



Rhizobium induces DNA damage in *Caenorhabditis elegans* intestinal cells

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In their natural habitat of rotting fruit, the nematode *Caenorhabditis elegans* feeds on the complex bacterial communities that thrive in this rich growth medium. Hundreds of diverse bacterial strains cultured from such rotting fruit allow *C. elegans* growth and reproduction when tested individually. In screens for *C. elegans* responses to single bacterial strains associated with nematodes in fruit, we found that *Rhizobium* causes a genome instability phenotype; we observed abnormally long or fragmented intestinal nuclei due to aberrant nuclear division, or defective karyokinesis. The karyokinesis defects were restricted to intestinal cells and required close proximity between bacteria and the worm. A genetic screen for *C. elegans* mutations that cause the same intestinal karyokinesis defect followed by genome sequencing of the isolated mutant strains identified mutations that disrupt DNA damage repair pathways, suggesting that *Rhizobium* may cause DNA damage in *C. elegans* intestinal cells. We hypothesized that such DNA damage is caused by reactive oxygen species produced by *Rhizobium* and found that hydrogen peroxide added to benign *Escherichia coli* can cause the same intestinal karyokinesis defects in WT *C. elegans*. Supporting this model, free radical scavengers suppressed the *Rhizobium*-induced *C. elegans* DNA damage. Thus, *Rhizobium* may signal to eukaryotic hosts via reactive oxygen species, and the host may respond with DNA damage repair pathways.

C. elegans | microbiota | DNA damage | ROS | *Rhizobium*

The free-living nematode *Caenorhabditis elegans* in the laboratory feeds on *Escherichia coli*. Cultivation of *C. elegans* on a diet of *E. coli* was natural for the *E. coli* geneticists who were the pioneers of *C. elegans* research. But the native habitat of *C. elegans* is not the laboratory, and, until recently, very little was known about its natural history (1–6). For many years, *C. elegans* has been called a soil nematode, which now appears to be inaccurate. In geographically diverse areas, *C. elegans* inhabits decaying plants and fruits, which fuel the proliferation of diverse species of bacteria which *C. elegans* finds scrumptious (5, 7). Félix and coworkers have systematically characterized the bacterial species that can be cultured from the same decaying fruits that foster the growth of *C. elegans* and other nematodes (5, 7). The potential influence on *C. elegans* of these bacterial species, either individually or in reconstructed communities of bacteria, has begun to emerge (8).

Here, we show that particular species of bacteria from the *C. elegans* natural habitat tested individually as the sole food source for *C. elegans* cultivation: For example, *Rhizobium huautlense*, inhibited normal karyokinesis during postembryonic intestinal development. Our analysis based on comprehensive *C. elegans* genetics screens, and free radical exposure or free radical scavenger treatments, show that *Rhizobium* produces reactive oxygen species (ROS), which induces DNA damage in the animal intestine to in turn cause defective karyokinesis. This ROS damage is likely to cause the formation of chromosome bridges preventing the proper separation of nuclei after a cell cycle. Chemical free radical scavengers can partially suppress this *Rhizobium*-induced DNA damage. Chemical addition of reactive

oxygen to an otherwise benign *E. coli* bacteria also induced such an aberrant nuclear division. Our findings provide an insight into the interactions between microbes and animal cells.

Results and Discussion

Feeding *C. elegans* on Particular Strains of Bacteria Cultured from Its Natural Habitat Causes Abnormal Intestinal Nuclei Divisions. In the laboratory, *C. elegans* is often cultivated on a diet of *E. coli* (OP50 strain) bacteria grown from a few drops of a saturated liquid overnight culture, which is spotted on nematode growth media agar plates (NGM-plates). We cultivated *C. elegans* on individual bacterial species isolated from the *C. elegans* natural habitat of rotting fruit (3, 4, 9) instead of *E. coli* and tested whether any of these strains induce the expression of GFP fusion reporter genes associated with particular developmental milestones or with stress and immunity (8). We tested these reporter genes on a cultured collection of 90 species that cohabit rotting fruits with *C. elegans* (the JUB bacterial library) (8) (*Materials and Methods*), which have been 16s rRNA gene sequenced for classification. A *sur-5::GFP* fusion gene (10), which expresses a nuclear-localized green fluorescent protein (GFP) in all somatic cells, allows many postembryonic cell divisions that occur during larval stages of *C. elegans* to be monitored. In *C. elegans* carrying the *sur-5::GFP* fusion cultivated on benign *E. coli*, bright GFP is observed in the intestine, where 20 large round intestinal nuclei line up in a regular pattern. The number and spatial distribution of these nuclei in larvae and adults is stereotyped. At the first larval (or L1) stage, larvae hatch with 20 mononucleated

Significance

An incredible diversity of microbial species inhabit the Earth. Many new interactions of bacteria with animals are only beginning to surface. We show that feeding a worm, *Caenorhabditis elegans*, with *Rhizobium* bacteria causes failed nuclear divisions in intestinal cells. *C. elegans* mutations compromising DNA damage repair pathways and exogenous reactive oxygen species cause similar failure in intestinal nuclear divisions, suggesting that *Rhizobium* may produce reactive oxygen species. In accordance, free radical scavenger chemicals suppress the response to *Rhizobium*. We propose that *Rhizobium* induces reactive oxygen species in its encounter with *C. elegans*. Our findings provide an insight into the interactions between microbes and animal cells.

