INTRODUCTION

In the course of ongoing global change, an ever-increasing number of species is threatened by habitat loss and fragmentation (Haddad et al., 2015; Murphy, Battocletti, Tinghitella, Wimp, & Ries, 2016; Swift & Hannon, 2010). This habitat change can entail a severe reduction in the size of natural populations and simultaneously increase the degree of isolation among them (Agnarsson, Avilés, & Maddison, 2013; Bates, Sadler, et al., 2014). Strong population declines are typically associated with genetic drift and a loss of genetic variation, which can be amplified by restricted gene flow among populations, if habitats exhibit low connectivity (Aguilar, Quesada, Ashworth, Herrerías-Diego, & Lobo, 2008; Murphy et al., 2016). Such demographic events can reduce the viability of populations, as they increase the likelihood of matings among closely related individuals (i.e., inbreeding), which can have highly detrimental effects on...
Inbreeding can induce manifold detriments in affected organisms, which typically culminate in reduced survival and/or reproductive success of inbred relative to outbred individuals (Fox, Scheibly, Smith, & Wallin, 2007; Harano, 2011; Lihoreau, Zimmer, & Rivault, 2007). This loss of individual fitness following inbreeding is referred to as inbreeding depression. Inbreeding depression arises from an increase in genome-wide homozygosity in the offspring generation, which results in the increased phenotypic expression of recessive deleterious mutations and the reduced expression of heterozygote advantage (Charlesworth & Willis, 2009; Keller & Waller, 2002). Inbreeding depression can become evident in the offspring of only one generation inbreeding (Müller & Müller, 2016) or after several generations inbreeding (Bilde, Maklakov, & Schilling, 2007). Moreover, inbreeding can also influence traits not closely linked to fitness, such as the chemical (Menzel, Radke, & Foitzik, 2016) or behavioral phenotype of insects (Müller & Juškauskas, 2018; Pilakouta & Smiseth, 2017; Richardson & Smiseth, 2017).

Given the high inbreeding costs (inbreeding depression) and the comparably low probability to benefit from inbreeding (reproductive assurance in the absence of non-related mating partners and inclusive fitness benefits; Peer & Taborsky, 2005; Kokko & Ots, 2006), many animals evolved mechanisms to avoid the mating with close relatives (Pusey & Wolf, 1996). These avoidance mechanisms come into action either before (precopulatory, e.g., Liu, Tu, He, Chen, & Xue, 2014) or after (postcopulatory, e.g., Simmons, Beveridge, Wedell, & Tregenza, 2006; Bretman, Newcombe, & Tregenza, 2009) inbreeding events. Precopulatory mechanisms base on the discrimination of kins as mating partners (Lihoreau & Rivault, 2009; Metzger, Bernstein, Hoffmeister, & Desouhant, 2010; Whitehorn, Tinsley, & Goulson, 2009). Postcopulatory mechanism can rest, for example, on kin sperm discrimination (Bretman et al., 2009; Welke & Schneider, 2009), that is, a lowered transfer of sperm if females mate with closely related males (Lewis & Wedell, 2009). Family-specific cuticular hydrocarbon (CHC) (Lihoreau & Rivault, 2009) or pheromone patterns (Herzner, Schmitt, Heckel, Schreier, & Strohm, 2006) partly serve as recognition cues on which inbreeding avoidance mechanisms are based on (Lihoreau & Rivault, 2009; Thomas & Simmons, 2011). However, pre- or postcopulatory kin discrimination as inbreeding avoidance mechanisms is not necessarily expressed in all species suffering from inbreeding depression (Bouchebti, Durier, Pasquaretta, Rivault, & Lihoreau, 2016; Edvardsson, Rodriguez-Munoz, & Tregenza, 2008). As alternative or in combination, the costs of inbreeding can be reduced by a polyandrous mating system (Bayoumy, Michaud, & Bain, 2015; Duthie, Bocedi, & Reid, 2016; Tregenza & Wedell, 2002). Moreover, a specific dispersal strategy can lead to inbreeding avoidance by preventing the encountering of closely related individuals (Pusey & Wolf, 1996).

