Effect of carcass contamination on necrophagous invertebrate performance

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Abstract

Background: The breakdown of dead organic matter is driven by a diverse array of organisms and is an important process increasingly impacted by a range of contaminants. While many studies have documented how contaminants affect food webs that are fueled by decaying plant litters, much less is known about how contaminants affect organisms that rely on dead animal material. Here, we begin to explore the effects of food contamination—using silver nanoparticles (AgNPs) as a model contaminant—on the carrion beetle Nicrophorus vespilloides that buries carcasses of small vertebrates in soils as food source and larval nursing grounds.

Results: Our data show that a single ingestion of a non-lethal dose of 1 μg mL−1 AgNPs by adult female beetles does not affect overall gut microbial activity but results in shifts in the gut microbial community composition towards pathogens including Alcaligenes, Morganella, and Pseudomonas. While no effects were observed in offspring clutch size, some reductions were visible in clutch weight, number of larvae, and number of eclosing pupae in exposed N. vespilloides in comparison with controls. Repeated ingestion of AgNPs over several weeks led to a decrease in survival of adult beetles, suggesting that more environmentally realistic exposure scenarios can directly affect the success of carcass-feeding animals.

Conclusions: Sub-lethal carcass contamination with a model pollutant can affect the gut microbial composition in female beetles and reduce offspring fitness. This encourages consideration of currently overlooked propagation routes of contaminants through necrophagous food webs and inherent consequences for ecological and evolutionary processes.

Keywords: Gut microbiome, Maternal transfer, Offspring fitness, DECOTAB, Nanoparticles, Bactericidal compounds, Carrion beetle, Nicrophorus vespilloides

Introduction

Dead organic matter fuels food webs by serving as a food source for a diverse array of organisms, ranging from microorganisms and macroinvertebrates to scavenging megafauna. While most of the dead organic matter originates from senescent plant materials, carcasses of organisms provide a particularly high-quality and seemingly preferred resource to a variety of organisms and create an intense, localized pulse of carbon and nutrients into the environment (e.g., Wilson and Wolkovich 2011; Schrama et al. 2017). The continuous release of chemicals into the environment by humans, however, can directly affect an organism’s health or indirectly affect organisms by reducing the quality of their resources as contaminants accumulate in organic matter (Theng and Yuan 2008). This potentially provides a substantial propagation route of contaminants through food webs. Indeed, many studies have documented how contaminants can accumulate in plant matter-fueled food webs and affect consumers that feed on living as well as dead plant material (e.g., Judy et al. 2011; Hunting et al. 2016). Yet, little is known about how contaminants affect food webs that rely on dead animal material.

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Some studies have addressed this issue and showed that anti-inflammatory drugs and antibiotics present in cattle carcasses can accumulate in organisms (e.g., vultures) that feed directly on dead organic matter (Oaks et al. 2004; Blanco et al. 2019, 2016). In addition to direct consumption, some animals are reliant on carcasses to rear their offspring, and it is conceivable that contaminants can adversely affect such life history strategies and characteristics. A relatively well-studied example of such a necrophagous organism is the carrion beetle *Nicrophorus vespilloides* that lives in forests where it seeks and buries small vertebrate carcasses (e.g., rodents or birds). These carcasses are scarce but essential for reproductive success. The carcasses are intricately prepared as a rearing ground for their larvae through the manipulation of the carcasses’ microbial community. This occurs through the excretion of antimicrobial compounds and the inoculation of the beetles’ well-defined gut microbiome that is transmitted both vertically from the parents and horizontally from the carcass (Kaltenpoth and Steiger 2014; Wang and Rozen 2017). Here, the microbiota of carrion beetles is particularly relevant for their offspring as it enhances defense against pathogens and preserves carrion, thereby aiding in the development of larvae (e.g., Kaltenpoth and Steiger 2014; Wang and Rozen 2017; Shukla et al. 2018). Provided that gut microorganisms are generally recognized to fulfill many pivotal roles in host functions essential for fitness (Bright and Bulgheresi 2010; Heijtz et al. 2011; Hsiao et al. 2013; Kau et al. 2011; Engel and Moran 2013; Funkhouser and Bordenstein 2013), alterations in the gut microbiota can conceivably have adverse effects on their host and offspring.