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Table 1. Mutations in the listed genes alter karyokinesis in the *C. elegans* developing intestine

| Gene | Name | Cellular function | EMS mutation/ alleles | Amino acid change | CGC strains | C-test* | CGC strain gut nuclei abnormal | Human homolog |
|--------------|---|----------------------------|---|--|---------------|-----------------|-----------------------------------|------------------------|
| <i>atm-1</i> | Ataxia telangiectasia mutated family | DDR | C578T | R/C | VC381 (gk186) | No | Yes | ATM |
| <i>dog-1</i> | Deletion of G-rich DNA | DNA repair/ maintenance | G118A [†] C15021247T position on chromosome I G759A G845A C346T | G/R Splice site donor C/Y G/D P/S | VC13 (gk10) | No No n/t | Yes | FANCI/ BRIP1/ BACH1 |

*C-test, complementation test. "No" means that a cross between the EMS mutant and the corresponding CGC strain does not rescue the abnormal phenotype, indicating that the same gene is mutated in both strains.

[†]Five alleles of *dog-1* were isolated in the screen; two were evaluated in a C-test. n/t, not tested. Amino acid abbreviations: R, arginine; C, cysteine; G, glycine; Y, tyrosine; D, aspartic acid; P, proline; S, serine. Sources: <https://wormbase.org/#012-34-5>.

redox status in living *Rhizobium* or *E. coli* cells (*Materials and Methods*). *Rhizobium* caused much more intense H₂DCFDA-AM fluorescence than *E. coli*, indicating the presence of more oxidizing metabolites and possibly a higher level of ROS (Fig. 5 A and B). One possible source of bacterial ROS is hydrogen peroxide (H₂O₂). We measured H₂O₂ concentrations in *E. coli* and *Rhizobium* using a fluorimetric hydrogen peroxide assay. The assay utilizes a substrate for hydrogen peroxidase, an enzyme which, in the presence of hydrogen peroxide, converts it into a fluorescent compound (*Materials and Methods*). The assay is applicable for the analysis of live cells. The measurement revealed that *Rhizobium* contains about five times more hydrogen peroxide than *E. coli* (Fig. 5 C and D).

Hydrogen peroxide is cytotoxic, but highly conserved catalase enzymes protect cells by metabolizing hydrogen peroxide to water and oxygen. A simple assay can detect catalase: A drop of 3% H₂O₂ onto a freshly grown *E. coli* lawn bubbles due to oxygen production by catalase (*Movies S1* and *S2*). A fresh *Rhizobium* lawn does not generate bubbles (*Movie S3*), indicating low or absent catalase activity. *C. elegans* L1 larvae exposed to 3% hydrogen peroxide on unseeded agar or on a *Rhizobium* lawn became paralyzed within minutes due to H₂O₂ toxicity. On *E. coli*, *C. elegans* larvae exposed to 3% hydrogen peroxide remained mobile. The catalase activity of *E. coli* may detoxify the H₂O₂ whereas the *Rhizobium* does not produce catalase and could not detoxify the H₂O₂ (Fig. 5 E–J). Treatment of *C. elegans* early L1 larvae with 3% hydrogen peroxide on an *E. coli* lawn induced the same defective intestinal karyokinesis as caused by *Rhizobium* feeding (Fig. 5 K and L and *SI Appendix, Table S1*): 63% of *C. elegans* fed *E. coli* and treated with 3% H₂O₂ showed defective karyokinesis.

Based on the facts that feeding on *Rhizobium*, but not *E. coli*, causes oxidative stress in *C. elegans*, that *Rhizobium* has higher cellular oxidative state and likely higher ROS than *E. coli*, that *Rhizobium* accumulate more hydrogen peroxide possibly because of low catalase activity, and that exogenous hydrogen peroxide causes intestinal karyokinesis defects similar to those induced by feeding with *Rhizobium*, we hypothesized that hydrogen peroxide produced by *Rhizobium* could be the oxidizing metabolite. *Rhizobium* could produce hydrogen peroxide in the *C. elegans* gut, the probable first *C. elegans* cells to encounter *Rhizobium* separated by only a cell membrane, as digestion ensues, causing, directly or indirectly, DNA damage in adjacent intestinal cells.

Hypoxia or Antioxidants Suppress the Karyokinesis Defect in *C. elegans* Grown on *Rhizobium*. If the oxidative stress model is correct, then karyokinesis defect caused by *Rhizobium* should depend on oxygen concentration. We found that hypoxia (1% oxygen) significantly reduced the defective gut karyokinesis phenotype in *C. elegans* grown on *Rhizobium* whereas high

oxygen (100% oxygen) enhanced the phenotype on *Rhizobium* (graph in Fig. 6, Fig. 6 A and B, and *SI Appendix, Table S1*). Moreover, 100% O₂ caused the defective gut karyokinesis phenotype in *C. elegans* fed on normally benign *E. coli* (Fig. 6C and

