Selfing-outcrossing as a gradient, not a dichotomy: propensity for

selfing varies within a population of hermaphroditic animals

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Abstract

For mating-system evolution, individual-level variation is essential. In self-compatible

hermaphrodites, individuals may vary in their lifetime propensity for selfing, which consists of

individual and environmental components. According to the reproductive assurance hypothesis

explaining partial selfing, a key environmental factor is mate availability, which fluctuates with

population density.

We quantified individual variation in selfing propensity in a hermaphroditic snail by

manipulating mate availability, recording mating behaviour, estimating selfing rates from

progeny arrays, and measuring female lifetime fitness. We found substantial among-individual

variation in selfing propensity, including pure outcrossers, pure selfers, and two types of plastic

individuals. This variation only manifested in the laboratory; for the highly dense field

population, data suggest full outcrossing. Meanwhile, experimental levels of mate availability

(low *versus* moderate) neither significantly affected selfing propensities nor selfing rates.

Instead, selfing propensities had an individual, environment-independent component.

Our results imply that selfing propensities are partially heritable and, when selected on, cause

mean selfing rates to evolve. We propose that genetic variation in selfing propensities offers a

reconciliation between the reproductive assurance hypothesis and its limited empirical support

in animals: distributions of selfing propensities vary temporally and spatially, thus obscuring

the relationship between population density and realised selfing rates.

Background

The mating system structures populations and species with respect to sexual behaviour. In self-

compatible hermaphrodites, it primarily describes the relative frequency of self-fertilisation

(hereafter selfing) *versus* outcrossing. The mating system adopted by such hermaphrodites

determines the quantity and genetic quality of offspring (1). Consequently, it has pervasive

effects on individual fitness and on the evolutionary and demographic properties of populations

and species (2, 3, 4).

Variation in selfing rates has often been studied at the species level (5, 6, 7, 8), where the

distribution of selfing rates has been found to be bimodal, with more outcrossers and selfers

than mixed maters (0.2 < selfing rate s < 0.8). In plants, many studies have documented

variability between populations of a single species (e.g., 9, 10, 11, 12). For example, in 51% of

105 surveyed plant species, populations fell into several mating-system categories (13). Our

knowledge on animal populations is sparser. Inter-population variation was observed in the

androdioecious nematode *C. elegans*, whose hermaphrodites cannot mate with each other and

where all outcrossing is due to males (14). In the few studies where selfing rates, or the delay in

the onset of reproduction when selfing (termed "waiting time", 15), have been estimated across

populations of self-compatible hermaphrodites capable of outcrossing, inter-population

variation was the norm (16, 17, 18, 19, 20, 21, 22, 23). It is therefore important to characterise

species by more than one population estimate, lest crucial differences between populations may

be overlooked.

Variation in selfing rates among individuals is equally essential. Evolutionary change primarily

happens at the population level; among-individual variation constitutes the raw material natural

selection acts upon. If individuals within isolated populations of simultaneous hermaphrodites

were uniform in their ability to self and had equal opportunities to outcross, the selection

gradient for the selfing rate would be zero: selection can only act on traits that have both

phenotypic and genetic variance (24).

Despite the importance of among-individual variation in selfing rates, empirical studies are not

usually designed to capture variation at this level. Reliable estimates of individual selfing rates

require a large number of rather large progenies to be genotyped with genetic markers of

sufficient resolution to unequivocally identify outcrossing events (25, 26). Existing studies have

shown that selfing rates varied within populations of plants (e.g., 27, 28, 29) and animals (e.g.,

18, 19, 20, 30, 31, 32, 33, 34). However, especially in studies of animals, many estimates rely

on small numbers of weakly polymorphic loci, which hampers the detection of outcrossing

events, and so may be biased (25, 26). Substantial differences among individuals were also

found in proxies of the selfing rate, such as in animals' waiting time (22, 35, 36) and in the

presence of seeds in self-pollinated plants (37, 38).

Individual selfing rates likely depend on genetic factors, environmental conditions, and

genotype-by-environment interactions. Genetic self-incompatibility systems are well-known in

plants (39, 40) and have recently been discovered in ascidians (41) and oomycetes (42).

Differences in waiting times among families, suggestive of genetic variation in selfing rates,

have been found in snails (15, 22, 35) and flatworms (36). A key environmental determinant of

individual selfing rates is the availability of mating partners. Selfing provides reproductive

assurance when mates are scarce, but due to inbreeding depression and other costs of selfing

(43, 44) may be unpreferred if outcrossing is possible. Yet, to date there are no direct tests of

how individual selfing rates in animals respond to differential mate availability; existing studies

(45, 46) lack selfing rates estimated using molecular markers. Here we aim to close this

knowledge gap.

Individual variation in selfing can be assessed by estimating individuals' lifetime propensity for

selfing, here defined as a biological property of an individual (the ability and readiness to self)

and its environment (mate availability), loosely following Brandon's propensity interpretation

of fitness (47). This definition allows a separate exploration of the individual (supposedly

genetic) *versus* ecological determinants of variation in selfing rates, the need for which recent

studies have abundantly shown (9, 13, 23, 48).

The freshwater snail *Radix balthica* is an ideal species to study within-population variation in

selfing propensity. Laboratory studies have confirmed that self-compatibility is geographically

widespread yet not omnipresent within populations (49, 50). There is also tentative evidence for

between- and within-population variation in selfing rates based on molecular genetic markers

(32, 33, 51). The population studied here is almost fully outcrossing in the field (26), in

accordance with its high density.

We derived individuals' lifetime propensity for selfing for two laboratory environments with

differential mate availability. We first ascertained how readily 274 laboratory-reared, unmated

snails selfed in isolation. We then assigned them to one of two mating treatments, simulating (i)

a low population density and scarcity of mating opportunities, or (ii) conditions in the ancestral

field population, where density was high and multiple paternity within egg clutches common

(52). We recorded individual mating behaviour, the lifetime production of eggs in isolation

(necessarily selfed) and thereafter (potentially outcrossed), the proportion of developing eggs,

and the proportion of surviving juveniles. For 56 individuals across both treatments (32 of them

once- and 24 repeatedly paired), we estimated individual selfing rates in eggs produced post-

isolation from progeny arrays.

