

Supporting Information

Jain et al. 10.1073/pnas.1211144109

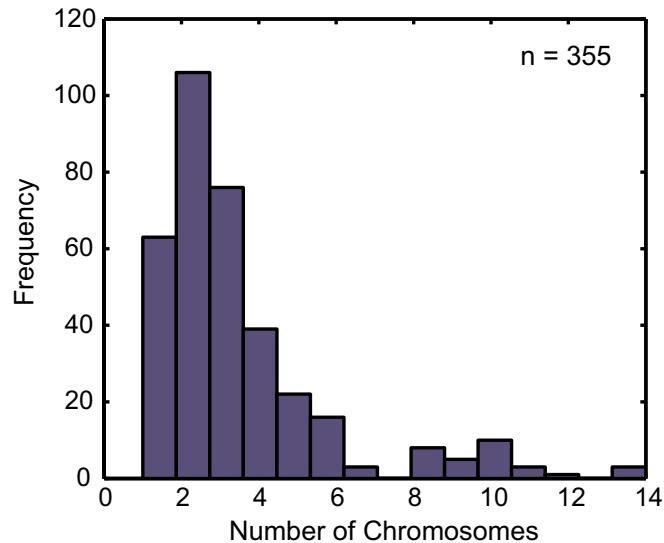


Fig. S1. Histogram of the number of chromosomes per cell in an exponentially growing wild-type population.