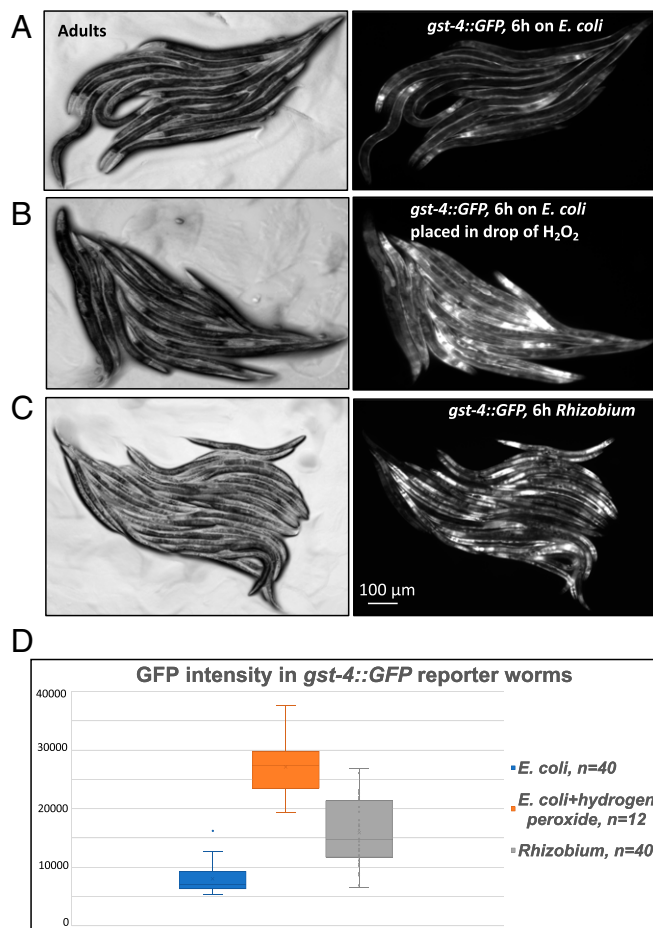


Fig. 4. Feeding on *Rhizobium* causes oxidative stress response. (A–C) Fluorescence and differential interference contrast *gst-4p::GFP* images of *C. elegans* adults transferred from an *E. coli* lawn to a new *E. coli* lawn (A), or transferred to a new *E. coli* lawn and treated with 30 μL of 3% H₂O₂ (B), or transferred to a *Rhizobium* lawn (C). Images were taken 6 h after the transfer of the *C. elegans*. (D) Graph showing the corresponding GFP intensities in the gut measured with ImageJ software. *t* test for *E. coli* vs. *E. coli* + hydrogen peroxide returns $P = 8.95503E-24$, and for *E. coli* vs. *Rhizobium*, $P = 1.29942E-12$.

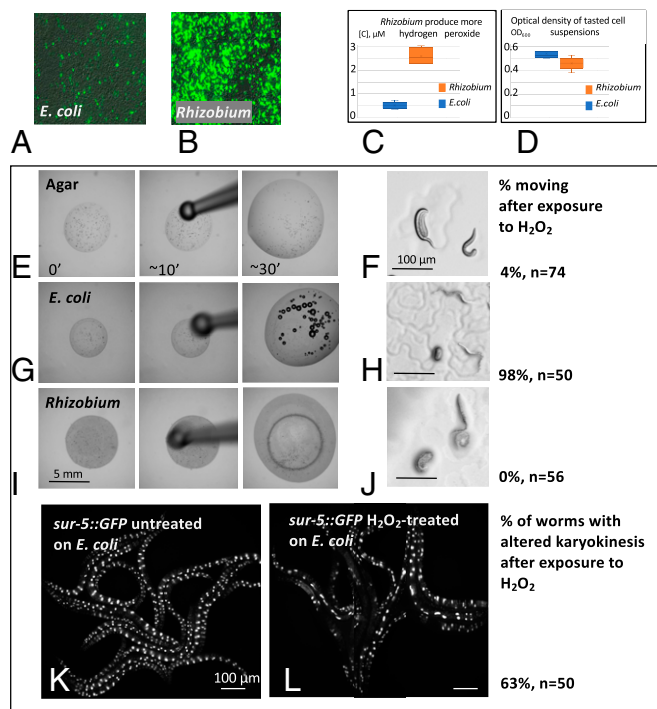


Fig. 5. *Rhizobium huautlense* cells are positive for oxidative stress, produce more hydrogen peroxide, and have less or no catalase activity. Hydrogen peroxide is sufficient to interfere with karyokinesis in the *C. elegans* gut. (A and B) A fluorescence test for oxidative stress indicates that *Rhizobium* has more redox activity (more cells have brighter fluorescence) than *E. coli* grown on NGM agar at room temperature. (C) A graph showing that *Rhizobium* produces more hydrogen peroxide than *E. coli* as measured by fluorimetric hydrogen peroxide assay. (D) Graph showing similar absorption (OD600) and therefore concentrations of *E. coli* and *Rhizobium* suspensions used in the assay. (E–J) Images of L1 larvae *C. elegans* treated with 3% hydrogen peroxide on agar (E and F), *E. coli* lawn (G and H), and *Rhizobium* lawn (I and J). Catalase activity of *E. coli* (G and H), but not *Rhizobium* (I and J) suppresses the paralysis of the L1. Numbers on the right of F, H, and J show percent of moving L1 larvae, and “n” is a number of scored larvae for each condition. (K and L) Fluorescence images of *sur-5::GFP* reporter *C. elegans* untreated (K) or treated with hydrogen peroxide (L) during L1 stage. Approximately 63% of treated animals developed the phenotype, $n = 50$.

SI Appendix, Table S1). These DNA damage responses to increased oxygen both on *E. coli* and *Rhizobium* are likely to be via increased production of ROS.

