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Bacterial metabolism of tryptophan causes toxicity in *Caenorhabditis elegans*

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Abstract

Tryptophan is an essential amino acid required not only for protein biosynthesis but also for the production of several physiologically important metabolites, including serotonin, kynurenine, and nicotinamide. Although dietary tryptophan is associated with various health benefits, excessive intake can result in adverse physiological effects. The specific tryptophan-derived metabolites responsible for such toxicity, however, remain incompletely characterized. Here, we investigate the mechanisms underlying tryptophan-induced toxicity in *Caenorhabditis elegans*. We observe that tryptophan concentrations of 1 mM or higher are highly toxic to *C. elegans*, blocking egg hatching. Notably, supplementation with various sugars alleviates this toxicity. Genetic analyses reveal that host tryptophan metabolism is dispensable for the observed effects. Instead, bacterial metabolism, particularly the conversion of tryptophan to indole, is essential for mediating toxicity. Bacterial strains deficient in indole production abolished tryptophan-induced toxicity, and all sugars that conferred protection also suppressed bacterial indole synthesis. These findings demonstrate that tryptophan toxicity in *C. elegans* is primarily mediated by microbial metabolism and underscore the pivotal role of host-microbiota interactions in shaping the physiological consequences of dietary amino acid intake.

Introduction

Tryptophan is an essential amino acid that is obtained exclusively through the diet. Beyond its fundamental role in protein biosynthesis, tryptophan regulates diverse physiological processes. It serves as a precursor for several bioactive molecules, including nicotinamide, serotonin, tryptamine, melatonin, and kynurenine, which participate in key metabolic and signaling pathways (Gao *et al*, 2020; Xue *et al*, 2023). As the sole precursor of the neurotransmitter serotonin, tryptophan is vital for maintaining optimal neurological and physiological function (Fernstrom, 1983). Systemic or oral administration of tryptophan enhances serotonin synthesis (Haleem, 2023), supporting its use as a dietary supplement to improve sleep quality and mood, and to manage psychiatric conditions such as depression and anxiety (Lindseth *et al*, 2015).

Emerging evidence suggests that dietary tryptophan and its metabolites may have therapeutic potential in various pathological conditions, including microbial infections, multiple sclerosis, inflammatory bowel disease, chronic kidney disease, and cardiovascular disease (Agus *et al*, 2018; Haleem, 2023; Xue *et al*, 2023). In rodent models, tryptophan supplementation has been shown to lower blood glucose levels in type 2 diabetic rats (Inubushi *et al*, 2012). In *Caenorhabditis elegans*, elevated tryptophan levels, either through direct supplementation or inhibition of its catabolism, have been linked to increased lifespan (van der Goot *et al*, 2012; Edwards *et al*, 2015; Dang *et al*, 2023). Additional studies suggest that high dietary intake of tryptophan may protect against intestinal inflammation (Zhang *et al*, 2021). Tryptophan fortification of foods has also been associated with improved sleep in infants with sleep disorders, weight gain in malnourished children, and enhanced mood and sleep quality in elderly individuals (Cubero *et al*, 2009; Friedman, 2018).

Despite these benefits, excessive tryptophan intake has also been associated with adverse outcomes (Hu *et al*, 2023; Izadi *et al*, 2024). In rodents, dietary restriction of tryptophan has been shown to extend lifespan and delay the onset of age-associated pathologies, including tumor development (Ooka *et al*, 1988; Segall & Timiras, 1976). Interestingly, the non-steroidal anti-inflammatory drug ibuprofen extends lifespan in yeast, worms, and flies by inhibiting tryptophan uptake (He *et al*, 2014). Although a minor fraction of dietary tryptophan is utilized for serotonin and tryptamine synthesis, over 95% is catabolized via the kynurenine pathway (Le Floc'h *et al*, 2011; Xue *et al*, 2023). Elevated levels of kynurenine pathway metabolites have been implicated in autoimmune diseases, neurodegenerative disorders, and various cancers (Schwarcz & Stone, 2017). Additionally,

gut microbiota metabolize tryptophan into indole and its derivatives, which can be further converted in the liver into indoxyl sulfate, a uremic toxin associated with oxidative stress and tissue injury (Agus *et al*, 2018; Graboski & Redinbo, 2020). Given that tryptophan has both beneficial and detrimental effects, it is critical to elucidate the mechanisms underlying its toxicity and to better understand the regulation of its metabolism in host-microbe interactions.

In this study, we investigated the mechanisms underlying tryptophan toxicity in *C. elegans*. We found that tryptophan was highly toxic at concentrations of 1 mM or greater, leading to a complete inhibition of egg hatching. Interestingly, supplementation with various sugars alleviated this toxicity. Genetic analyses revealed that host tryptophan metabolic pathways were not required for the manifestation of toxic effects. Instead, microbial metabolism of tryptophan played a central role in mediating toxicity. Specifically, the bacterial conversion of tryptophan into indole was responsible for the deleterious effects, and bacterial diets lacking the ability to produce indole abolished tryptophan toxicity. Notably, all sugars that alleviated tryptophan toxicity also inhibited bacterial indole production. Our findings reveal that tryptophan toxicity in *C. elegans* is driven by microbial metabolism and demonstrate the critical role of the host-microbiota axis in shaping the physiological outcomes of dietary amino acid intake.

Results

Tryptophan is highly toxic to *C. elegans* and inhibits egg hatching

While rearing *C. elegans* on *Escherichia coli* OP50-seeded nematode growth medium (NGM) plates supplemented with various amino acids, we observed that tryptophan was highly toxic. At concentrations of 2 mM or higher, tryptophan completely inhibited egg hatching (Figure 1A, B). This observation is consistent with a recent study (Ke *et al*, 2020), which reported that tryptophan exhibited the highest toxicity among the 20 standard amino acids, significantly impairing *C. elegans* development. In our assays, toxic effects were evident at 1 mM tryptophan, where more than half of the eggs failed to hatch, and at 2 mM, no hatching occurred (Figure 1A, B).

To investigate the mechanisms underlying tryptophan toxicity, we performed RNA sequencing to profile transcriptional changes following a 6-hour exposure of adult wild-type

98 N2 worms to 2 mM tryptophan. Transcriptome analysis revealed 1,625 differentially
99 expressed genes, including 963 upregulated and 662 downregulated transcripts (Figure 1C,
100 Table S1). Gene Ontology (GO) analysis of the upregulated genes revealed significant
101 enrichment in biological processes related to innate immune and xenobiotic responses (Figure
102 1D). In terms of molecular function, the upregulated genes were enriched for monooxygenase
103 and oxidoreductase activities, including several members of the cytochrome P450 and UDP-
104 glucuronosyltransferase (UGT) families (Figure S1A). These findings suggested activation of
105 xenobiotic detoxification pathways in response to tryptophan exposure. Conversely, GO
106 analysis of downregulated genes indicated enrichment in terms associated with cuticle
107 structure and transmembrane transport (Figure S1B, C).

108

109 **Sugars rescue tryptophan toxicity**

110 Interestingly, GO analysis also revealed that genes involved in monosaccharide and hexose
111 transport were upregulated upon tryptophan exposure (Figures 1D and S1A), suggesting a
112 possible interaction between sugar metabolism and tryptophan toxicity. To explore this, we
113 supplemented NGM plates containing 10 mM tryptophan with various monosaccharides at a
114 concentration of 50 mM. Notably, while these monosaccharides alone did not affect *C.*
115 *elegans* development in the absence of tryptophan (Figure S2A), they fully rescued the toxic
116 effects of tryptophan (Figure 2A, B). Egg hatching was restored on plates containing 10 mM
117 tryptophan supplemented with D-glucose, D-fructose, D-mannose, D-galactose, or L-
118 rhamnose, whereas no hatching occurred on plates with 10 mM tryptophan alone.

119 We further tested whether disaccharides could similarly suppress tryptophan toxicity.
120 To this end, we supplemented tryptophan-containing plates with lactose, sucrose, or maltose.
121 Interestingly, only lactose supplementation rescued tryptophan-induced toxicity (Figure 2A,
122 B). Neither sucrose nor maltose restored egg hatching in the presence of tryptophan.
123 Importantly, all three disaccharides supported normal development in the absence of
124 tryptophan (Figure S2A), indicating that the inability of sucrose and maltose to rescue was
125 not due to inherent developmental toxicity.

126 To determine whether lower concentrations of sugar could also rescue tryptophan
127 toxicity, we supplemented 10 mM tryptophan plates with varying concentrations of glucose
128 and assessed egg hatching. No rescue was observed at 2 mM glucose, whereas glucose

concentrations of 5 mM and above fully restored egg hatching (Figures 2C and S2B), demonstrating a dose-dependent protective effect.

