1 **Identifying the fitness consequences of sex in complex natural environments**
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<u>Abstract</u>

In the natural world, sex prevails, despite its costs. While much effort has been dedicated

to identifying the intrinsic costs of sex (e.g. the cost of males), few studies have identified

the ecological fitness consequences of sex. Furthermore, correlated biological traits that

differ between sexuals and asexuals may alter these costs, or even render the typical costs

of sex irrelevant. Here we use a large-scale transplant experiment of a North American

wildflower to show that sex is associated with reduced lifetime fitness, despite lower

herbivory. We separate the effects of sex from hybridity, finding that over-winter survival

is elevated in asexuals regardless of hybridity, but herbivores target hybrid asexuals.

Survival is lowest in homozygous sexual lineages, implicating inbreeding depression as a

cost of sex. Our results show that the consequences of sex are shaped by complex natural

environments, correlated traits, and the identity of mates, rather than sex itself.

Introduction

Despite over a century of research, sex remains a fundamental mystery in evolutionary biology. Although recombination removes deleterious alleles and unites beneficial ones (Hill & Robertson 1966; Keightley & Otto 2006), enabling organisms to adapt (Felsenstein 1974) and evolve with antagonists (Jaenike 1978), sex has many costs. These typically include the "twofold cost," wherein sexual organisms produce half as many offspring as asexuals due to their investment in male offspring (Maynard Smith 1978), as well as the disruption of co-adapted beneficial loci via recombination (Lehtonen *et al.* 2012). While theory has identified many costs and benefits of sex, these are likely mediated by the genomic composition of asexuals, the patterns of mating among sexuals, and the biotic and abiotic selective environment.

Despite widespread attention to the genetic processes maintaining sex, the ecological context in which sex occurs has received less attention. This is a crucial gap in our knowledge, as the processes governing sexual/asexual dynamics occur in complex natural environments. Our understanding of the ecology of sex stems largely from decades of experimental work on the freshwater snail, which has established sterilizing trematode parasites as the primary selective agent maintaining sex via frequency-dependent selection. This system offers a key example of Red Queen dynamics, wherein sexual reproduction generates novel lineages that avoid infection. Water depth moderates parasite frequency, leading to a striking ecological pattern: sex predominates in shallow water, where secondary hosts occur, while asexuality is widespread in deeper sites (Lively 1987; Jokela & Lively 1995; Lively & Dybdahl 2000; Vergara *et al.* 2014). This

research shows that the abiotic and biotic environment operate in concert to shape

sexual/asexual dynamics in the natural world. Critically, both ecology and its interactions

with a system's biological features provide the context necessary to pinpoint the real-life

costs and benefits of sex (Neiman *et al.* 2018).

Similarly, very few studies have accounted for the widespread observation that

sex and asex are often differently associated with fundamental biological and life history

traits, despite the critical role these traits play in shaping the costs of sex (Meirmans *et al.*

2012). For example, the cost of males will be minimized in plants with hermaphroditic,

self-compatible flowers (Charlesworth 1980; Mogie 1996); as such, frequent inbreeding

will further minimize this cost. Additional costs of asexuality may be rooted in the origin

and composition of asexual lineages. For example, while polyploidy and hybridity are

overrepresented in asexual plants and vertebrates as compared to sexuals (e.g.,

Vrijenhoek 1978; Asker & Jerling 1992; Lutes *et al.* 2010), the impact of these key traits

is rarely considered. Further costs of sex may be related to mate choice (inbreeding or

outcrossing) and mate availability for sexual organisms, which have profound phenotypic

and genetic consequences (Meirmans *et al.* 2012). Hybridization, or outcrossing between

widely divergent taxa, alters numerous fitness-related traits (Yakimowski & Rieseberg

2014), which has repercussions for hybrid asexuals. The genetic impacts of mate choice

will similarly influence the evolution of sex. For example, sex via inbreeding can expose

a recessive genetic load (Charlesworth & Willis 2009), while asexual reproduction may

maintain heterozygosity. Such life history traits likely contribute substantially to

sexual/asexual dynamics, but are rarely discussed in studies of the costs and benefits of

sex.