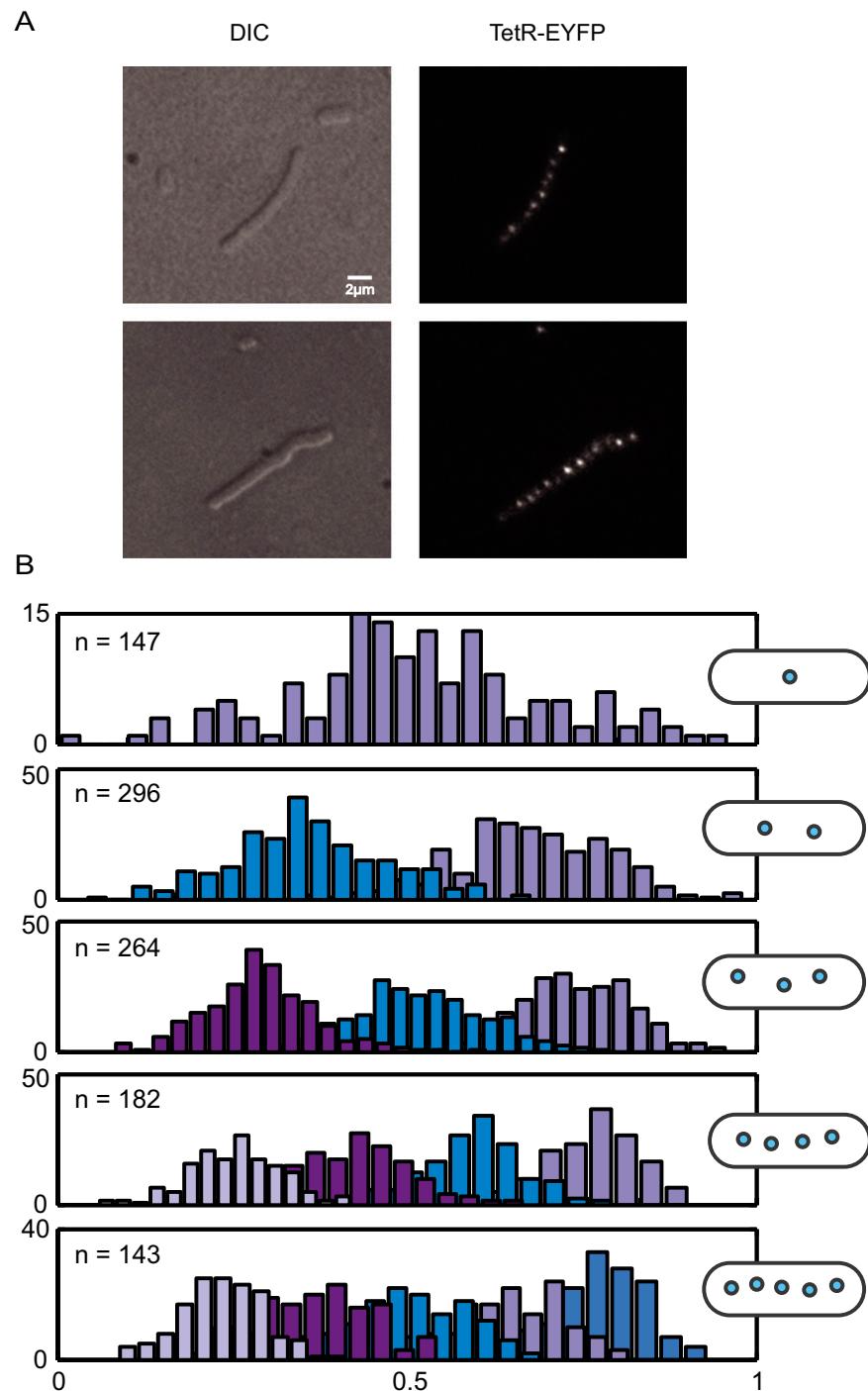


Fig. S2. Cells maintain chromosome ordering along the long axis in a $\Delta minD$ strain. (A) Examples of representative $\Delta minD$ cells. A single z-section is shown. (B) Cells maintain chromosome ordering along the long axis in a $\Delta minD$ strain. Sample size (n) refers to the number of cells analyzed with the given number of chromosomes.