Next, we examined whether glucose supplementation could reverse the transcriptional changes induced by tryptophan exposure. RNA sequencing analysis of adult N2 worms exposed to 2 mM tryptophan plus 50 mM glucose for 6 hours showed a marked reduction in transcriptomic perturbation. (Table S2). Only 244 genes were upregulated under the tryptophan-plus-glucose condition, compared to 963 genes upregulated in response to tryptophan alone (Figure 2D). Notably, 806 of the genes upregulated in the tryptophan-only condition were no longer upregulated when glucose was present (Table S3). Moreover, most of the 157 genes upregulated in both conditions exhibited higher expression levels under tryptophan-only exposure than with tryptophan plus glucose (Figure 2E). A similar trend was observed among downregulated genes, with a sharp reduction in the number of genes downregulated under the tryptophan-plus-glucose condition compared to tryptophan alone (Figure S3, Table S4). Collectively, these findings demonstrated that glucose supplementation effectively rescues tryptophan-induced toxicity at both the developmental and transcriptional levels.

Blocking tryptophan catabolism in *C. elegans* does not alleviate tryptophan toxicity

To understand how sugars mitigate tryptophan-induced toxicity, we first investigated whether excess tryptophan is metabolized into a toxic downstream product, and whether sugars act by suppressing the production or activity of this metabolite. In *C. elegans*, as in other eukaryotes, tryptophan is catabolized via three primary pathways: the kynurenine, serotonin, and indole branches (Figure 3A). The kynurenine pathway accounts for over 95% of dietary tryptophan metabolism, and elevated kynurenine/tryptophan ratios have been associated with a variety of pathological conditions (Xue *et al,* 2023). We therefore hypothesized that a toxic metabolite produced through the kynurenine pathway may underlie the observed developmental toxicity.

To test this hypothesis, we examined *tdo-2* mutants, which lack tryptophan-2,3-dioxygenase, the rate-limiting enzyme that initiates kynurenine synthesis. If toxicity were mediated by kynurenine-derived metabolites, *tdo-2* loss-of-function mutants would be expected to exhibit resistance to tryptophan toxicity. However, *tdo-2(ve552)*

160 mutants failed to hatch on 2 mM tryptophan, displaying sensitivity comparable to that of
161 wild-type N2 animals (Figure 3B, C).

162 We next asked whether the serotonin or indole branches of tryptophan metabolism
163 contribute to toxicity. Mutants defective in serotonin synthesis (*tph-1(mg280)*) or indole
164 production (*amx-2(ok1235)*) were exposed to 2 mM tryptophan. Eggs from both mutant
165 strains failed to hatch, showing no resistance relative to N2 (Figure 3B, C). Taken together,
166 these results indicated that blocking any single host-mediated tryptophan catabolic pathway
167 does not rescue tryptophan toxicity in *C. elegans*.

168

169 **Bacterial metabolism of tryptophan is essential for its toxicity in *C. elegans***

170 Since host-mediated tryptophan catabolism did not appear to underlie tryptophan toxicity, we
171 investigated whether bacterial metabolism of tryptophan contributes to the toxic effects
172 observed in *C. elegans*. To test this, we transferred *C. elegans* eggs to unseeded NGM plates
173 containing 10 mM tryptophan and monitored their hatching. It is well established that *C.*
174 *elegans* eggs can hatch in the absence of food and arrest at the L1 larval stage. While eggs
175 fail to hatch on *E. coli* OP50-seeded plates containing 10 mM tryptophan, the use of unseeded
176 plates enables assessment of tryptophan toxicity in the absence of bacterial metabolism.
177 Strikingly, all eggs transferred to unseeded plates containing 10 mM tryptophan hatched
178 successfully (Figure S4A, B). This observation strongly suggested that active bacterial
179 metabolism is required for tryptophan-induced toxicity.

180 To validate this finding, we monitored the hatching of *C. elegans* eggs on 10 mM
181 tryptophan plates seeded with kanamycin-killed *E. coli* OP50. Notably, in the presence of
182 dead bacteria, tryptophan did not impair egg hatching or subsequent development; all worms
183 developed to become fertile adults (Figure 4A, B). Together, these results indicated that active
184 bacterial metabolism of tryptophan generates a toxic product that mediates the observed
185 effects in *C. elegans*.

186 In *E. coli*, tryptophan is primarily metabolized via the indole pathway (Lee & Lee,
187 2010). Specifically, the bacterial enzyme tryptophanase (TnaA) catalyzes the conversion of
188 tryptophan into indole, pyruvate, and ammonia (Smith, 1897). We therefore hypothesized that
189 indole production by *E. coli* mediates the observed toxicity. To test this, we utilized an *E. coli*
190 Δ*tnaA* mutant, which lacks tryptophanase, and assessed *C. elegans* egg hatching on 10 mM
191 tryptophan plates seeded with this mutant strain. Strikingly, all eggs hatched and developed

192 normally on 10 mM tryptophan plates seeded with *E. coli* Δ*tnaA* (Figure 4C, D), confirming
193 that the toxic effect of tryptophan required its conversion to indole by bacterial TnaA.

194 To further substantiate this conclusion, we examined whether tryptophan was toxic
195 when metabolized by a bacterium that does not produce indole. *Pseudomonas aeruginosa*
196 catabolizes tryptophan via the oxidative kynurenine pathway, generating anthranilate rather
197 than indole (Bortolotti *et al*, 2016). When *C. elegans* eggs were cultured on 10 mM
198 tryptophan plates seeded with *P. aeruginosa*, all eggs hatched, and no toxicity was observed
199 (Figure 4E, F). Together, these data demonstrated that the production of indole by bacterial
200 tryptophanase is responsible for the toxic effects of tryptophan in *C. elegans*.

201

202 **Indole causes toxicity in *C. elegans***

203 To directly assess whether indole causes toxicity in *C. elegans*, we supplemented NGM plates
204 with indole and examined its effect on egg hatching. Similar to the effects observed with
205 tryptophan, indole supplementation on *E. coli* OP50-seeded plates inhibited egg hatching,
206 with toxic effects evident at concentrations of 1 mM and above (Figure 5A, B). Since *P.*
207 *aeruginosa* rescued the toxic effects of tryptophan, we next investigated whether it could
208 similarly attenuate indole-induced toxicity. However, indole remained toxic on *P. aeruginosa*-
209 seeded plates, and no eggs hatched at concentrations of 2 mM or higher (Figure 5A, B). At
210 1 mM indole, a significantly greater number of eggs hatched on *P. aeruginosa*-seeded plates
211 compared to those seeded with *E. coli*. This difference is likely attributable to the endogenous
212 production of indole by *E. coli*, which *P. aeruginosa* does not produce, resulting in a higher
213 effective concentration of indole in the *E. coli*-seeded condition.

214 To determine whether bacterial presence is required for indole toxicity, we examined
215 egg hatching on both unseeded and *E. coli*-seeded indole plates. Notably, egg hatching was
216 similarly inhibited on unseeded plates containing 2 mM indole (Figures 5C and S5A),
217 indicating that indole exerts its toxic effects independently of live bacteria. Similarly, indole
218 inhibited egg hatching on plates seeded with kanamycin-killed *E. coli* (Figure 5D, E), further
219 confirming that bacterial viability is not required for indole-induced toxicity.

220 We also tested whether the *E. coli* Δ*tnaA* mutant, which lacks tryptophanase and
221 cannot convert tryptophan to indole, could alter indole toxicity. Importantly, the Δ*tnaA*
222 mutant did not mitigate indole toxicity, as eggs failed to hatch on 2 mM indole plates seeded

with this strain (Figure 5F, G). Collectively, these findings demonstrate that indole itself is inherently toxic to *C. elegans*, and this toxicity is independent of bacterial metabolism. While *E. coli*-derived indole mediates the toxic effects of tryptophan, the presence of indole alone is sufficient to exert toxicity, irrespective of bacterial species or metabolic activity.

Sugars do not alleviate indole-induced toxicity in *C. elegans*

We next sought to understand the mechanism by which glucose and other sugars suppress tryptophan toxicity. Transcriptomic profiling (Table S1) revealed that tryptophan exposure leads to the upregulation of numerous UGT genes. Notably, the transcriptional response to tryptophan closely resembles that elicited by indole exposure, as reported previously (Lee *et al*, 2017), suggesting that UGT induction may constitute a host defense mechanism against indole-mediated toxicity. UGT enzymes play a central role in phase II xenobiotic detoxification by conjugating glucose moieties to xenobiotic substrates, thereby increasing their solubility and facilitating excretion. Consistent with this, *C. elegans* has been shown to glycosylate indoles to reduce their toxicity (Stupp *et al*, 2013).

We hypothesized that glucose supplementation may enhance UGT-mediated detoxification by increasing the availability of glucose substrates for conjugation. Under this model, the glycosylation of indoles could be responsible for the observed rescue of tryptophan toxicity when glucose is added. To test this hypothesis, we supplemented indole-containing plates with glucose and assessed egg hatching. Unexpectedly, even 50 mM glucose failed to promote hatching on plates containing 2 mM indole (Figures 6A and S5B), indicating that glucose supplementation does not alleviate indole toxicity. To further evaluate this, we supplemented indole plates with other sugars that were tested for mitigating tryptophan toxicity. However, none of these sugars ameliorated the toxic effects of indole (Figure S5C). These results suggested that sugar-mediated rescue of tryptophan toxicity does not occur via enhanced UGT-mediated glycosylation of indole, and that sugars do not protect *C. elegans* from indole toxicity.