Contaminants of particular interest are pollutants with antimicrobial properties, including antibiotics, heavy metals, persistent organic pollutants, and pesticides (Jin et al. 2017). Although most of these environmental pollutants do not directly target the gut microbiome, their major route of uptake is through ingestion associated with food items that indirectly target the gut microbiota. Previous studies indeed confirm that contaminants can accumulate in the guts of animals, thereby potentially affecting the gut microbiota of the host (Ding et al. 2019) which could translate to adverse effects on invertebrate fitness. The dysbiosis of the maternal microbiome potentially translates to fewer resources being invested in producing offspring or adverse effects on offspring in growth and survival (Arce et al. 2012; Kaltenpoth and Steiger 2014; Wang and Rozen 2017). However, studies hitherto report contradictory effects with respect to the significance of microbiomes for their host’s health (e.g., mice, zebrafish, and daphnids), ranging from no compositional changes of the microbiome to severe adverse effects on reproduction and growth (Merrifield et al. 2013; Sison-Mangus et al. 2014; van den Brule et al. 2015; Wilding et al. 2016).

This study centers on the hypothesis that the accumulation of contaminants in organisms can have postmortem effects on the organisms that use these carcasses as food and nursery grounds. To begin to test this assumption, we used the carrion beetle *N. vespilloides* in well-controlled laboratory incubations and assessed the effects of one feeding episode on a mimicked carcass with sub-lethal contaminant concentrations on (1) maternal microbiome composition and metabolic activity and (2) fitness parameters of their offspring. Subsequently, the effect of chronic consumption of a contaminated food source on the survival of adults was assessed.

**Materials and methods**

**Experimental setup**

The carrion beetles *Nicrophorus vespilloides* (Herbst, 1783) used in this study were caught from a wild population in 2014 (May to June) from Warmond (The Netherlands) and ever since maintained in the laboratory in unsterilized commercially available garden soil (baseline). Mating between sibling pairs was avoided to maintain an outbred population (Mattey et al. 2013). Beetles were reared in climate chambers at 20 °C under a constant light-dark cycle (15 h light-9 h dark) and were fed chicken liver at weekly intervals.

Exposure diet was prepared using Decomposition and Consumption TABlets, or DECOTABs in short, that can embed a homogenized and standardized mixture of contaminants in a standardized substrate (Kampfraath et al. 2012; Van der Lee et al. 2020). DECOTABs were prepared as 1-mL frozen tablets using an acrylic mold with a mixture of chicken liver and cornstarch (Kampfraath et al. 2012). Silver nanoparticles (AgNPs) were used as a model toxic compound as they represent an emerging class of toxicants (Fabrega et al. 2011; Gottschalk et al. 2013; Pillai et al. 2014). AgNPs have known antimicrobial properties in which they affect the permeability and respiration of the bacterial cell, ultimately leading to cell death (Dakal et al. 2016) and shifts in bacterial community composition (Zhai et al. 2016; Tili et al. 2017; Zhai et al. 2018). AgNPs with a nominal particle size of 15 nm were purchased from Nanostructured & Amorphous Materials (Houston, USA). Characterization of the particle morphology of the AgNP is described elsewhere (Zhai et al. 2018). It is technically challenging to disentangle the relative contributions of nanoparticles and ions within complex systems such as the food item served here (chicken liver); hence, the agglomeration dynamics and ratio of AgNPs and ions were not assessed in the DECOTABs. To confirm that AgNPs were actually ingested, the guts (control n = 16; AgNP 1 μg mL⁻¹, n = 17) were extracted and homogenized using a conical plastic pestle in 2.0-mL Eppendorf with 1.5 mL sterile phosphate buffer saline (PBS; 100 nM; pH
7.2). The samples of the gut mixture were acidified with a mixture of 5% HNO₃ and PBS adding up to 3 mL. The concentration of silver was then assessed by graphite furnace atomic absorption spectrometry (GF-AAS), and standards of silver nitrate (0.1, 1, 10, 100 µgL⁻¹) were used to establish a calibration curve.