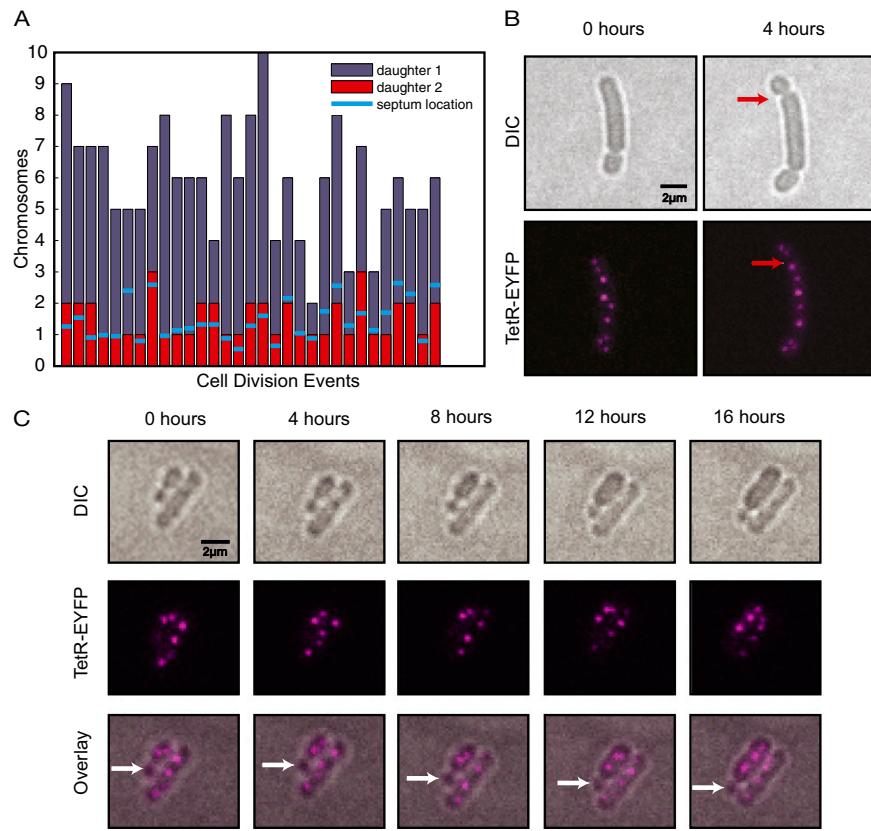


Fig. S3. Cell division in $\Delta minD$ cells. (A) The number of chromosomes partitioned to daughter cells in $\Delta minD$ cells. Each stacked bar represents a single cell division event with the height of red and dark blue bars representing chromosomes partitioned to each daughter cell. Light blue lines represent the actual septum location as a fraction of total chromosome number (bar height). (B) An example of chromosome partitioning in $\Delta minD$ cells based on the position of the septum (red arrow). A single z-section is shown. (C) Misplaced septum formation in $\Delta minD$ cells can result in anucleate daughter cells, shown by white arrows. A single z-section is shown.

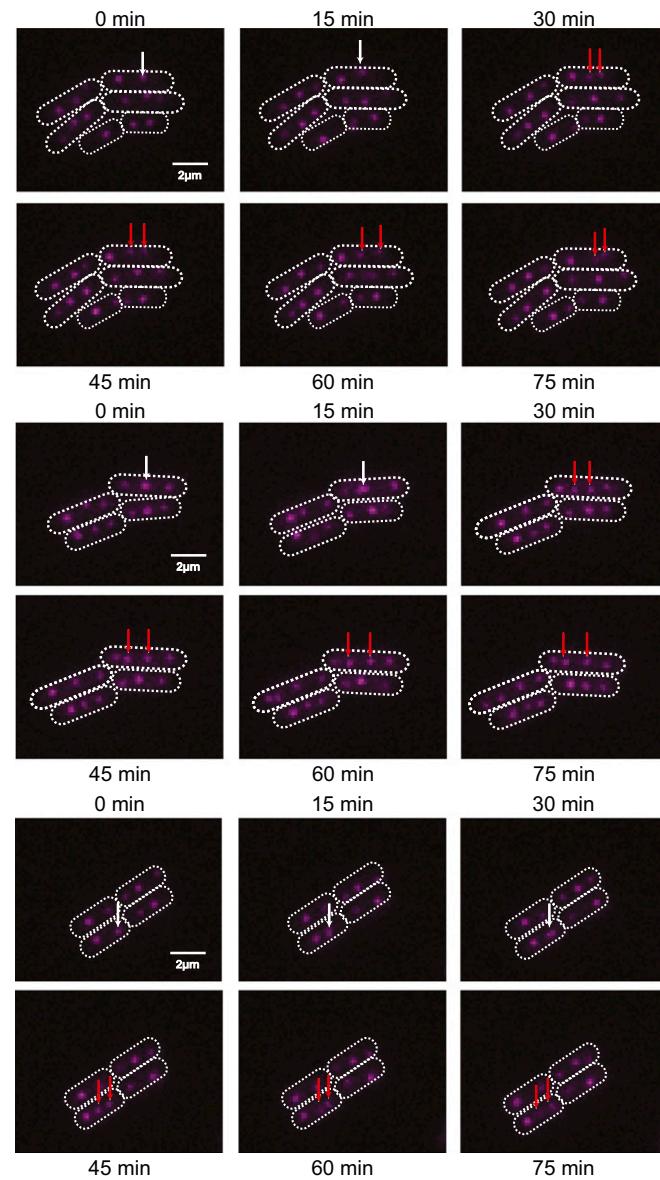


Fig. S4. Three time courses of wild-type cells. A single genomic locus proximal to the origin is labeled by using tet operator arrays (pink dots). White arrows point to a replicating chromosome and red arrows point to the resulting, replicated chromosomes. A single z-section is shown for each time course.