Sugars alleviate tryptophan toxicity by suppressing bacterial indole production

We observed that several sugars alleviate the toxic effects of tryptophan but not those of indole. This suggests that these sugars may act upstream of indole production, potentially

254 preventing tryptophan from being converted into its toxic derivative. We therefore asked
255 whether sugar supplementation affects bacterial indole production. Previous studies have
256 shown that glucose inhibits the activity of tryptophanase, the enzyme responsible for
257 converting tryptophan to indole in bacteria (Botsford & DeMoss, 1971; Li & Young, 2014).
258 Based on these findings, we hypothesized that supplementation of sugars might inhibit
259 tryptophanase activity, thereby reducing indole production even in the presence of excess
260 tryptophan.

261 To test this, we quantified indole production by bacteria using Kovac's reagent. As
262 expected, *E. coli* cultured in LB medium produced indole under basal conditions, and indole
263 levels were further elevated upon tryptophan supplementation (Figure 6B). Notably, glucose
264 supplementation significantly reduced indole production in cultures exposed to high
265 concentrations of tryptophan (Figure 6B). We extended this analysis to other
266 monosaccharides and disaccharides. Strikingly, only those sugars that rescued tryptophan
267 toxicity also reduced indole production. In contrast, sucrose and maltose, both ineffective at
268 rescuing tryptophan toxicity, did not significantly lower indole levels compared to the
269 tryptophan-only control (Figure 6B). Furthermore, the sugars that alleviated tryptophan
270 toxicity also diminished basal indole production in the absence of exogenous tryptophan
271 (Figure 6C).

272 Collectively, these findings support a model in which *E. coli* converts tryptophan to
273 indole, which is then taken up by *C. elegans*. At high concentrations, indole is toxic and
274 impairs egg hatching. Various sugars, including multiple monosaccharides and the
275 disaccharide lactose, inhibit indole biosynthesis in *E. coli*, thereby preventing the downstream
276 toxic effects of tryptophan in *C. elegans* (Figure 6D).

277

278 **Discussion**

279 Our study demonstrates that bacterial metabolism of tryptophan into indole can profoundly
280 impact host physiology and contribute to toxicity. Over 85 bacterial species, including both
281 Gram-positive and Gram-negative, have been reported to express functional tryptophanase
282 and convert tryptophan to indole (Lee & Lee, 2010). Many of these species reside in the
283 human gut, where they efficiently produce indole and its derivatives (Roager & Licht, 2018).
284 Indeed, indole is the most abundant tryptophan-derived metabolite detected in human feces
285 and mouse cecal contents, accounting for 50-75% of total tryptophan catabolites, with

concentrations of 2.6-2.7 mM (Darkoh *et al*, 2015; Dong *et al*, 2020). The indole concentrations could increase further during enhanced tryptophan uptake, as indole concentrations as high as 6.6 mM have been observed in human faeces (Darkoh *et al*, 2015). Therefore, indoles produced by gut microbiota might lead to toxicity under specific circumstances.

Indole and its derivatives exhibit pleiotropic effects on host biology, ranging from beneficial to detrimental, depending on the context (Jaglin *et al*, 2018; Tennoune *et al*, 2022; Yang *et al*, 2023; Perdijk *et al*, 2024; Seymour *et al*, 2024). At moderate concentrations, indole can promote host health. For example, 250 μM indole has been shown to extend healthspan in *C. elegans*, *Drosophila*, and mice via conserved molecular pathways (Sonowal *et al*, 2017, 2023). Indole also supports intestinal homeostasis by enhancing epithelial barrier function, modulating immune responses, and suppressing the colonization of pathogenic bacteria (Bansal *et al*, 2010; Shimada *et al*, 2013; Alexeev *et al*, 2018; Scott *et al*, 2020). Furthermore, indole and its derivatives have been implicated in ameliorating inflammatory bowel disease, hemorrhagic colitis, colorectal cancer, and type 2 diabetes mellitus (Ye *et al*, 2022).

However, at elevated concentrations or in the context of gut dysbiosis, indole and its metabolites can become deleterious. In the liver, indole is converted to indoxyl sulfate, a known uremic toxin linked to systemic toxicity. Indoxyl sulfate has been implicated in the pathogenesis of cardiovascular and chronic kidney diseases (Banoglu *et al*, 2001; Hung *et al*, 2017; Hoibian *et al*, 2018), and is capable of inducing neuroinflammation and neuronal damage (Adesso *et al*, 2017, 2018). Additionally, indole exacerbates portal hypertension in a rat model of liver cirrhosis (Huc *et al*, 2018) and can promote the development of collagen-induced arthritis in mice via microbiota-dependent mechanisms (Seymour *et al*, 2024). In *C. elegans*, high levels of indole reduce lifespan (Lee *et al*, 2017). These observations underscore that the pathological consequences of indole are context-dependent, influenced by its concentration and specific derivatives.

Our findings reveal that several monosaccharides, as well as the disaccharide lactose, inhibit indole production in *E. coli*. Previous studies have shown that glucose and arabinose suppress indole production by inhibiting tryptophanase activity (Botsford & DeMoss, 1971; Li & Young, 2014). Although catabolite repression via cyclic AMP (cAMP) was initially proposed as the underlying mechanism (Isaacs *et al*, 1994), subsequent studies have shown

that tryptophanase regulation also involves cAMP-independent pathways (Li & Young, 2014). Tryptophanase activity is regulated through complex and multilayered networks (Stewart & Yanofsky, 1985; Isaacs *et al*, 1994; Gong & Yanofsky, 2003; Li & Young, 2014), suggesting that indole production may offer evolutionary advantages to bacteria (Boya *et al*, 2021). Indeed, indole serves as a critical signaling molecule in microbial communities and can influence microbial composition (Lee & Lee, 2010; Lee *et al*, 2015). Diets rich in protein and tryptophan may thus promote the proliferation of indole-producing bacteria, potentially leading to gut dysbiosis.

Our results suggest that indole production by gut microbiota can be modulated through dietary interventions. Indeed, dietary components such as fiber, non-starch polysaccharides, and high-fat diets have been shown to reduce gut indole production, shifting tryptophan metabolism toward more beneficial pathways (Knarreborg *et al*, 2002; Krishnan *et al*, 2018; Huang *et al*, 2023; Sinha *et al*, 2024). We identified additional sugars capable of suppressing indole synthesis in *E. coli* and possibly in other bacterial species. Targeted delivery or regulated release of these sugars in the gut may provide a strategy to fine-tune microbial tryptophan metabolism and mitigate the pathological effects associated with excessive indole production.

Materials and methods

Bacterial strains

The following bacterial strains were used in this study: *Escherichia coli* OP50, *E. coli* BW25113, *E. coli* BW25113 Δ*tnaA* mutant, and *Pseudomonas aeruginosa* PA14 Δ*gacA* mutant. Bacterial cultures were grown in Luria-Bertani (LB) broth at 37°C. The *E. coli* BW25113 Δ*tnaA* mutant was from the *E. coli* Keio collection (Baba *et al*, 2006), and was cultured in LB broth supplemented with 25 μg/mL kanamycin. Because *P. aeruginosa* virulence might affect the hatching and development of *C. elegans*, we used the *gacA* mutant, which has drastically reduced virulence (Tan *et al*, 1999). Nematode growth medium (NGM) plates were seeded with the respective bacterial cultures and incubated at room temperature for at least two days prior to use in experiments.

C. elegans strains and growth conditions

C. elegans hermaphrodites were maintained at 20°C on NGM plates seeded with *E. coli* OP50. The Bristol N2 strain was used as the wild-type control. The following strains were used in this study: RG3052 *tdo-2(ve552[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])*, RB1190 *amx-2(ok1235)*, and MT15434 *tph-1(mg280)*. Some of the strains were obtained from the Caenorhabditis Genetics Center (University of Minnesota, Minneapolis, MN).

Preparation of NGM plates with different supplements

The following supplements, indicated with their product numbers, were obtained from HiMedia BioSciences: L-tryptophan (#GRM067), D-(+)-glucose (#TC130), D-(-)-fructose (#GRM196), D-(+)-mannose (#RM104), L-(+)-rhamnose (#RM062), D-(+)-galactose (#GRM101), lactose (#GRM017), sucrose (#GRM134), D-(+)-maltose (#GRM3050), and indole (#GRM824). All the supplements were added to the NGM immediately prior to pouring into plates to achieve the desired final concentrations. Indole was dissolved in methanol, and control plates were supplemented with equivalent volumes of methanol for assays involving indole.

***C. elegans* egg hatching assays**

A synchronized egg population was obtained by transferring 15-20 gravid adult hermaphrodites onto NGM plates and allowing them to lay eggs for 1.5-2 hours. After that, the adults were removed, and the number of eggs laid was counted. The plates were then incubated at 20°C, and the number of hatched eggs/L1 larvae was quantified after 24 hours. Representative images of the plates were captured after 96 hours of incubation at 20°C. For each experimental condition, at least three independent biological replicates were performed.