The mustard genus *Boechera* offers a rare opportunity to study the ecological

costs of sex while accounting for multiple sex-associated biological traits. Apomixis

(asexual reproduction via seed) is widespread in the genus (Böcher 1951; Schranz *et al.*

2005; Aliyu *et al.* 2010). These asexuals may be polyploid or diploid, enabling separate

assessment of the effects of polyploidy and asexuality (Kantama *et al.* 2007; Beck *et al.*

2012; Mau *et al.* 2015). Furthermore, lineages that reproduce asexually are derived from

outcrossing events, which may be either interspecific or intraspecific (Li *et al.* 2017;

Rushworth *et al.* 2018), allowing separable examination of hybridity and asexuality.

Finally, similar to other asexual systems (e.g., Lutes *et al.* 2010; Fradin *et al.* 2017),

asexuals are highly heterozygous, whereas sexual *Boechera* are highly self-fertilizing and

thus highly homozygous (Roy 1995; Song *et al.* 2006; Li *et al.* 2017). This particular trait

is of broad relevance to all sexual/asexual systems, as all sexual organisms undergo shifts

in population size and variable biparental inbreeding. In other words, the consequences of

sex will always depend in large part on the identity of available sexual partners. For this

reason, the well-characterized mating system of *Boechera* offers an ideal system for

studying the evolution of sex.

Using a reciprocal transplant experiment in intact native habitat, we identified the

fitness components underlying the costs of sex in *Boechera*. We examined female

lifetime fitness and insect herbivory of sexual and asexual plants, finding strong over-

winter viability selection against sexual lineages, which is lessened in heterozygous

sexuals. This survival difference resulted in increased lifetime fitness for asexual

lineages, despite elevated insect herbivory on hybrid asexuals. Importantly, the negative

impacts of herbivory appeared in later life stages, reducing the probability of survival to a

second year, consistent with a cost of hybridization. Numerous lines of evidence suggest

that the cost of sex in this system is due at least in part to inbreeding depression in highly

homozygous sexual lineages.

Materials and Methods

Study system

Boechera retrofracta (Brassicaceae) is a short-lived perennial wildflower native to

western North America. Multiple *Boechera* species are highly self-fertilizing (Hamilton

& Mitchell-Olds 1994; Roy 1995; Song *et al.* 2006), and *B. retrofracta*'s high

microsatellite homozygosity (mean homozygosity=0.94, Rushworth *et al.* 2018) and

small flowers that self-pollinate before anthesis (C. Rushworth, pers. obs.) suggest

frequent self-fertilization. Nonetheless, *B. retrofracta* forms hybrids with at least 12

species (Windham & Al-Shehbaz 2006). Asexuality is tightly associated with outcrossing

and hybridization in *Boechera* (Schranz *et al.* 2005; Kantama *et al.* 2007; Beck *et al.*

2012; Rushworth *et al.* 2018). Apomixis in this system is pseudogamous, requiring

endosperm (but not embryo) fertilization (Böcher 1951). Because sexuals are highly

selfing and asexuals must also produce a small amount of viable pollen, several of the

traditional costs of sex are minimized in this system, allowing investigation of broader

ecological and biological factors maintaining sex.