We tested if a treatment of L1 larvae with *N*-acetyl-cysteine (NAC), a potent antioxidant that reacts with hydrogen peroxide (33), could alleviate the *Rhizobium* gut karyokinesis defect. We treated L1 larvae with NAC by letting them hatch overnight in a solution of NAC. The treated and untreated *C. elegans* were plated on *Rhizobium* or *Rhizobium* supplemented with NAC, as well as on control *E. coli* lawns, and scored for the intestinal karyokinesis defect (*Materials and Methods*). Treatment of L1 larvae with NAC before exposure to *Rhizobium* suppressed the karyokinesis defect, but a supplementation of a *Rhizobium* lawn with NAC could not rescue the *Rhizobium*-induced karyokinesis defect in untreated larvae (*SI Appendix, Fig. S4A and Table S1*). The data indicate that NAC promotes a resistance to oxidative stress in larvae rather than reduced ROS levels in *Rhizobium*. These results support the hypothesis that bacterial-induced oxidative stress and ROS cause abnormal karyokinesis in intestinal nuclei.

In a related experiment, we found that growth of *C. elegans* on *Rhizobium* in the presence of an apple slice (*Materials and Methods*), likely to contain a natural antioxidant, also inhibited

the intestinal karyokinesis defect (*SI Appendix, Fig. S4B and Table S1*). A reduction of ROS by antioxidants in the apple or indirectly through a change in *Rhizobium* metabolism is possible. The suppression of ROS damage by the apple could explain why proliferating *C. elegans* are almost always found on fallen fruits in nature, but not in soil harboring similar bacterial strains (34, 35).

***Rhizobium* Induces *C. elegans* Intestinal Karyokinesis Defects over a Short Range.** Since there were no obvious changes in karyokinesis outside of the intestine (*SI Appendix, Fig. S5*), we hypothesized that ingested *Rhizobium* disrupts intestinal karyokinesis over a short range from the lumen of the gut to the gut nucleus. We tested this hypothesis by obstructing direct contact between *Rhizobium* and *C. elegans* with a 0.22- μ m-pore-size filter. The filter was placed on the *Rhizobium* lawn, and the *C. elegans* eggs were spotted on top of the filter along with a small aliquot of *E. coli* as a food to stimulate postembryonic development (36). A second filter prepared with a mixture of *E. coli* with *Rhizobium* without filter served as a control. We found that a presence of the filter between *Rhizobium* and the *C. elegans* inhibited the intestinal karyokinesis defects whereas control *C. elegans* in contact with *Rhizobium* showed the gut karyokinesis defect (*Materials and Methods* and *SI Appendix, Fig. S6*). These results suggest that the intestinal karyokinesis defect could be caused by *Rhizobium* cells contacting cells in the nematode. Such a short-range agent of DNA damage is consistent with hydrogen peroxide.

We also measured formation of 8-hydroxydeoxyguanosine (8-OHdG), one of the hallmarks of oxidative DNA damage, in chromosomal DNA isolated from *C. elegans* grown on *E. coli* and *Rhizobium* (*Materials and Methods*). We did not detect a significant difference in the concentration of 8-OHdG between the *Rhizobium*- or *E. coli*-grown *C. elegans* (*SI Appendix, Fig. S7*).

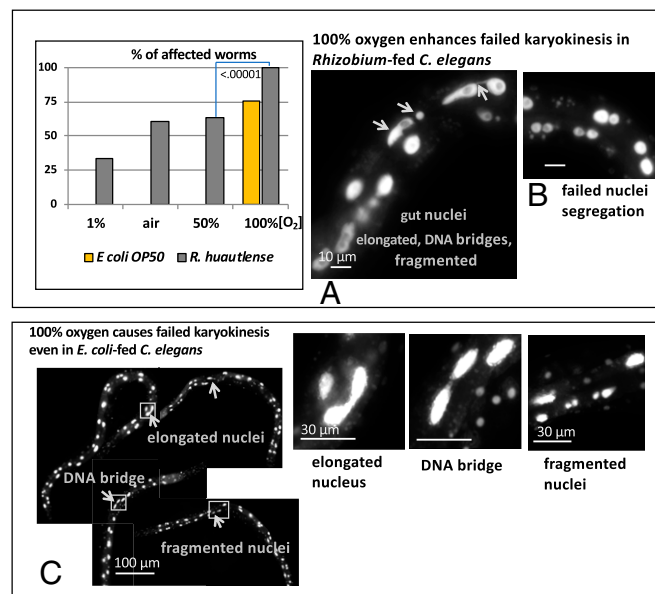


Fig. 6. Abnormal intestinal karyokinesis in *C. elegans* fed with *Rhizobium* depends on oxygen. (A) The graph shows the percentage of *C. elegans* with defective gut nuclear division when grown on *Rhizobium* from hatching in a chamber with indicated concentration of oxygen; $n > 50$, P value is shown. (A and B) Fluorescence images of gut nuclei in adult *C. elegans* fed on *Rhizobium* and maintained at 100% oxygen; (A) arrows point to elongated nuclei, fragmented nuclei, and nuclei connected with DNA bridges. (B) Closely positioned nuclei indicate a failed nuclei segregation. (C) Images of *C. elegans* fed with *E. coli* at 100% oxygen from the time of hatching illustrate abnormal gut nuclei pattern (arrowheads and *Insets*).