RNA isolation

RNA isolation was performed as described previously (Ravi *et al*, 2023; Rao & Singh, 2025). Briefly, wild-type N2 animals were synchronized by egg laying. Approximately 50 gravid adult hermaphrodites were transferred to 9-cm NGM plates seeded with *E. coli* OP50 and allowed to lay eggs for 4 hours. After removing the adults, the eggs were incubated at 20°C

379 for 72 hours to allow development into adult worms. The adult animals were then collected,
380 washed with M9 buffer, and transferred to *E. coli* OP50-seeded plates containing the
381 following three conditions: control plates without supplements, 2 mM tryptophan, and 2 mM
382 tryptophan plus 50 mM glucose. Following a 6-hour incubation at 20°C, worms were
383 collected, washed with M9 buffer, and snap-frozen in TRIzol reagent (Life Technologies,
384 Carlsbad, CA). Total RNA was extracted using the RNeasy Plus Universal Kit (Qiagen,
385 Netherlands). The RNA samples from three independent biological replicates were isolated
386 for each condition.

387

388 **RNA sequencing and data analysis**

389 Total RNA was isolated from N2 worms grown under three conditions: control, 2 mM
390 tryptophan, and 2 mM tryptophan plus 50 mM glucose, using three biological replicates per
391 condition, as described above. Library preparation and sequencing were conducted at Unipath
392 Specialty Laboratory Ltd., India. The cDNA libraries were sequenced on the NovaSeq 6000
393 platform using 150-bp paired-end reads.

394 The RNA sequencing data were analyzed using the web platform Galaxy
395 (https://usegalaxy.org/), following previously described workflows (Ghosh & Singh, 2024;
396 Rao *et al*, 2024). Briefly, paired-end reads were first trimmed using the Trimmomatic tool.
397 The resulting reads obtained for each sample were mapped to the *C. elegans* reference
398 genome (WS220) using the STAR aligner. Read counts per gene were quantified using
399 the htseq-count tool. Differential gene expression analysis was then performed using
400 DESeq2. Genes exhibiting at least a two-fold change and a P-value <0.01 were considered
401 differentially expressed. Gene ontology enrichment analysis was carried out using the DAVID
402 Bioinformatics Resource (https://david.ncifcrf.gov/tools.jsp). Venn diagrams were generated
403 using the BioVenn web tool (https://www.biovenn.nl/), and heatmaps were created using the
404 pheatmap package in R version 4.5.0 (R Statistical Software).

405

406 **Bacterial killing**

407 Bacterial killing was performed as described previously (Lee *et al*, 2024). Briefly, *E. coli*
408 OP50 was cultured in 50 mL of LB broth at 37°C for 24 hours. The culture was then
409 centrifuged at 3000 × *g* for 30 minutes. The supernatant was discarded, and the bacterial

pellet was resuspended in 100 mL of autoclaved water containing 250 μg/mL kanamycin. The suspension was distributed equally into two 250-mL flasks and incubated with shaking at 37°C for an additional 24 hours. To confirm bacterial killing, 1 mL of the culture was aliquoted, washed three times with sterile water, and plated on LB agar. Plates were incubated at 37°C for 24 hours and checked for viable colonies. If no colonies were detected, the bacteria were considered effectively killed and stored at 4°C. Before use, the killed bacterial culture was washed three times with 100 mL of sterile water and concentrated 20-fold before seeding onto NGM plates.

Indole measurement using Kovac's reagent

To prepare samples for indole quantification, various sugars were added to LB broth at a final concentration of 50 mM. Each sugar was added to 10 mL of LB broth in 50 mL tubes, with or without added tryptophan. After that, 20 μL of the same primary culture of *E. coli* OP50 was added into each tube, which were then incubated at 37°C for 12 hours. The optical density at 600 nm (OD_{600}) was measured for all cultures, and the cultures were adjusted to the same OD_{600} prior to indole quantification. As a control, 10 mL of LB broth containing *E. coli* OP50 without added sugar or tryptophan was similarly cultured. Indole standards (0, 0.1, 0.5, 1, 2, 5, and 10 mM) were prepared in LB broth and processed in parallel with the samples to generate a calibration curve.

Indole levels were measured as described previously (Mueller *et al*, 2009), with minor modifications. Briefly, 1 mL of each culture was transferred to a 1.5 mL microcentrifuge tube and centrifuged at 11,300 × *g* for 20 seconds. Then, 250 μL of the supernatant was mixed with 250 μL of 20% (w/v) trichloroacetic acid (Sigma-Aldrich #T6399) and incubated on ice for 15 minutes. The mixture was centrifuged at 13,000 × *g* for 10 minutes to remove precipitated proteins. Subsequently, 450 μL of the resulting supernatant was combined with 500 μL of Kovac's reagent (Sigma-Aldrich #67309) and vortexed. The samples were incubated at room temperature for 2 minutes to allow phase separation. Then, 200 μL of the upper (colored) layer was transferred to 800 μL of ethanol, and absorbance was measured at 540 nm. Indole concentrations were calculated using a standard curve generated on the same day.

Quantification and statistical analysis

The statistical analysis was performed with Prism 8 (GraphPad). All error bars represent the mean ± standard deviation. The unpaired, two-tailed, two-sample *t*-test was employed when needed, and statistical significance was determined when $P < 0.05$. For more than two samples, ordinary one-way ANOVA followed by Dunnett's multiple comparisons test was used. In the figures, asterisks (*) denote statistical significance as follows: $*P < 0.05$; $**P < 0.01$; and $***P < 0.001$, as compared with the appropriate controls. All experiments were performed at least three times.

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Author Contributions

S.G., Subodh, and J.S. conceived and designed the experiments. S.G. and Subodh performed the experiments. S.G. and J.S. analyzed the data and wrote the paper.

Declaration of interests

470 The authors declare no competing interests.

471

472 **Data availability**

473 The RNA sequencing data for N2 worms exposed to tryptophan alone and tryptophan plus
474 glucose, along with the control, have been submitted to the public repository, the Sequence
475 Read Archive, with BioProject ID PRJNA1271509. All data generated or analyzed during this
476 study are included in the manuscript and supporting files.

477

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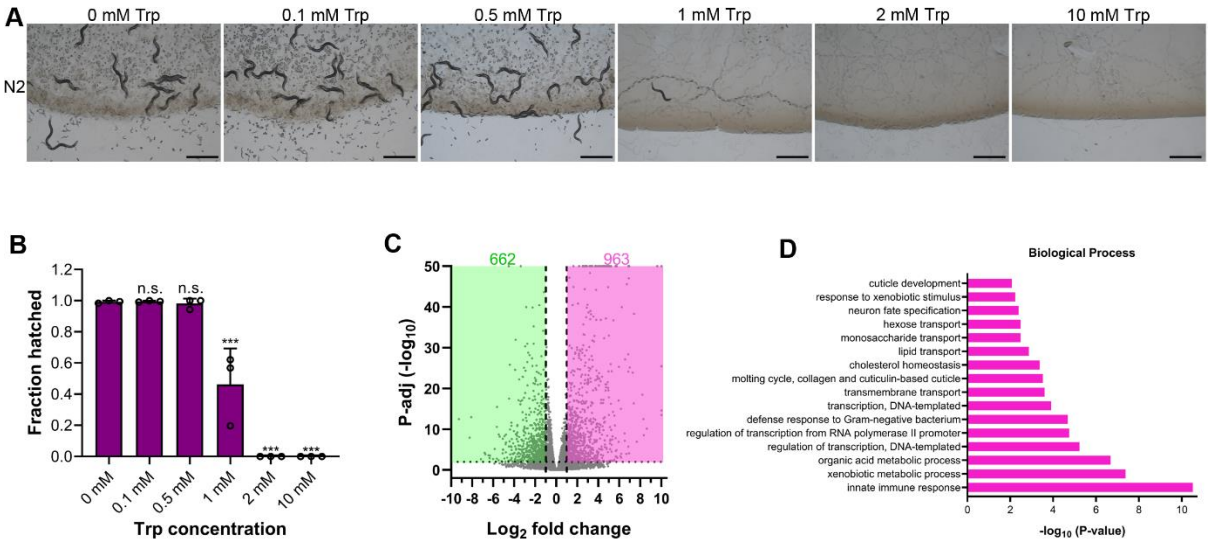
646

647 **Figures**

648



649

650 **Figure 1. Tryptophan is highly toxic to *C. elegans* and inhibits egg hatching**

651 (A) Representative images of N2 worms on various concentrations of tryptophan (Trp) on *E.*
652 *coli* OP50 diet after 96 hours of hatching at 20°C. Scale bar = 1 mm.

653 (B) Quantification of hatched eggs of N2 worms on various concentrations of tryptophan
654 on *E. coli* OP50 after 24 hours of hatching at 20°C. ***$P < 0.001$ and n.s., nonsignificant via
655 ordinary one-way ANOVA followed by Dunnett's multiple comparisons test (n = 3 biological
656 replicates).

657 (C) Volcano plot of upregulated and downregulated genes in N2 worms exposed to 2 mM
658 tryptophan.

659 (D) Gene ontology enrichment analysis of genes upregulated upon tryptophan exposure for
660 biological processes.