Assuming variation in post-isolation selfing rates, we had two *a priori* hypotheses: Selfing

post-isolation is more common under low mate availability (hypothesis 1) and in snails that

selfed in isolation (hypothesis 2). Hypothesis 1 posits that selfing provides reproductive

assurance at low density but is unfavoured when outcrossing opportunities are abundant (43,

44). If supported, it suggests selfing propensities have a strong ecological component.

Hypothesis 2 postulates that a period of isolation reveals classes of individuals with different

propensity for selfing, which remain largely unaffected by environmental change. It thus

assumes selfing propensities to be stable, fundamental properties based in genetic differences.

The hypotheses are not mutually exclusive; ecological and genetic factors can jointly shape

individual selfing rates. We further explored how variation in selfing propensities is maintained

by investigating their association with individuals' sexual drive, female lifetime reproductive

success, and their progenies' embryonic development and juvenile survival.

Methods

Study system

R. balthica is a simultaneously hermaphroditic snail inhabiting the shallow littoral zone of

European lakes (51, 53). In Lake Zurich, Switzerland, it is an annual with non-overlapping

generations. Snails hatch from eggs in spring, reach sexual maturity in winter, reproduce from

March to May, and then die (52). During the single breeding season, individuals may copulate

repeatedly in both sexual roles and lay hundreds of eggs in distinct egg clutches (52).

Copulation is unilateral. Lymnaeid snails can store allosperm and should not be allosperm-

limited even if they do not mate continuously (54).

Experimental snails

We caught 86 adult snails (P0 mothers) at peak breeding season (24 April 2013) in Uerikon,

Lake Zurich, Switzerland. In the laboratory at Eawag-Duebendorf, Switzerland (room

temperature 18°C), they were kept individually in 200 ml plastic cups filled with aged tap water

and fed *ad libitum* with organic lettuce. Water was changed once a week. In these cups, P0

mothers laid egg clutches until they died of natural causes in late spring 2013 (details in

reference 49). P0 mothers were not allowed to mate in the laboratory; their offspring (F1 snails)

were outcrossed using allosperm stored from copulations in the field, as shown by genotyping

P0 mothers and F1 snails (see Results). Previous work showed that egg clutches collected from

the ancestral field population throughout the breeding season often had multiple fathers and

contained many unique parental genotypes (52). This suggests that the fathers of F1 snails,

while unknown, came from a large and diverse population, and that F1 offspring of the same P0

mother were often just half siblings. Hence, an overrepresentation of certain fathers among F1

snails is unlikely. Each egg clutch laid by P0 mothers was placed in a separate water-filled 40

ml plastic cup. After 17 days, when hatching was imminent, each clutch was transferred to a

larger 200 ml cup. F1 snails were reared in sibling groups and individually from age 19.4 ± 4.1

weeks onwards (mean ± SD) to preserve their virginity (details in 49). This start of isolation is

very early; paired control snails only began to mate at age 41.3-45.7 weeks. F1 snails were fed

Spirulina powder mixed with finely ground chalk and flakes of fish food as juveniles, and

organic lettuce from age 37.0-42.3 weeks onwards. In spring 2014, mating trials (see next

section) were conducted with 274 F1 snails derived from 38 P0 mothers and 108 sibling groups.

The effects of mating trials on mating behaviour, fecundity, and the population growth rate are

described elsewhere (49). For a subset of 56 of these 274 F1 snails, derived from 22 P0 mothers

and 38 sibling groups, we estimated selfing rates (see Genetic analysis).

147   Mating trials

Mating trials began in early May 2014 when F1 snails were 52.1 ± 1.0 weeks old. By that time,

all F1 snails had experienced a prolonged period of isolation and 38.7% had begun to reproduce

through selfing. In the ancestral field population, breeding started in early March (52), and

laboratory control snails had started to copulate several weeks earlier (see above).

A detailed description of the design of mating trials is provided in (49). In brief, two types of

mating trials were conducted. In the first type, 137 snails had one mating opportunity with one

partner, simulating low population density and a scarcity of mating opportunities. In the second

type, 137 snails had six mating opportunities over the course of six weeks, each with a different

partner, simulating moderate population density. Hence, no F1 snail remained permanently

isolated; they were all paired at least once. Each mating opportunity lasted for 10.3 ± 0.8 hours,

long enough for copulations in both sexual roles, and happened during daylight hours. Once-

paired snails were thus allowed to mate for only ~0.5% of the length of their natural, three-

month breeding season. Additionally, having just a single mating partner creates substantial

obstacles for outcrossing (*e.g.*, unsuccessful sperm transfer, cryptic female choice, sperm

competition with autosperm, genetic incompatibility between sperm and egg). So, if a snail's

sole mating partner happened to be sterile or incompatible, then this was part of the low-density

environment we wanted to simulate. Hence, once-paired snails were clearly mate-limited. By

contrast, a choice of six mating partners may come close to conditions in the ancestral, fairly

dense field population, where field-collected egg clutches had 2.1 fathers on average (range: 1-

9; 52). Given that not all mate encounters will lead to copulations, and the above-mentioned

obstacles to outcrossing, the average number of potential mating partners of a free-living snail

throughout its life is probably a multiple of 2.1. The contrast between once- and six-times-

paired snails thus enables a rigorous test of hypothesis 1 ("selfing post-isolation is more

common under low mate availability"). F1 snails were euthanised one week after repeatedly

paired snails had had their sixth mating opportunity, when mortality began to increase reflecting

the snails' natural, annual life cycle.