Table S1. Table of plasmids

Plasmid	Description	Resistance	Source
pAM1303	Neutral site 1 integration vector	Sp/Sm	(1)
pAM1579	Neutral site 2.1 integration vector	Kan (Amp)	(1)
pAM1573	Neutral site 2.1 integration vector	Cm (Amp)	(1)
EB2065	Neutral site 2.2 integration vector	Cm (Amp)	This work
EB2066	Site A integration vector (between chromosomal position 1595934 and 1595935 relative to GenBank CP000100)	Cm (Amp)	This work
EB2067	Site B integration vector (between chromosomal position 1758400 and 1758401 relative to GenBank CP000100)	Cm (Amp)	This work
eBB110	Source of 120 Tet operator repeats	Amp	(2)
PLAU43	Source of 120 Lac operator repeats	Amp	(3)
EB2068	120 Tet operators PCR'd using primers pBBSall and pBBXbal from eBB110 cloned between Sall and XbaI of pAM1579	Kan (Amp)	This work
EB2069	120 Lac operators from XbaI, SmaI fragment of pLAU43 cloned between XbaI and SmaI of pAM1573	Cm (Amp)	This work
EB2070	120 Lac operators from SmaI, Sall fragment of pLAU43 cloned between SmaI and Xhol of EB2066	Cm (Amp)	This work
EB2071	120 Lac operators from SmaI, Sall fragment of pLAU43 cloned between SmaI and Xhol of EB2067	Cm (Amp)	This work
pJRC23	Source of ECFP	Sp/Sm	(4)
EB2072	PkaiBC::tetR::ECFP PkaiBC::EYFP::lacI cloned between NotI and SacI of pAM1303	Sp/Sm	This work
EB2073	PkaiBC::tetR::EYFP cloned between NotI and SacI of pAM1303	Sp/Sm	This work
EB2074	PapcA::rbcl::ECFP cloned between SmaI and Xhol of EB2065	Cm (Amp)	This work
pAM2055	Source of gentamycin cassette for deletion vectors	Gm (Amp)	(1)
EB2075	mreB (Synpcc7942_0300) deletion vector with gentamycin cassette inserted between BamHI and SacI of pUC18	Gm (Amp)	This work
EB2076	ftsZ (Synpcc7942_2378) deletion vector with gentamycin cassette inserted between HindIII and SacI of pUC18	Gm (Amp)	This work
EB2077	parA1 (Synpcc7942_0220) deletion vector with gentamycin cassette inserted between BamHI and SacI of pUC18	Gm (Amp)	This work
EB2078	minD (Synpcc7942_0896) deletion vector with gentamycin cassette inserted between BamHI and SacI of pUC18	Gm (Amp)	This work
EB2079	parA2 (Synpcc7942_1833) deletion vector with gentamycin cassette inserted between BamHI and SacI of pUC18	Gm (Amp)	This work

Markers shown in parenthesis are additional markers used for selection of plasmids in *E. coli*. Cm, chloramphenicol; Gm, gentamycin; Kan, kanamycin; Sp/Sm, spectinomycin/streptomycin.

1. Mackey SR, Ditty JL, Clerico EM, Golden SS (2007) Detection of rhythmic bioluminescence from luciferase reporters in cyanobacteria. *Methods Mol Biol* 362:115–129.

2. Marquis KA, et al. (2008) SpollIE strips proteins off the DNA during chromosome translocation. *Genes Dev* 22:1786–1795.

3. Lau IF, et al. (2003) Spatial and temporal organization of replicating *Escherichia coli* chromosomes. *Mol Microbiol* 49:731–743.

4. Chabot JR, Pedraza JM, Luitel P, van Oudenaarden A (2007) Stochastic gene expression out-of-steady-state in the cyanobacterial circadian clock. *Nature* 450:1249–1252.