In this study, we investigated the effects of inbreeding over the lifetime of *Phaedon cochleariae* F. (Coleoptera: Chrysomelidae, Figure 1). We determined the larval development time, larval and adult survival, adult body mass, and reproductive output of outbred versus inbred mustard leaf beetles. Moreover, we conducted a mating assay in order to test whether postcopulatory inbreeding avoidance mechanisms occur in our study species. We sequentially mated females with two nonsiblings, two siblings, a nonsibling and a sibling, and a sibling and a nonsibling. The reproductive output of these distinct mating combinations was detected by determining the larval hatching rate of eggs laid by the females.

In a previous study on *P. cochleariae*, we disclosed that the larval hatching rate of inbred offspring is decreased in comparison with outbred offspring (Müller & Müller, 2016). Thus, this beetle species suffers from inbreeding depression and we consequently expected negative inbreeding effects on other fitness-related traits throughout its ontogeny. Moreover, the CHC profile, which is decisive for mate choice in *P. cochleariae* (Geiselhardt, Otte, & Hilkner, 2012) and other insect species (Howard & Blomquist, 2005), differs between families (Müller & Müller, 2016). Although CHCs can function as mechanism to discriminate between closely related and unrelated individuals (Weddle, Hunt, & Sakaluk, 2013), the family-specific CHC patterns in *P. cochleariae* do not mediate precopulatory inbreeding avoidance by a faster mating of nonsiblings compared to siblings (Müller & Müller, 2016). Thus, family-specific CHC profiles might serve as a recognition cue for postcopulatory inbreeding avoidance. Here, we specifically tested whether such postcopulatory inbreeding avoidance mechanisms exist in *P. cochleariae*. In detail, we hypothesized a lower hatching rate of larvae from eggs laid by females that were sequentially mated with two siblings compared to a mating with two nonsiblings or a nonsibling and a sibling (or vice versa). Such patterns would indicate the existence of sibling sperm recognition and discrimination (Tregenza & Wedell, 2002).
2 | MATERIALS AND METHODS

2.1 | Study organisms and rearing

*Phaedon cochleariae* individuals of our laboratory strain descended from different regions in Germany, that is, side arms of the river Main, Botanical Garden of Berlin-Dahlem and the south of Bielefeld along the Furlbach. This mixed strain, which consisted of 500–600 individuals, was reared for approximately 25 generations under laboratory conditions, where individuals mated randomly. Every year, 100–150 wild *P. cochleariae* individuals were collected at the Bielefeld location and were integrated into the strain in order to refresh the gene pool. Beetles were reared in separate ventilated plastic boxes (20 × 20 × 6.5 cm) in a climate cabinet (20°C, 16:8 hr light:dark, 65% relative humidity) with a density of about 100–200 individuals per box and a nearly balanced sex ratio. The beetles were fed ad libitum with leaves of 8–10 weeks old *L. pekinensis* var. Michihili plants (seeds received from Kiepenkerl; Bruno Nebelung GmbH, Konken, Germany). Plants were grown in a greenhouse (16:8 hr light:dark, 60% relative humidity) in pots (12 cm diameter) filled with composted soil.

2.2 | Experiment 1: measurements of larval development time, larval and adult survival, adult body mass, and reproductive output

Pupae originating from three different rearing boxes were collected and each pupa and later each young adult was reared separately in one small petri dish (5.5 cm diameter). Thus, unintended mating combinations and sibling matings were prevented prior to the experiment. After adult emergence and sex determination, pairs of one female and one male, which descended from different rearing boxes, were transferred into small petri dishes (one pair per petri dish). These individuals served as parental (P) generation. In total, six different breeding pairs (corresponding to six families) were used. Eggs laid by the females were separately collected and larvae (F1 generation) were reared in large petri dishes (9 cm diameter) until pupation. Pupae were separated and the emerging F1 adults were again reared in a density of one individual per small petri dish. The F1 generation was used to examine the effects of inbreeding on various fitness-related traits over the lifetime of *P. cochleariae*. For this purpose, we performed experimental outbreeding and inbreeding with the F1 generation. Here, we set up pairs of one female and one male, which were either nonsiblings (i.e., outbreeding) or siblings (i.e., inbreeding) for each of the six families. We used two to four pairs per family × breeding treatment combination, depending on the availability of *P. cochleariae* individuals. All pairs were separately reared in small petri dishes for 14 days. During this time interval, each pair mated repeatedly and all females laid eggs. Subsequently, we collected the eggs produced by nonsibling pairs (i.e., outbred offspring) and sibling pairs (i.e., inbred offspring) and transferred them to new large petri dishes (one separate dish for eggs of each mating pair). Based on the performed breeding scheme, the parents of outbred offspring were neither full nor half siblings nor cousins and the parents of inbred offspring were all full siblings.

