

1 **The ParB homologs, Spo0J and Noc, together prevent premature midcell Z ring**
2 **assembly when the early stages of replication are blocked in *Bacillus subtilis*.**

3

4

SUPPLEMENTARY MATERIALS

5

6

7 Isabella V. Hajduk, Riti Mann, Christopher D. A. Rodrigues, Elizabeth J. Harry

8

9

10

11 ¹The ithree institute, University of Technology Sydney, PO Box 123 Broadway NSW 2007
12 Australia

13

14

15 **Running title:** ParB homologs block premature midcell Z ring assembly

16

17

18

19

20 For correspondence:

21

22 Elizabeth.Harry@uts.edu.au

23

24

25

26

27

28 The ithree institute

29 Faculty of Science

30 University of Technology Sydney

31 T. +61 (02) 9514 4173

32 M. +61 (0) 404 643 181

33 PO Box 123 Broadway NSW 2007 Australia

34

1 **Table S1: *Bacillus subtilis* strain list used in this study**

| Strain | Genotype | Source |
|--------|---|---------------------------------|
| SU5 | 168 <i>trpC2</i> | E. Nester |
| SU492 | 168 <i>trpC2 amyE::(spc P_{xyl}-ftsZ-yfp)</i> | Lab stock |
| SU504 | 168 <i>trpC2 amyE::Pspachy-ftsZ (cat)</i> | Rodrigues & Harry, 2012 |
| SU533 | 168 <i>trpC2 Δnoc::cat</i> | Lab stock |
| SU629 | 168 <i>trpC2 Δnoc::tet amyE::(spc P_{xyl}-noc-yfp)</i> | Lab stock |
| SU656 | 168 <i>trpC2 Δnoc::tet</i> | Lab stock |
| SU661 | 168 <i>trpC2⁺ dna-1</i> | Rodrigues & Harry, 2012 |
| SU682 | 168 <i>trpC2 amyE::(spc P_{xyl}-ftsZ-yfp) ΔminCD::cat</i> | Lab stock |
| SU746 | 168 <i>trpC2⁺ dna-1 amyE::(spc P_{xyl}-ftsZ-yfp)</i> | This work |
| SU747 | <i>trpC2 Δsoj::neo</i> | Scholefield <i>et al</i> , 2011 |
| SU748 | <i>trpC2 Δspo0J::neo, amyE::spo0J (cat)</i> | Gruber & Errington, 2009 |
| SU765 | 168 <i>trpC2 Δsoj-spo0J::tet</i> | This work |
| SU766 | 168 <i>trpC2⁺ dna-1 Δsoj-spo0J::tet</i> | This work |
| SU767 | 168 <i>trpC2 amyE::(spc P_{xyl}-ftsZ-yfp) Δsoj-spo0J::tet</i> | This work |
| SU768 | 168 <i>trpC2⁺ dna-1 amyE::(spc P_{xyl}-ftsZ-yfp) Δsoj-spo0J::tet</i> | This work |
| SU769 | 168 <i>trpC2 Δspo0J::kan</i> | This work |
| SU770 | 168 <i>trpC2⁺ dna-1 Δspo0J::kan</i> | This work |
| SU771 | 168 <i>trpC2 Δsoj::kan</i> | This work |
| SU772 | 168 <i>trpC2⁺ dna-1 Δsoj::kan</i> | This work |
| SU802 | 168 <i>trpC2⁺ dna-1 Δnoc::cat</i> | This work |
| SU803 | 168 <i>trpC2 Δnoc::cat Δsoj-spo0J::tet</i> | This work |
| SU804 | 168 <i>trpC2⁺ dna-1 Δnoc::cat Δsoj-spo0J::tet</i> | This work |
| SU823 | 168 <i>trpC2 Δsoj-spo0J::tet hutM(345°)::lacO(cat), thrC (283°)::lacI-cfp(erm) amyE::(spc P_{xyl}-ftsZ-yfp)</i> | This work |
| SU824 | 168 <i>trpC2⁺ dna-1 Δsoj-spo0J::tet hutM(345°)::lacO(cat), thrC (283°)::lacI-cfp(erm) amyE::(spc P_{xyl}-ftsZ-yfp)</i> | This work |
| SU827 | 168 <i>trpC2 Δnoc::cat Δsoj::kan</i> | This work |
| SU828 | 168 <i>trpC2⁺ dna-1 Δnoc::cat Δsoj::kan</i> | This work |
| SU829 | 168 <i>trpC2 Δnoc::cat Δspo0J::kan</i> | This work |
| SU830 | 168 <i>trpC2⁺ dna-1 Δnoc::cat Δspo0J::kan</i> | This work |
| SU831 | 168 <i>trpC2 Δnoc::cat amyE::(spc P_{xyl}-noc-yfp)</i> | This work |
| SU832 | 168 <i>trpC2⁺ dna-1 Δnoc::cat amyE::(spc P_{xyl}-noc-yfp)</i> | This work |
| SU833 | 168 <i>trpC2 Δnoc::cat Δspo0J::kan amyE::(spc P_{xyl}-noc-yfp)</i> | This work |
| SU834 | 168 <i>trpC2⁺ dna-1 Δnoc::cat Δspo0J::kan amyE::(spc P_{xyl}-noc-yfp)</i> | This work |
| SU835 | 168 <i>trpC2 Δnoc::cat Δsoj-spo0J::tet amyE::(spc P_{xyl}-ftsZ-yfp)</i> | This work |
| SU836 | 168 <i>trpC2⁺ dna-1 Δnoc::cat Δsoj-spo0J::tet amyE::(spc P_{xyl}-ftsZ-yfp)</i> | This work |
| SU849 | <i>smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | Wang <i>et al.</i> 2014a |
| SU850 | 168 <i>trpC2 smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | This work |
| SU851 | 168 <i>trpC2⁺ dna-1 smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | This work |
| SU874 | 168 <i>trpC2 Δnoc::cat, smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | This work |
| SU875 | 168 <i>trpC2⁺ dna-1 Δnoc::cat, smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | This work |
| SU876 | 168 <i>trpC2 Δsoj-spo0J::tet, smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | This work |
| SU877 | 168 <i>trpC2⁺ dna-1 Δsoj-spo0J::tet, smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | This work |
| SU878 | 168 <i>trpC2 Δnoc::cat, Δsoj-spo0J::tet, smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | This work |
| SU879 | 168 <i>trpC2⁺ dna-1 Δnoc::cat, Δsoj-spo0J::tet, smc-ssrA loxP (kan), lacA::PxylA Ec sspB loxP (erm)</i> | This work |
| SU887 | 168 <i>trpC2 amyE::Pspachy-ftsZ (cat) Δspo0J::kan</i> | This work |
| SU888 | 168 <i>trpC2 amyE::Pspachy-ftsZ (cat) Δnoc::tet</i> | This work |
| SU889 | 168 <i>trpC2 amyE::Pspachy-ftsZ (cat) Δspo0J::kan Δnoc::tet</i> | This work |
| SU890 | 168 <i>trpC2 amyE::(spc P_{xyl}-ftsZ-yfp) Δspo0J::kan</i> | This work |
| SU891 | 168 <i>trpC2 amyE::(spc P_{xyl}-ftsZ-yfp) Δnoc::cat</i> | This work |
| SU892 | 168 <i>trpC2 amyE::(spc P_{xyl}-ftsZ-yfp) Δspo0J::kan Δnoc::cat</i> | This work |
| SU893 | 168 <i>trpC2 amyE::(spc P_{xyl}-ftsZ-yfp) Δspo0J::kan ΔminCD::cat</i> | This work |
| SU894 | <i>yuxG(-87°)::lacO48 (phleo), yhdG(+87°)::tetO48 (cat), ycgO::P_{ftsW} tetR-cfp (spec) terminators P_{ftsW} lacI-mypet</i> | Wang <i>et al.</i> 2014a |
| SU895 | 168 <i>trpC2⁺ dna-1 yuxG(-87°)::lacO48 (phleo), yhdG(+87°)::tetO48 (cat), ycgO::P_{ftsW} tetR-cfp (spec) terminators P_{ftsW} lacI-mypet</i> | This work |
| SU897 | 168 <i>trpC2⁺ dna-1 soj-spo0J::tet, yuxG(-87°)::lacO48 (phleo), yhdG(+87°)::tetO48 (cat), ycgO::P_{ftsW} tetR-cfp (spec) terminators P_{ftsW} lacI-mypet</i> | This work |
| SU899 | 168 <i>trpC2⁺ dna-1 Δspo0J::kan, ycgO::spo0J (erm)</i> | This work |
| SU901 | 168 <i>trpC2⁺ dna-1 Δnoc::cat Δspo0J::kan, ycgO::spo0J (erm)</i> | This work |
| SU903 | 168 <i>trpC2⁺ dna-1 Δsoj-spo0J::tet, ycgO::spo0J (erm)</i> | This work |
| SU905 | 168 <i>trpC2⁺ dna-1 Δnoc::cat Δsoj-spo0J::tet, ycgO::spo0J (erm)</i> | This work |

