset of chromosome segregation. Black

897 arrow indicates timing of the addition of IAA. Colors represent mClover3-mAID-Lem2
898 (green), RFP (magenta) and Hoechst 33342 (white). Orange arrows indicate timing of
899 RFP expression. Bar, 10 μ m.

900 **b.** Timing of RFP expression in control parental (control) and Lem2-depleted HeLa
901 (IAA) cells. Timing of RFP expression was determined for individual cells as shown in
902 panel (a). Time 0 is the onset of chromosome segregation. Cell numbers tested are
903 indicated in the graph.

904 **c.** Time-lapse images of transfected HeLa cells treated with siRNA targeting *BAFNI*
905 (iBAF) and control siRNA (iLuc). Cells were treated with siRNA (iBAF or iLuc) and
906 transfected with pEF1 α -mRFP plasmid. DNA was stained with Hoechst 33342 during
907 transfection. Immediately after removal of the transfection reagent, time-lapse images
908 for each wavelength were acquired at 10-min intervals (2 μ m x 5 z-stacks for a single
909 time-point) using DeltaVision. Images projected by the maximum intensity projection
910 for z-stacks are shown. RFP expression from transfected DNA and Hoechst 33342 are
911 shown in magenta and white, respectively. The numbers represent time (h:min). Time 0
912 represents the onset of chromosome segregation. Bar, 10 μ m.

913 **d.** Western blotting of lysates of HeLa cells treated with siRNA targeting *BAFNI*
914 (iBAF) and control siRNA (iLuc). Tubulin was used as a loading control.

915 **e.** Timing of RFP expression in HeLa cells treated with iBAF and iLuc. Timing of RFP
916 expression was determined for individual cells as shown in panel (c). Time 0 represents
917 the onset of chromosome segregation. Cell numbers tested are indicated in the graph.

918

919 **Fig. 6. NE rupture causes premitotic gene expression.**

920 **a.** HeLa cells were transfected with pEF1 α -mRFP plasmid. NE was ruptured with a
921 microinjection needle by microinjecting FITC-dextran into the nucleus. Left,
922 microinjected cells (green) marked with FITC-dextran. Right, representative image of
923 microinjection needle used for NE rupture. Bar, 10 μ m.

924 **b.** Time-lapse images of cells with NE rupture shown in (a). Immediately after NE
925 rupture, time-lapse images for each wavelength were captured every 10 min (2 μ m x 5
926 z-stacks for a single time-point) using DeltaVision. Images projected by the maximum
927 intensity projection for z-stacks are shown. From the left, FITC-dextran, RFP, and
928 Hoechst images are shown. Black-and-white inverted images are shown for each
929 wavelength. Merged images (second column from the right), FITC-dextran (green),
930 RFP (red), and Hoechst 33342 (blue). The cell in the red-boxed region is shown in the
931 lower panels. The numbers represent time (h:min) from the start of imaging. Red arrow
932 indicates the start of RFP expression. G1, S, G2, and M represent the cell cycle phases
933 estimated based on our previous report³⁸. Bars, 10 μ m.

934 **c.** Western blotting of HeLa cells treated with siRNA targeting *EMD* (iemerin) and the
935 control (iLuc), showing that emerin expression in the cells treated with siRNA was
936 reduced to 17% of control levels.

937 **d.** Time-lapse images of the cells treated with iemerin. HeLa cells were treated with
938 siRNA targeting *EMD* (iemerin). The cells were transfected with pEF1 α -mRFP
939 plasmid. After transfection reagent was replaced with the culture medium, time-lapse
940 images for each wavelength were collected every 10 min (2 μ m x 5 z-stacks for a single
941 time-point) using DeltaVision. Images projected by the maximum intensity projection
942 for z-stacks are shown. The numbers represent time (h:min) from the start of imaging.
943 Expression of RFP started in pre-mitotic cells (white arrow). White arrow indicates the

944 start of RFP expression. S, G2 and M represent the S, G2 and M phases of the cell
945 cycle. Bar, 10 μm .