**Experimental design**

Intergenerational effects of consumption of AgNPs on offspring fitness were evaluated by serving DECOTABs with 1 µgL⁻¹ AgNPs to adult female beetles and compared these with a control group (n = 20 per group) that were served DECOTABs without AgNPs. In a preliminary study, three AgNP exposure concentrations (0.1, 1, and 5 µgL⁻¹) were tested for lethality within 7 days (n = 20, data not shown). While none of the exposures resulted in mortality, the highest concentration reduced the beetle activity, and therefore, all subsequent experiments were performed using 1 µgL⁻¹. Under natural conditions, carrion beetles give bi-parental care to their brood at the pre-hatching stage (Steiger et al. 2011). However, to reduce variability, the setup includes maternal care only throughout all stages. Before exposure to AgNPs, we ensured that the weight of the female beetles did not differ between the control and exposure groups (control vs exposure, 0.194 ± 0.037 (STD) vs 0.197 ± 0.044 (STD) µg; Student t-test: n = 20, p = 0.810). After female beetles were assigned to a treatment, the unmated female beetles were put in Petri dishes containing a layer of filter paper soaked with 1-mL water to keep the air moist and were then left at 20°C for 2 days to consume the DECOTABs (see Fig. 1 for a conceptual presentation of the experimental outline). After consuming the DECOTABs, the female was placed in a Petri dish with an adult male to mate for 24 h, after which the female was placed in a plastic container of 15 × 10 × 5 cm containing 30-cm³ unsterilized potting soil and a mouse carcass weighing between 16 and 31 g. Mouse carcasses were kindly provided by the laboratories of Leiden University Hospital.

**Effect of carcass contamination on parental microbiome composition and metabolic activity**

The overall respiratory capacity of the gut microbiome of the control and AgNP-exposed beetles was approximated by dehydrogenase enzyme activity via the reduction of 2-(p-iodophenyl)-3-(p-nitrophenyl)-5-phenyl tetrazolium chloride (INT) to formazan (INTF). INT successfully competes with NADH⁺ and NADPH⁺ and therefore INT reduction is a common assay of respiratory activities under both anaerobic and aerobic conditions (Smith and McFeters 1996). The guts (n = 10 per group) were extracted and homogenized in a 100-µL PBS solution using an autoclaved plastic pestle. This gut mixture was filtered through a cell strainer (40 µm Nylon) to obtain a clear solution containing microbial cells without gut tissue. To measure enzymatic activity, a 5-µL aqueous INT solution (200 µL; 1 mM with 0.05% dimethyl sulfoxide (DMSO)) was added to each gut extract, consisting of 5 µL of strained gut solution from
individual beetles. Samples were incubated in a dark room at 22 °C for 3 h, and thereafter, enzyme activity was stopped and INT formazan crystals were solubilized by adding 5 μL DMSO. Absorbance reflecting the dehydrogenase activity was measured using a NanoDrop 2000/2000c at 585 nm and corrected for microbial cell density measured at 600 nm (gut solution without INT), in which both were baseline-corrected using the aforementioned incubation solution without microbial cells (Hunting et al. 2010). Microbial density was found to be variable among the guts of individual beetles, and therefore, the microbial activity of the exposed beetles and the control group was expressed as dehydrogenase activity measured at an absorbance at 585 nm relative to microbial cell density measured at 600 nm to obtain a measure of metabolic activity per unit cell density, in which relative activity equals 0.1 × (A_{585}/A_{600}).

To obtain a qualitative structural account of the effect of AgNPs on the microbial gut composition, we used CHROMagar Orientation. CHROMagar allows differentiation and presumptive identification of bacteria. CHROMagar Orientation was purchased from the manufacturer CHROMagar (Paris, France) in powder base (agar, 15 g L\(^{-1}\); peptone and yeast extract, 17 g L\(^{-1}\); chromogenix mix, 1 g L\(^{-1}\)). The plates were made with 33 g of CHROMagar Orientation powder base dissolved in 1 L of purified water. The guts of the exposed beetles (n = 4) and the control (n = 5) were extracted and homogenized within a 750-μL PBS solution and then diluted with autoclaved ultrapure water to obtain a final volume of 5 mL. Each gut was then plated on 3 CHROMagar plates and incubated for 24 h at 32 °C, after which the colony-forming units (CFU) were determined and counted, and the mean of the three plates per gut was calculated.