Table S2. Table of *S. elongatus* strains

Strain	Description	Plasmids used to create this strain from WT	Resistance	Source
EOC200	TetO (NS 2.1)	EB2068	Kan	This work
EOC201	TetO (NS 2.1), LacO (A)	EB2068, EB2070	Kan, Cm	This work
EOC202	TetO (NS 2.1), LacO (B)	EB2068, EB2071	Kan, Cm	This work
EOC203	TetO (NS 2.1), <i>tetR::EYFP</i> (NS 1)	EB2068, EB2073	Kan, Sp/Sm	This work
EOC204	TetO (NS 2.1), LacO (A), <i>tetR::ECFP EYFP::lacI</i> (NS 1)	EB2068, EB2070, EB2072	Kan, Cm, Sp/Sm	This work
EOC205	TetO (NS 2.1), LacO (B), <i>tetR::ECFP EYFP::lacI</i> (NS 1)	EB2068, EB2071, EB2072	Kan, Cm, Sp/Sm	This work
EOC206	TetO (NS 2.1), <i>tetR::EYFP</i> (NS 1), <i>rbcL::ECFP</i> (NS 2.2)	EB2068, EB2073, EB2074	Kan, Cm, Sp/Sm	This work
EOC207	$\Delta mreB$ incomplete segregation	EB2075	Gm	This work
EOC208	$\Delta ftsZ$ incomplete segregation	EB2076	Gm	This work
EOC209	$\Delta parA1$	EB2077	Gm	This work
EOC210	$\Delta minD$	EB2078	Gm	This work
EOC211	$\Delta parA2$	EB2079	Gm	This work
EOC212	$\Delta mreB$, TetO (NS 2.1)	EB2075, EB2068	Gm, Kan	This work
EOC213	$\Delta ftsZ$, TetO (NS 2.1)	EB2076, EB2068	Gm, Kan	This work
EOC214	$\Delta parA1$, TetO (NS 2.1)	EB2077, EB2068	Gm, Kan	This work
EOC215	$\Delta minD$, TetO (NS 2.1)	EB2078, EB2068	Gm, Kan	This work
EOC216	$\Delta parA2$, TetO (NS 2.1)	EB2079, EB2068	Gm, Kan	This work
EOC217	$\Delta mreB$ incomplete segregation, TetO (NS 2.1), <i>tetR::EYFP</i> (NS 1), <i>rbcL::ECFP</i> (NS 2.2)	EB2075, EB2068, EB2073, EB2074	Gm, Kan, Sp/Sm, Cm	This work
EOC218	$\Delta ftsZ$ incomplete segregation, TetO (NS 2.1), <i>tetR::EYFP</i> (NS 1), <i>rbcL::ECFP</i> (NS 2.2)	EB2076, EB2068, EB2073, EB2074	Gm, Kan, Sp/Sm, Cm	This work
EOC219	$\Delta parA1$, TetO (NS 2.1), <i>tetR::EYFP</i> (NS 1), <i>rbcL::ECFP</i> (NS 2.2)	EB2077, EB2068, EB2073, EB2074	Gm, Kan, Sp/Sm, Cm	This work
EOC220	$\Delta minD$, TetO (NS 2.1), <i>tetR::EYFP</i> (NS 1), <i>rbcL::ECFP</i> (NS 2.2)	EB2078, EB2068, EB2073, EB2074	Gm, Kan, Sp/Sm, Cm	This work
EOC221	$\Delta parA2$, TetO (NS 2.1), <i>tetR::EYFP</i> (NS 1), <i>rbcL::ECFP</i> (NS 2.2)	EB2079, EB2068, EB2073, EB2074	Gm, Kan, Sp/Sm, Cm	This work

