Supplemental movie legends

Supplemental movie 1. Replication and segregation dynamics during novobiocin treatment. TLMM movies of representative cells of the Alpha-EYFP/DnaN-mCherry (top panel) and the DnaN-mCherry/ParB-mNeon (bottom panel) strains. After 5 hours (300 min) of growth under optimal conditions, the novobiocin (10xIC₅₀) was added (+Novo) and then removed 5 hours later (600 min). Images were acquired every 10-min. Scale bar 5 μm.

Supplemental movie 2. Replication and segregation dynamics during Ndx treatment. TLMM movies of representative cells of the Alpha-EYFP/DnaN-mCherry (top panel) and the DnaN-mCherry/ParB-mNeon (bottom panel) strains. After 5 hours (300 min) of growth under optimal conditions, the Ndx (10xIC₅₀) was added (+Ndx) and then removed 5 hours later (600 min). Images were acquired every 10-min. Scale bar 5 μm.

Supplemental movie 3. Replication and segregation dynamics during GM treatment. TLMM movies of representative cells of the Alpha-EYFP/DnaN-mCherry (left panel) and the DnaN-mCherry/ParB-mNeon (right top panel) and the Alpha-EYFP/ParB-mCherry (right bottom panel) strains. After 5 hours (300 min) of growth under optimal conditions, the GM (10xIC₅₀) was added (+GM) and then removed 5 hours later (600 min). Images were acquired every 10-min. Scale bar 5 μm.

Supplemental movie 4. Chromosome and replisome dynamics during exposure to replicationaffecting drugs. TLMM movies of representative cells of the HupB-EGFP/DnaN-mCherry strain during Ndx (left panel), novobiocin (middle panel) and GM (right panel) treatment. After 5 hours (300 min) of growth under optimal conditions, the antibiotics (10xIC₅₀) were added (+Ndx, +Novo, +GM) and then removed 5 hours later (600 min). Images were acquired every 10-min. Scale bar 5 μ m.

Supplemental Tables

Strain name	Relevant genotype	Reference
Alpha-EYFP/DnaN- mCherry	<i>M. smegmatis Mc² 155 alpha-eyfp, dnaN-mCherry</i>	(1)
DnaN- mCherry/ParB- mNeon	M. smegmatis Mc ² 155 dnaN-mcherry , parB- mNeon,	(2)
Alpha-EYFP/ParB- mCherry	<i>M. smegmatis Mc² 155 alpha-eyfp, parB-mCherry</i>	(1)
DnaN-mCherry/HupB- EGFP	<i>M. smegmatis Mc² dnaN-mcherry, hupB-egfp</i>	(3)
RRL-19B	E. coli AB1157 (thr-1, araC14, leuB6(Am), Δ(gpt- proA)62, lacY1, tsx-33, qsr'-0, glnV44(AS), galK2(Oc), LAM-, Rac-0, hisG4(Oc), rfbC1, mgl- 51, rpoS396(Am), rpsL31(strR), kdgK51, xylA5, mtl-1, argE3(Oc), thi-1) ypet-dnaN	(4)

Table S1. Characteristics of strains used in this study

Supplemental Figures



Fig. S1. Growth curves of the *M. smegmatis* strains used in this study in the presence of the different antibiotics. The right bottom panel presents the growth of all strains in antibiotic-free medium.



Fig. S2. Inhibition curves calculated from the growth curves plotted in Figure S1. Top panel, griselimycin; middle panel, nalidixic acid; bottom panel, novobiocin. IC_{50} is marked with a red dotted line.



Fig. S3. The binding of HupB-EGFP to the chromosome is not affected under GM treatment, and reflects the nucleoid structure. Micrographs and corresponding fluorescence intensity profiles of DnaN-mCherry/HupB-EGFP cells stained with Hoechst 33342 before (A) and after 5 hours of GM treatment (B and C). During GM treatment, the chromosome is decondensed in some cells (B), while diffuse fluorescence is observed in others (C). Scale bar, $2 \mu m$.

Supplemental references

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