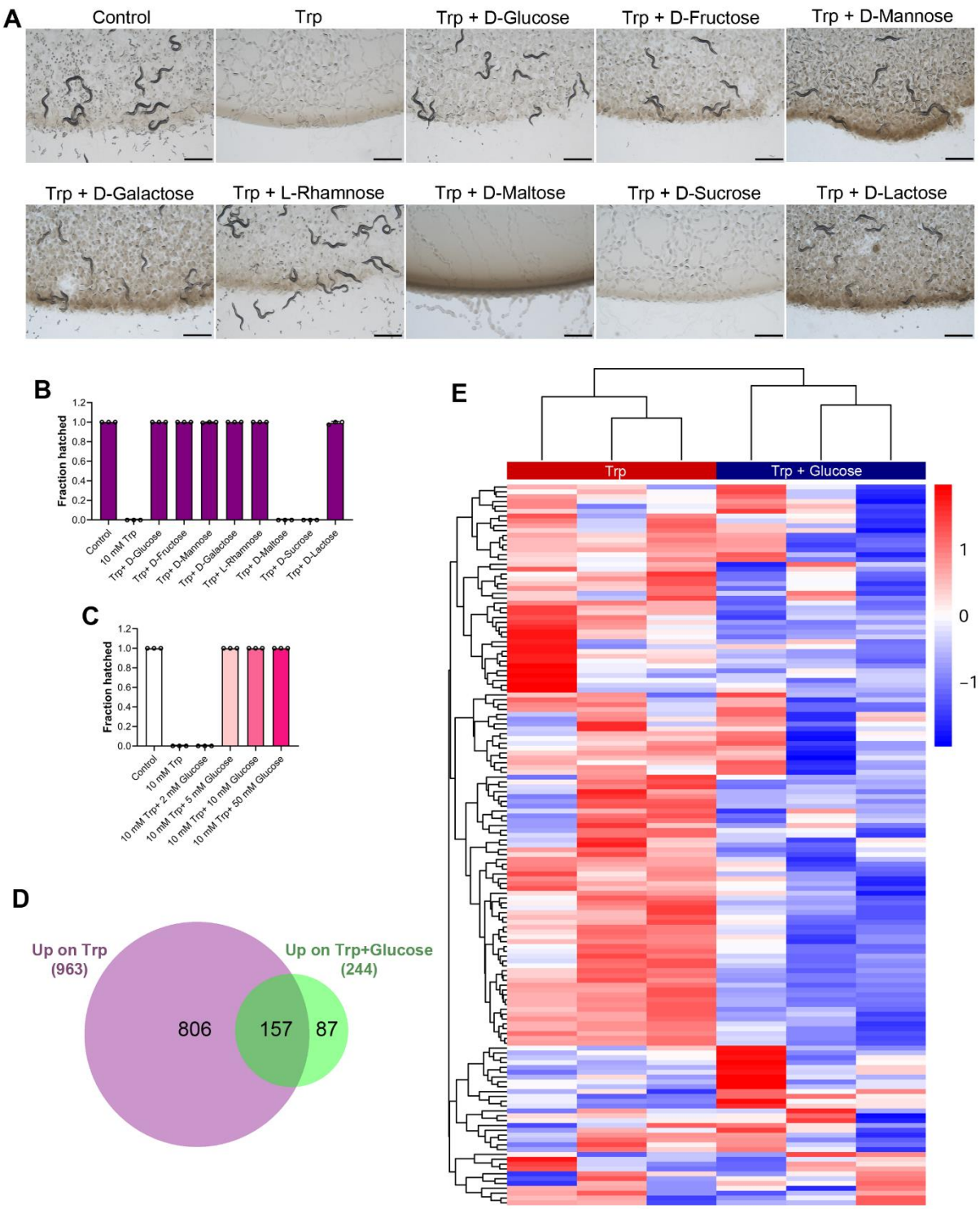


Figure 2. Sugars rescue tryptophan toxicity

(A) Representative images of N2 worms grown on *E. coli* OP50 diet for 96 hours at 20°C containing 0 mM tryptophan (Trp), 10 mM Trp, and 10 mM Trp supplemented with 50 mM of different sugars. Scale bar = 1 mm.

(B) Quantification of hatched eggs of N2 worms grown on *E. coli* OP50 diet for 24 hours at 20°C containing 0 mM Trp, 10 mM Trp, and 10 mM Trp supplemented with 50 mM of different sugars. (n = 3 biological replicates).

669 (C) Quantification of hatched eggs of N2 worms grown on *E. coli* OP50 diet for 24 hours at
670 20°C containing 0 mM Trp, 10 mM Trp, and 10 mM Trp supplemented with different
671 concentrations of glucose. (n = 3 biological replicates).

672 (D) Venn diagram showing the overlap between the genes upregulated upon exposure to 2
673 mM Trp alone and 2 mM Trp plus 50 mM glucose.

674 (E) Heat map showing the expression of upregulated genes common between Trp alone and
675 Trp plus glucose.

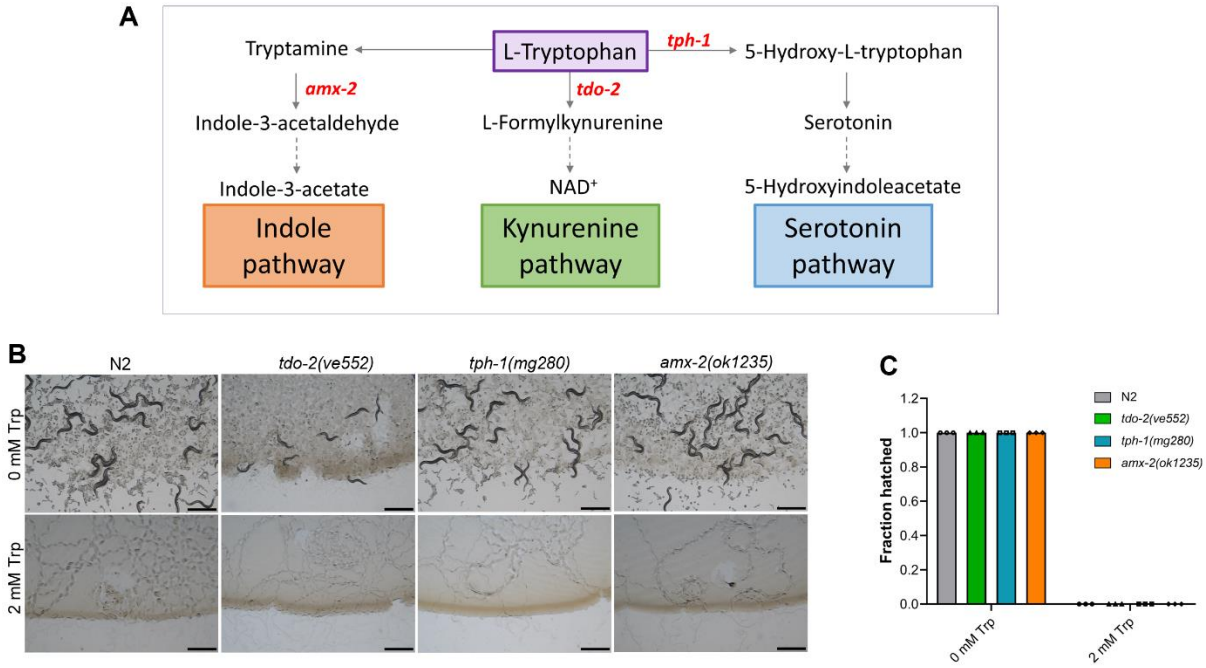


Figure 3. Blocking tryptophan catabolism in *C. elegans* does not alleviate tryptophan toxicity

(A) Simplified representation of tryptophan metabolism pathways. L-tryptophan is metabolized in *C. elegans* via three main pathways: the kynurenine, indole, and serotonin pathways.

(B) Representative images of N2, *tdo-2(ve552)*, *tph-1(mg280)*, and *amx-2(ok1235)* worms after 96 hours of hatching at 20°C on *E. coli* OP50 diet containing 0 mM and 2 mM tryptophan (Trp). Scale bar = 1 mm.

(C) Quantification of hatched eggs of N2, *tdo-2(ve552)*, *tph-1(mg280)*, and *amx-2(ok1235)* worms after 24 hours of hatching at 20°C on *E. coli* OP50 diet containing 0 mM and 2 mM Trp. (n = 3 biological replicates).