Table S3. Table of primers

Primer	Sequence
Primers for site A (EB2066), B (EB2067), and NS 2.2 (EB2065) integration vectors	
MCS_F	5' GTGCTTCTGGCTATAGCTGACTcgtaacggtaaccga 3'
MCS_R	5' ccgcgtgggttaccag 3'
A-Up-F	5' GACCACACCCGTCTGTGGATCCTGCAACTATCCCTCGATC 3'
A-Up-R	5' AGTCAGCTATAAGCCAGAACGACTGGGAGGCATTAGAACG 3'
A-Dn-F	5' ctggtaaccccaagcgcggATCGGATTGAAATTGTTTC 3'
A-Dn-R	5' TGCCTCCGGCGTAGAGGATCCGTCTCGACCACCCATTAG 3'
B-Up-F	5' GACCACACCCGTCTGTGGATCCGTGTTGTAGGCCACGAT 3'
B-Up-R	5' AGTCAGCTATAAGCCAGAACGACCGCTAACGGATGAGCG 3'
B-Dn-F	5' ctggtaaccccaagcgcggGACTGTCTGAATTACCTTCAG 3'
B-Dn-R	5' TGCCTCCGGCGTAGAGGATCCGTAGAAGCACGTGGC 3'
NS22-Up-F	5' GACCACACCCGTCTGTGCAGGCTCAGTGTGGTTCG 3'
NS22-Up-R	5' AGTCAGCTATAAGCCAGAACGAC 3'
NS22-Dn-F	5' ctggtaaccccaagcgcggAGATCTCGCAGCGTAAAGCCGT 3'
NS22-Dn-R	5' TGCCTCCGGCGTAGAGGATCCGTAGGCCATCCACGCAC 3'
Primers for tet operator vector (EB2068)	
pBBSall	5' TAAACTATGTCGACCTTCTTATCTTGATAATAAGGGTAAC 3'
pBBXbal	5' ATAGTTTATCTAGAccgtccttgaacatgact 3'
Primers for PapcA::RbcL::ECFP vector (EB2074)	
Smal_ApcA	5' CATGCCCGGGTACGAGCGCTATATCACCCC 3'
ApcA_RBS	5' TAATGGATTCTCCAAGACTAGATTGAAACCAGACTGGCCTCCACC 3'
RBS_RbcL	5' TCTAGTCTGGAGGAATCCATTAATGCCAAGACGCAATCTG 3'
RbcL_linker	5' ACTAGAACCGAAACTACCAACTAGAGAGCTGTCCATGTTCAATTG 3'
Linker_ECFP	5' TCTAGTGGTAGTTCTGGTTAGTATGGTGGAGCAAGGGCGAGGAG 3'
ECFP_Xhol	5' CATGCTCGAGTTACTGTACAGCTCGCATGC 3'
Primers for deletion vectors (EB2075 to EB2079)	
Gent_F	5' gacgcacacccgtgga 3'
Gent_R	5' gccgcgttgtgacaa 3'
1833KO-Up-F	5' GATCGATCGGATCCACAAAAAGGGGGCTGGTTAG 3'
1833KO-Up-R	5' GttccacgggtgcgtcACTAGT CGACAACCTCCAAAGCG 3'
1833KO-Dn-F	5' AaattgtcacaacgcgcTGACTGACGCCATTGACC 3'
1833KO-Dn-R_Sac	5' GATCGATCGAGCTCAAACAAAATGCCCAAAGT 3'
0300KO-Up-F	5' GATCGATCGGATCCGGTAGCCGAATCACTCCGA 3'
0300KO-Up-R	5' GttccacgggtgcgtcACTAGTCCGCCATTGATGACGTG 3'
0300KO-Dn-F	5' AaattgtcacaacgcgcAGCGGTGCGGAGCAG 3'
0300KO-Dn-R_Sac	5' GATCGATCGAGCTCAGAGTGTACTAGCGCTAGAAGTG 3'
0220KO-Up-F	5' GATCGATCGGATCCACCGAGATATGCTCGATTG 3'
0220KO-Up-R	5' GttccacgggtgcgtcACTAGTGGTGGATCTCAAGTCAGGG 3'
0220KO-Dn-F	5' AaattgtcacaacgcgcCAGCTGCCAAATCTG 3'
0220KO-Dn-R_Sac	5' GATCGATCGAGCTCGAACACTGAGCTTGGCTTGCT 3'
2378KO-Up-F_HindIII	5' GATCGATCAAGCTTAACGGCTGAGCGCTT 3'
2378KO-Up-R	5' GttccacgggtgcgtcACTAGTGGGGTGAGTAGTAGCA 3'
2378KO-Dn-F	5' AaattgtcacaacgcgcTGGCTGTTCCGATCGCC 3'
2378KO-Dn-R_Sac	5' GATCGATCGAGCTCTGTCAATGGTCCCGCT 3'
0896KO-Up-F	5' GATCGATCGGATCCGCCAGTAACCGTCACAGAT 3'
0896KO-Up-R	5' GttccacgggtgcgtcACTAGTAGGGTCCGAAGAGCAGGAG 3'
0896KO-Dn-F	5' AaattgtcacaacgcgcGGGTCTTGCAGCAATGCT 3'
0896KO-Dn-R_Sac	5' GATCGATCGAGCTCTGACCACCTAGCAAATGTTCC 3'

