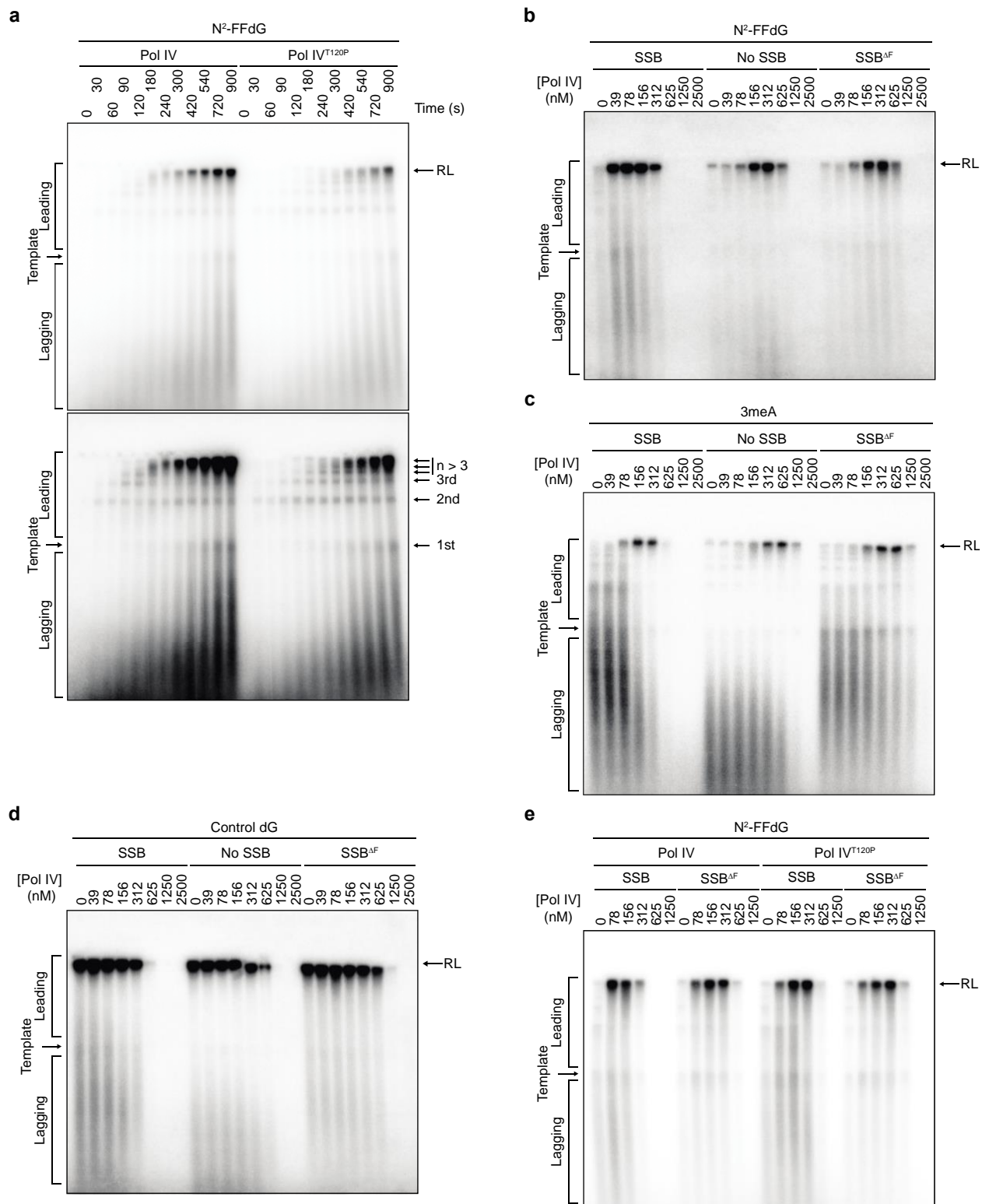


Supplementary Figure 1.

a. Sensitivity of various Thr¹²⁰ mutants of *dinB* to NFZ and MMS.

b. (Left) Changes in the ellipticity of Pol IV or Pol IV^{T120P} at 222 nm upon increasing temperature. (Right) 1st derivatives of the unfolding curves shown in the left panel. The maxima correspond to the midpoints (T_M) of the unfolding curve. Similar results were reproduced twice.