1 **Table S2: Plasmids and oligonucleotides used in this study**

2

| Plasmid | Description | Source |
|------------------|--|-----------|
| pKM084 | <i>ycgO::cat</i> | D. Rudner |
| pIH003 | <i>ycgO::Psoj-optRBS-spo0J (erm)</i> | This work |
| Oligonucleotides | Sequence ^a | Source |
| olH004 | cgc GAATTC AAACCATTTTCTCACCATCCTG | This work |
| olH005 | cgc GCTAGC CTTTCACATGAACATGTAATC | This work |
| olH008 | cgc GCTAGC ACATAAGGAGGAACTACTATGGCtaaAGGCCTTGAAAAAGGGAT | This work |
| olH0010 | gcg GGATCC TTATGATTCTCGTTCAGACA | This work |

3 ^a Bold letters indicate the recognition sequence for restriction enzymes

4

5 **Plasmid construction**

6

7 **pIH003** [*ycgO::Psoj-optRBS-spo0J (erm)*] was generated by a two-way ligation of a
 8 EcoRI-NheI PCR product containing the promoter region for the *soj-spo0J* operon
 9 (amplified with oligonucleotide primers olH004 and olH005), and a NheI-BamHI PCR
 10 product containing the *spo0J* gene (amplified with oligonucleotide primers olH008 and
 11 olH0010) into pKM084 cut with EcoRI-BamHI. Both PCR products were amplified from
 12 SU5 genomic DNA as the template. pKM084 is an ectopic integration vector for
 13 recombination into the *ycgO* locus (David Rudner).

14

15 **SUPPLEMENTARY FIGURE LEGENDS**

16

17 **Figure S1: Z ring positioning when initiation of DNA replication is blocked in**
 18 **vegetatively grown fixed cells.** (A-D) Vegetative cells were grown in PAB to the mid-
 19 exponential phase at the permissive temperature (34°C), then shifted to the non-
 20 permissive temperature (48°C) for a further 60 min: (A) wild-type; SU5, (B) *dna-1*;
 21 SU661, (C) Δ *soj-spo0J*; SU765 and (D) *dna-1* Δ *soj-spo0J*; SU766. Percentages
 22 shown are the frequencies of Z rings occurring at midcell in the range of 0.45 – 0.5 on
 23 the x-axis.

24 **Figure S2: Z ring positioning when initiation of DNA replication is blocked in**
 25 **vegetatively grown fixed cells of individual *soj* or *spo0J* mutants.** (A-F)
 26 Vegetative cells were grown in PAB to the mid-exponential phase at the permissive
 27 temperature (34°C), then shifted to the non-permissive temperature (48°C) for a
 28 further 60 min: (i) wild-type; SU5, (ii) *dna-1*; SU661, (iii) Δ *soj*; SU769, (iv) *dna-1* Δ *soj*;
 29 SU770, (v) Δ *spo0J*; SU771 and, (vi) *dna-1* Δ *spo0J*; SU772. Percentages shown are
 30 the frequencies of Z rings occurring at midcell in the range of 0.45 – 0.5 on the x-axis.

1 **Figure S3: Z ring positioning in outgrown spore cells at the permissive**
2 **temperature.** Z ring positioning was examined in outgrown spores at the permissive
3 temperature (34°C) for 120 min in strains: (A) wild-type; SU492, (B) *dna-1*; SU746, (C)
4 *Δsoj-spo0J*; SU767 and (D) *dna-1 Δsoj-spo0J*; SU768. Percentages shown are the
5 frequencies of Z rings occurring at midcell in the range of 0.45 – 0.5 on the x-axis.

6 **Figure S4: Whole field of view of Z rings overlayed with DAPI and phase contrast**
7 **when initiation of DNA replication is blocked in live outgrown spores.**
8 Microscopic images of an overlay of phase contrast, DAPI (red) and FtsZ-YFP (green)
9 in (A) *dna-1*; SU746, and (B) *dna-1 Δsoj-spo0J*; SU768. Right side images show closer
10 details of selected cells with different nucleoid morphologies and varying Z ring
11 positions. Scale bar represents 2 μm.

12 **Figure S5: Flow cytometry profiles of *Δsoj-spo0J* strains when initiation of DNA**
13 **replication is blocked via *dna-1* mutant or addition of HPUra.** Flow cytometry
14 profiles in controls where initiation of DNA replication is progressing normally (wild-
15 type; SU492 and *Δsoj-spo0J*; SU767) or blocked (controls: *dna-1*; SU746 and wild-
16 type +HPUra; and test strains: *dna-1 Δsoj-spo0J*; SU768 and *Δsoj-spo0J* +HPUra. n
17 = 10,000.