All except two mating partners were from among the 274 snails, so snails could act both as

mothers and fathers. The first mating partner of each snail was size-matched, to ensure that

anatomical or developmental differences did not constrain copulation. All snails were

individually marked (details in 49). Mating pairs were set up randomly with respect to

relatedness. Consequently, in 22/465 pairings (4.7%) involving 42 F1 snails, mating partners

had the same P0 mother (*i.e.*, full or half siblings). Of these 42 snails, 37 were paired

repeatedly, and so were also paired with four to five unrelated snails each, reducing potential

effects of mating partner relatedness. The five snails solely paired with a sibling were excluded

from all analyses, as was a snail with missing data on female reproductive output. Hence, our

sample sizes are 131 snails paired once (32/56 snails with selfing rate estimates) and 137 snails

paired repeatedly (24/56).

Mating trials were recorded on time-lapse movies (one frame/30 sec). From these we extracted

the number of partners each snail had copulated with as a male and as a female. We also

collected all egg clutches F1 snails laid both before and after being paired to measure female

lifetime reproductive success. We placed clutches in individual 40 ml cups and counted the

number of eggs and how many of these contained developed embryos 17 days after oviposition

(details in 49). Then each F1 snail's clutches were transferred to two 2 l plastic containers each,

where selfed (clutches laid in isolation) and potentially outcrossed (clutches laid post-isolation)

F2 family groups were reared separately. F2 snails were fed a mixture of *Spirulina* algae, finely

ground chalk, and flakes of fish food.

Genetic analysis

We selected those F1 snails (n = 56) for selfing rate estimation that post-isolation produced at

least twelve F2 offspring that reached 12.3 ± 0.5 weeks of age. At that time, some F2 juveniles

were used in a field experiment and the rest frozen for genetic analysis. As by then 53.4 ±

18.1% of F2 offspring born post-isolation had died, we here estimate "secondary" selfing rates

after the expression of potential in- or outbreeding depression in early development. For logistic

reasons, we did not genotype any juveniles from eggs laid shortly before the experiment was

terminated. Hence, the genotyped offspring of repeatedly paired snails originate from eggs laid

before mating opportunity 4 (n = 6 snails), 5 (n = 15 snails), and 6 (n = 3 snails), respectively.

Offspring of these snails can thus be sired by three to five mating partners.

Of 40 F1 snails, 15 or more F2 juveniles were available for genotyping; of these we genotyped

at least 15, resulting in 16.0 ± 1.9 genotyped juveniles per F1 snail (range: 15-22). Of the 16 F1

snails with fewer than 15 juveniles available for genotyping, all juveniles were genotyped. This

resulted in 8.1 ± 3.6 genotyped juveniles per F1 snail (range: 3-14). We also genotyped all 86

P0 and 274 F1 snails to estimate population-level inbreeding coefficients using Genetix version

4.05.2 (55) and to ascertain whether F1 snails themselves were selfed (Table S1). Snails were

genotyped for ten highly polymorphic microsatellite loci (GenBank Accession No. KX830983-

KX830992) developed specifically for this population by Ecogenics GmbH (Zurich,

Switzerland). The genotyping and scoring protocol is described elsewhere (26). One locus

showed a non-negligible frequency of null alleles (locus Rb_3, KX830985) and was excluded

from all analyses.

Paternity analysis

Paternity analyses were performed using a custom-built R routine. We excluded 41 juveniles

(5.3% of 768 juveniles genotyped) from the analysis because they did not meet the inclusion

criteria (fewer than six loci of both mother and offspring genotyped successfully, OR genotypes

lacking a maternal allele at more than two loci, OR genotypes with six successfully genotyped

loci but two loci without a maternal allele). This left us with 13.0 ± 4.1 successfully genotyped

juveniles per family (range: 3-18). For these 727 juveniles, we counted the loci consistent with

being outcrossed, *i.e.*, those with one non-maternal allele. We then compared each juvenile's

genotype to those of all its candidate fathers, defined as all the potential mates of a juvenile's

mother, irrespective of whether copulations had been observed. A candidate father was

considered a perfect match if he possessed a juvenile's paternal allele at every locus successfully

genotyped in the juvenile, the mother, and himself. Candidate fathers lacking a juvenile's paternal

allele at one or several loci were considered non-perfect matches and deemed increasingly

unlikely fathers the higher the number of father-offspring mismatches. Additionally, we repeated

all paternity analyses using COLONY version 2.0.5.9 (56, see Supporting Methods), known for

its highly accurate parentage assignments (57).

Using these procedures, paternity was assigned to 717 (98.6%) juveniles with high-quality

genotypes, originating from 55 F1 snails (13.0 ± 3.9 assigned offspring per F1 snail). Paternity

assignments were inconclusive for all of the offspring of one F1 snail (see Table S2 for numbers

of genotyped, successfully genotyped, and assigned juveniles per family). For 705 juveniles, the

R routine and COLONY produced identical assignments. Of the remaining twelve juveniles, five

were left unassigned by the R routine, four by COLONY, and three were assigned by both but to

different fathers; here we gave preference to the R routine (details in Table S2). Allelic diversity

was high across loci (Table S1), yielding powerful paternity analyses. The mean probability of

being selfed for offspring identified as selfed was 0.79 ± 0.22, while that of offspring identified

as outcrossed was 0.00 ± 0.00 (Table S2). Outcrossed offspring were assigned to a father with a

probability of 0.93 ± 0.16 (Table S2).

Statistical analysis

We tested whether F1 snails were more likely to produce at least some selfed juveniles post-

isolation (no/yes) under low mate availability (test of hypothesis 1) and upon successful selfing

in isolation (test of hypothesis 2) using a GLMM with binomial errors ($n = 55$). Fixed effects

were the experimental treatment (once- vs. repeatedly paired), the number of developed

embryos produced in isolation, and the number of male and female partners snails copulated

with. P0 mother identity was used as a random intercept to account for potential effects of

relatedness.