After hatching of the outbred and inbred larvae (F2 generation), the days until adult emergence were counted (i.e., development time). During that time, all larvae descending from identical parents were captured in one large petri dish with up to 18 individuals. Moreover, we determined the proportion of surviving larva until pupation (larval survival) as well as the proportion of viable adults directly after adult emergence (adult survival) per petri dish. Adult beetles were then transferred into new small petri dishes (one beetle per petri dish). At day 8–10 of adult life, the body mass of virgin males and females was determined, using a microbalance (ME36S, accuracy 0.001 mg; Sartorius AG, Göttingen, Germany). Afterwards, outbred and inbred females were mated with an outbred male of a different family. The number of eggs laid per female was counted over a period of 4 days. At the first day of counting, the females were 11–16 days old (predictor variable age). All F1 and F2 beetles were reared ad libitum on middle-aged cabbage leaf disks, which were replaced every to other day. All petri dishes were lined with a moistened filter paper to prevent desiccation.

2.3 | Experiment 2: effects of nonsibling and sibling mating partner order on larval hatching rate

In experiment 2, we investigated whether postcopulatory inbreeding avoidance mechanisms occur in *P. cochleariae*. For this purpose, we set up four mating assay treatments, using either nonsiblings (i.e., outbreeding) or siblings (i.e., inbreeding) from the F1 generation (see previous section) as mating partners. In the first treatment group, females were sequentially mated with two nonsibling males, in the second group first with a nonsibling and second with a sibling, in the third group first with a sibling and second with a nonsibling and in the fourth group sequentially with two sibling males. We used beetles of all families for each mating combination. For each treatment, F1 females and males were separately captured in small petri dishes until day 10 of adult life. Thereafter, the first virgin male was added to the petri dish of the virgin female at about 10.00 a.m. After 6 hr, this first male was removed and after additional 18 hr the second virgin male was added for 6 hr from 10.00 a.m. to 16.00 p.m. To prevent confounding effects of daytime on mating success (matings might be avoided in the evening hours, as beetles are mainly active from 10 a.m. to 16 p.m.), we did not add the second male at the same day after 16.00 p.m. We checked whether beetles mated during the time spent together in the petri dish. Three, 5 and 7 days after the last mating occurred, all eggs laid by a specific female in 24 hr were counted and removed from the petri dish. We determined the proportion of larvae hatched from all eggs laid per female individual (i.e., hatching rate). Beetles for experiment 2 were reared in the same way as for experiment 1 (see above).
2.4 Statistical analyses

All data on the performance of *P. cochleariae* from both experiments were statistically analyzed with mixed effects models (R-package: *lme4*, Bates, Maechler, Bolker, & Walker, 2014) in R version 3.2.3 (R Development Core Team, 2015). For responses with normal error distribution, we used linear mixed effects models (LMM), whereas for responses with Poisson or binomial error distribution, we used generalized linear mixed effects models (GLMM).