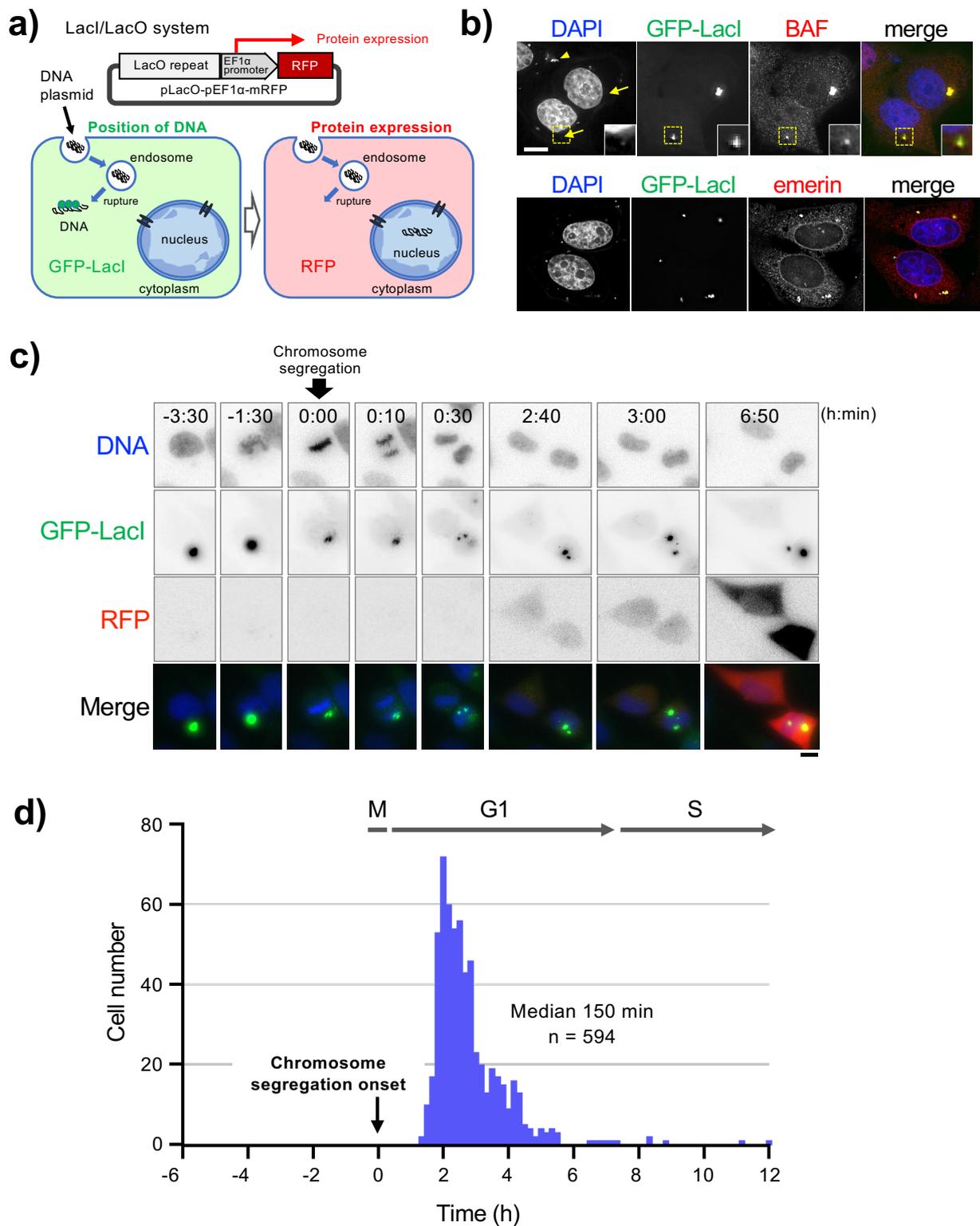


Fig. 1. Gene expression from foreign DNA occurs after mitosis.