Sampling and growth methods

We used 53 wild-collected diploid genotypes (24 sexual, 29 asexual) in a multi-site

reciprocal transplant experiment (see Table S1 in Supporting Information). Species (or

parental species, in the case of asexuals) were determined based on microsatellites and

morphology (Rushworth *et al.* 2018) using the *Boechera* Microsatellite Website (Li *et al.*

2017). Heterozygosity, estimated from microsatellites, was calculated for each maternal

lineage. 21 sexual genotypes were entirely homozygous, and three were not

(heterozygosity ranging from 0.08 – 0.42); heterozygosity of all but one asexual

genotypes was >0.5 (Figure S1). Eighteen asexual genotypes are "hybrid asexuals,"

resulting from interspecific hybridization events, while eleven are "non-hybrid asexuals,"

resulting from within-species outcrossing. One sexual genotype was not *B. retrofracta*,

and was not analyzed; all asexual genotypes but one ("A5") involved *B. retrofracta* as a

parental species. Other parental species represented in hybrids were *Boechera*

sparsiflora, *Boechera exilis*, *Boechera pendulocarpa*, *Boechera stricta*, and *Boechera*

puberula. To ensure that experimental selection pressures approximated those of each

genotype's native environment, we selected lines based on proximity or similarity of

collection and garden environments. All but two genotypes were collected within 10km

of an experimental garden.

To minimize maternal environmental effects, genotypes were grown in controlled

conditions for one generation before the experiment. Seeds were germinated on wet filter

paper in petri dishes and transplanted as seedlings onto a blend of Fafard 4P Mix and

MetroMix 200 (Sun Gro Horticulture, Agawam, MA, USA) in Ray Leach Cone-tainers

(Steuwe and Sons, Tangent, OR, USA) in the Duke University greenhouses, and shipped

as rosettes for transplanting at six weeks of age. Transplantation into replicated,

randomized blocks occurred in fall for data collection the following season, allowing

synchronization of flowering time with the native vegetation.

Transplant experiment

Subsets of the 53 genotypes were planted in two cohorts of replicated, randomized blocks

in the fall of 2012 and 2013. Data was collected for two growth seasons for each planting

year, totaling three years of data collection. Planting year one consisted of 2000

individuals representing 19 asexual and 22 sexual genotypes, planted in five gardens with

400 plants/site (ALD, CAP, CUM, MIL, and SIL; Table S1). In year two, 2080

individuals representing 18 asexual and 21 sexual lineages were transplanted at the same

sites (520 plants/site; Figure S2). However, the SIL site was excluded from year two

because its dense vegetation and high humidity are atypical *B. retrofracta* habitat. Over-

winter survival was scored the following spring. Plants that survived were measured for

width, height, flower number, and fruit number. Due to low reproduction, MIL in year

one (N=34 reproducing plants) and CUM in year two (N=2) were excluded from

analyses. Leaf damage was quantified visually by trained observers, as in Prasad et al.

(2012). The CUM site was excluded in herbivory analyses, as no herbivory

measurements were made there. Prior to fruit dehiscence, 1 – 4 fruits were collected from

2 – 3 replicates of each genotype within each garden. Seeds were counted from each fruit

and averaged across replicates to estimate average seed set per genotype per garden.

Fecundity was estimated as average seed set per fruit multiplied by individual fruit

number. This average was rounded to the nearest integer to allow us to model seed set as

a negative binomially distributed random variable.

Pollen viability

To evaluate how sexual system impacted pollen viability, we grew 25 sexual and 26

asexual genotypes in a growth chamber at the Duke University Phytotron. We collected

mature developing buds (stage 12, Smyth *et al.* 1990) from 1 – 3 individuals per

genotype (N=112 individual plants), and stained pollen to differentiate viable from

inviable grains (Peterson *et al.* 2010), averaging pollen viabilities across replicates. For

each individual, 100 pollen grains were randomly counted from a microscopy slide

visualized on a Leica MZ7.5 Stereozoom dissecting microscope and imaged using a

Canon EOS Rebel T3i digital camera.

Statistical analyses

Our primary aim was to evaluate the association between Reproductive System

(*RS*, a factor which takes the value of sexual or asexual), and group (*group*, a factor

which can take the value of sexual, hybrid asexual, or non-hybrid asexual) with fitness

components over time and space. Because sexuals and asexuals differ intrinsically in

heterozygosity, any observed fitness differences may stem from inbreeding or

outbreeding depression. We conducted additional analyses to address this possibility.