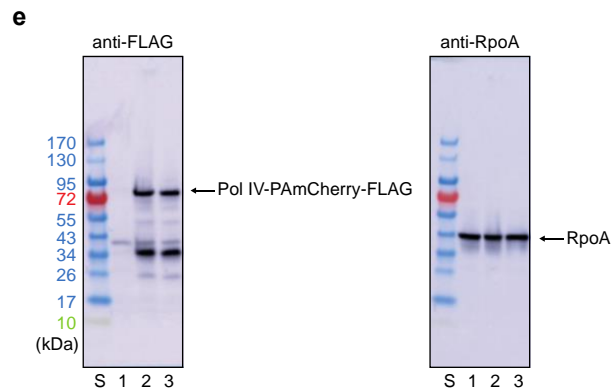
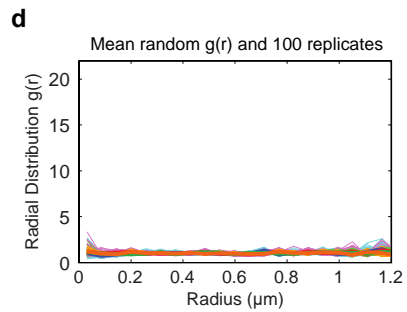
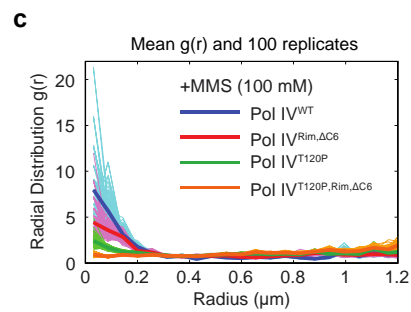
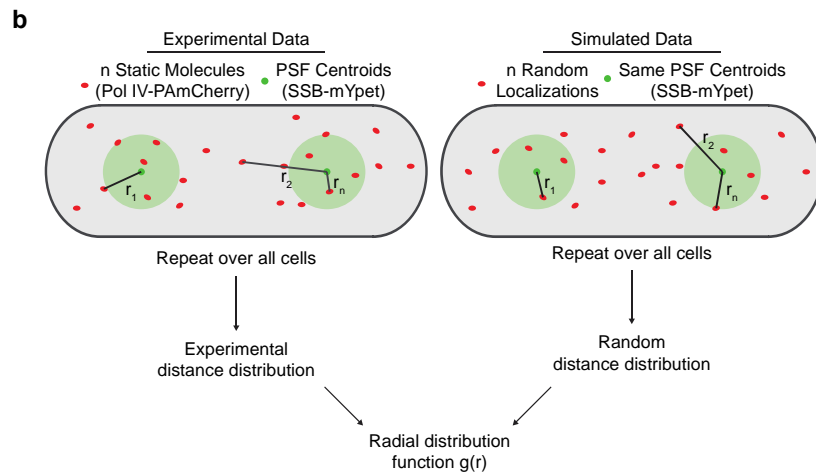
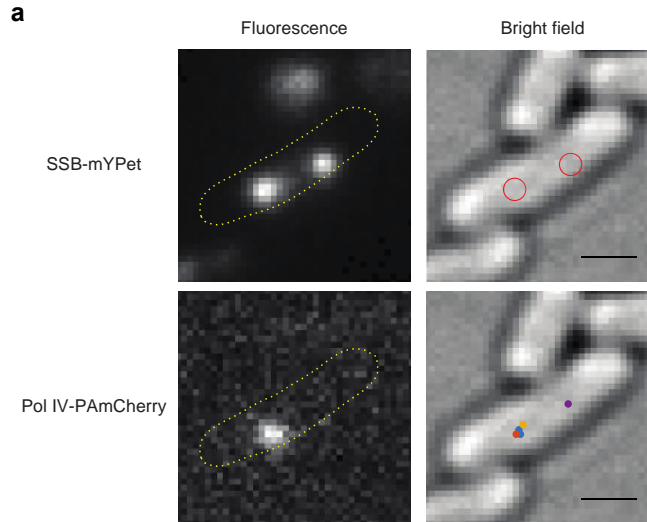
Supplementary Figure 2.

a. (Top) The entire gel for the replication products at RL shown in Fig. 2c. (Bottom) The same gel shown in the top was log-transformed to clearly show stalled replication. Numbers, stalling/collapse of replisomes through multiple passages through N²-FFdG during rolling circle replication.

b, c. The entire gels for the replication products at RL shown in Fig. 2d.

d. SSB weakly potentiates replication inhibitory activity of Pol IV through SSB-Ct. A lesion-free control substrate was replicated with SSB or SSB^{ΔF} or without SSB. As previously reported^{1,2}, replication of both leading and lagging strands was reduced in the absence of SSB or presence of SSB^{ΔF} with lagging strands products becoming shorter. The effects of SSB^{ΔF} on replication were less compared with omitting SSB. The reductions in the amount of replication products from lesion-containing templates in the absence of SSB or the presence of SSB^{ΔF} are attributable to this general reduction in processive replication. Similar results were reproduced twice.

e. The entire gel for the replication products at RL shown in Fig. 2e.