To assess whether the gut microbial communities were different depending on treatment, we assessed substrate utilization profiles using commercial EcoPlates (Biolog, Hayward, USA). Each well plate contains 3 replicates comprising 31 different carbon substrates which are ecologically relevant and structurally diverse compounds. Since it does not include specific compounds typical of the food items used in this study, it remains impossible to directly relate substrate utilization profiles to the actual functioning of the microbiome. Nonetheless, the number of substrates used can serve as a proxy for the (metabolic) diversity and resource niche breadth of the microbial community (Garland 1999; Hunting et al. 2013; Krause et al. 2014; Hunting et al. 2017). For this experiment, the gut microbial community of beetles that ingested chicken liver laced with AgNP (1 μg mL\(^{-1}\), n = 4) was compared to the microbial community of control beetles (n = 5) that ingested chicken liver only. The guts were extracted and homogenized in 700 μL PBS and 4300 μL purified water, which then was dispersed between 3 replicates with 50 μL in each well. The Eco-Plates were then incubated at 20 °C for 5 days in dark conditions and thereafter measured at 600 OD using a standard plate reader.

**Offspring fitness**

After exposure and mating, the female was left to take care of her young until the offspring started to disperse from the mouse carcass to go into the pupal stage (Mattey et al. 2013). The time of dispersal varied ranging from 9 to 10 days post-hatching, which was similar between treatments. We collected the following four fitness measures after mating control and exposed female beetles (n = 20 per treatment group) with unexposed males: (1) the number of eggs produced by the female beetles after the burial of the mouse, as counted at the bottom of the plastic container; (2) the number of larvae present after the larvae dispersed (by sorting through the soil); (3) the total brood weight of the larva at dispersal; and (4) the number of eclosing pupae (Fig. 1).

**Survival of adult beetles after chronic exposure**

A repeated exposure experiment was set up to assess chronic, and thereby more environmentally realistic, effects on survival. Female beetles were fed DECO TABs with or without 1 μg mL\(^{-1}\) AgNPs as described above, and the treatment was repeated every 7 days. The weight of the beetles in the first experiment was 0.230 ± 0.043 g (STD) for the control and 0.210 ± 0.051 μg (STD) for the exposed group (Student t-test p = 0.365), and in the second experiment, the weight was 0.280 ± 0.037 μg (STD) for the control and 0.270 ± 0.049 μg (STD) for the exposed group (Student t-test p = 0.786). The first experiment lasted for 36 days (n = 10 per group), and the second experiment had to be terminated due to unforeseen events at the laboratory after 27 days (n = 10 per group). The beetles were checked daily for survival.

**Statistical analysis**

To assess meaningful differences between the control and treatment groups taking into account the effect size, Ag concentrations in the gut, enzyme activity, and beetle offspring parameters were analyzed using estimation statistics and depicted as Gardner-Altman plots (Ho et al. 2019). For comparison, a two-tailed permutation t-test was performed. Both tests build upon resampling the measured data, and therefore, the data will approach a normal distribution even if the underlying population is not normally distributed. Differences in bacterial community substrate utilization profiles in exposed and control treatments were visualized using a principal component analysis (PCA) and tested performing a Euclidean-based ANOSIM in PAST (Villéger et al. 2013).
The differentiation of microbial species determined using CHROMagar was depicted in pie charts for visual comparison, and each species was tested for proportional differences between treatment and control using a Z-test. The survival of the female beetles in the chronic exposure was drawn using the Kaplan-Meier estimates of the survival functions and analyzed with a Mantel-Cox log-rank test. All statistics are summarized in Table 1.

Results

Effect of carcass contamination on parental gut microbial composition and activity

Silver was measured to accumulate in the guts of the exposed beetles where all concentrations of exposed beetle guts were clearly above the detection limit (0.026 ppb) and the concentrations of control beetle gut below the detection limit (Fig. 2). Microbial activity as shown using the results of the relative dehydrogenase activity was in a similar range in the exposed beetles compared to the control beetles (permutation t-test, \( p = 0.084 \); Fig. 3A). The (metabolic) composition of the gut microbial community was different between exposed and control treatments (Fig. 3B, one-way ANOSIM, Euclidean-based similarity, \( n = 4 \), \( R = 0.969 \), \( p = 0.030 \)). The composition of the colony-forming units between both the treatments was different for a number of taxa, as shown using the results of the CHROMagar plates (Fig. 3C). Exposure to nanoparticles decreased the abundance of Candida and Serratia (Fig. 3C) while promoting the growth of Alcaligenes, Morganella, and Pseudomonas (Z-test, \( p < 0.05 \)).