All analyses were performed using R version 3.5.2 (R Core Team 2018). Outliers

were visually identified using an adjusted boxplot analysis suitable for far-right skewed

data (Hubert & Vandervieren 2008), with robustbase (Maechler *et al.* 2019). To analyze

fitness data, we used generalized linear mixed models (GLMMs) in glmmTMB (Brooks

et al. 2017). Model diagnostics were conducted using DHARMa (Hartig 2019) and

marginal means were estimated using the package ggeffects (Lüdecke 2018). Lifetime

fitness was estimated using a zero-inflated negative binomial distribution with the

canonical link functions, where structural zeros represent plants that did not survive the

winter and the conditional portion of the model represents fecundity of survivors.

Separate fecundity and survival models were also run, using the negative binomial and

binomial error distributions, respectively. The model used was:

220                         $y \sim RS + garden + RS{*}garden + year + RS{*}year$

where y indicates a fitness component, lifetime fitness, or herbivory. Two random effects

were included in all models: block nested in site and genotype nested in *RS*. The same

structure was used for the zero-inflation and conditional portions of the zero-inflated

negative binomial model for lifetime fitness. In all estimable models, a random effect of

genotype nested in reproductive system, crossed with garden, accounted for essentially

no variance, and was therefore removed from analyses. A scaled covariate for width at

planting was included in survival models, while a scaled covariate of plant height was

added to all herbivory models to account for a link between plant size and insect

attraction.

231   We next tested for fitness differences between hybrid asexual genotypes, non-

hybrid asexual genotypes, and sexual genotypes, by running the same models as above,

replacing the *RS* term with the *group* term. To assess the effects of heterozygosity on

survival, we used fixed-effects only models to analyze survival of heterozygous and

homozygous sexual lineages, and complementary analyses (heterozygous sexuals vs.

asexuals; homozygous sexuals vs. asexuals) were also conducted. Significance testing for

fitness analyses was conducted via likelihood ratio tests (Bolker *et al.* 2009) with Holm

corrections (Holm 1979).

Leaf damage was log-transformed and analyzed using a linear mixed model

(LMM) in lme4 (Bates *et al.* 2015). Significance testing was conducted using *F* tests with

Kenward-Roger approximation of denominator degrees of freedom (Kenward & Roger

1997) using pbkrtest (Halekoh & Højsgaard 2014).

To understand the relationship between herbivory and fitness, and how it differs

among groups, two additional models were estimated with log herbivory and its

interaction with group and garden as predictors. Each model examined a fitness

component: fecundity and survival to a second year, both of which are conditional upon

survival to year one and non-zero herbivory. The model specified was:

y ~ group + garden + year + group*garden + group*year + herbivory +

250   *group*herbivory + garden*herbivory*

Results

Higher asexual lifetime fitness due to survival advantage

Higher asexual lifetime fitness: Lifetime fitness, the product of survival and

reproduction, was higher in asexuals than sexuals (χ^2=11.90, df=2, *P*=7.82×10^{-3}). The

effect of RS depended on year (χ^2=34.74, df=2, *P*=1.72×10^{-7}) and garden (χ^2=23.45,

df=8, *P*=0.016; Table S2). To disentangle the effects of hybridization and sex, we

analyzed the same data with the group term. This model similarly showed an effect of

group (χ^2=13.33, df=4, P=0.039), and its interaction with both year (χ^2=36.87, df=4,

P=1.53×10^{-6}) and garden (χ^2=66.70, df=16, P=3.00×10^{-7}; Table 1, Table S2). Estimated

lifetime fitness for each group varied across year and site, but was elevated in both

asexual groups, with consistently high hybrid asexual fitness (Figure S3). While our

model design did not allow us to predict overall lifetime fitness for each group without

accounting for interactions, raw summaries showed higher overall fitness in both groups

of asexuals, although years varied slightly (Year 1 means: sexual, 211.54; hybrid asexual,

289.40; non-hybrid asexual, 299.93; Year 2 means: sexual, 76.29; hybrid asexual, 181.57;

non-hybrid asexual, 145.55; Figure S4). Estimated marginal means for all models are in

Appendix S1.