The model for the response development time (GLMM, Poisson) from experiment 1 comprised the fixed effects of breeding treatment and the random effects of family in P-generation and petri dish nested within family. The models for the responses larval survival (GLMM, binomial) and adult survival (GLMM, binomial) from experiment 1 included the fixed effect of breeding treatment and the random effect of family in P-generation. The model for the response body mass (LMM, Gaussian, log-transformed) from experiment 1 comprised the fixed effects of breeding treatment, sex (covariate) and the interaction among these factors as well as the random effect of family in P-generation. The model for the response egg number (GLMM, Poisson) from experiment 1 included the fixed effects of breeding treatment, female age (covariate) and their interaction as well as the random effect of family in P-generation. The model for the response larval hatching (GLMM, binomial) from experiment 2 included the fixed effects of mating treatment and the random effect family in P-generation.

All of the described models were fitted with a maximum likelihood approach. After ensuring that models exhibit variance homogeneity and normal distribution of residuals by the means of visual inspection, we applied stepwise backward model selection to obtain the minimal adequate model. Here, we removed fixed effect terms with *p* > 0.05 based on chi-square likelihood ratio tests (R-package: MASS, Venables & Ripley, 2000). In case a minimal adequate mixed model included fixed effect factors with more than two levels (mating treatment from experiment 2), we additionally performed multiple comparisons (Tukey post hoc tests) of all factor levels on the respective model (R-package: *multcomp*, Hothorn, Bretz, & Westfall, 2008). For illustration of the effects of breeding treatment (experiment 1) and mating treatment (experiment 2), we generated box-whisker plots, which additionally included least square means with their standard errors extracted from the respective minimal adequate mixed effects models (R-package: *lsmeans*, Lenth, 2016). In contrast to raw data means and their standard errors, these model estimates account for the specific error distribution of the response, for the effects of covariates as well as for random effects.

3 RESULTS

3.1 Inbreeding effects on fitness-related traits

The larval development time, larval survival rate, viability of freshly hatched adults, and the reproductive output of *P. cochleariae* were significantly affected by inbreeding (Figure 2, Tables 1, 2). The development time of inbred larvae was on average 1.5 days prolonged compared to outbred larvae (Figure 2a). The larval survival rate until adulthood was 37.6% lower in inbred (53.8% larvae survived) than in outbred individuals (91.4% larvae survived; Figure 2b). Moreover, only 81.5% of the inbred beetles were viable after adult emergence, compared to 96.9% of the outbred beetles (Figure 2c). Likewise, the number of eggs laid by inbred females was 36.2% lower than in outbred females (Figure 2d). Adult body mass was significantly affected by the covariate sex, with females being heavier than males (Table 1).

3.2 Effects of nonsibling versus sibling male mating partner order on larval hatching rate

All female individuals mated with the first and the second male partner during experiment 2. The hatching rate of larvae descending from these matings was significantly affected by the mating treatment, which combined different breeding levels (outbreeding versus inbreeding) and a different succession of the two breeding levels in a sequential mating with different male partners (Figure 3, Tables 1, 2). Larvae descending from females mated with two nonsibling males
TABLE 1  Effects of breeding treatment [outbred (O) versus inbred (I)], covariates, and their interaction on development time ($N_O = 437, N_I = 131$), larval survival ($N_O = 37, N_I = 28$), adult survival ($N_O = 20–30, N_I = 15–17$), and egg number ($N_O = 25, N_I = 28$) of *Phaedon cochleariae* from experiment 1 (E1), as well as effect of mating treatment [male order: nonsibling, nonsibling (N/N) versus nonsibling, sibling (N/S) versus sibling, nonsibling (S/N) versus sibling, sibling (S/S)] on larval hatching rate ($N_{N/N} = 24, N_{N/S} = 25, N_{S/N} = 25, N_{S/S} = 25$) of *Phaedon cochleariae* from experiment 2 (E2).