We used eight GLMMs to test if snails with different propensies for selfing differed in their

lifetime number of eggs (female LRS) and developed embryos, in their lifetime proportion of

undeveloped embryos, and in the proportion of developed embryos that died before reaching

the juvenile age of 12.3 ± 0.5 weeks, when surviving F2 juveniles were genotyped. We

estimated these models for F1 snails with successfully estimated selfing rates (four models, n =

50; the five outcrossers were excluded due to low sample size) and for all female fertile snails,

here defined as snails with developed embryos (four models, n = 123 or 113, depending on the

trait analysed). We fitted GLMMs using negative binomial errors (nbinom 1 family) for the

number of eggs and developed embryos, and using Gaussian errors for the proportion of

undeveloped embryos and dead juveniles. Despite being proportions, the latter two response

variables did not deviate significantly from normality (one-sample Kolmogorov-Smirnov tests:

$D \geq 0.06$, $p \geq 0.33$). Furthermore, model assumptions were not violated, as shown by both

conventional diagnostic plots and those produced by DHARMa, a simulation-based approach to

assess the model fit (58). In all eight models, adult body size was used as a continuous

covariate, the propensity for selfing as a factor with three levels ("selfer", "plastic mixer", and

"plastic switcher" when n = 50, and "apparent selfer", "apparent plastic snail", and "apparent

outcrosser" when n = 123 or 113), and P0 mother identity as a random intercept. In all models

except those of the number of eggs, we also included the "pair identity" on mating opportunity

1 as a random intercept. This accounts for the non-independence of sexual functions within

pairs of once-paired snails, where it was, *e.g.*, impossible for one snail to remain unmated while

its sole mating partner mated. In models of the number of eggs, the likelihood of selfing post-

isolation (see above), and the likelihood of mating in both sexual roles among snails with

estimated selfing rates (see below), we did not add the pair identity on mating opportunity 1

because doing so resulted in convergence problems. However, of 31 once-paired snails with

estimated selfing rate, only six were paired with one another, while 25 were paired with

repeatedly paired snails or with once-paired snails whose selfing rate was not estimated. Any

non-independence in selfing rate estimates is thus likely small.

Furthermore, we tested if snails that mated in both sexual roles (no/yes) differed in their

propensity for selfing using a GLMM with binomial errors and including snails with

successfully estimated selfing rates only (n = 50, see above). The propensity for selfing (three

levels: "selfer", "plastic mixer", "plastic switcher"), adult body size (continuous covariate) and

P0 mother identity (random intercept) were used as predictors.

Finally, we used two-tailed Pearson's Chi-square tests to investigate effects of the experimental

treatment of F1 snails (once- vs. repeatedly paired) on multiple paternity (supporting analysis),

selfing (test of hypothesis 1), and the propensity for selfing (test of hypothesis 1). Specifically,

2x2 contingency tables were used to test for treatment effects on the number of F2 progenies

with multiple paternity, the number of F2 progenies with selfed offspring, and the overall

number of selfed F2 offspring. Treatment effects on the propensity for selfing in all female

fertile F1 snails (categories: "apparent selfer", "apparent plastic snail", and "apparent

outcrosser") and only in snails with selfing rate estimate (categories: "selfer", "plastic mixer",

"plastic switcher", and "outcrosser") were analysed using 3x2 and 4x2 contingency tables,

respectively.

Analyses were performed using R v. 4.2.2 (59). GLMMs were fitted using "glmmTMB" (60).

The model fit was assessed using "DHARMa" (58). We tested the significance of random

effects with log-likelihood ratio tests (full model vs. model without random effect in question).

Values are given as mean ± SD. Scatter plots were prepared using "beeswarm" (61) and depict

data points superimposed on boxplots. The entire code and model output is provided as an R

Markdown document (62).

Results

Frequency of selfing and multiple paternity

Selfed offspring were found in 43.6% of genotyped families (24/55; Fig. 1a). Nine families

showed low selfing rates of 20% at most and 15 families high ones of 83% or more. Families

with intermediate selfing rates ($0.20 < s < 0.80$) were absent. Overall, 23.2% of offspring were

identified as selfed (166/717).

There was no evidence of selfing in either the F1 or P0 generation. All 274 F1 snails were

outcrossed, *i.e.*, they possessed a non-maternal allele at one or more loci (mean: 4.6 ± 1.4 loci;

Table S1). P0 mothers of F1 snails with genotyped offspring were likely outcrossed too: their

genotypes did not show reduced heterozygosity ($F_{IS} = 0.009$, 95% CI: -0.010, 0.068; Table S1).

P0 mother identity did not explain any variance in the probability that, post-isolation, F1 snails

reproduced via selfing (Table 1).

Multiple paternity was significantly more common among the offspring of repeatedly paired

(45.8%) than once-paired snails (12.9%, Pearson's Chi-squared test with Yates' continuity

correction: observed counts: 4, 27, 11, 13, expected counts: 8.5, 22.5, 6.5, 17.5, $\chi^2_1 = 5.83$, $p =$

0.0158; Fig. 1a); note that progenies of once-paired snails show multiple paternity when they

contain both selfed and outcrossed offspring. On average, the progenies of repeatedly and once-

paired snails had 1.7 ± 0.9 (max. = 4) and 1.1 ± 0.3 (max. = 2) fathers, respectively.

Only subtle effect of mate availability on selfing

Mean selfing rates did not differ significantly among once- (0.37 ± 0.48) and repeatedly paired

snails (0.19 ± 0.36; Wilcoxon rank sum test with continuity correction: $W = 433.5$, $p = 0.25$), nor

did their variance (Levene's test for homogeneity of variance: $F_{1,53} = 2.44$, $p = 0.12$; Fig. 1b).