But, if DNA damage is localized to only intestinal cells (less than 1% of the DNA content of a fertile hermaphrodite), we would not expect to see a dramatic change in 8-OHdG as measured for the entire DNA content of a *C. elegans*. In published experiments, treatment of *C. elegans* with hydrogen peroxide, a known agent promoting formation of 8-OHdG (37), resulted only in a modest increase in [8-OHdG], from 20 to 25 pg per microgram of chromosomal DNA, despite high dosage of hydrogen peroxide and lengthy treatment (38). Thus, our results should not be interpreted as an absence of a change in 8-OHdG as this assay might not be sensitive enough to reveal the change in chromosomal DNA of intestinal cells.

Does Higher ROS Level Associated with Aberrant Karyokinesis in *C. elegans* Come Exclusively from Ingested *Rhizobium*? The pathogenic bacteria *Enterobacteria faecalis* does not produce ROS during infection but induces ROS production by the worm itself through activation of its dual oxidase, BLI-3 (39–41). BLI-3 has a NADPH oxidase (NOX) domain responsible for the ROS production and a peroxidase-like domain utilizing hydrogen peroxide (40, 42–44). We tested if NOX contributes to aberrant karyokinesis in worms fed on *Rhizobium*. An inhibition of NOX function in *bli-3(im10)* mutants decreases hydrogen peroxide level during *E. faecalis* infection (40). We thus crossed *bli-3(im10)* worms with *sur-5::GFP* reporter strain and compared the karyokinesis phenotype in the worms grown on *E. coli* and *Rhizobium* (Fig. 7). The characteristic *Rhizobium*-induced elongated, double and fragmented nuclei phenotype could still be seen in these worms. This result indicated that a NOX-mediated increase in hydrogen peroxide level upon infection is not essential for the karyokinesis defect in worms grown on *Rhizobium*.

Other Phenotypes Associated with Feeding on *Rhizobium*. *C. elegans* grown on *Rhizobium* from the time of hatching were thinner and smaller in body size (SI Appendix, Fig. S8 A and B). We tested a hypothesis that altered karyokinesis in some intestinal cells might impair intestinal function, resulting in the body mass deficiency. We supplemented *Rhizobium* food with *E. coli* by mixing *Rhizobium* with *E. coli* 1:10, 1:1, 10:1, and 100:1 by volume and evaluated both body size and gut nuclei morphology in *C. elegans* maintained on the mixed bacterial lawns from the time of hatching. We found that supplementation with *E. coli* at any dilution fully rescued the small, scrawny phenotype, but not the intestinal karyokinesis defect, although reducing its penetrance from 79 to 40% (SI Appendix, Fig. S8 C and D). This result indicates that *Rhizobium* may generate less of an essential nutrient or micronutrient for rapid *C. elegans* growth that is produced by *E. coli*. Because only a small fraction of *E. coli* could suppress the slow growth phenotype, but not the intestinal karyokinesis defect, the bacterial mixing experiment also suggested that a micronutrient malnutrition rather than a gut function defect prevented normal growth in *C. elegans* fed on *Rhizobium*. Considering that the abnormal gut nuclei morphology was observed only in a few nuclei in a 1,000-celled worm, it is not surprising that the physiological consequence of it was insignificant.

A test for bacterial aversion behavior revealed that *Rhizobium* does not repel *C. elegans* (SI Appendix, Fig. S9), indicating that the *C. elegans* pathogen surveillance system was not activated (45). In nature, *Rhizobium* forms a symbiotic relationship with particular legumes, fixing atmospheric nitrogen for the plant in return for carbon (46, 47). The *Rhizobium*–*C. elegans* ROS interaction may be mechanistically related to this more intimate plant bacterial symbiosis.

Concluding Remarks

We have shown that the *Rhizobium* that *C. elegans* may encounter in nature in rotting fruits inhibits normal karyokinesis in intestinal cells. We could cause a similar intestinal cell pheno-