| Responses                  | Development time (E1) | Larval survival (E1) | Adult survival (E1) | Body mass (E1) | Egg number (E1) | Larval hatching (E2) |
|----------------------------|-----------------------|----------------------|--------------------|---------------|----------------|----------------------|
| GLMM (Poisson)             | GLMM (Binomial)       | GLMM (Binomial)      | LMM (Gaussian)     | GLMM (Poisson) | GLMM (Binomial)  |                       |
| Fixed effects:             | $\chi^2$              | $p$                  | $\chi^2$          | $p$           | $\chi^2$      | $p$                  |
| Breeding treatment         | 9.350                 | 0.002                | 127.950            | <0.001        | 30.774         | <0.001               |
| Sex                        | Not tested            | Not tested           | Not tested         | 3.305         | 0.069          | Not tested           |
| Age                        | Not tested            | Not tested           | Not tested         | Not tested    | 133.540        | <0.001               |
| Breeding x sex             | Not tested            | Not tested           | Not tested         | 1.347         | 0.246          | Not tested           |
| Breeding x age             | Not tested            | Not tested           | Not tested         | 0.893         | 0.345          | Not tested           |
| Mating treatment           | Not tested            | Not tested           | Not tested         | Not tested    | 203.700        | <0.001               |
| Random effects:            | Variance              | Variance             | Variance           | Variance      | Variance       | Variance             |
| Petri dish                 | 0.000                 | Not tested           | Not tested         | 0.018         | Not tested     | 0.009                |
| Family in P-generation     | 0.000                 | 0.046                | 0.353              | 0.004         | Not tested     | Not tested           |

Note. The table gives $p$ and $\chi^2$-Values from chi-square likelihood ratio tests for all fixed effects in the respective (G)LMM as well as the amount of variance explained by the random effects in each model (extracted from the minimal adequate mixed effects models). Significant $p$-Values are indicated in bold.
The average over all families is shown. Note. The coefficient of inbreeding depression ($\delta = \hat{\omega}_{\text{Outbred}} - \hat{\omega}_{\text{Inbred}} / \hat{\omega}_{\text{Outbred}}$) was calculated according to Hedrick and Kalinowski (2000). The average over all families is shown.

### DISCUSSION

#### 4.1 Lifetime inbreeding depression in *P. cochleariae*

The present study revealed that mustard leaf beetles suffer from strong inbreeding depression throughout their ontogeny. Inbreeding does not only reduce the larval hatching rate of the beetle as detected in a previous study (Müller & Müller, 2016), but also crucially increases the mortality of larvae prior adult emergence and of young adults. Moreover, the reproductive output of inbred females was reduced compared to outbred females, although their body mass was not decreased by inbreeding. Comparable effects were previously detected in the beetle *Propylea quatuordecimpunctata* (Morjan, Obrycki, & Krafsur, 1999), which suggests that body mass might be more robust against inbreeding impacts than other fitness-related traits. Possible effects of the males’ breeding status on the reproductive success remain to be tested in *P. cochleariae*. Given the comprehensive detrimental effects of inbreeding detected in this and in previous studies on *P. cochleariae* (Müller & Juškauskas, 2018; Müller & Müller, 2016), it can be expected that the reproduction abilities of inbred males are lower relative to outbred males. Other studies illustrate that sperm competiveness and/or amount are depressed by inbreeding in other arthropod species (Ala-Honkola et al., 2013; Fox, Xu, Wallin, & Curtis, 2012; Konior, Keller, & Radwan, 2005; Michalczyk, Martin, Millard, Emerson, & Gage, 2010). In summary, our results for lifetime inbreeding depression in *P. cochleariae* suggest that natural populations of the species, in which inbreeding depression is also detectable (Müller & Juškauskas, 2018), have a risk to become extinct within a few generations. This scenario can be relevant for a wide range of species that suffer from strong inbreeding depression after only one or few generations of nonrandom mating (Bilde et al., 2007; Fox et al., 2007; Roff, 1998).

This applies as long as the purging of deleterious recessive mutations does not mitigate or eliminate inbreeding depression under nonrandom mating conditions (Crnokrak & Barrett, 2002; Hedrick & García-Dorado, 2016). In addition, inbreeding effects on the behavioral phenotype, that is, personality, might either amplify or counteract the reduced viability of inbred *P. cochleariae* individuals, depending on the environment (Müller & Juškauskas, 2018). Such inbreeding by environment interaction effects on the behavioral phenotype represents a so far underestimated field of evolutionary ecology and might be relevant over a wide range of animal species (Ala-Honkola, Uddström, Díaz Pauli, & Lindström, 2009; Aspi, 2000; Briskie & Mackintosh, 2004). Taken together, our studies on the inbreeding effects on lifetime fitness and the behavior in *P. cochleariae* depict a comprehensive scenario of possible costs associated with inbreeding in insect populations. Such laboratory studies are a first important step to achieve a better understanding of the manifold inbreeding effects suffered in wild populations. However, the extent