Effects of carcass contamination on offspring fitness

There was no difference in clutch sizes between the exposed and control treatments (permutation t-test, \( p = 0.239 \); Fig. 4A). The development into larvae within 10 days showed a tendency of being reduced in the exposed group (permutation t-test, \( p = 0.050 \); Fig. 4B) as did the total larval brood weight in the exposed group in comparison with the control (permutation t-test, \( p = 0.037 \); Fig. 4C). In connection with those results, the number of larvae reaching the pupal stage was similarly reduced (permutation t-test, \( p = 0.111 \); Fig. 4D). The larval weight did not appear to vary with carcass weight (\( y = 0.0008713x + 0.15753, R^2 = 0.0063, p = 0.618 \)).

Effects of carcass contamination on adult survival

Continuous exposure to AgNP over the course of 4 weeks impaired the survival of the beetles (Fig. 5; pool of two experiments, Mantel-Cox log-rank test, \( p = 0.016 \)).

Discussion

This study explored the direct and indirect effects of carrion contamination with a bacteriocidal compound (AgNP) on the gut microbiome and offspring fitness of the carrion beetle N. vespilloides. Our findings indicate that, at levels that do not result in acute toxicity, a single feeding event on an AgNP-contaminated food source can influence the composition of the beetle microbiome without clearly affecting its overall metabolic activity. The resulting alteration of the gut microbiome potentially translates to the observed subsequent effects on their offspring which rely on the maternal transmission of the microbiome and may be related to the observed increased mortality of adult beetles in chronic exposure.

While a single exposure to silver nanoparticles did not appear to alter the overall gut microbial activity, a compositional change was observed in gut microbial communities in N. vespilloides. The abundance of certain strains such as Serratia sp., a group of gram-negative bacteria that includes opportunistic insect pathogens

**Table 1** Summary of statistical test results

| Assay                         | \( n \) (control) | \( n \) (exposed) | Effect size [CI width lower bound; upper bound] | Two-sided permutation t-test | Euclidean-based ANOSIM | Log-rank (Mantel-Cox) test |
|-------------------------------|-------------------|-------------------|-----------------------------------------------|-----------------------------|-------------------------|----------------------------|
| Gut Ag concentration          | 16                | 17                | 1.36 [95CI 1.07; 1.65]                           | \( p = 0.000 \)             |                         |                             |
| Dehydrogenase activity        | 10                | 10                | \(-0.00724 [95CI −0.0147; −0.000163]\)          | \( p = 0.084 \)             |                         |                             |
| Biolog EcoPlate               | 5                 | 4                 |                                               | \( p = 0.030 \)             |                         |                             |
| Offspring fitness             |                   |                   |                                               |                             |                         |                             |
| Eggs per clutch               | 20                | 20                | \(-3.5 [95CI −8.9; 2.46]\)                     | \( p = 0.239 \)             |                         |                             |
| Larvae emerging               | 16                | 19                | \(-6.82 [95CI −13.3; −0.694]\)                 | \( p = 0.050 \)             |                         |                             |
| Clutch weight                 | 16                | 19                | \(-0.973 [95CI −1.9; −0.169]\)                 | \( p = 0.037 \)             |                         |                             |
| Pupal stage                   | 16                | 19                | \(-6.82 [95CI −13.3; −0.694]\)                 | \( p = 0.111 \)             |                         |                             |
| Chronic mortality             | 20                | 20                |                                               |                             |                         | \( p = 0.016 \)             |
(Flyg et al. 1980; Motta et al. 2018), was decreased while the growth of other strains exhibited a relative increase. Such differences are potentially due to the sensitivity of certain bacterial strains to AgNPs, particularly gram-negative bacteria (Choi et al. 2008; Ip et al. 2006). This result is in line with an increasing number of studies that demonstrate the consequences of environmental contaminant exposure can include shifts in symbiont and pathogen abundances (Motta et al. 2018). This can have potential implications for the metabolism as an altered bacterial composition may affect the success in extracting nutrients from the food provided and have subsequent knock-on effects on fitness (Cornman et al. 2016; Gupta et al. 2011; Kaltenpoth and Steiger 2014; Lehman et al. 2009). However, the actual underlying mechanisms often remain poorly understood, and this remains a key challenge for future research. Since the gut microbiome is likely more diverse in wild specimens.