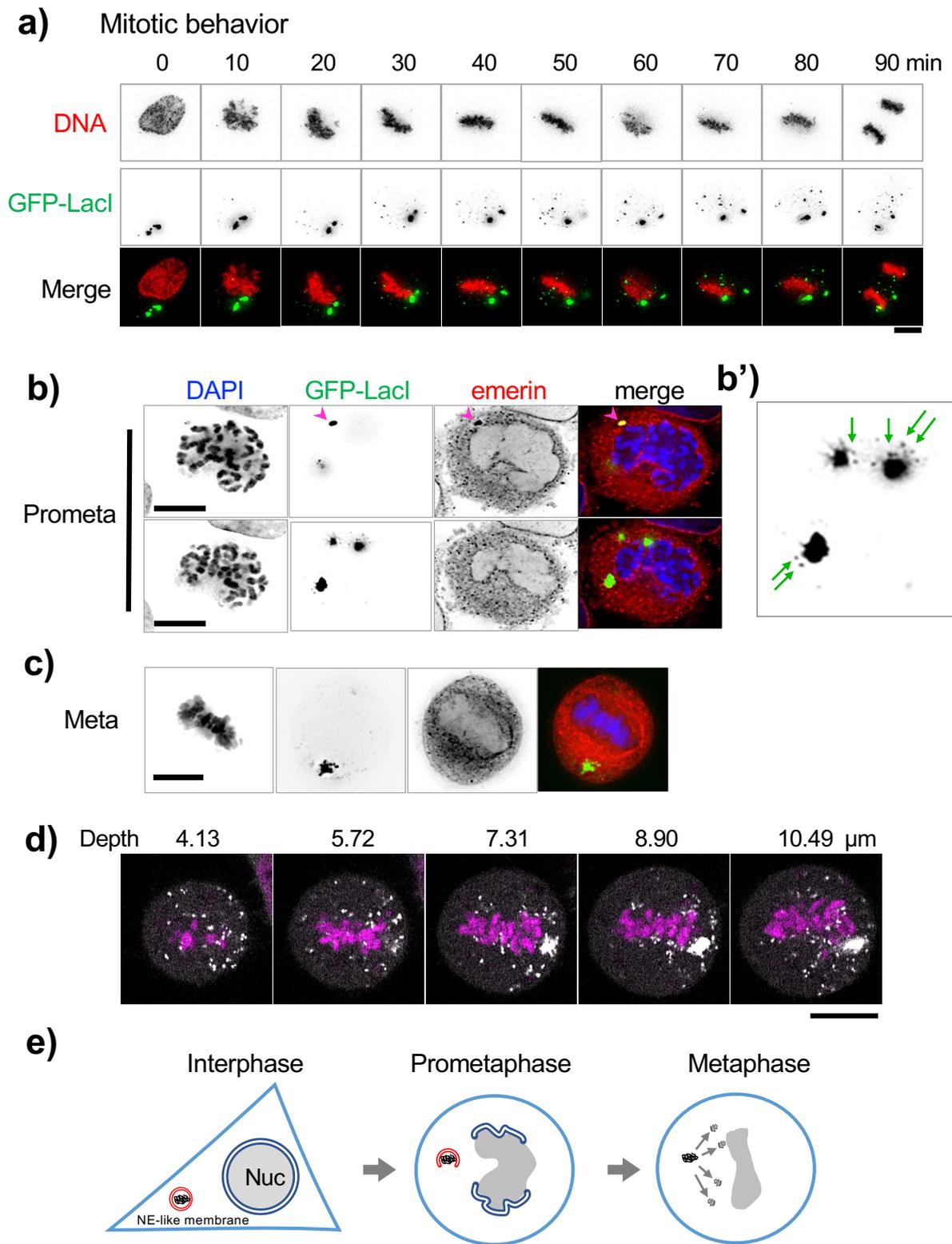


Fig. 2. DNA puncta disperse during mitosis.