Supplementary Figure 3.

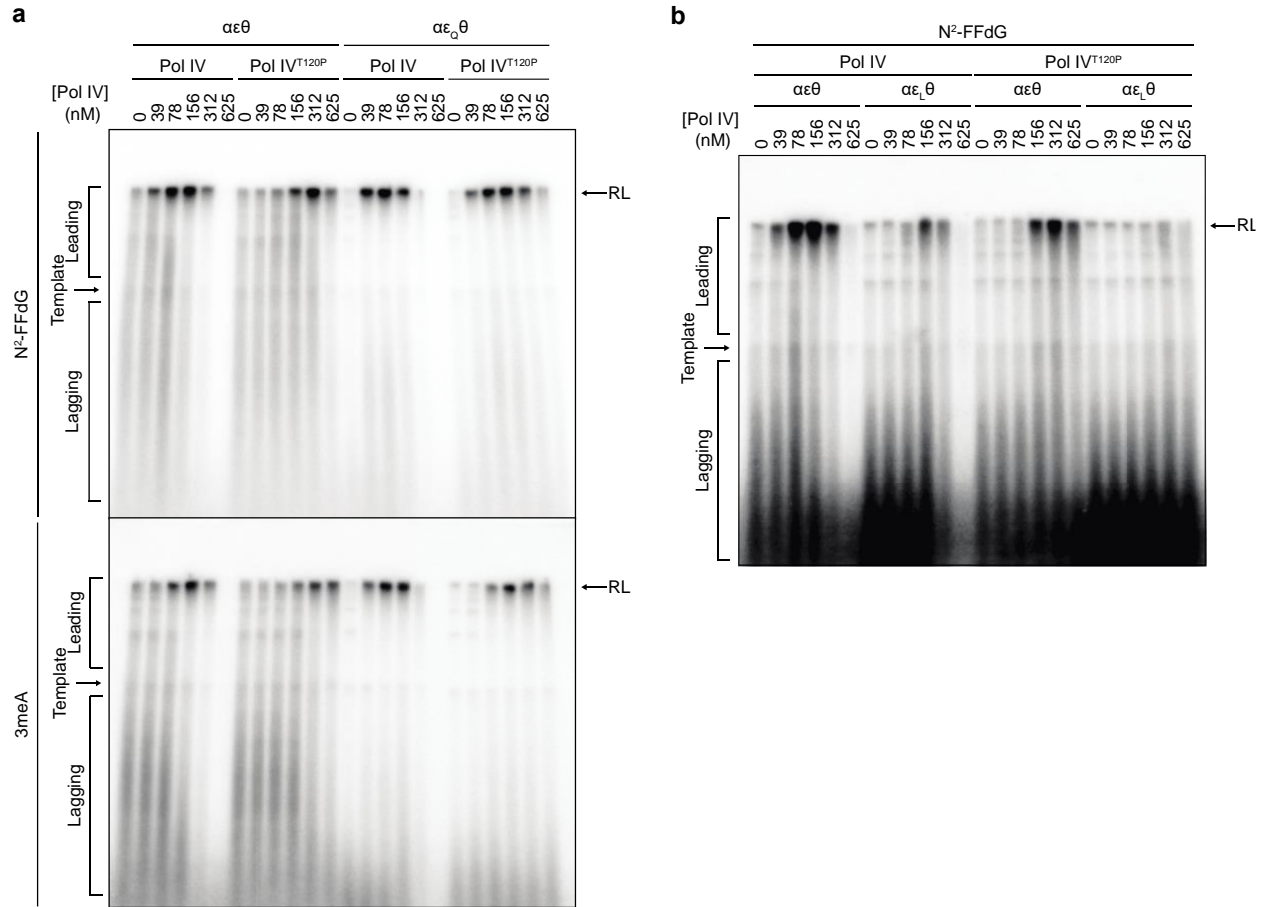
a. (Left) Representative fluorescence micrographs of a cell containing two SSB-mYPet foci (top) and a single photoactivated Pol IV-PAmCherry molecule (bottom) with overlays showing the cell outlines. (Right) Corresponding brightfield micrographs with overlays showing SSB foci (red circle, top) or all detected Pol IV tracks (colored dots, bottom) (Scale bars: 1 μ M).