Repeated mating opportunities did not significantly reduce the likelihood of a snail producing at

least some selfed offspring (Table 1), nor the proportion of families with non-zero selfing rates

(once-paired: 48.4%, repeatedly paired: 37.5%, Pearson's Chi-squared test: observed counts: 15,

16, 9, 15, expected counts: 13.5, 17.5, 10.5, 13.5, $\chi_1^2 = 0.65$, $p = 0.42$; Fig. 1c). We thus did not

find strong support for hypothesis 1, which predicted that selfing is more common at low mate

availability. However, as a group, once-paired snails produced significantly more selfed offspring

than repeatedly paired snails: 34.2% vs. 8.9% (Pearson's Chi-squared test: observed counts: 138,

265, 28, 286, expected counts: 93.3, 309.7, 72.7, 241.3, $\chi_1^2 = 63.63$, $p < 0.0001$; Fig. 1d).

Nevertheless, the distribution of selfing rates was similarly bimodal among both treatment groups

(Fig. 2a).

Selfing in isolation linked to selfing post-isolation

Predominant selfing after mating trials had started was restricted to snails that had selfed

successfully in isolation, while snails without prior selfing exclusively outcrossed post-isolation

(Fig. 2b). Consequently, selfing in isolation predicted continued selfing (Table 1); its effect

remained statistically significant when an extremely prolific snail was excluded ($b = 0.01$, $z =$

2.04, $p = 0.0418$). We therefore conclude that our data support hypothesis 2, which posited that

selfing post-isolation is more common in snails with a history of selfing.

Individual-level variation in propensity for selfing

Snails varied in their propensity for selfing. For snails with estimated selfing rates (n = 55), we

ascertained the selfing propensity from the presence/absence of selfing (*i.e.*, developed

embryos) in isolation and the post-isolation selfing rate. This revealed four groups of snails:

9.1% "outcrossers" that never selfed, 47.3% "plastic switchers" that only selfed in isolation,

16.4% "plastic mixers" that combined selfing in isolation with predominant yet incomplete

outcrossing post-isolation, and 27.3% "selfers" that selfed fully or mostly throughout (Fig. 2c).

Most outcrossers did not produce any undeveloped embryos in isolation either. Consequently,

their proportion (9.1%) was largely unchanged when based on the number of undeveloped

(10.9%; Fig. S1a), or developed and undeveloped embryos (5.5%; Fig. S1b). Numbers of

developed and undeveloped embryos were positively correlated across the whole dataset

(Pearson's product-moment correlation: $r = 0.61$, $t_{266} = 12.53$, $p < 0.0001$; Fig. S1c).

The subsample with estimated selfing rate underrepresents snails that reproduced only before or

only after mating trials started (Fig. S2). To estimate the frequency of selfing propensities more

accurately, we thus included all female fertile snails; the selfing propensity of female infertile

snails is unknown. We ascertained the apparent selfing propensity of snails without selfing rate

estimates based on when they laid eggs and how they mated. Apparent outcrossers only

reproduced post-isolation and mated as a female, apparent plastic snails reproduced both before

and after being paired and mated as a female, and apparent selfers only reproduced in isolation,

or reproduced in isolation and post-isolation yet never mated as a female. Across all female

fertile snails (n = 124), this revealed 30.6% apparent outcrossers, 37.1% apparent plastic snails,

and 32.3% apparent selfers

The pairing treatment did not influence the selfing propensity of snails with estimated selfing

rates (Pearson's Chi-squared test: observed counts: 11, 4, 15, 1, 4, 5, 11, 4, expected counts:

8.5, 5.1, 14.7, 2.8, 6.5, 3.9, 11.3, 2.2, $\chi^2_3 = 4.98$, $p = 0.17$). However, when using the full dataset,

selfing propensities differed among experimental groups (observed counts: 29, 26, 11, 11, 20,

27, expected counts: 21.3, 24.5, 20.2, 18.7, 21.5, 17.8, $\chi^2_2 = 15.17$, $p = 0.0005$). Low mate

availability led to more apparent selfers (43.9% vs. 19.0%) and fewer apparent outcrossers

(16.7% vs. 46.6%) yet did not affect apparent plastic snails (39.4% vs. 34.5%), lending some

support to hypothesis 1.

367 **Reduced female LRS and increased offspring mortality in selfers**

368 Selfing was associated with a significant decrease in female LRS (number of developed

369 embryos) of 48.9% compared to plastic mixers and of 26.8% compared to plastic switchers

370 (Fig. S3a, Table S3). On the one hand, this was because selfers produced significantly fewer

371 eggs – 39.9% and 19.1% fewer than plastic mixers and switchers, respectively (Table S3). On

372 the other hand, selfers laid more eggs that failed to develop (28.3%) than plastic mixers

373 (16.2%); the difference to plastic switchers (21.4%) was non-significant (Fig. S3b, Table S3).

374 Outcrossers were excluded because of low sample size (n = 5). Across all female fertile snails,

375 female LRS in apparent selfers was not only reduced compared to apparent plastic snails (by

376 62.9%), but also compared to apparent outcrossers (by 52.0%; Fig. 3a, Table 2). Again, this was

377 both due to a reduced fecundity (reductions by 57.0% and 44.9%, respectively) and increased

378 proportion of undeveloped embryos in apparent selfers (36.8% vs. 25.4% and 28.2%,

379 respectively; Fig. 3b, Table 2).

380 Additionally, selfing resulted in inbreeding depression in juvenile survival. In snails with

381 selfing rate estimates, the proportion of dead F2 juveniles was significantly higher among the

382 progenies of selfers (69.5%) than among those of plastic switchers (53.4%); the difference to

383 plastic mixers (64.5%) was non-significant (Fig. S2c, Table S3). Within all female fertile snails,

384 apparent selfers had more dead juveniles (65.5%) than apparent outcrossers (51.6%), but not

385 significantly more than apparent plastic snails (60.9%; Fig. 3c, Table 2).

386 These results were all corrected for effects of body size, P0 mother identity, and, where

387 possible, pair identity on mating opportunity 1 (Table 2, Table S3).