18 **Figure S6: Noc localisation in the absence of *spo0J* when initiation of DNA**
19 **replication is blocked.** Strains (A) wild type; SU831, (B) *dna-1*; SU832, (C) *Δspo0J*
20 *Δnoc*; SU833, and (D) *dna-1 Δspo0J Δnoc*; SU834, possessing Noc tagged with a
21 yellow fluorescent protein (*noc-yfp*; falsely coloured cyan) were grown vegetatively
22 supplemented with 0.3% xylose. DNA stained with DAPI (falsely coloured red). Scale
23 bar = 2 μm.

24 **Figure S7: Z ring positioning when initiation of DNA replication is blocked**
25 **during spore outgrowth.** Z ring positioning was examined in two conditions: in the
26 temperature-sensitive *dna-1* background (A) and with the addition of the DNA
27 polymerase III inhibitor HPUra (B), in spore outgrown spore cells. A: (i) wild-type;
28 SU492, (ii) *dna-1*; SU746, (iii) *Δnoc*; SU831, (iv) *dna-1 Δnoc*; SU832, (v) *Δsoj-spo0J*
29 *Δnoc*; SU835, and (vi) *dna-1 Δsoj-spo0J Δnoc*; SU836; B: (i) wild-type +HPUra and
30 (ii) *Δsoj-spo0J Δnoc* +HPUra.

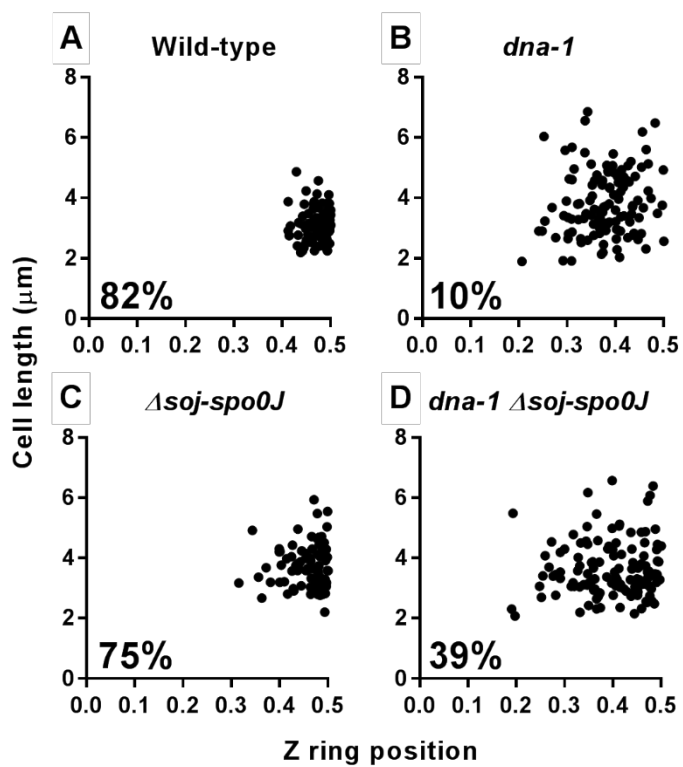
31 **Figure S8: Z ring positioning in the individual mutants, *soj* or *spo0J*, in the *dna-***
32 **1 *Δnoc* mutant at the non-permissive temperature.** Scatter plots showing Z ring

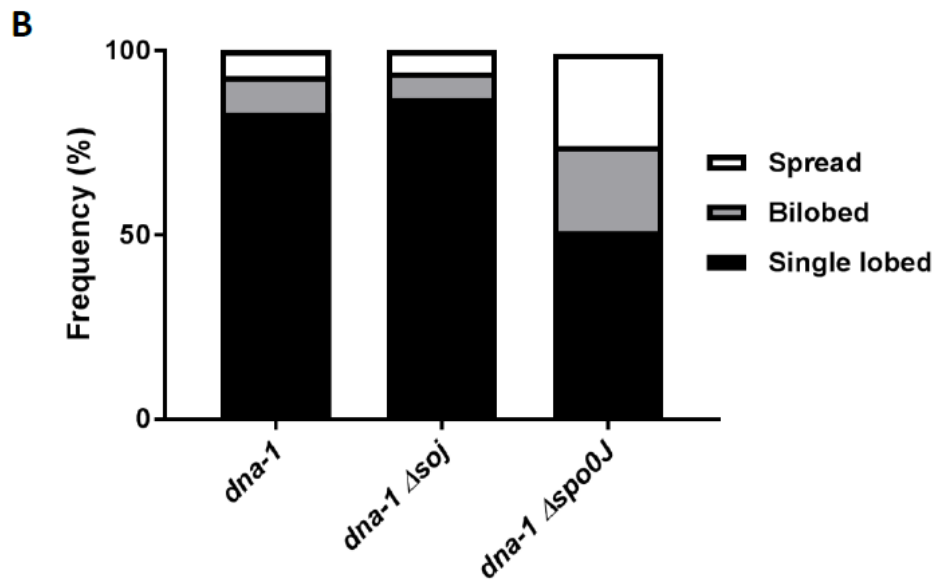
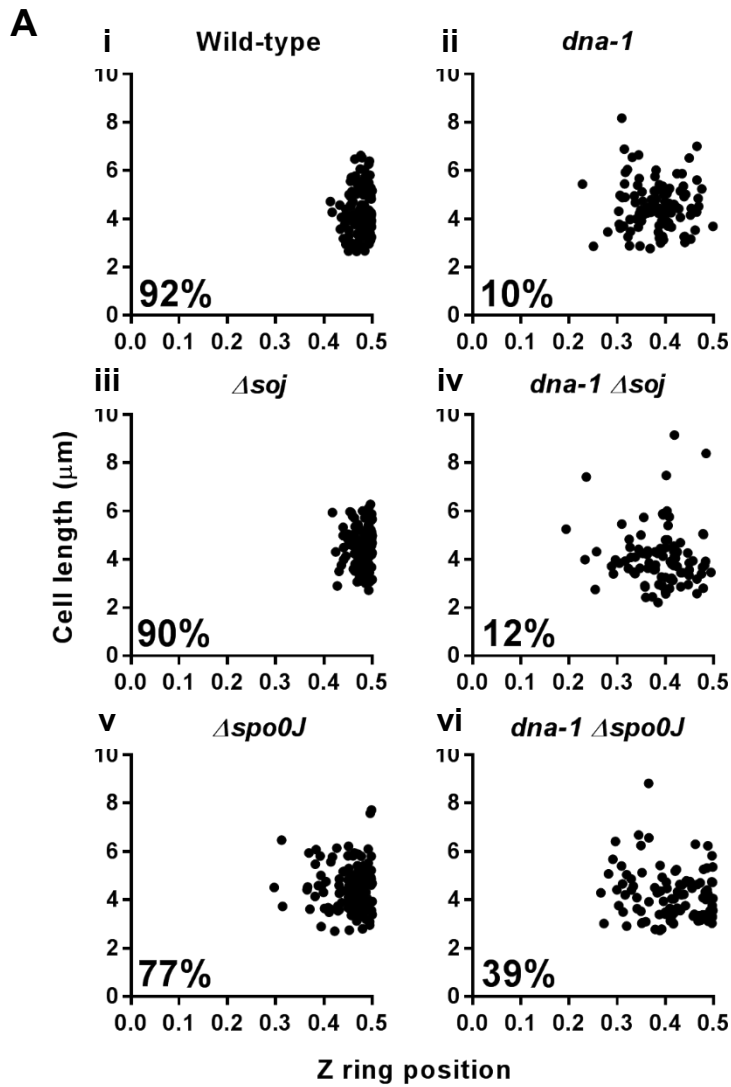
1 positioning and average cell length of vegetatively grown, fixed cells of (A) wild-type;
2 SU5, (B) *dna-1*; SU661, (C) Δ *soj* Δ *noc*; SU829, (D) *dna-1* Δ *soj* Δ *noc*; SU830, (E)
3 Δ *spo0J* Δ *noc*; SU827, and (F) *dna-1* Δ *spo0J* Δ *noc*; SU828.