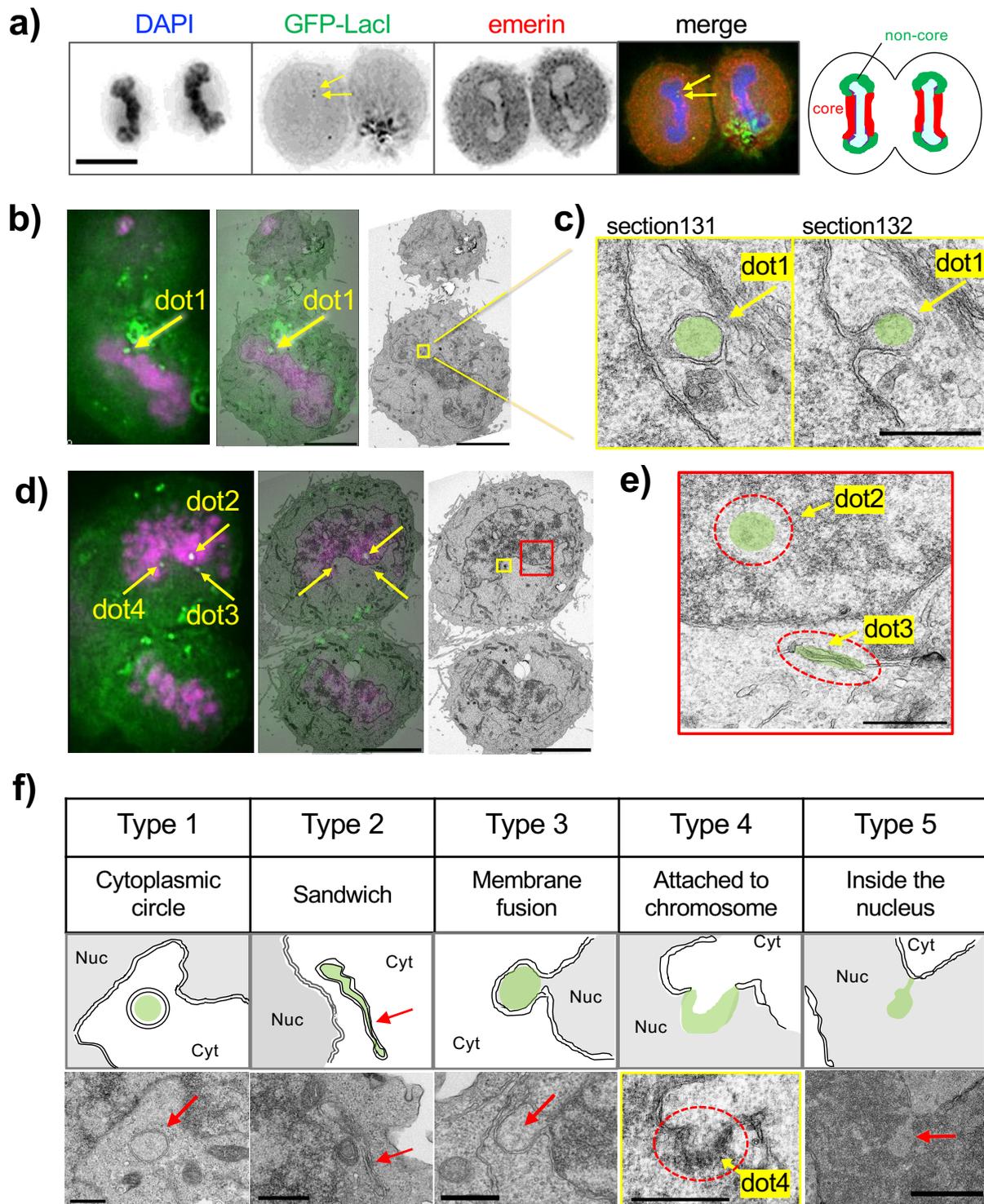


Fig. 3. CLEM images showing small puncta of transfected DNA in telophase cells.