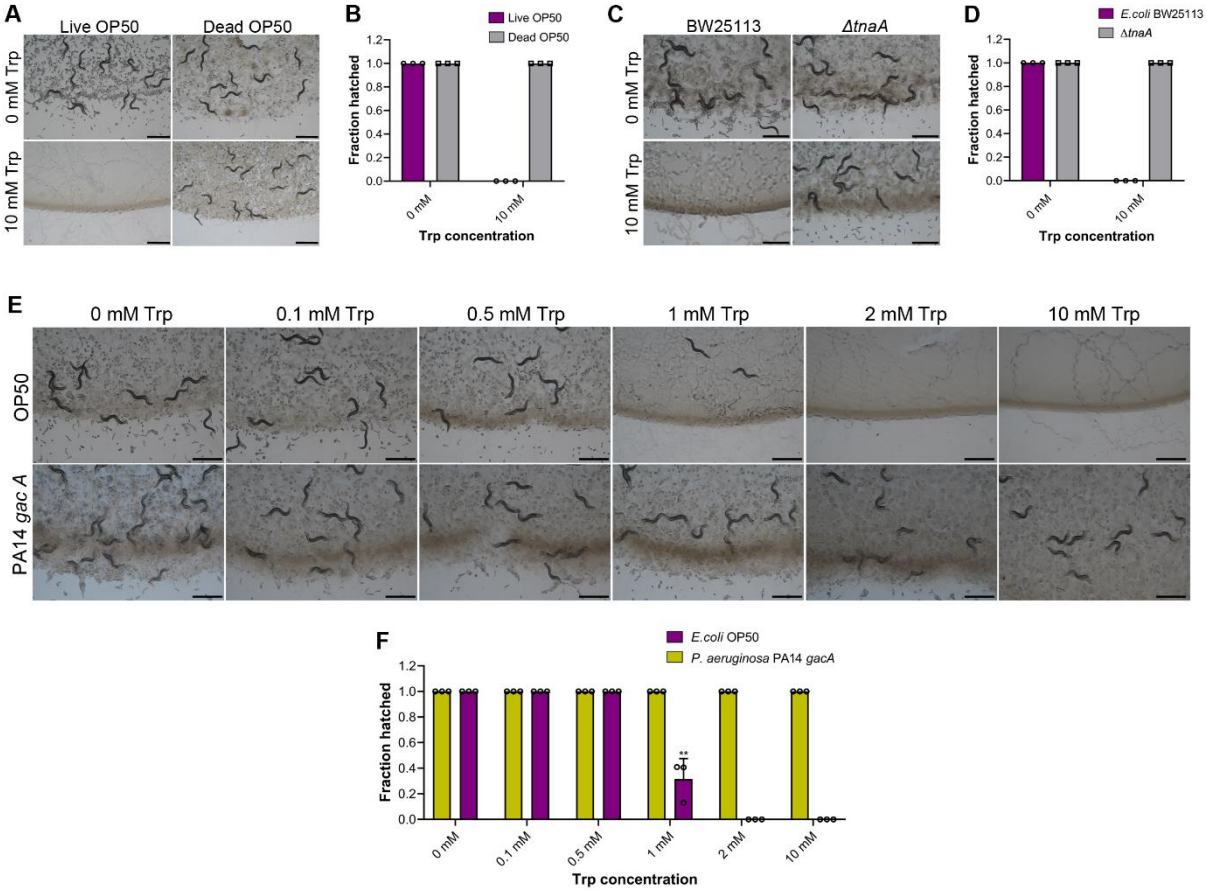


Figure 4. Bacterial metabolism of tryptophan is essential for its toxicity in *C. elegans*

(A) Representative images of N2 worms after 96 hours of hatching at 20°C on live and dead *E. coli* OP50 diet containing 0 mM and 10 mM tryptophan (Trp). Scale bar = 1 mm.

(B) Quantification of hatched eggs of N2 worms after 24 hours of hatching at 20°C on live and dead *E. coli* OP50 diet containing 0 mM and 10 mM Trp. (n = 3 biological replicates).

(C) Representative images of N2 worms after 96 hours of hatching at 20°C on wild-type *E. coli* BW25113 and Δ*tnaA* mutant diet containing 0 mM and 10 mM Trp. Scale bar = 1 mm.

(D) Quantification of hatched eggs of N2 worms after 24 hours of hatching at 20°C on wild-type *E. coli* BW25113 and Δ*tnaA* mutant diet containing 0 mM and 10 mM Trp. (n = 3 biological replicates).

(E) Representative images of N2 worms after 96 hours of hatching at 20°C on various concentrations of Trp on *E. coli* OP50 and *P. aeruginosa* PA14 diets. Scale bar = 1 mm.

(F) Quantification of hatched eggs of N2 worms after 24 hours of hatching at 20°C on various concentrations of Trp on *E. coli* OP50 and *P. aeruginosa* PA14 diets. ***$P < 0.001$ via the t-test (n = 3 biological replicates).

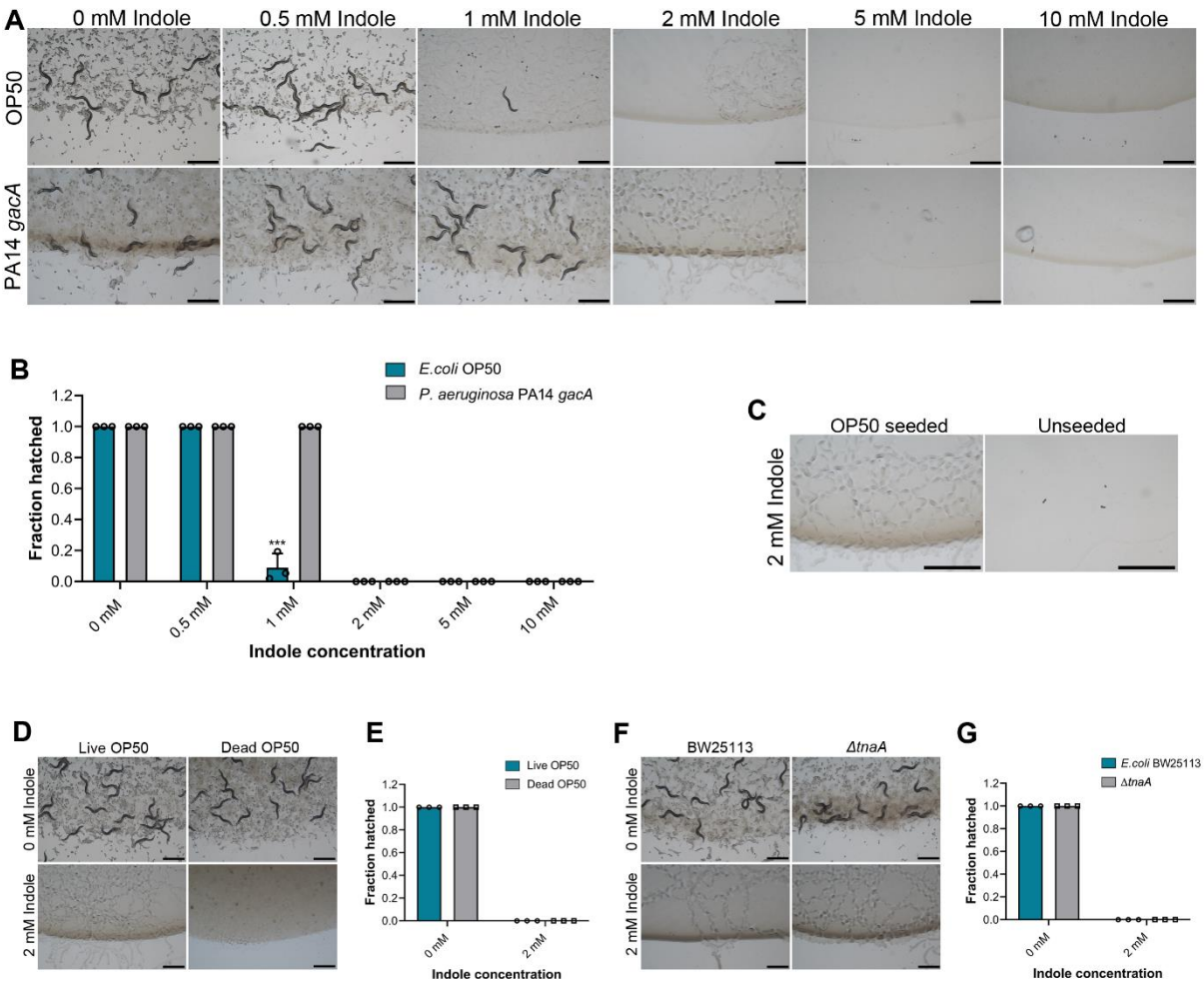


Figure 5. Indole causes toxicity in *C. elegans*

(A) Representative images of N2 worms after 96 hours of hatching at 20°C on various concentrations of indole on *E. coli* OP50 and *P. aeruginosa* PA14 diets. Scale bar = 1 mm.

(B) Quantification of hatched eggs of N2 worms after 24 hours of hatching at 20°C on various concentrations of indole on *E. coli* OP50 and *P. aeruginosa* PA14 diets. ***$P < 0.001$ via the *t*-test ($n = 3$ biological replicates).

(C) Representative images of N2 worms after 24 hours of hatching at 20°C on unseeded and *E. coli* OP50-seeded plates supplemented with 2 mM indole. Scale bar = 1 mm.

(D) Representative images of N2 worms after 96 hours of hatching at 20°C on live and dead *E. coli* OP50 diet containing 0 mM and 2 mM indole. Scale bar = 1 mm.

(E) Quantification of hatched eggs of N2 worms after 24 hours of hatching at 20°C on live and dead *E. coli* OP50 diet containing 0 mM and 2 mM indole. ($n = 3$ biological replicates).

(F) Representative images of N2 worms after 96 hours of hatching at 20°C on wild-type *E. coli* BW25113 and Δ*tnaA* mutant diet containing 0 mM and 2 mM indole. Scale bar = 1 mm.

(G) Quantification of hatched eggs of N2 worms after 24 hours of hatching at 20°C on wild-type *E. coli* BW25113 and Δ*tnaA* mutant diet containing 0 mM and 2 mM indole. ($n = 3$ biological replicates).