b. Cartoon of radial distribution function analysis. (Left) Experimental distances between static Pol IV-PAmCherry molecules and the nearest SSB-mYPet foci are calculated. (Right) For each cell, the same number of random localizations are generated and the distances to the same SSB-mYPet foci are determined. This calculation is performed over all cells and the resulting experimental distribution is normalized by the random distribution to give the radial distribution function $g(r)$. This procedure is repeated 100 times to generate 100 $g(r)$ curves, which are averaged to give the final mean $g(r)$ curve.

c. Variability of calculated Pol IV-SSB radial distribution function curves for Pol IV and mutants in MMS-treated cells. The 100 $g(r)$ curves (thin lines) and mean $g(r)$ curves (thick lines) are plotted for each data set.

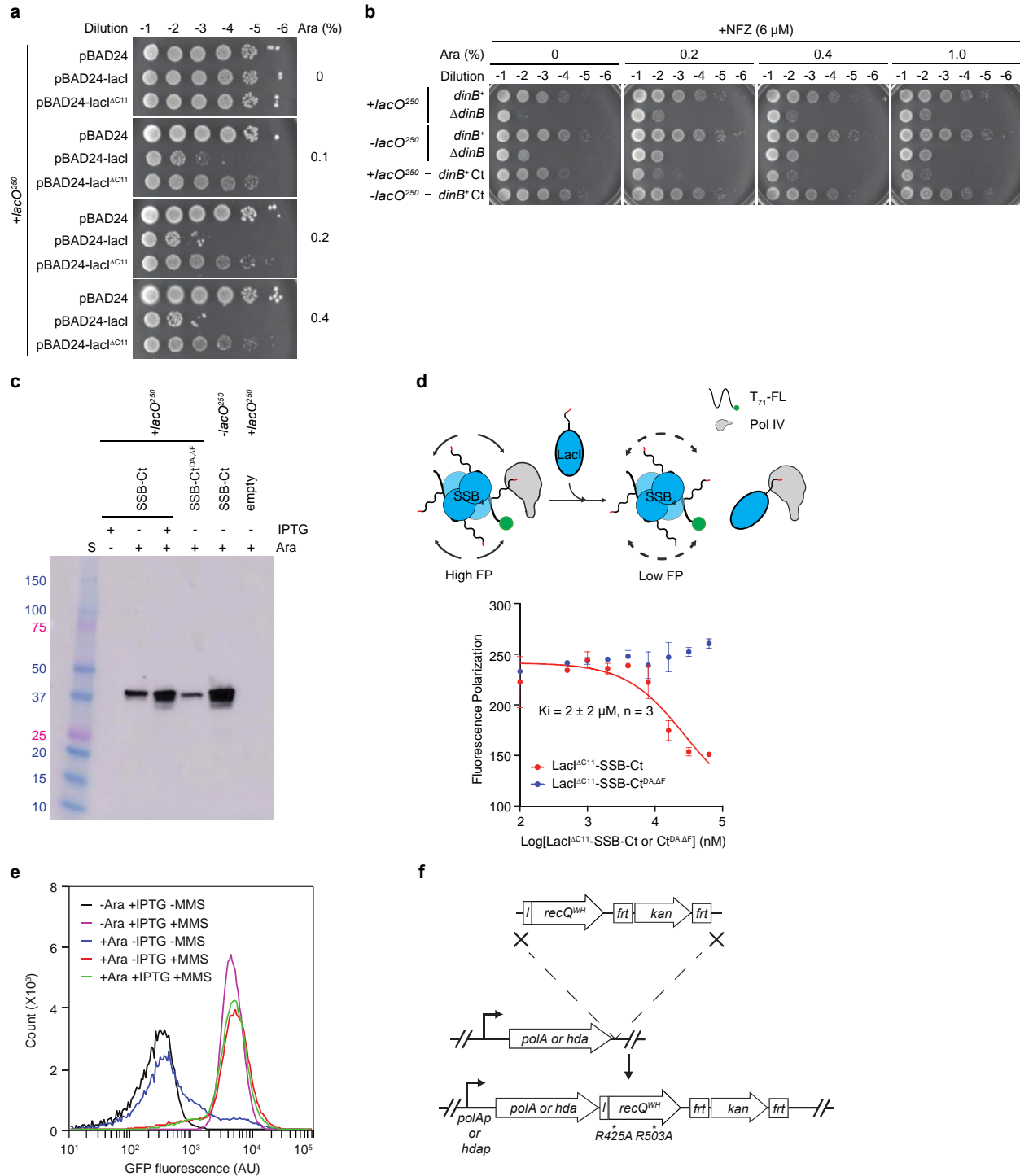
d. As in B, but for the random $g(r)$ curves.