| Trait                        | $\delta$ |
|------------------------------|----------|
| Development time             | 0.066    |
| Larval survival              | 0.404    |
| Adult survival               | 0.189    |
| Body mass females            | 0.077    |
| Body mass males              | 0.007    |
| Egg number                   | 0.394    |
| Larval hatching rate         | 0.254    |

Larvae descending from eggs laid by females, which were first mated with a sibling and consecutively with a nonsibling, had an intermediate hatching rate (79.9%).

![FIGURE 3](image.png)

**FIGURE 3** Effects of the mating treatment [male order: nonsibling, nonsibling (Non/Non) versus nonsibling, sibling (Non/Sib) versus sibling, nonsibling (Sib/Non) versus sibling, sibling (Sib/Sib)] on the larval hatching rate of eggs laid per female of *Phaedon cochleariae* ($N = 24–25$). The box-whisker plots show the following statistics: medians (solid gray lines), interquartile range (boxes), 1.5 * lower/upper quartile (whiskers), outliers (white dots), and least square means with their standard error (black dots with error bars) extracted from the minimal adequate (G)LMM. The letters indicate significant differences between the treatment groups and are based on a Tukey post hoc test on the minimal adequate GLMM had the highest hatching rate (90.1%) and larvae descending from females either mated with two siblings (65.4%) or first with a nonsibling and consecutively with a sibling (69.4%) had the lowest hatching rate.
of inbreeding depression under benign laboratory conditions may not precisely predict the magnitude of inbreeding depression under more stressful field conditions in natural populations (Armbruster & Reed, 2005). Some studies illustrated that results gained from laboratory-reared populations can predict the performance of wild populations under field conditions (Coelho, Rugman-Jones, Reigada, Stouthamer, & Parra, 2016; Fox et al., 2007). Nevertheless, there is broad evidence that the magnitude of inbreeding depression is higher under stressful field conditions relative to benign laboratory conditions (Fox & Reed, 2011). As global change not only increases inbreeding rates through habitat loss and fragmentation, but simultaneously raises the levels of abiotic and biotic stress in remaining habitats (Davis, 2003; Kingsolver, Diamond, & Buckley, 2013), inbreeding depression is a huge recent and future challenge for many species (Andersen, Fog, & Damgaard, 2004).

4.2 | Evidence for second male sperm precedence in *P. cochleariae*

In our study, no female-mediated sibling sperm discrimination was detectable in *P. cochleariae*. If postcopulatory inbreeding avoidance mechanisms would exist, one would expect a similar larval hatching rate resulting from a mating with a nonsibling and a sibling male in both orders, because females might be able to preferentially choose sperm of nonsiblings. This phenomenon is postulated as cryptic female choice (Eberhard, 1996) and is supposed to be realized in various arthropod species (Fedina, 2007; Peretti & Eberhard, 2010). However, in the present study, it was not detected. Contrary, the hatching rate resulting from mating with a sibling followed by a nonsibling was significantly higher compared to a mating in the opposite order. Moreover, the hatching rate was highest in the mating with two nonsiblings. This points to second male sperm precedence, a phenomenon in polyandrous species in which the second/last male mating partner sires more offspring than the first (Kehl, Karl, & Fischer, 2013; Xu & Wang, 2010). Paternity analyses must be conducted to clarify this assumption in *P. cochleariae*. If the break between the first male and second male during the sequential mating would be shorter than the applied 18 hr, the impact of the second male may become even more pronounced than it already was. A potential mechanism behind second male sperm precedence can be associated with either females or males. Females can release sperm of the first mating from their reproductive tract (Snook & Hosken, 2004). Alternatively, the second male might displace the sperm of the first male (Xu & Wang, 2010). The realized mechanisms underlying second male sperm precedence in *P. cochleariae* remain to be clarified in future studies. Second male sperm precedence would shed a new light on a specific behavior previously observed in *P. cochleariae*. Couples of the beetle often mate several hours. During that time, no other male has the possibility to mate with the female. This long mating duration is potentially a mate guarding behavior or, more precisely, a mounting behavior practiced by males. In doing so, males may prevent fertilization by other males to improve their own paternity chances, which was also observed in the West Indian sweetpotato weevil (Sato & Kohama, 2007). A mate guarding behavior combined with second male sperm precedence and a possibility of remating, can enhance male fitness (Arnqvist, 1988; Baxter, Barnett, & Dukas, 2015; Calbacho-Rosa, Cordoba-Aguilar, & Peretti, 2010).