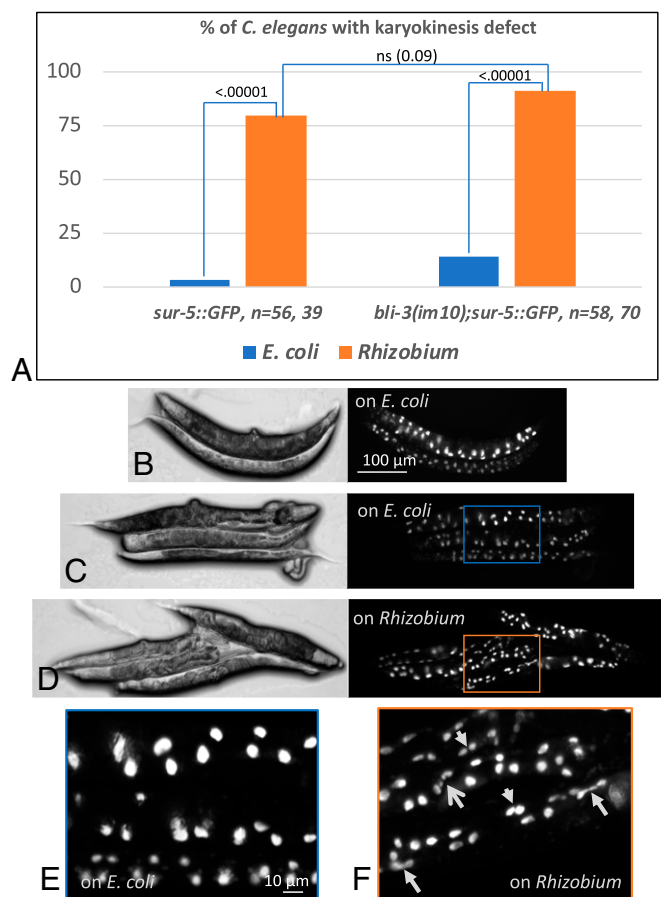


Fig. 7. Inhibition of ROS production by NOX in *bli-3(im10)* mutants does not rescue aberrant karyokinesis in gut nuclei of *C. elegans* grown on *Rhizobium*. (A) Graph showing percentage of worms with karyokinesis defect on *E. coli* and *Rhizobium*. P values are shown. ns, not significant. (B–F) differential interference contrast and fluorescence *bli-3(im10);sur-5::GFP* images of adult worms grown on *E. coli* (B, C, and E) and *Rhizobium* (D and F). (E and F) Enlarged Insets shown in C and D, correspondingly. (F) Arrows point to elongated (closed arrows), fragmented (open arrow), and double nuclei (short arrows). *bli-3* mutants have “blistered” phenotype and small and deformed body shapes due to its role in collagen crosslinking of the cuticle (43, 44). Because of that, the gut nuclei pattern is somewhat disturbed in comparison with WT worms but the karyokinesis defect can be observed in the intestine.

type via genetic lesions in *C. elegans* DNA damage response genes, *atm-1* and *dog-1*, suggesting that, indeed, DNA damage is a likely reason for abnormal nuclei divisions in the gut of *C. elegans* fed on *Rhizobium*. How does feeding on *Rhizobium* cause DNA damage in *C. elegans*? A common cause of DNA damage is reactive oxygen species (ROS). We showed that the *Rhizobium* diet evoked an oxidative stress response in *C. elegans*. A measurement of intracellular oxidative activity in *Rhizobium* revealed that it is higher than *E. coli*. The suppression of the karyokinesis defect on *Rhizobium* by growth under hypoxia conditions or with free radical scavengers also supports the model of ROS as a cause. What could be the *Rhizobium*-produced ROS? We favor the model that the ROS is hydrogen peroxide. Treatment with exogenous hydrogen peroxide causes abnormal karyokinesis similar to that caused by *Rhizobium*, and the concentration of hydrogen peroxide is higher in *Rhizobium* than in *E. coli*. In addition, it is feasible that relatively stable (half-life in milliseconds) and polar hydrogen peroxide can transit from bacteria inside the intestinal lumen, across two bacterial and one eukaryotic cell membrane, into the intestinal cell where it is converted into a hydroxyl radical in the presence of ferrous (Fe^{2+})

iron via the Fenton reaction (48) to damage DNA. The short half-life of this ROS would explain why a proximity between gut and *Rhizobium* cells is required for DNA damage to occur. It would also explain why most nuclei are not affected, which would be expected from a diffusion of more stable toxic product along intestinal tract. Why does the *Rhizobium* impair intestinal karyokinesis but not noticeably affect the function of the gut as might be expected from DNA damage mutagenesis? One possible explanation is that, in *C. elegans*, intestinal cells endoreduplicate genomic DNA during postembryonic development to generate the 32C per nucleus (11). The high copy number of these genomes may decrease the probability of a protein deficiency caused by DNA damage.

The altered gut nuclei phenotype is an easy readout; it could be used to further understand consequences of a high oxygen environment on animal cells. An interesting by-product of our EMS screen was a finding that the altered intestinal nuclei phenotype in *sur-5::GFP* reporter strain could be used as a discovery tool in a search for genes required for genome stability. Some of the candidate mutations selected in the screen are in the genes not previously linked to DNA damage response, repair, or chromosome segregation and, therefore, may reveal new genes involved in these processes (*SI Appendix, Fig. S10*).

In bacteria, ROS [superoxide (O_2^-), hydrogen peroxide (H_2O_2), and hydroxyl ion ($HO\cdot$)] are naturally generated during aerobic growth when molecular oxygen acquires electrons from reduced cofactors of various flavoproteins (49) and from side reactions of the electron transport chain (50). ROS inhibits enzymes containing iron or iron-sulfur clusters, including multiple enzymes involved in DNA replication and repair. To resist ROS, bacteria have evolved a number of strategies, such as the ROS efflux, compartmentalization, scavenging, and DNA repair mechanisms (49, 51). Some bacteria use ROS in fighting prokaryotic competitors less protected from oxidative stress, by either releasing ROS or inducing ROS production within neighboring cells by secreting redox cycling drugs (51–56). ROS-producing *Helicobacter pylori* and *Enterococcus faecalis* in the mammalian intestine may cause double strand breaks in DNA of epithelial cells, thus contributing to gastric carcinogenesis (57, 58). Killing of *C. elegans* with streptococcal species and *Enterococcus faecium* via hydrogen peroxide has been reported (59–61). It would be fascinating to explore the potential countermeasures that eukaryotes marshal to respond to bacteria and the counter countermeasures in turn used by bacteria.