Table S4. Sequence of TetR and LacI fluorescent fusion protein constructs

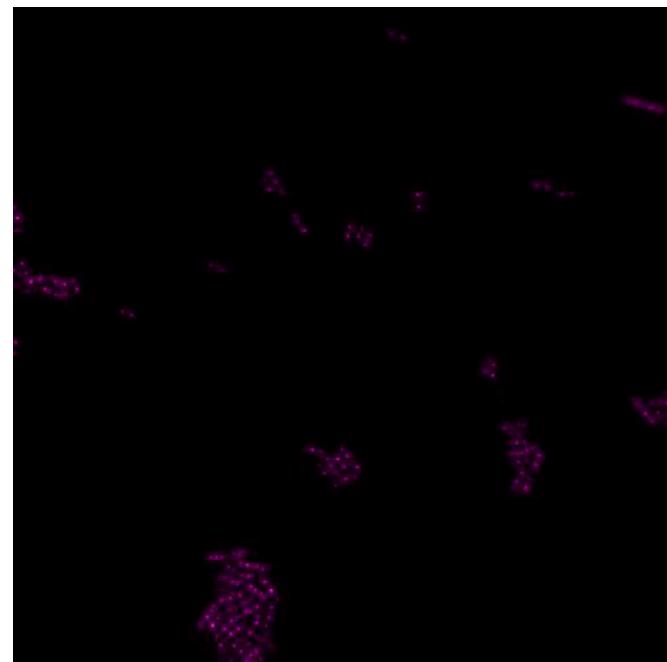
EB2072: *PkaiBC::RBS::tetR::ECFP PkaiBC::RBS::EYFP::linker::lacI* cloned between *NotI* and *SacI* of pAM1303. Note that the beginning of *PkaiBC* contains the *kaiA* terminator which will serve to decouple TetR and LacI expression.

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Table S4. Cont.

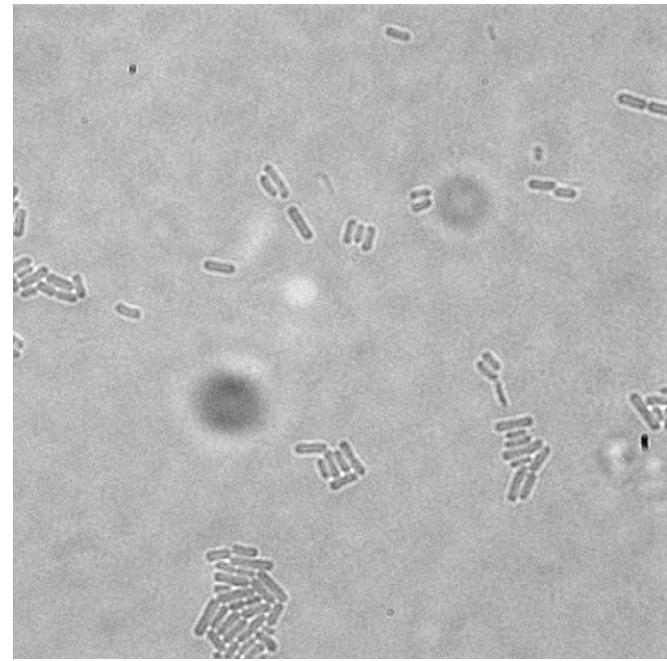
EB2073: *PkaiBC*::*RBS*::*teIR*::*linker*::*EYFP* cloned between *NotI* and *SacI* of pAM1303