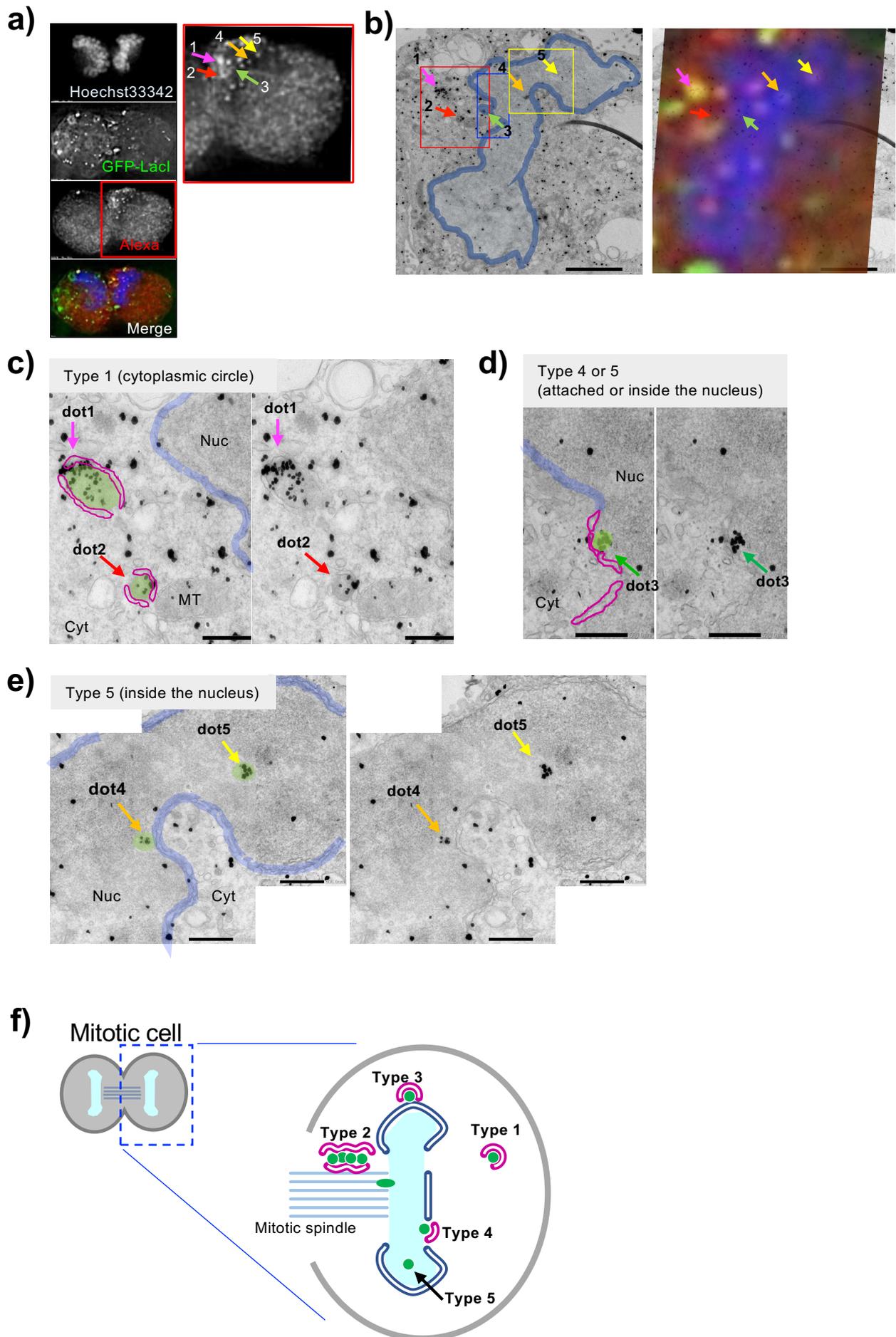
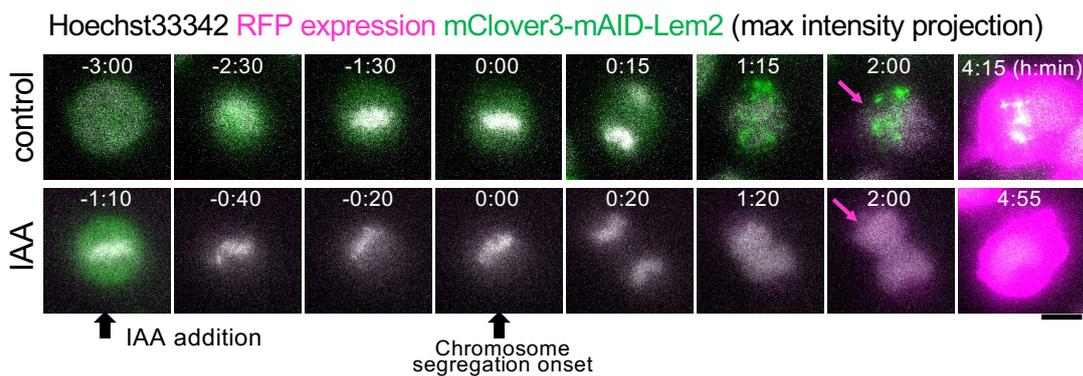
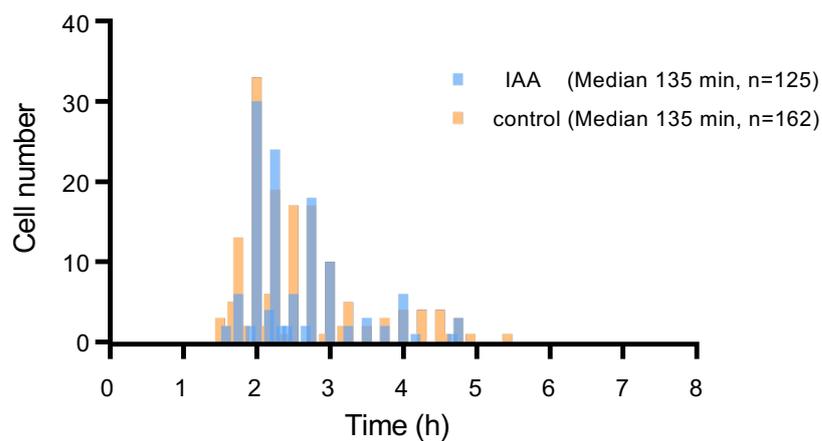