![Fig. 2](image_url) Concentration of silver in control and AgNP-exposed carrion beetle guts. All values for controls were below the detection limit.

![Fig. 3](image_url) Effect of carcass contamination on gut microbial community activity and composition. A Microbial activity (expressed as the relative dehydrogenase activity) for control and the exposed beetles (data point in parentheses was excluded from analysis; n = 9 for control, n = 10 for exposed, p = 0.084). B Statistical separation of bacterial community (metabolic) composition in exposed and control treatments as illustrated by a PCA analysis of the Biolog substrate utilization profiles of the bacterial communities (n = 5 for control, n = 4 for exposed, ANOSIM p = 0.030). C Gut microbial composition of control beetles (left, n = 5) and exposed beetles (right, n = 4) presented as percentages of the overall number of colony-forming units for each of the treatments on CHROMagar plates. Asterisks indicate that no unit of that taxa was found. Assessment of the differences in species between control (left, n = 5) and treatments (right, n = 4) suggests nanoparticles decrease the abundance of Candida and Serratia while promoting the growth of Alcaligenes, Morganella, and Pseudomonas (Z-tests scoring p < 0.05).
of *N. vespilloides*, they may compensate for shifts in abundances and metabolic activities. Despite this, the observed shift towards pathogenic microorganisms hints that contaminated food items could potentially have adverse health effects for individual beetles and their offspring or other scavenging organisms.

Beetles accumulated nanoparticles and associated free silver species in their intestinal tract (Fig. 2), and several tendencies of reduced fitness of the subsequent generation were observed (Fig. 4). Since we only used a single feeding event, it should be considered that an environmentally more realistic scenario includes the repetitive ingestion of contaminated food items. This could result in more severe effects, as visible in the observed increased mortality of adult beetles in chronic exposure (Fig. 5). The mechanisms driving the adverse effects on the offspring remain uncertain and could offer avenues for further research. For instance, particles might be excreted when the mother treats the carcass with anal secretions where silver could accumulate to potentially toxic levels for the larvae. Alternatively, a change in the

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**Fig. 4** Fitness measures in the offspring of *N. vespilloides* in AgNP-exposed and control beetles. A Number of eggs (*n* = 20 female beetles per treatment group, *p* = 0.239). B Number of larvae per clutch (parentheses indicate 4 broods of controls and 1 brood of exposed that have failed and were excluded from the analysis, *p* = 0.050). C Weight of the clutch (failed broods excluded, *p* = 0.037). D Number of larvae pupating (failed broods excluded, *p* = 0.111)

**Fig. 5** Survival after chronic exposure. Survival for control and exposed beetles was assessed over 36 (experiment 1) and 27 (experiment 2) days. Shown is the pooled data of both experiments (*n* = 20, *p* = 0.016)
microbiota may lead to behavioral changes in the female beetle which can potentially affect brood care. Likewise, silver nanoparticles could have affected the mother’s health via mechanisms other than dysbiosis of the gut microbiome, or an inhibited or impaired microbiota might have prevented the maternal transfer of microorganisms critical to the healthy development of the offspring (Jacobs et al. 2014). It would be further interesting to directly expose the carcass that is used as rearing ground and assess its effect on offspring fitness independent of the exposed parents.

Using a simplified model setup mimicking a carcass-dependent food web, we observed that food contamination with a model antibacterial compound (AgNP) can alter the metabolic diversity of the burying beetle *Nicrophorus vespilloides* microbiome. The observed patterns on gut microbiomes likely hold true for myriad contaminants as many have antibacterial and antifungal properties. The resulting alteration of the gut microbiome could potentially be the cause of increased mortality of adult beetles in chronic exposure. A single feeding event was observed to have some apparent, albeit minor effects on offspring, but it should be considered that a more realistic repetitive exposure to contaminants could potentially coincide with a stronger negative effect on the fitness of their offspring. The observed effects in our model necrophagous food web may be relevant for other scavengers that rely on the maternal transfer of microbiomes and encourage consideration of currently overlooked propagation routes of contaminants through food webs.

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Authors' contributions
EH: conceptualization, experimental design, data analysis, writing, and editing. MS: data analysis, writing, and editing. DR: experimental design and editing. EH: conceptualization, experimental design, data analysis, writing, and editing. The authors read and approved the final manuscript.

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Competing interests
The authors declare that they have no competing interests.

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