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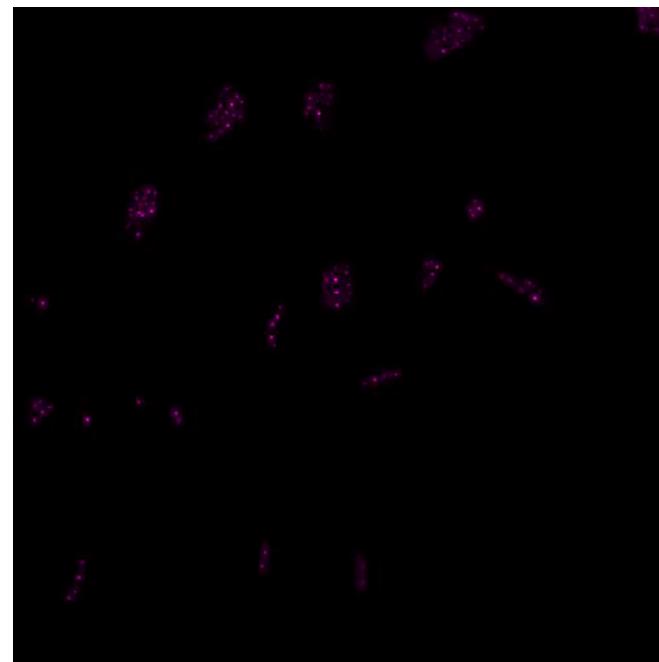
Movie S1. Time-lapse movies of wild-type cells with labeled chromosomes (TetR-EYFP). Individual frames are separated by 1 h. A single z-plane is shown.

[Movie S1](#)



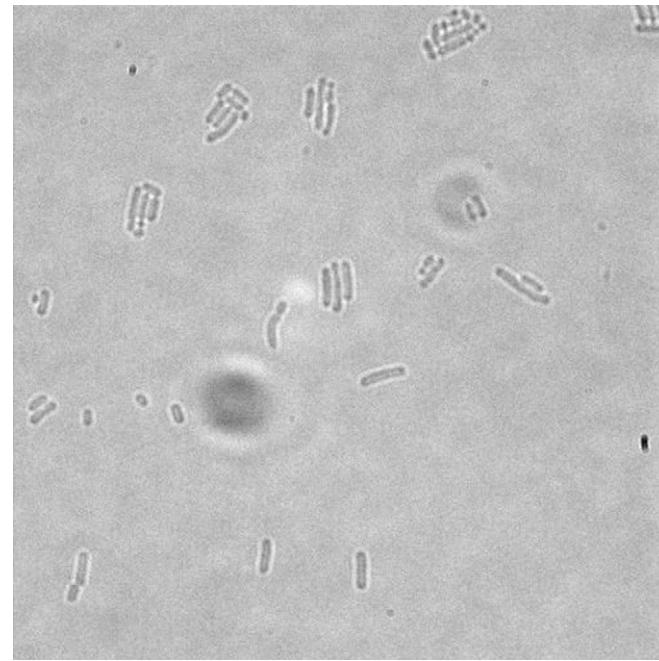
Movie S2. Corresponding DIC images for [Movie S1](#).

[Movie S2](#)



Movie S3. Time-lapse movies of $\Delta minD$ cells with labeled chromosomes (TetR-EYFP). Individual frames are separated by 1 h. A single z-plane is shown.

[Movie S3](#)



Movie S4. Corresponding DIC images for [Movie S3](#).

[Movie S4](#)