Mating as a female did not prevent selfing

A strong propensity for selfing was not linked to a lack of sexual drive. Neither the number of male (potential sperm donors) nor female mating partners (potential sperm recipients) was significantly associated with the likelihood of selfing post-isolation (Table 1). Many selfers (53.3%) copulated in the female role and so might have received allosperm. Four out of five selfers (80.0%) mated as a male, potentially siring outcrossed offspring. Of snails that selfed post-isolation, 70.8% mated as a female.

However, the selfing propensity and sexual activity were not entirely independent. Significantly fewer selfers (40.0%) than plastic switchers (92.3%) copulated in both sexual roles ($b = 2.88$, $z = 3.18$, $p = 0.0015$; Fig. 4); the difference to plastic mixers was non-significant (66.7%, $b = 1.12$, $z = 1.26$, $p = 0.21$). Outcrossers (80.0%) were excluded from the model due to low sample size (n = 5, Fig. 4). Body size ($b = -0.05$, $z = -0.24$, $p = 0.81$) and P0 mother identity ($\chi^2_1 = 0.0$, $p = 1.00$) did not affect the likelihood of mating in both sexual roles.

Discussion

We found that propensities for selfing differed substantially within a snail population.

Individual-level variation in selfing propensity has not received much attention in the mating-

system literature, which has traditionally been focused on the level of species (5, 6, 7, 8) or

populations (10, 13).

We controlled for several possible sources of error that could hamper the estimation of

individual selfing propensities. Focal individuals were outcrossed, sexually mature, and paired

with at least one size-matched partner that was not a close relative. The pairing treatments

contrasted two biologically meaningful levels of mate availability. We estimated individual

selfing rates post-isolation from progeny arrays, which are almost free of problematic

assumptions (26, 63, 64). Our molecular markers were highly polymorphic, yielding reliable

paternity assignments with sufficient statistical power (26). Nonetheless, it is important to note

that we estimated "secondary" selfing rates, which might be biased if there is in- or outbreeding

depression prior to sampling. Here, we found inbreeding depression: juvenile mortality was

highest when progenies were selfed, intermediate when mixed, and lowest when outcrossed.

Most pre-sampling mortality thus likely affected selfed juveniles. Accordingly, there should be

no bias in high selfing rates, while low selfing rates ($0 < s < 0.21$) and those of zero might be

slightly too low. This does not change the high prevalence of self-fertility in our population (in

fact, it underestimates it), nor the bimodal distribution of selfing rates, nor the substantial

variation in selfing propensities among individuals.

The mating-system variation was cryptic in two ways. First, it only became apparent in the

laboratory. By contrast, both direct and indirect selfing rate estimates of the ancestral field

population were near-zero across three years and throughout the breeding season (26). Using

the field results only, the population would have been deemed outcrossing and its considerable

potential for selfing missed. Second, much variation would have been overlooked had we only

recorded selfing during an isolation treatment, without estimating post-isolation selfing rates.

This approach, used in pioneering earlier studies (e.g., 65, 66), merely reveals individuals'

ability to self but not their propensity for selfing, which must be assessed in the face of

outcrossing opportunities and using genetic markers. Here we show just how variable the

propensity is, comprising two types of plastic individuals, selfers, and outcrossers. Selfing-

outcrossing thus is a gradient, not a dichotomy – also, and perhaps especially, at the individual

level.

We found that variation in selfing propensity had a stable, environment-independent

component: the better an individual was at selfing in isolation, the more likely it was to self

post-isolation (hypothesis 2). In fact, those individuals that did not self while isolated invariably

outcrossed thereafter. These pure outcrossers were numerous, representing nearly a third of

female fertile snails. Notably, most of them did not produce any undeveloped embryos in

isolation either, showing that they did not try to self and failed (perhaps due to early-acting

inbreeding depression), but rather did not try at all. Furthermore, given the deliberately late start

of mating trials, it is unlikely that many outcrossers would have begun to self if isolated even

longer. These results hint at a genetic background to selfing propensities, which is in line with

findings in other hermaphroditic animals (15, 22, 35, 36, 41). An aversion to selfing so strong

as to risk reproductive failure when deprived of mates is so clearly maladaptive that a genetic

self-incompatibility mechanism appears plausible. While common in plants (39, 40), such

mechanisms have only recently been discovered in animals (41). Our results imply that a

detailed investigation of the genetic basis of variation in self-compatibility would be

worthwhile in this population.

By contrast, we found only limited support for a strong ecological component of selfing

propensities: selfing post-isolation was not substantially more common under low mate

availability (hypothesis 1). While selfing was indeed most prevalent when snails were isolated,

it often persisted post-isolation. Moreover, pairing snails repeatedly rather than once had a

surprisingly subtle effect, significantly affecting neither the mean selfing rate, the proportion of

selfing individuals, nor the likelihood of selfing. This was despite large effect sizes and

sufficient statistical power: doubling our sample size would not have rendered any of these

effects significant (see R Markdown document). Instead, treatment effects were non-significant

because variation among snails was large even within treatment groups. The only significant

consequences of pairing snails repeatedly were a lower overall number of selfed offspring,

showing that increased mate availability reduced selfing in high-fitness individuals, and an

overabundance of apparent outcrossers and scarcity of apparent selfers when considering the

full dataset.

Superficially, the subtlety of these effects contradicts ecological models of selfing, which

assume an advantage and thus higher prevalence of selfing at low population density (43, 44).

So far, empirical evidence for reproductive assurance in animals is mixed. While some support

was found in androdioecious species such as the clam shrimp *Eulimnadia texana* (67) and the

nematode *C. elegans* (68), studies of hermaphroditic snails struggled to find any. Selfing rates

were not increased in low-density, disturbed, or temporary habitats (16, 19, 21, 31), neither the

frequency nor duration of outcrossing depended on the length of a mating opportunity (45), and

individuals did not adjust their waiting time to the perceived density of conspecifics (46). And

yet, it is hard to imagine that the capacity for independent reproduction does not feature among

the primary evolutionary drivers of selfing.