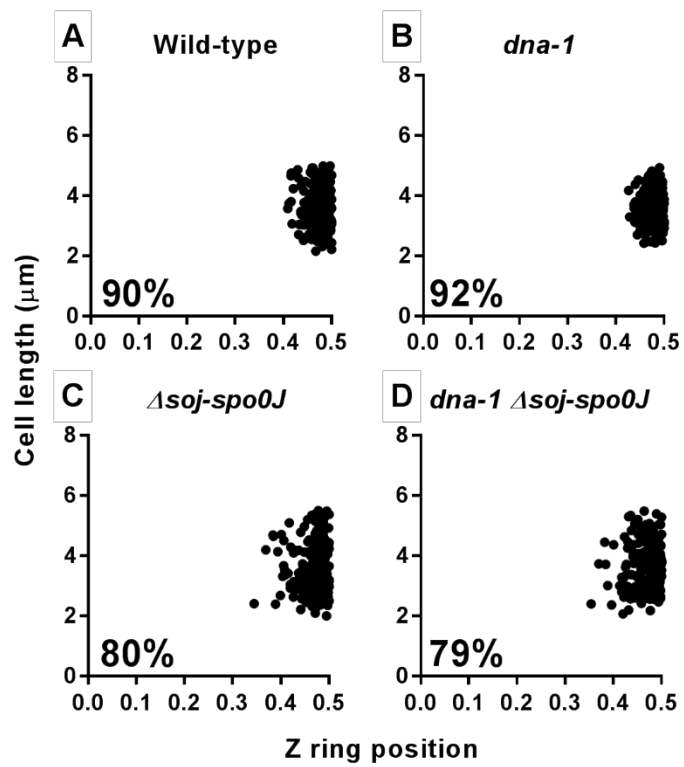
4 **Fig S9. Complementation of *spo0J* in the temperature sensitive strains.** (A) Z ring
5 positioning, and (B) nucleoid morphologies in strains (i) *dna-1 spo0J ycg::spo0J*
6 (SU899), (ii) *dna-1 spo0J noc ycg::spo0J* (SU901), (iii) *dna-1 soj-spo0J ycg::spo0J*
7 (SU903), and (iv) *dna-1 soj-spo0J noc ycg::spo0J* (SU905), grown vegetatively at the
8 non-permissive temperature and examined via IFM and ethanol fixation, respectively.
9 n = 200.

10 **Figure S10: Z ring positioning when initiation of DNA replication is blocked**
11 **during spore outgrowth in the *dna-1* temperature-sensitive SMC-depleted**
12 **mutant.** Scatter plots showing Z ring positioning and average cell length of outgrown
13 spores grown in the presence of xylose (1% v/v) to induce SMC degradation in
14 replicating cells (left column) and non-replicating cells (*dna-1* mutant; right column). Z
15 ring positioning was examined in strains (A) wild-type; SU850, (B) *dna-1*; SU851, (C)
16 Δ *noc*; SU874, (D) *dna-1* Δ *noc*; SU875, (E) Δ *soj-spo0J*; SU876, (F) *dna-1* Δ *soj-spo0J*;
17 SU877, (G) Δ *noc* Δ *soj-spo0J*; SU878, and (H) *dna-1* Δ *noc* Δ *soj-spo0J*; SU879.

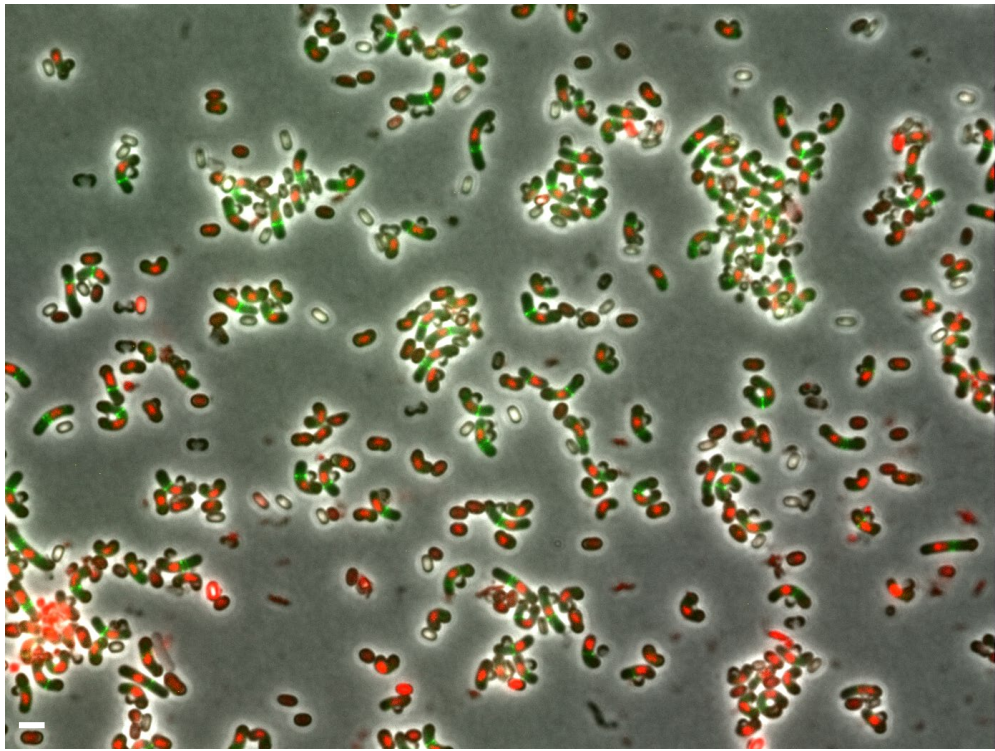
18 **Figure S11: Cell division is not inhibited in a *minCD spo0J* double mutant.** Phase
19 contrast images of mid-exponentially growing cells of strains (A) wild-type; SU492, (B)
20 Δ *spo0J*; SU890, (C) Δ *minCD*; SU682 and (D) Δ *spo0J* Δ *minCD*; SU893 grown at 30°C
21 and 37°C. Scale bar represents 2 μ m.