Fig. 4 Immuno-CLEM images in telophase cells

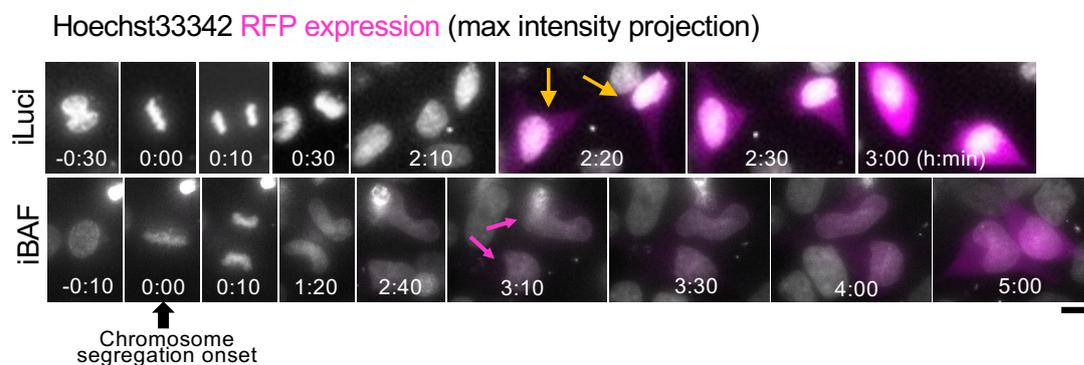
a) Lem2 AID



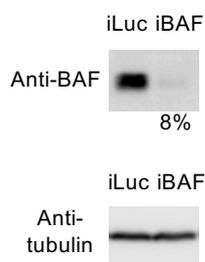
b)



c) iBAF



d) iBAF



e)

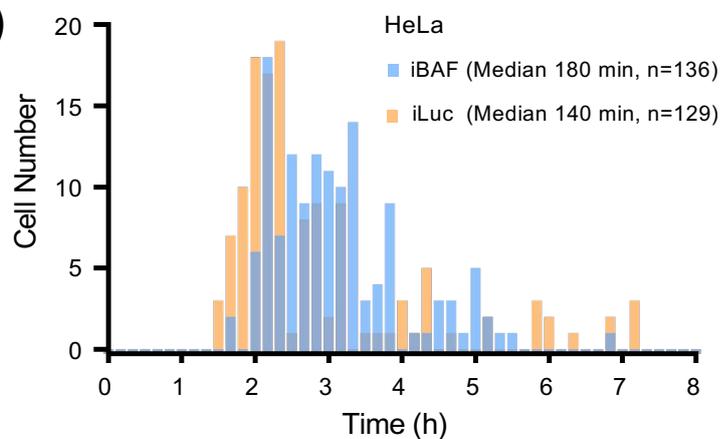


Fig. 5. Depletion of BAF delays plasmid gene expression.