Our study offers a potential reconciliation between the reproductive assurance hypothesis and

its limited empirical support in animals. First, reproductive assurance matters, but only in

extreme situations. Consequently, there is an ascertainment bias: if a population is dense

enough to be sampled, it might be too dense for selfing to be common. Second, it is important

to understand that a population's selfing rate depends on the distribution of individual selfing

propensities among its members. As selfing propensities consist of an individual (*e.g.*,

genotype) and environmental component (*e.g.*, population density), the relationship between

density and population-level selfing rates is complicated. It depends on the relative frequency of

individuals that can respond to density variation with increased or decreased selfing, *i.e.*, on the

frequency of plastic individuals and selfers compared to outcrossers. Hence, population-level

selfing rates can be low because the population mainly consists of outcrossers, or because

density is high enough for outcrossing to occur even in plastic individuals and selfers.

Interpreting high population-level selfing rates is perhaps easier, as they necessitate a sizeable

proportion of self-fertile individuals. However, also in highly selfing populations density is

difficult to predict, because the threshold density beneath which selfing occurs may be

population- or species-specific.

These considerations might explain the apparent conflict between our field (near-zero selfing

rate; 26) and laboratory results (widespread self-fertility). We think it likely that the high

density of our field population prompts even selfers to reproduce through outcrossing.

Consequently, the evident self-fertility of large parts of the population does not manifest under

the current conditions. Interestingly, the commonness of selfing in the laboratory also among

repeatedly paired snails suggests that, compared to their free-living conspecifics, these snails

were still mate-limited. If so, then a future experiment with massively increased mate

availability should reduce selfing to zero.

So, what maintains self-compatibility alleles in our study population, and why have they not

gone to fixation? The cost of lacking self-compatibility is obvious – an occasional crash of

population size will quickly wipe out any individuals without self-compatibility alleles. The

benefits of lacking self-compatibility alleles, as demonstrated here, include a higher female

LRS, resulting from increased fecundity and decreased embryo mortality. Also juvenile

mortality was lower among outcrossed progenies. Hence, selfing did not only lead to inbreeding

depression in embryo development, but also reduced juvenile survival. The presence of

inbreeding depression in laboratory settings tallies with findings in other populations of *R.*

balthica (69) and in a close relative (65, 66), and presumably represents a major obstacle for the

spread of self-compatibility alleles. The reduced female lifetime fecundity of selfers (45%

fewer eggs than outcrossers) is interesting as well. We previously found that selfed egg clutches

were only half the size of (supposedly outcrossed) clutches laid after a female mating (49). Our

study shows that potential compensation mechanisms of selfers, such as a faster clutch-laying

rate, were – if at all present – clearly insufficient. Another benefit of lacking self-compatibility

alleles might be an increased male LRS, particularly if selfers have sperm discounting (70) or

reduced energy allocation to the male function (71). However, the strong sexual drive of selfers

documented here (80% mated in the male role) suggests that reductions in siring success when

selfing might be more modest. The ability to self while also acting as a paternal parent is a main

requirement of another hypothesis for the evolution of selfing: the 50% transmission advantage

of a selfer's genes over those of an outcrosser (43, 72). If future work finds this ability to be

common, then it provides a mechanism for maintaining self-compatibility alleles also at high

population density.

Finally, it is worth mentioning the dangers of assuming that pairing individuals results in their

offspring being outcrossed, as done in some early work (e.g., 65, 66). In our study, 44% of

paired snails selfed at least partially, and 71% of snails that selfed post-isolation copulated as a

female. Hence, outcrossing is neither guaranteed by pairing individuals once or repeatedly, nor

by observing female copulations. Potential reasons for a failure to outcross include, *e.g.*,

unsuccessful sperm transfer, cryptic female choice, sperm competition with autosperm, and

genetic incompatibility between sperm and egg. Hence, to verify outcrossing events, genetic

paternity analysis is unavoidable. Risks of failing to do so include underestimating the

frequency and extent of selfing, and underestimating the strength of inbreeding in studies that

compare the offspring of isolated parents (obligately selfed) with those of paired parents

(purportedly outcrossed though potentially selfed; e.g., 48, 73, 74, 75). We here add to a

growing number of studies that tested the assumption of (near-)exclusive outcrossing after

pairing empirically, and found it occasionally confirmed (76), yet more often refuted (34, 54,

77, 78).

Ethics

532   We performed our study in accordance with national laws.

Data accessibility

534   Microsatellite markers: GenBank Accession nos. KX830983-KX830992. All data are available

535   from Dryad X.

Authors' contributions

537   A.F. designed the study, performed experiments, carried out molecular work, and analysed the

538   data. A.F. and A.S. drafted the manuscript, with input from J.J. All authors contributed to

539   manuscript revision and gave final approval for publication.

Competing interests

541   We declare we have no competing interests.

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Tables

Table 1

GLMM on the probability of selfing post-isolation. Results are shown for F1 snails (n = 55)

with genotyped offspring that could be assigned to fathers. The response variable was the

likelihood of F1 snails producing at least one selfed juvenile after having been paired (no/yes),

modelled using binomial errors. Fixed effects were the pairing treatment (once- vs. repeatedly;

test of hypothesis 1), the number of developed embryos produced in isolation (test of

hypothesis 2), and the number of male (potential sperm donors) and female partners (potential

sperm recipients) a snail copulated with. P0 mother identity was included as a random intercept;

its effect was non-significant ($\chi^2_1 = 0.0$, $p = 1.0$). Results are provided on a logit scale. β:

estimate, S.E.: standard error, z: z-value, p: p-value.