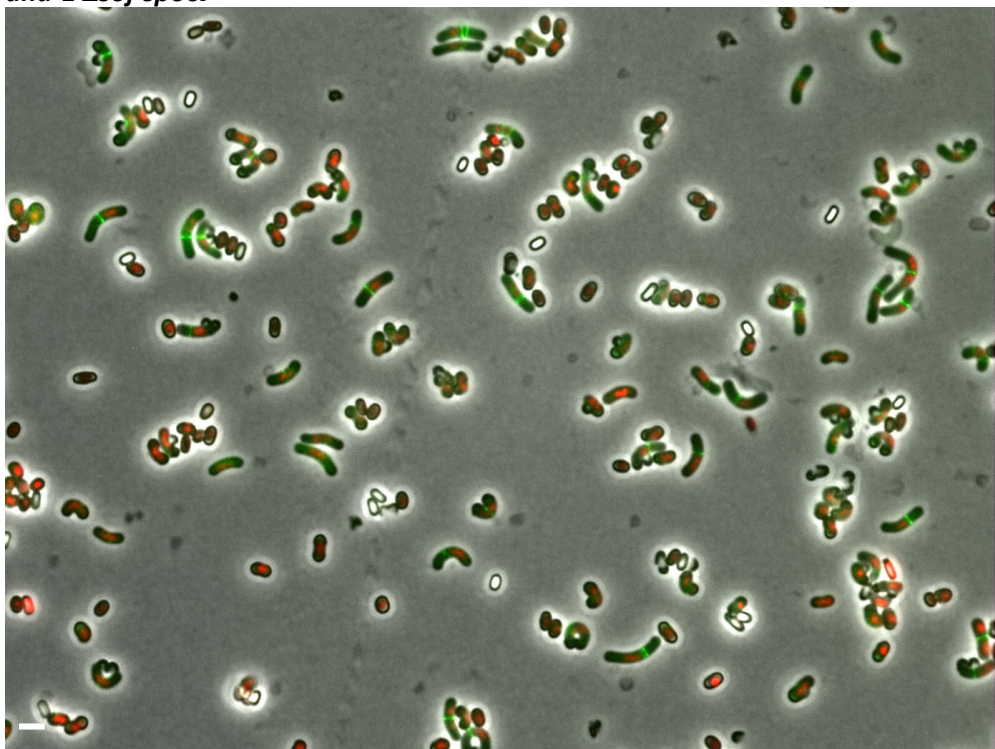


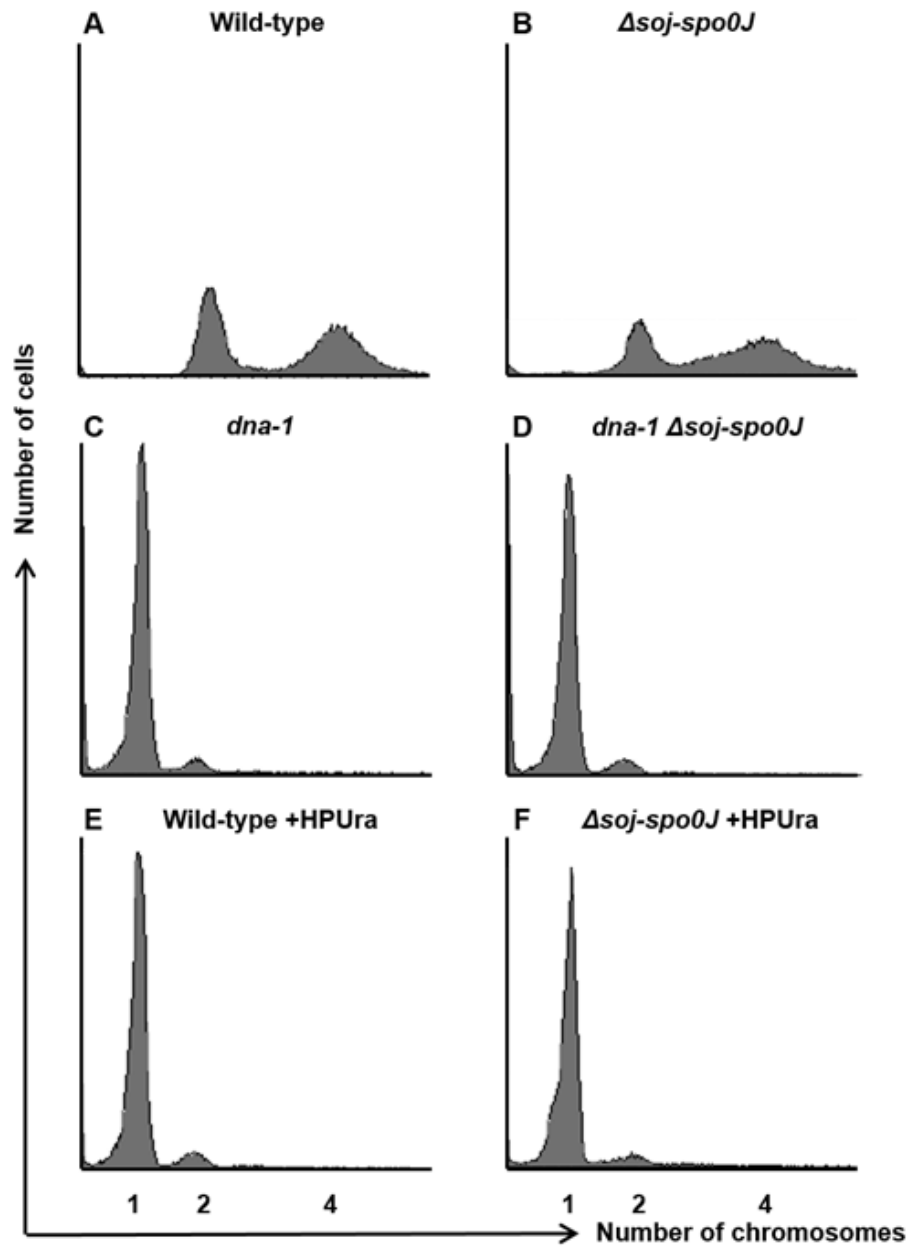


A *dna-1*

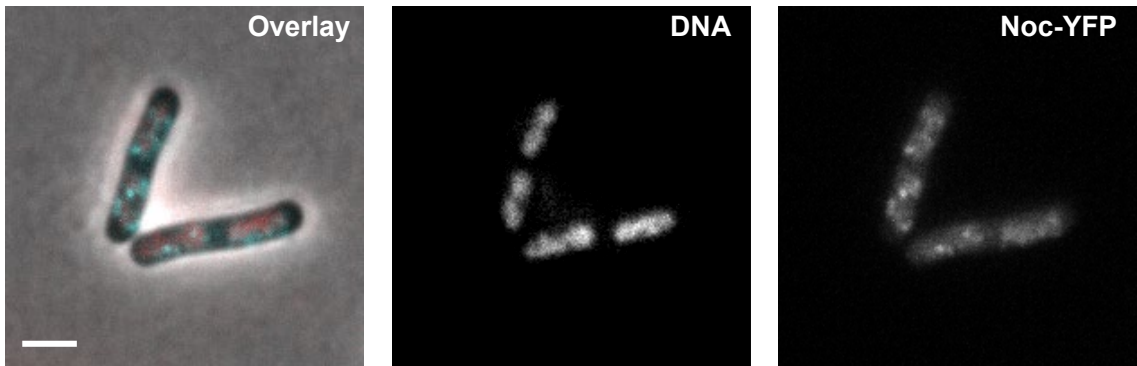


B *dna-1 Δsoj-spo0J*

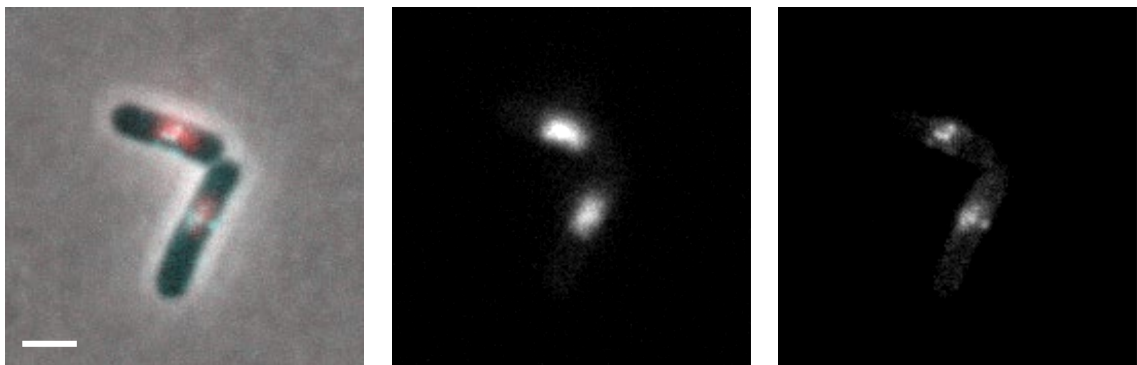