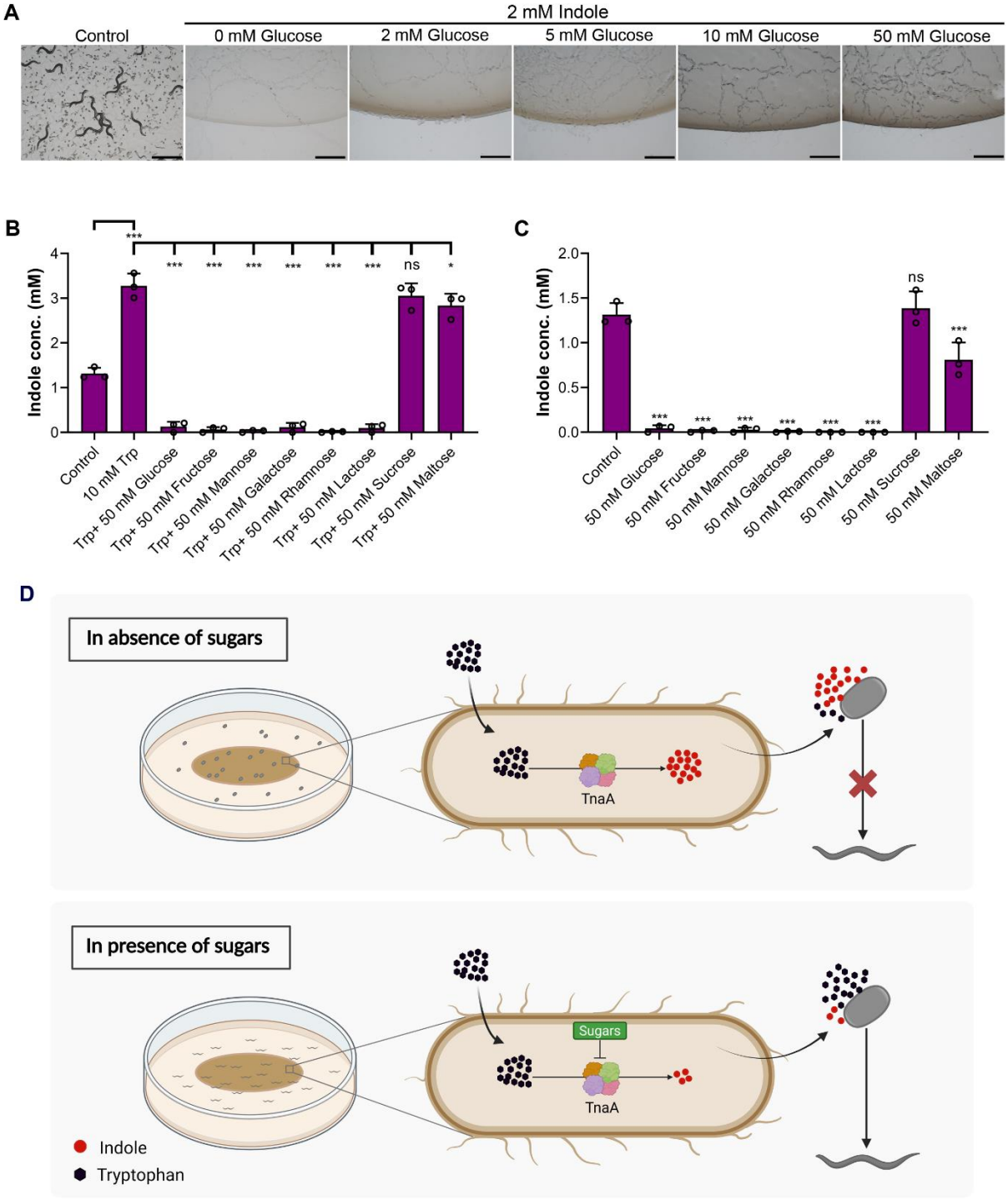


Figure 6. Sugars alleviate tryptophan toxicity by suppressing bacterial indole production

(A) Representative images of N2 worms grown on *E. coli* OP50 diet for 96 hours at 20°C containing 0 mM indole, 2 mM indole, and 2 mM indole supplemented with increasing concentration of glucose. Scale bar = 1 mm.

(B) Quantification of indole produced by *E. coli* OP50 cultured without any supplement, with 10 mM tryptophan (Trp), and with 10 mM Trp plus 50 mM of various sugars. *$P < 0.05$,

731 ***$P < 0.001$, and n.s., nonsignificant via ordinary one-way ANOVA followed by Dunnett's
732 multiple comparisons test ($n = 3$ biological replicates).

733 (C) Quantification of indole produced by *E. coli* OP50 cultured without any supplement and
734 with 50 mM of various sugars. ***$P < 0.001$ and n.s., nonsignificant via ordinary one-way
735 ANOVA followed by Dunnett's multiple comparisons test ($n = 3$ biological replicates). The
736 control data in (B) and (C) are the same.

737 (D) Model showing that the bacterial metabolism of tryptophan causes toxicity in *C elegans*.
738 *E. coli* takes up tryptophan and catabolizes it to indole via tryptophanase (TnaA). Indole can
739 freely diffuse through lipid membranes and prevents hatching of worm eggs at high
740 concentrations. The addition of sugars decreases indole production from tryptophan by
741 inhibiting TnaA in *E. coli*, resulting in the hatching and development of *C. elegans* in the
742 presence of excessive tryptophan.

743 **Supplementary Figures**

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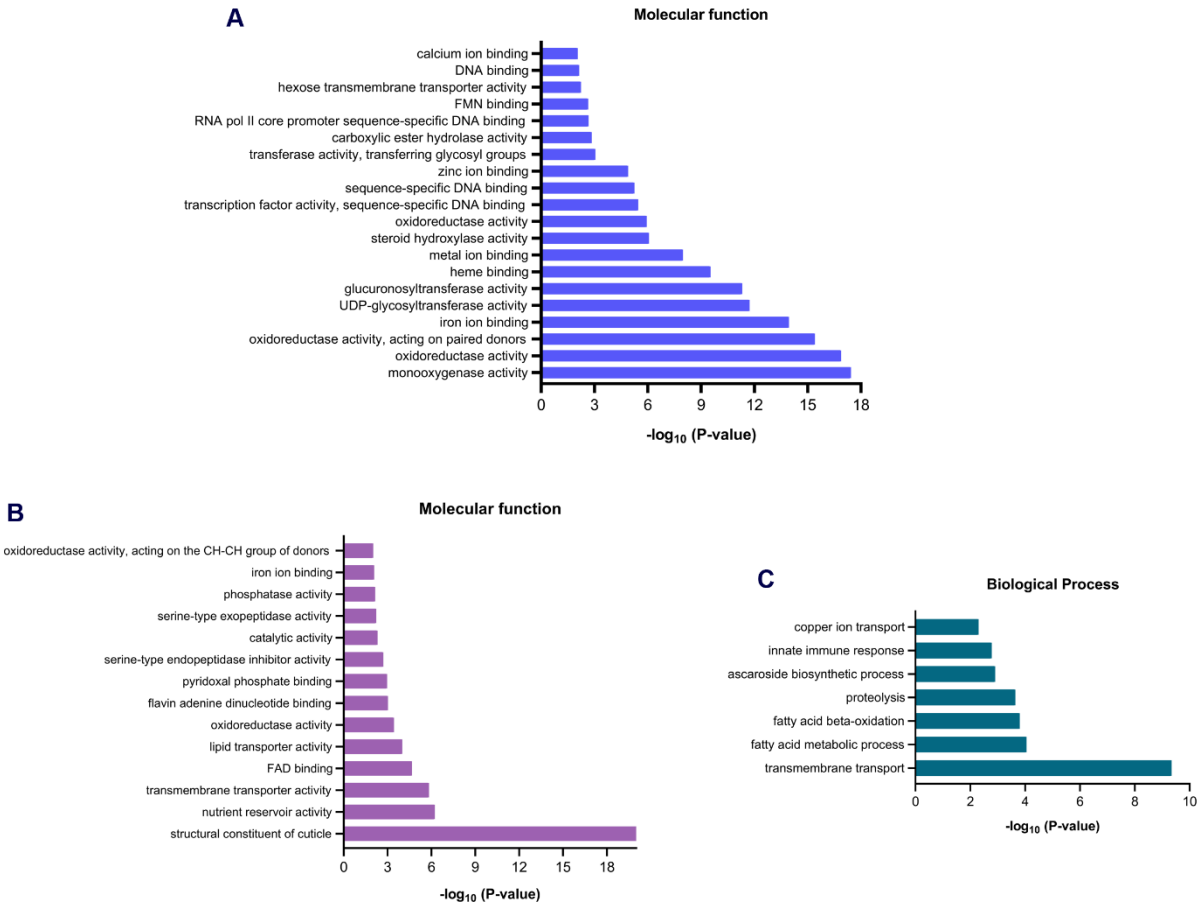

746

747 **Figure S1. Gene ontology analysis for genes differentially regulated upon tryptophan**
748 **exposure**

749 (A) Gene ontology enrichment analysis of genes upregulated upon tryptophan exposure for
750 molecular function.

751 (B)-(C) Gene ontology enrichment analysis of genes downregulated upon tryptophan
752 exposure for molecular function (B) and (C) biological processes.