Concluding from a previous (Müller & Müller, 2016) and the present study, neither pre- nor postcopulatory inbreeding avoidance mechanisms could be detected in *P. cochleariae*. This is surprising because *P. cochleariae* suffers from severe inbreeding depression in a laboratory (this study) and a wild population (Müller & Juškauskas, 2018). Simultaneously, these beetles have distinct family-specific CHC patterns, which could function as recognition cue for inbreeding avoidance. However, the absence of inbreeding avoidance in spite of the expression of inbreeding depression was also observed for burying beetles (Mattey & Smiseth, 2015; Pilakouta & Smiseth, 2016). Hence, the question arises whether any other mechanisms to avoid inbreeding costs exist in these species. One possibility could be a specific dispersal behavior, which was detected in many species throughout the animal kingdom (Pusey & Wolf, 1996). Indeed, in a previous study, we found that wild *P. cochleariae* males are more active than females and cover longer distances (Müller & Juškauskas, 2018). This result suggests that the dispersal of males may avoid the mating among close relatives in natural populations of the species. Sex-specific dispersal can be observed across a wide range of animal species and is often expressed to avoid inbreeding and/or intra-specific competition (Pusey, 1987; Pusey & Wolf, 1996). Which sex disperses is determined by the ratio of costs (increased predation or less mates in the new habitat) and benefits (less competition and relatives as mates) that are associated with dispersal (Motro, 1991; Perrin & Mazalov, 1999; Yoder, Marschall, & Swanson, 2004).

Moreover, a polyandrous mating system might exist in *P. cochleariae*. As postulated by the fertility restoration hypothesis (Bayoumy et al., 2015), such mating systems can evolve in species, if the reproductive success of females is decreased by inbreeding, but increased by outbreeding. A further mating assay, in which females have the possibility to mate with single nonsiblings or siblings or with groups of either nonsiblings, siblings or a mixed group of nonsiblings and siblings, can clarify if polyandry is beneficial for *P. cochleariae* in the context of inbreeding. Females of polyandrous species may use CHC patterns and a CHC based chemosensory self-referencing mechanisms to increase the diversity of their mating partners (Weddle, Hunt, et al., 2013; Weddle, Steiger, et al., 2013) and consequently the genetic diversity of their offspring. However, even for polyandrous species, an increasing degree of relatedness in a given population following habitat loss and fragmentation will decrease the efficiency of inbreeding avoidance by multiple paternity.

Overall, the present study highlights the need to measure inbreeding depression throughout the entire ontogeny of a study organism. Focusing on single traits or restricted periods in ontogeny could potentially lead to an under-estimation of the inbreeding effects on fitness-related traits. Although we found pronounced lifetime inbreeding depression, a postcopulatory inbreeding avoidance mechanism was not detected, which supports the existence of a polyandrous mating system or specific dispersal strategies for
inbreeding avoidance. Instead, we disclosed evidence for second male sperm precedence, which can potentially explain specific behavioral traits of our study species, that is, a mounting behavior of males after mating.

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CONFLICT OF INTEREST

None declared.

AUTHOR CONTRIBUTIONS

TM designed the study. TDL and TM conducted the breeding experiment and the bioassays and collected the data. KS and TM analyzed the data and prepared the figures. TM and KS wrote and revised the manuscript.

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