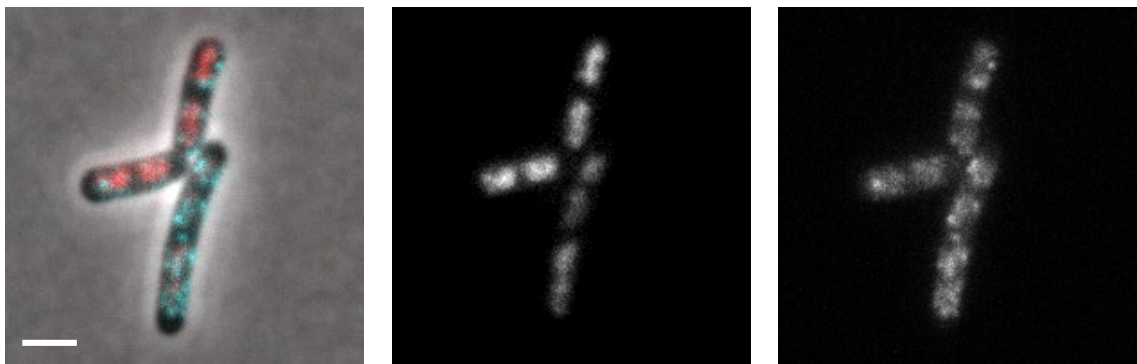
A Wild-type



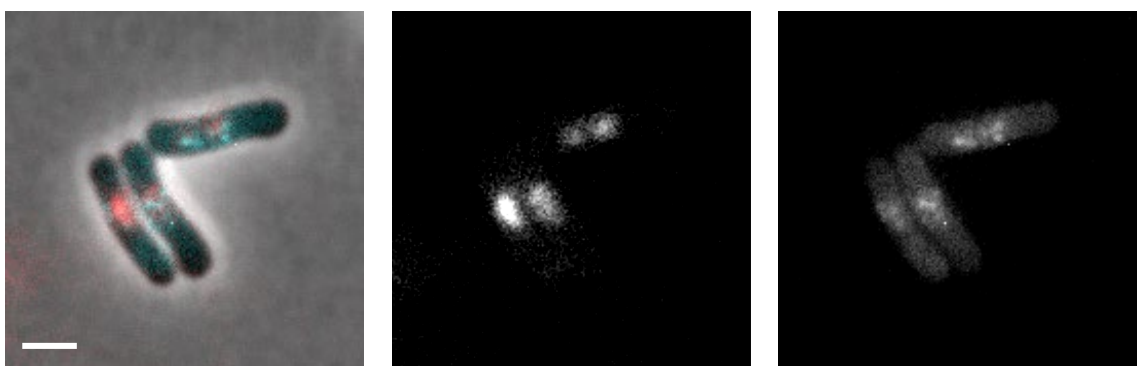
B *dna-1*

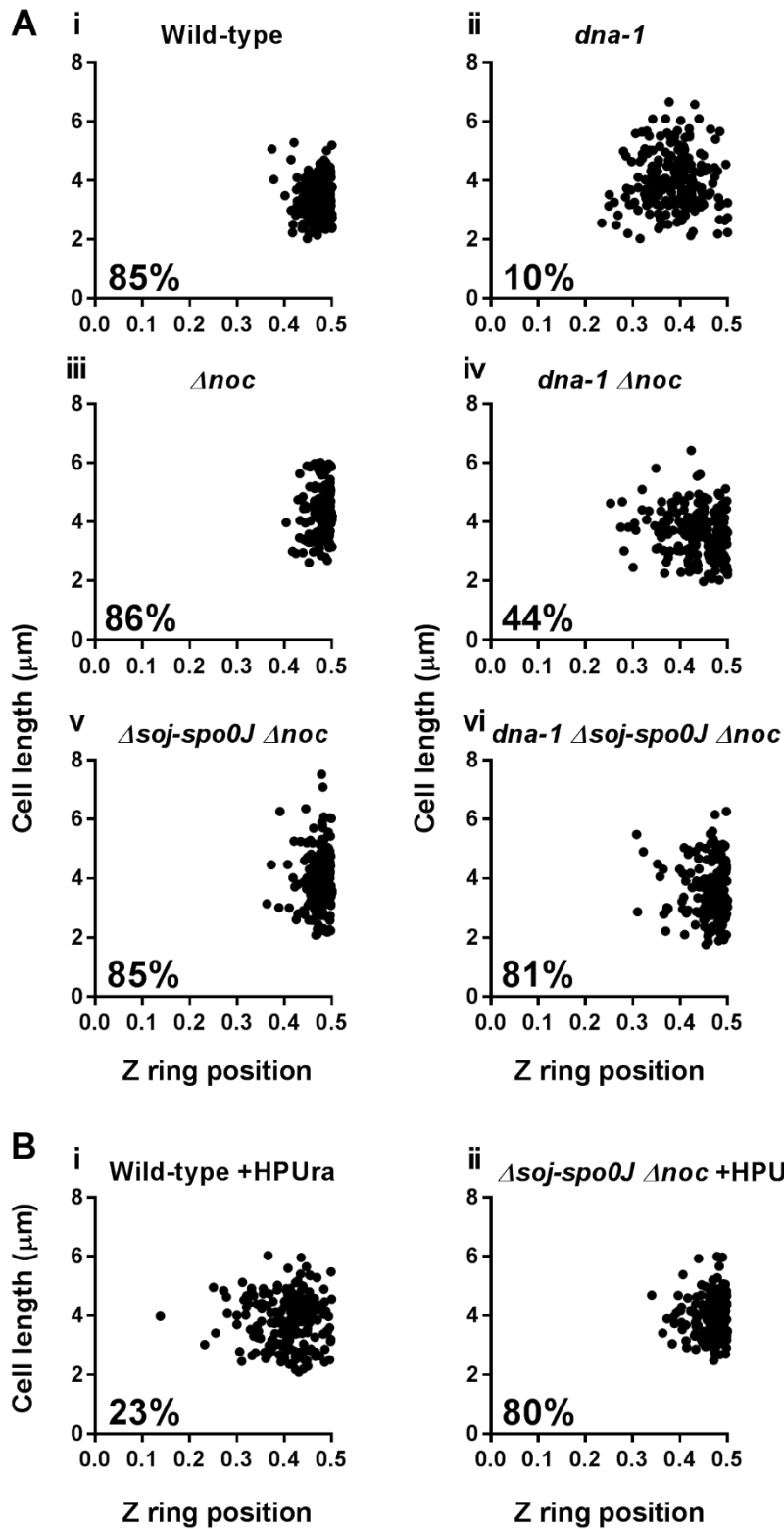


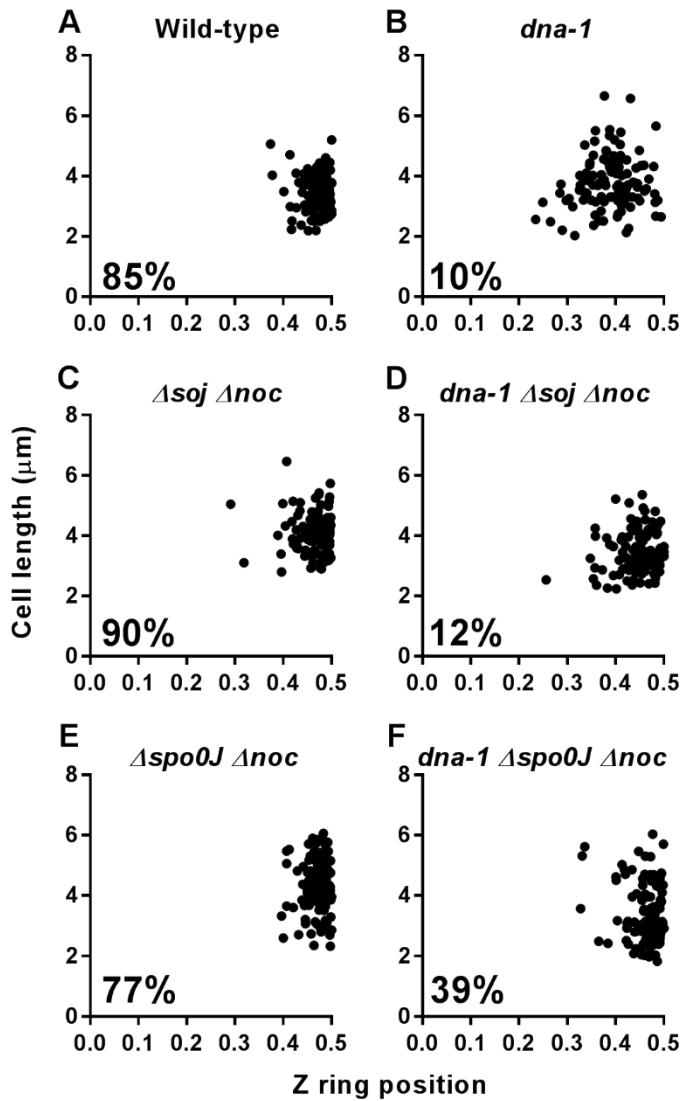