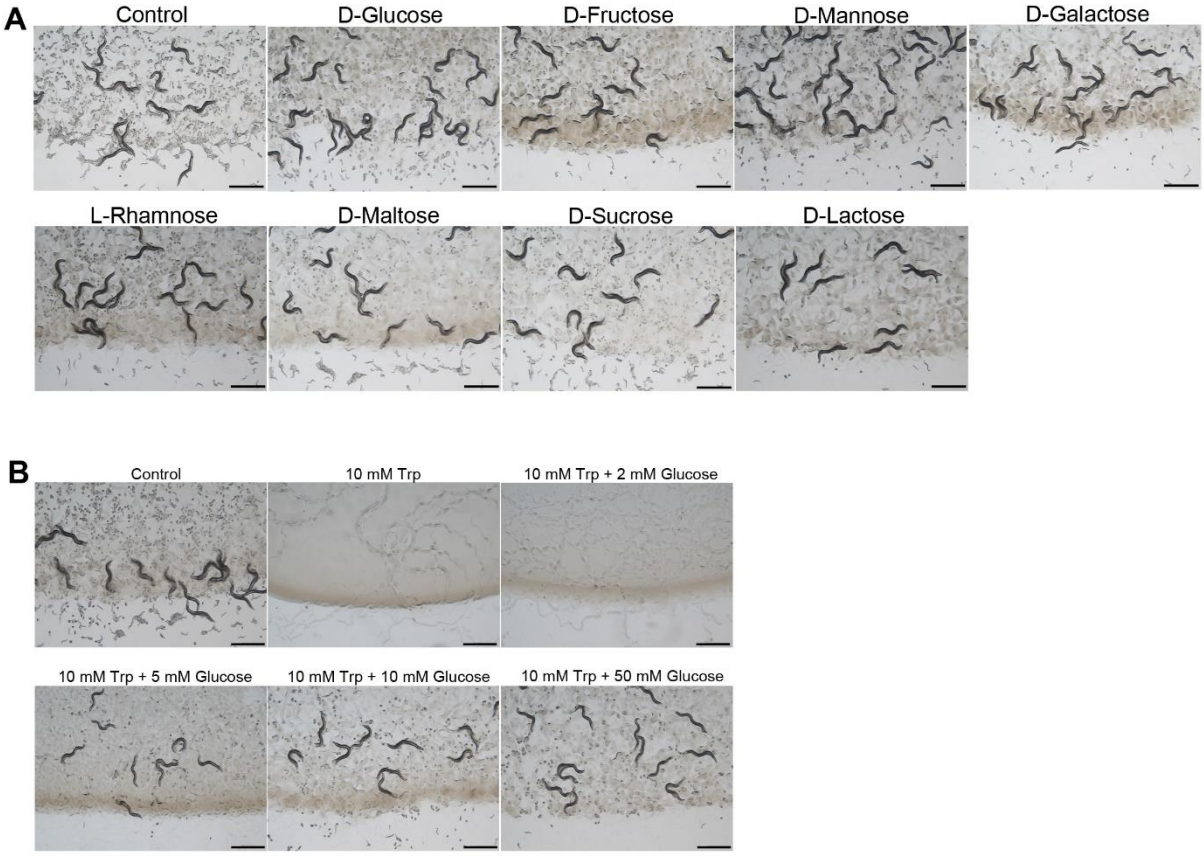

Figure S2. Sugars rescue tryptophan toxicity

(A) Representative images of N2 worms grown on *E. coli* OP50 diet for 96 hours at 20°C supplemented with 50 mM of different sugars. Scale bar = 1 mm.

(B) Representative images of N2 worms grown on *E. coli* OP50 diet for 96 hours at 20°C containing 0 mM Trp, 10 mM Trp, and 10 mM Trp supplemented with different concentrations of glucose. Scale bar = 1 mm.

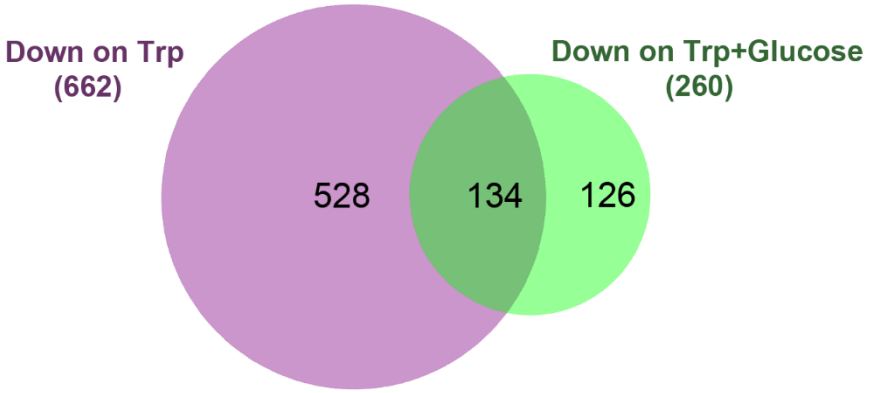

Figure S3. Sugars rescue tryptophan toxicity

Venn diagram showing the overlap between the genes downregulated upon exposure to 2 mM Trp alone and 2 mM Trp plus 50 mM glucose.

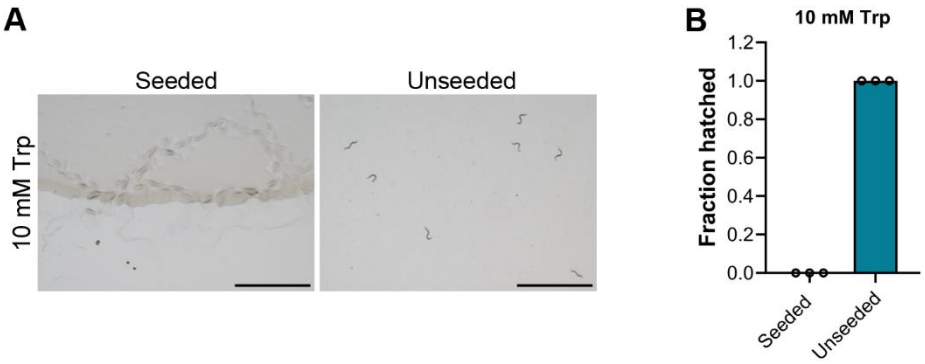

Figure S4. Bacterial metabolism of tryptophan is essential for its toxicity in *C. elegans*

(A) Representative images of N2 worms after 24 hours of hatching at 20°C on unseeded and *E. coli* OP50-seeded plates supplemented with 10 mM tryptophan (Trp). Scale bar = 1 mm.

(B) Quantification of hatched eggs of N2 worms after 24 hours of hatching at 20°C on unseeded and *E. coli* OP50-seeded plates supplemented with 10 mM Trp. (n = 3 biological replicates).

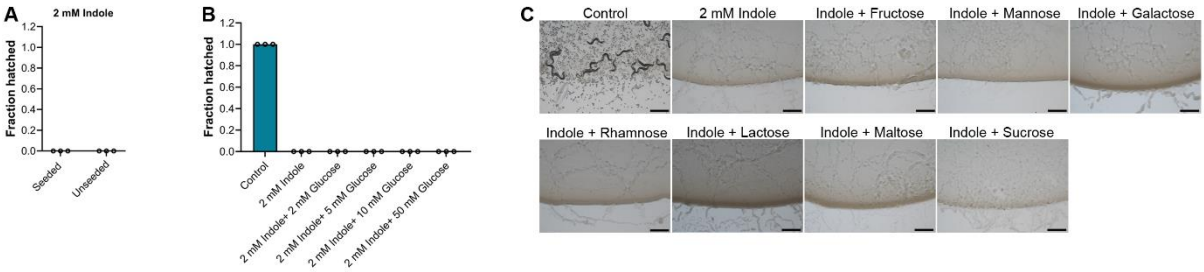

Figure S5. Sugars do not alleviate indole-induced toxicity in *C. elegans*

(A) Quantification of hatched eggs of N2 worms after 24 hours of hatching at 20°C on unseeded and *E. coli* OP50-seeded plates supplemented with 2 mM indole. (n = 3 biological replicates).

(B) Quantification of hatched eggs of N2 worms grown on *E. coli* OP50 diet for 24 hours at 20°C containing 0 mM indole, 2 mM indole, and 2 mM indole supplemented with different concentrations of glucose. (n = 3 biological replicates).

(C) Representative images of N2 worms grown on *E. coli* OP50 diet for 96 hours at 20°C containing 0 mM indole, 2 mM indole, and 2 mM Indole supplemented with 50 mM of different sugars. Scale bar = 1 mm.

Supplementary Table

Table S1. Differentially-regulated genes in adult N2 worms exposed to 2 mM tryptophan for 6 hours. A separate Excel file is provided.

Table S2. Differentially-regulated genes in adult N2 worms exposed to 2 mM tryptophan plus 50 mM glucose for 6 hours. A separate Excel file is provided.

Table S3. Comparison of genes upregulated in N2 worms upon exposure to tryptophan alone and tryptophan plus glucose. A separate Excel file is provided.

Table S4. Comparison of genes downregulated in N2 worms upon exposure to tryptophan alone and tryptophan plus glucose. A separate Excel file is provided.