e. Comparable expression of Pol IV-PAmCherry-FLAG and Pol IV^{T120P}-PAmCherry-FLAG in the imaging strains. FLAG-tagged proteins were detected by Western blotting with an anti-FLAG antibody. S. protein size markers; 1, the parent imaging strain; 2, an imaging strain expressing Pol IV-PAmCherry-FLAG; 3, an imaging strain expressing Pol IV^{T120P}-PAmCherry-FLAG. RpoA, a loading control, was probed by Western blotting with an anti-RpoA antibody. Similar results were reproduced twice.



Supplementary Figure 4.

- a. The entire gels for the replication products at RL shown in Fig. 4a.
 b. The entire gel for the replication products at RL shown in Fig. 4b.



Supplementary Figure 5.

a. Deleting the 11 C-terminal residues of LacI (LacI^{ΔC11}) eliminates the cell-killing activity of over-expressed LacI in the +lacO²⁵⁰ strain. Over-expressing LacI^{ΔC11} slightly slowed down cell growth presumably because the LacI^{ΔC11} cluster temporarily attenuates the movement of replisomes. Similar results were reproduced twice.

- b. Expression of LacI^{ΔC11}-SSB-Ct in the absence of the genomic *lacO*²⁵⁰ array does not sensitize cells to NFZ. Assay strains bear *pBAD24* or *pBAD24-lacI^{ΔC11}-ssb-Ct* (Ct) where indicated; + or -*lacO*²⁵⁰, presence or absence of the genomic *lacO*²⁵⁰ array.
- c. Expression of LacI^{ΔC11}-SSB-Ct was not reduced by either addition of IPTG or the lack of the *lacO*²⁵⁰ array. Expression of LacI^{ΔC11} with indicated SSB-Ct appended to the C-terminal end was probed by Western blotting with an anti-LacI antibody. S, size markers; empty, insert-free *pBAD24*. Similar results were reproduced twice.
- d. SSB-Ct in LacI^{ΔC11}-SSB-Ct retains wild-type binding to Pol IV. Interaction between Pol IV and LacI^{ΔC11}-SSB-Ct or SSB-Ct^{DA,ΔF} was determined by measuring FP. (Top) A competition binding assay scheme. (Bottom) FP changes in the presence of indicated concentrations of His₆-LacI^{ΔC11}-SSB-Ct or Ct^{DA,ΔF}. His₆-LacI^{ΔC11}-SSB-Ct competitively inhibits the Pol IV-SSB interaction whereas His₆-LacI^{ΔC11}-SSB-Ct^{DA,ΔF} does not. Means ± ranges of three duplicate measurements.
- e. Nearly wild-type SOS responses were induced upon MMS treatment in assay strains. The +*lacO*²⁵⁰ strain bearing *pBAD24-lacI^{ΔC11}-ssb-Ct* was treated with indicated agents, and the MMS-induced SOS response was measured 90 minutes after the addition of MMS. To eliminate the delay in the expression of LacI^{ΔC11}-SSB-Ct, cells were pre-treated with arabinose overnight and the same concentration of arabinose was present throughout the experiment. Cells were also treated with IPTG in the same way. In subpopulations of cells, induction of LacI^{ΔC11}-SSB-Ct induced varying levels of the SOS response in the absence of MMS. This induction is presumably due to the slowing of replisomes upon encountering the LacI^{ΔC11}-SSB-Ct cluster. In a small fraction of cells (~10%) expressing LacI^{ΔC11}-SSB-Ct (+Ara), the MMS-induced SOS response was only weakly induced. Notably, there is no difference in the MMS-induced SOS response between IPTG-untreated and treated cells. Therefore, IPTG does not eliminate the increased sensitivity to damaging agents of the +*lacO*²⁵⁰ strains by influencing the damage-induced SOS response. Similar results were reproduced twice.
- f. Addition of the coding sequence for wild-type (RecQ^{WH}) or SSB-binding defective RecQ^{WH} (RecQ^{WH(R425A,R503A)}) to the 3' end of native *polA* or *hda* gene. *l*, linker; *polAp* or *hdap*, promoter of *polA* or *hda*; *frt*, flippase recognition site; *kan*, kanamycin resistance gene.