Predictor	β (S.E.)	z	p
Intercept	-0.22 (0.82)	-0.26	0.79
Treatment (once- vs. repeatedly paired)	1.75 (1.77)	0.99	0.32
# dev. embryos produced in isolation	0.01 (0.01)	2.24	0.0251
# male mating partners	-0.82 (0.56)	-1.48	0.14
# female mating partners	-0.19 (0.41)	-0.46	0.64

Table 2

GLMMs on female LRS, its components, and juvenile mortality. Results are shown for four

models using all female fertile F1 snails. Response variables were the lifetime number of

developed embryos (female LRS) and eggs, the lifetime proportion of undeveloped embryos,

and the proportion of developed embryos that died before they could be genotyped (*i.e.*, before

reaching the juvenile age of 12.3 ± 0.5 weeks). For response variables 1 and 2, negative

binomial errors were fitted, and results are provided on the log scale. For the two response

variables that are proportions, Gaussian errors were fitted, as they did not deviate significantly

from normality and as model assumptions were fulfilled. Fixed effects were propensity for

selfing (reference level: apparent selfers) and body size (shell length). As random intercepts we

included P0 mother identity (significant: $\chi_1^2 \geq 5.59$, $p \leq 0.0181$, except for % dead juveniles: χ_1^2

$= 0$, $p = 1$) and, in all models except that of the number of eggs, pair identity on mating

opportunity 1 (non-significant: $\chi_1^2 \leq 2.51$, $p \geq 0.11$). Although there were 124 female fertile F1

snails, sample size in models was reduced because one snail lacked data on body size and ten

snails lacked developed embryos that were reared to the juvenile age. Results were very similar

when computed for snails with estimated selfing rates only (Table S3). S.E.: standard error, PS.:

propensity for selfing, app.: apparent.

Response	Sample size	Predictor	Estimate (S.E.)	*z*-value	*p*-value
# dev. embryos (LRS)	123	Intercept	2.59 (0.63)	4.11	<0.0001
		PS (app. plastic snail)	1.08 (0.18)	6.13	<0.0001
		PS (app. outcrosser)	0.89 (0.19)	4.81	<0.0001
		Body size	0.14 (0.04)	3.27	0.0011
# eggs	123	Intercept	2.75 (0.59)	4.64	<0.0001
		PS (app. plastic snail)	1.02 (0.17)	6.11	<0.0001
		PS (app. outcrosser)	0.83 (0.17)	4.79	<0.0001
		Body size	0.15 (0.04)	3.81	0.0001

Response	Sample size	Predictor	Estimate (S.E.)	*z*-value	*p*-value
% undev. embryos	123	Intercept	30.78 (14.87)	2.07	0.0384
		PS (app. plastic snail)	-12.77 (3.80)	-3.36	0.0008
		PS (app. outcrosser)	-11.14 (3.92)	-2.84	0.0045
		Body size	0.47 (1.07)	0.44	0.66
% dead juveniles	113	Intercept	106.09 (19.51)	5.44	<0.0001
		PS (app. plastic snail)	-7.80 (4.94)	-1.58	0.11
		PS (app. outcrosser)	-16.69 (5.12)	-3.26	0.0011
		Body size	-2.71 (1.38)	-1.96	0.0496

Figure legends

Figure 1

Experimental variation in mate availability did not affect individual-level selfing rates.

Snails had either one mating opportunity with one partner, or three to five, each with a different

partner. (a) Paternity distributions among the progenies of 56 F1 snails. Each vertical bar

depicts one F2 family, with differently coloured sections corresponding to juveniles sired by

different fathers. Father 1 refers to F1 snails' first and father 5 to their last mating partner.

Identical colours in different F2 families do not indicate identical father identities; all in all, F2

juveniles were sired by 68 unique fathers, with 10.5 ± 6.4 (mean ± SD) juveniles per father

(min = 1, max = 30). Within pairing treatments, families are arranged according to selfing rate

and number of genotyped juveniles. (b) Differences in mean selfing rates between once- (n =

31) and repeatedly paired snails (n = 24) were not statistically significant. White triangles on

boxplots show group means. (c) The proportion of F2 families with non-zero selfing rates was

similar among once- and repeatedly paired snails. (d) However, once-paired snails produced

significantly more selfed juveniles overall. Error bars are 95% confidence intervals (79).

Figure 2

Variation in individual propensity for selfing. We ascertained the selfing propensity of snails

with offspring that were successfully genotyped and assigned to a father (n = 55) based on the

presence/absence of selfing in isolation and the post-isolation selfing rate. (a) The distribution

of post-isolation selfing rates was bimodal among both once- and repeatedly paired snails. (b)

Selfing in isolation predicted continued selfing: predominant selfing after mating trials had

started only occurred in snails that had selfed successfully in isolation. (c) An a-posteriori

categorisation revealed four classes of snails with differential selfing propensity: outcrossers

that never reproduced via selfing, plastic switchers that selfed in isolation but outcrossed after

being paired, plastic mixers that mixed selfing in isolation with a low level of selfing post-

isolation, and selfers that selfed fully or predominantly throughout their lives. These categories

were very similar when based on the number of undeveloped embryos, or undeveloped and

developed embryos combined (see Fig. S1).

Figure 3

Reduced female LRS and increased offspring mortality in preferential selfers. Individuals

with a strong propensity for selfing had the fewest developed embryos (a, n = 123), the highest

proportion of embryos that failed to develop (b, n = 123), and the highest mortality rate among

their juvenile offspring (c, n = 113). Shown are snails with (black circles) and without

genotyped offspring (empty circles). We ascertained the apparent propensity for selfing of

snails without genotyped offspring based on the timing of egg production and the mating

behaviour. White triangles on boxplots show group means.

Figure 4

Poor correspondence between propensity for selfing and sexual drive. Although selfers (n =

15) were less likely to mate in both sexual roles than plastic switchers (n = 26), they showed

considerable sexual activity, with 53.3% mating as a female and 80.0% as a male. Error bars are

95% confidence intervals (79).

Figures

Figure 1

Figure 2

Figure 3

Figure 4