C $\Delta spo0J$

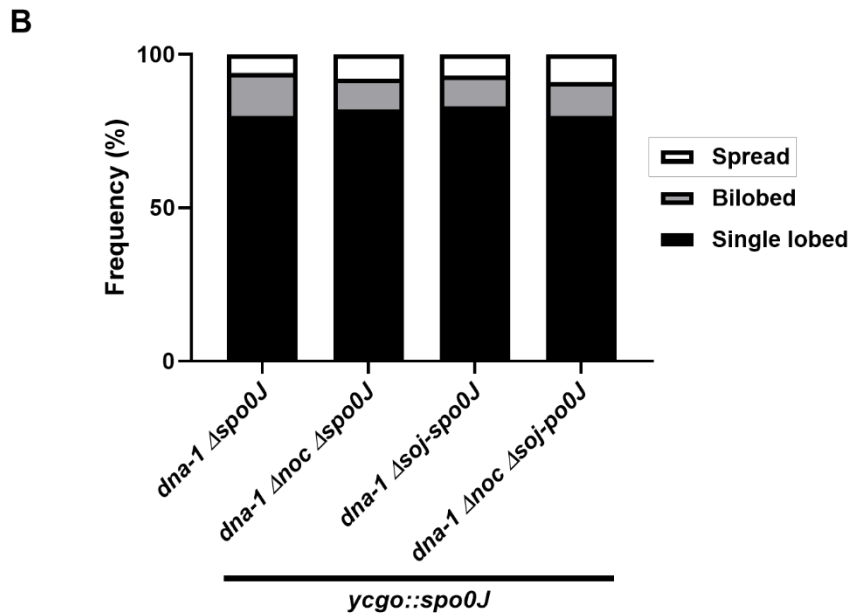
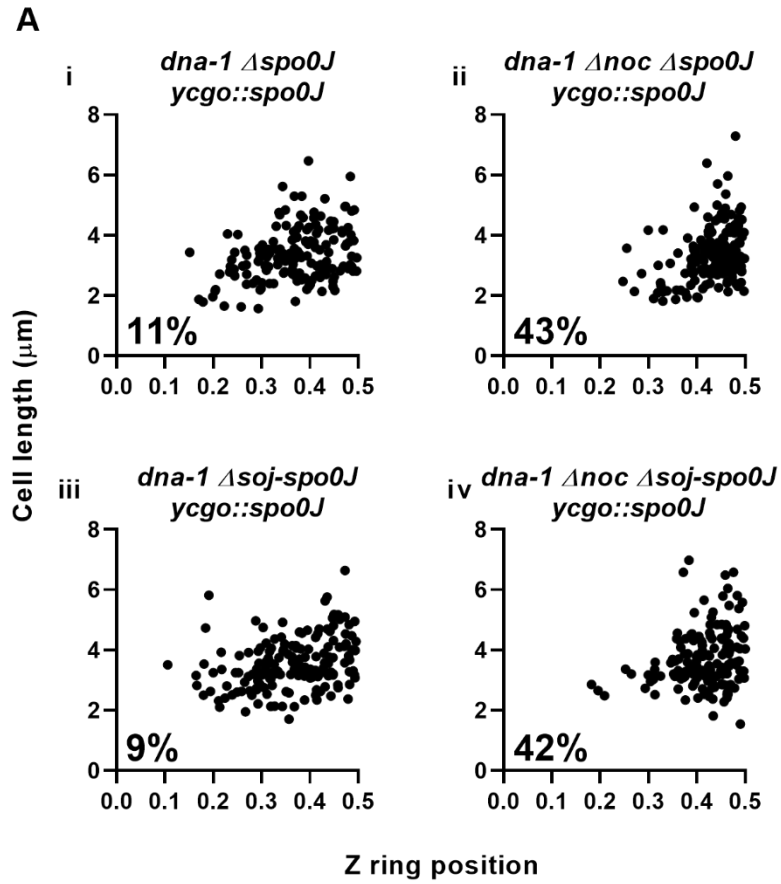


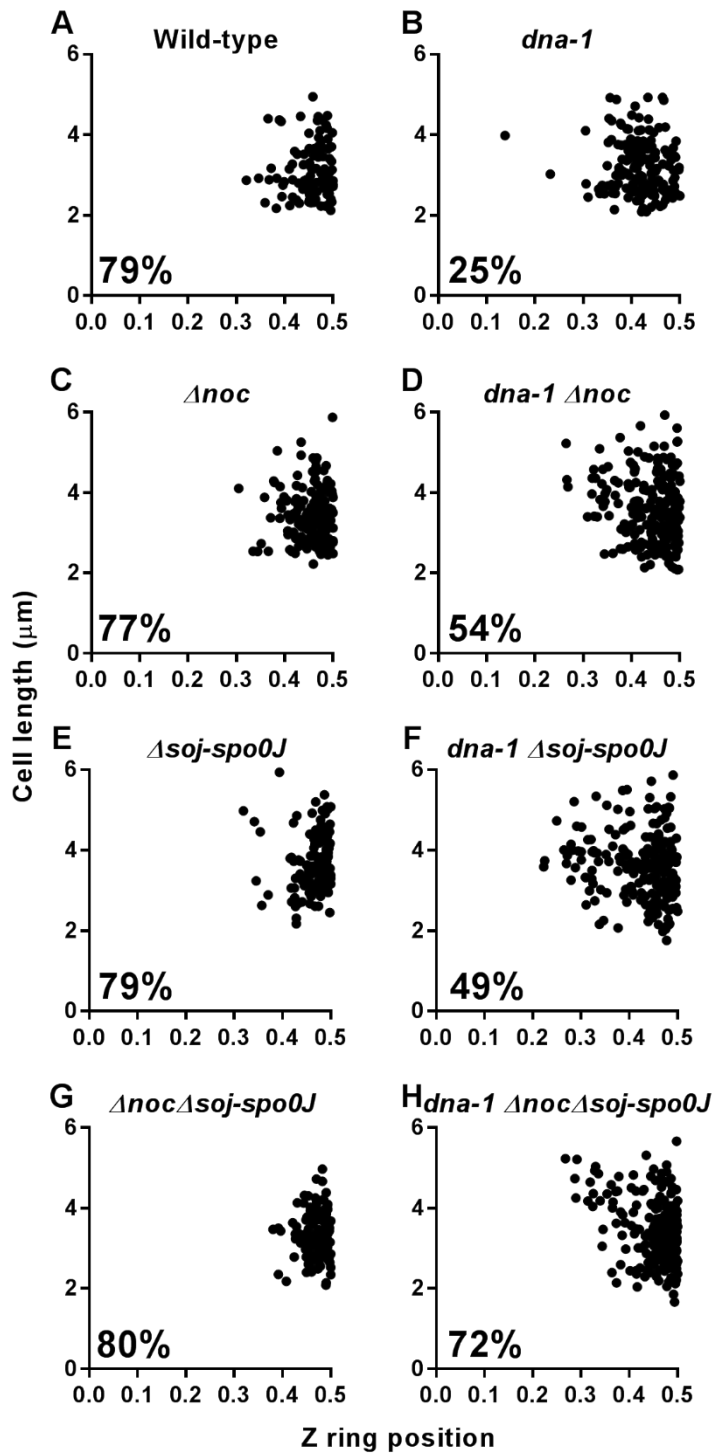
D *dna-1* $\Delta spo0J$