Materials and Methods

The *C. elegans* and Bacteria Strains and Maintenance. The *C. elegans* and bacteria strains and maintenance were as follows: N2 Bristol, GR3065: translational *sur-5::GFP* from laboratory stock; VC381: *atm-1(gk186)*, VC13: *dog-1(gk10)*, CL2166: *gst-4p::GFP*, GS3798: *arls99 [dpy-7p::2Xnls::YFP]*, and LW699: GFP-tagged LMN-1 were obtained from CGC; GR3066: *atp-1(mg665)*; *sur-5::GFP*, GR3067: *dog-1(mg666);sur-5::GFP*, GR3068: *dog-1(mg667);sur-5::GFP*, GR3069: *dog-1(mg668);sur-5::GFP*, GR3195: *dog-1(mg694);sur-5::GFP*, and GR3196: *dog-1(mg695);sur-5::GFP* were generated by EMS in the course of this study. *bli-3(im10)* was a gift from Hiroki Moribe, Kurume University School of Medicine, Fukuoka, Japan. These mutants were crossed with *sur-5::GFP* reporter strain. All strains were maintained at 15 °C on nematode growth media (NGM) (62) agar plates in the presence of streptomycin (250 µg/mL final concentration) and seeded with OP50-1 *E. coli* carrying an integrated streptomycin resistance gene (CGC). Throughout the text, it is referred to as *E. coli*. RNAi was performed as described (63).

The library of the wild bacterial isolates (JUB) was created by, and obtained from, the M.-A. Félix laboratory (3, 8, 64). *Rhizobium hualtense* (ATCCBAA-115) and *Rhizobium galegae* (ATCC43677) were obtained from ATCC. Agrobacterium was a gift from the Jen Sheen laboratory (Department of Molecular Biology, Massachusetts General Hospital, Harvard Medical School, Boston).

Screening the JUB Library. Bacteria were grown in liquid LB for 3 to 4 d at room temperature until suspensions appeared cloudy. Then, 300 µL of the liquid

cultures were spotted on 3-cm NGM agar plates without antibiotic. The plates with the bacteria were maintained at room temperature for 2 d before setting up experiments. *C. elegans sur-5::GFP* eggs were obtained by bleaching gravid adults (62) and were spotted on the top of the bacterial lawn. The plates with *C. elegans* were maintained at 15 °C. The phenotypes were scored at various time points. Identity of all bacterial strains mentioned in this work was confirmed by 16S sequencing (MacroGenUSA).

Microscopy and DAPI Staining. For staining with DAPI, 30 *C. elegans* were transferred to a 1.5-mL Eppendorf tube filled with 20 µL of cold methanol and kept at –20 °C for 5 min. Methanol was aspirated, and 20 µL of S-basal (62) was added and pipetted up and down to separate bacteria stuck to *C. elegans*. The bacteria suspended in S-basal were removed. Finally, 10 µL of S-basal with 0.5 µL of DAPI solution (200×) was added to the worm pellet. After 5 min of incubation with DAPI, the worm's corpses were mounted on a microscopic agarose pad.

Live and fixed *C. elegans* were visualized with AxioZoom.V16 (Zeiss) and AxioImagerZ.1 (Zeiss) microscopes equipped with OrkaFlash 4.0 (Hamamatsu) and AxioCam HRc (Zeiss) cameras, respectively. Images were processed with ImageJ software.

Feeding *C. elegans* with *Rhizobium*. The *Rhizobium* cultures and plates were prepared as described above for the JUB library clones. Control *E. coli* strain was grown the same way and plated on NGM without antibiotics. Typically, two to three adults or 100 eggs or L1 larvae hatched in S-basal buffer overnight were plated on the lawns.

Testing a Requirement for a Contact Between the *Rhizobium* and *C. elegans* for Inducing Intestinal Nuclei Phenotype. A small, 10-mm-diameter, *Rhizobium* lawn was covered with a polyethersulfone (PES) 0.2-µm, 25-mm membrane filter (Sterlitech Corp.). Then, 100 µL of overnight suspension of *E. coli*, was placed on the top of the filter and was allowed to form a lawn. This was needed to stimulate the worm's development, which requires consumption of the live bacterial food (36). As a control, 100 µL of LB without *E. coli* was added to the *Rhizobium* lawn and to the *Rhizobium* covered with the filter. Analogously, 100 µL of *E. coli* was added to the *Rhizobium* lawn without the filter. Bleached *sur-5::GFP* eggs were plated on the top. The gut phenotype was scored in the grown adults ($n = 50$).

EMS Mutagenesis and Selection of Mutants. Mutagenesis of *sur-5::GFP* was done according to the published protocol (6). The aliquots of treated *C. elegans* were distributed on dozens of 10-cm NGM agar plates seeded with 10× concentrated *E. coli* bacterial lawn. Second generation (F1) progeny were bleached and plated in aliquots on 10 10-cm NGM agar plates seeded with 10× concentrated *E. coli* bacterial lawn. Adults (F2) with the desired phenotype were cloned, and homozygote progeny were outcrossed with a parental strain more than two times. Fewer than 10,000 haploid genomes were screened; the screen was not performed to saturation.

Identification of EMS mutants.

Next-generation sequencing of EMS mutants. DNA isolation, library construction, and whole genome sequencing with Illumina were carried out according to the published protocol (65).

Complementation test. Available alleles of the candidate genes were obtained from CGC and crossed with corresponding EMS mutants. An appearance of the altered gut nuclei phenotype indicated failed complementation and suggested that corresponding genes are the causative genes.

Evaluation of the gut nuclei phenotype with *sur-5::GFP* reporter. The candidate mutants obtained from CGC were outcrossed two times and crossed with *sur-5::GFP*, and their gut nuclei phenotype was evaluated on *E. coli* bacterial lawn.

RNAi Screen. *C. elegans sur-5::GFP* adults were bleached, and eggs were plated on bacterial lawn expressing RNAi (63) targeting a subset of DDR genes. Adult *C. elegans* were evaluated for abnormal gut nuclei.