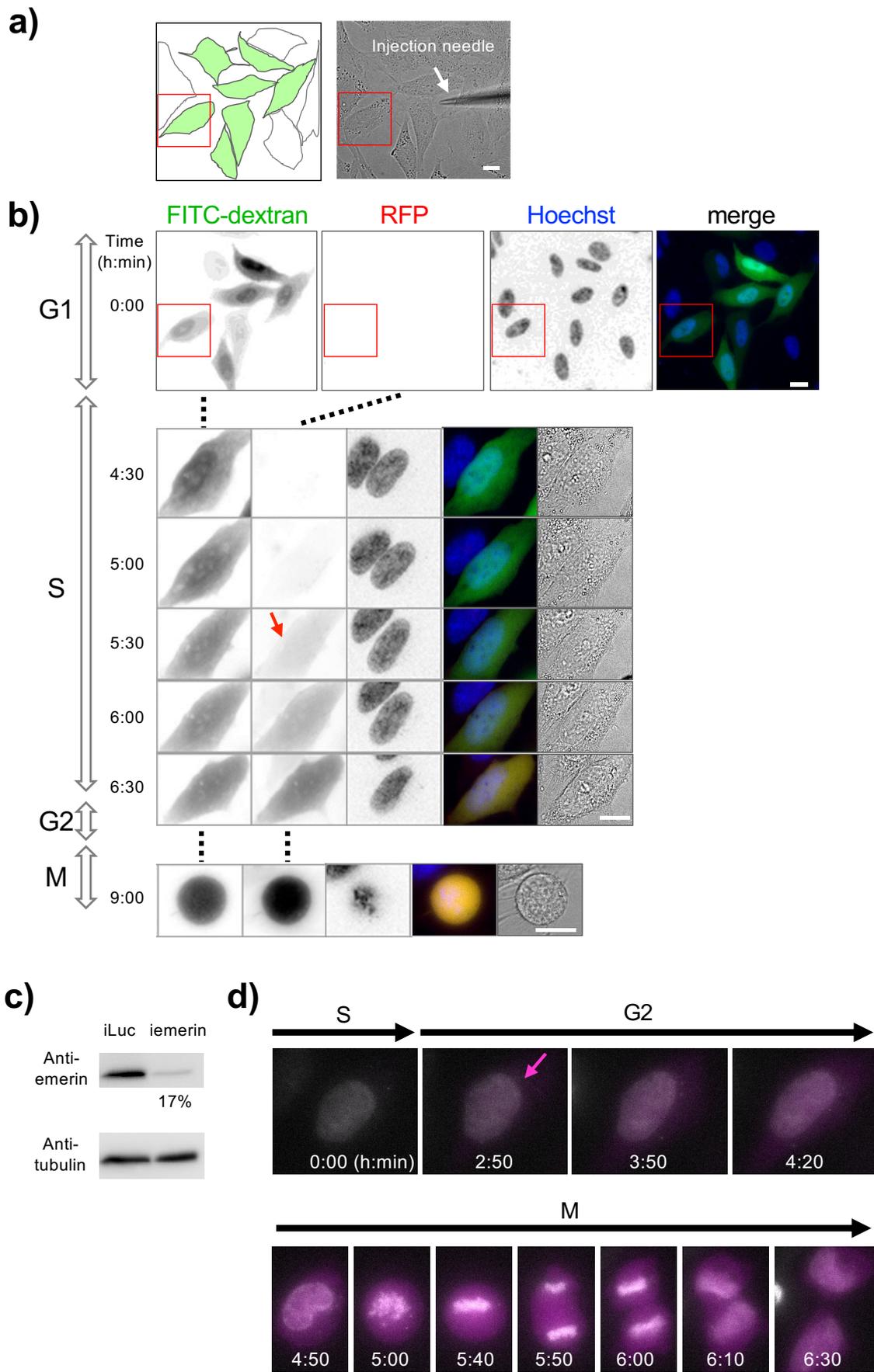


Fig. 6. NE rupture causes premitotic gene expression.

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