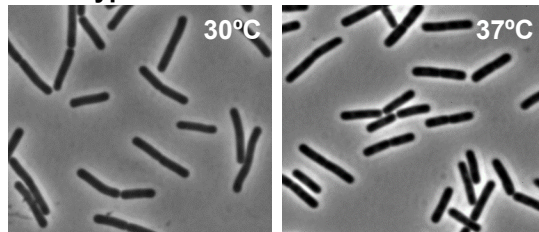




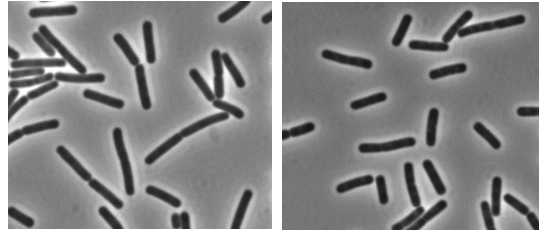




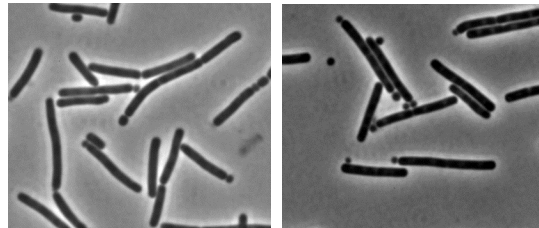
A Wild-type



B $\Delta spo0J$



C $\Delta minCD$



D $\Delta spo0J \Delta minCD$