Oxidative Stress Detection in *E. coli* and *Rhizobium* with H2DCFDA-AM Reagent.

The reagent dichlorodihydrofluorescein diacetate allowed the oxidative stress detection inside living cells. The procedure was according to the provided protocol: 10 µL of H2DCFDA-AM reagent (Molecular Probes) [10× stock solution in DMSO] spotted on the top of bacterial lawns. After 1 h at room temperature, cells were transferred on microscopic agarose and evaluated under the UV scope.

Hydrogen Peroxide Detection in *E. coli* and *Rhizobium*. Bacteria were grown in LB liquid media at room temperature to saturation ($OD_{600} \sim 0.5$). The suspensions were spun down and resuspended in the same volume of S-basal media. Then, 50 μ L of the suspension in triplicates were tested for hydrogen peroxide concentration with a Fluorimetric Hydrogen Peroxide Assay Kit according to the manufacturer's protocol (Sigma-Aldrich). Fluorescence Ex $\lambda 540$ /Em $\lambda 590$ was measured with microplate reader SpectraMax M5^e (Molecular Dynamics).

Hydrogen Peroxide Treatment. Early L1 larvae hatched in S-basal overnight were spotted on the small (~ 1 cm in diameter) *E. coli* lawn on regular NGM plates. Then, 50 μ L of 3% hydrogen peroxide was added so that all of the lawn area was covered. An appearance of bubbles (oxygen) indicated the presence of hydrogen peroxide and catalase activity: $2H_2O_2 \xrightarrow{\text{catalase}} 2H_2O + O_2$. The plates were kept at room temperature. The adult *C. elegans* grown from the treated L1 were inspected on day 3.

Catalase Tests. To test catalase activity, 10 μ L of 3% hydrogen peroxide was dropped on *E. coli* and *Rhizobium* lawns; a generation of bubbles was observed and recorded. To test the efficiency of bacterial catalase on reducing hydrogen peroxide concentration, 5 μ L of synchronized L1 larvae in S-basal were plated on agar, *E. coli* and *Rhizobium* lawns and immediately covered with hydrogen peroxide at a toxic dose of 30 μ L of 3%. The process and generation of bubbles on *E. coli*, but not on agar or *Rhizobium*, was recorded. Two hours later, the *C. elegans* were visually examined for their ability to move.

Oxygen Treatment. *Rhizobium* and *E. coli* lawns on NGM plates without antibiotics prepared as described above were seeded with *C. elegans* eggs and placed in oxygen chambers (Hypoxia Incubator Chamber; Stemcell technologies) with 1% (hypoxia) and high oxygen (50% and 100%) at 20 °C and kept under the conditions for 2 d. The phenotype was scored in adults.

Evaluation of DNA Damage with 8-OHdG ELISA. Chromosomal DNA was extracted from adult *C. elegans* grown on *E. coli* and *Rhizobium* according to the manufacturer's protocol (Puregene Tissue kit; Qiagen). Isolated DNA was treated with nuclease P1 and shrimp alkaline phosphatase (Sigma-Aldrich) to prepare for the assay as instructed in the protocol (OxiSelect Oxidative DNA Damage ELISA Kit; Cell Biolabs). Three biological replicates (DNA isolated from different batches of *C. elegans* on different days) and two technical replicates (split DNA solution from the same isolate) were obtained from *C.*

elegans grown on *E. coli* and two biological and two technical replicates from the *C. elegans* grown on *Rhizobium*. The samples and standards were treated according to the protocol (OxiSelect Oxidative DNA Damage ELISA Kit; Cell Biolabs). Microplate reader SpectraMax M5^e (Molecular Dynamics) was used for measurement absorption at 450 nm. Concentration of 8-hydroxydeoxyguanosine (8-OHdG) in the samples was determined according to the standard curve in Excel.

Treating *C. elegans* with *N*-acetyl-cysteine. Stock solution of NAC (Sigma-Aldrich) was prepared by dissolving 80 mg of NAC crystals in 10 mL of sterile water (~ 50 mM). Then, 1 mL of NAC solution or sterile water (as a control) was added on the top of 3-d-old *E. coli* or *Rhizobium* lawn on NGM agar without antibiotics, making roughly 5 mM final concentration of NAC. Eggs obtained by bleaching were spotted immediately after addition of NAC or water to the plates. Also, NAC (1/10 by volume) was added to the rest of the egg prep and incubated overnight at room temperature. L1 larvae hatched in NAC solution were plated on the *E. coli* and *Rhizobium* plates treated with NAC as above. Adult *C. elegans* were scored for the abnormal gut nuclei phenotype.

Feeding *C. elegans* with an Apple Supplement. *Rhizobium* plates were prepared as described above. Red apple cultivars, MacIntosh and Macoun, obtained from a local grocery store, were tested. The apples were wiped out with ethanol-soaked paper and rinsed with sterile water three times. Apple slices were cut out with a sterile blade and placed on the top of bacterial lawn. *C. elegans* eggs were plated on the top and near the slices (one slice per plate). No contaminating bacteria or fungi were visually detected on the apple plates within 5 to 6 d of the experiment.

Statistics. Whenever applicable, the experiments were done at least in three biological triplicates with 50 to 100 *C. elegans* at each condition when not indicated otherwise. *P* values were calculated with *t* test using Excel (Microsoft office) or calculating *z* score for the difference between two datasets.

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