Higher asexual survival: We next asked if fitness disparities were attributable to

differences in survival or fecundity. The zero-inflation portion of a zero-inflated negative

binomial GLMM addresses structural zeros, or the probability of not surviving to

reproduce. Over-winter survival was significantly elevated in asexual genotypes

compared to sexuals. In year 1, 81% of asexuals survived, while 64.3% of sexuals

survived. In year 2, 56.5% of asexuals survived, compared with only 23.8% of sexuals.

Survival differed by reproductive system (χ^2=11.82, df=1, P=2.35×10^{-3}), and

across year (χ^2=9.16, df=1, P=0.04) and garden (χ^2=46.73, df=4, P=1.21×10^{-8}; Table

S3A). In both years asexuals had higher survival; the interactions between year and RS,

and garden and RS, were also significant (year: χ^2=22.88, df=1, P=8.60×10^{-6}; garden:

χ^2=16.23, df=4, P=0.016; Table S3A). Similar results were seen in a separate model of

survival (Table S4A). Plant size also increased plant survival (χ^2=54.20, df=1,

P=7.24×10^{-13}).

Over-winter survival differed by group (χ^2=12.22, df=2, P=0.01; Table 1, Table

S3B), with both types of asexuals surviving better than sexuals in both cohorts (Figure

1A). The effects of group differed temporally (group × year interaction, χ^2=23.56, df=2,

P=5.36×10^{-5}) and spatially (group × site interaction, χ^2=42.82, df=8, P=6.66×10^{-6}; Figure

1B, Table 1, Table S3B). A separate model of survival showed similar results (Table

S4B). Hybrid asexuals were larger than other groups at planting; rosette size (χ^2=53.79,

df=1, P=7.24×10^{-13}; Table S4B). Across gardens, hybrid asexuals had higher predicted

survival than sexuals, while non-hybrid asexual survival was more variable (Figure 1B).

In three of five sites, survival was roughly equivalent in both asexual groups. In two sites,

survival was similarly low for sexuals and non-hybrid asexuals.

Heterozygosity also impacted survival. Survival of heterozygous sexuals

exceeded that of homozygous sexuals (χ^2=7.07, df=1, P=0.024; Figure S5, Table S5), but

did not result in increased total fitness for heterozygous sexuals (Table S6). Asexual and

heterozygous sexual survival did not differ (χ^2=0.47, df=1, P=1). Together, these results

are consistent with inbreeding depression manifesting as reduced over-winter survival in

homozygous sexual lineages.

No differences in fecundity: We next examined the conditional portion of the zero-

inflated negative binomial GLMM, which analyzed fecundity (seed set conditional on

survival). In total, 995 individuals set seed (597 asexuals and 398 sexuals); removing

nine outliers left a dataset of 986 individuals. Seed set did not differ intrinsically between

sexuals and asexuals, but the influence of RS varied by year (χ^2=11.86, df=1, P=1.72×10$^-$

^{3}; Table S3A). The conditional group model produced similar results: although the main

effect of group was not significant, non-hybrid asexual fecundity was lowest in both

years (Figure S6A). The influence of group varied by year (χ^2=13.32, df=2, P=5.13×10^{-3};

Table 1, Table S3B) and by garden (χ^2=23.70, df=8, P=0.01; Table 1, Table S3B, Figure

S6B). There were substantial environmental effects on seed set (Table S3). Similar results

were found in the separate fecundity models (Table S7). Pollen viability assays revealed

no difference between sexuals and asexuals (pooled t-test, t=0.599 on 49 df, P=0.55),

suggesting that any differences in seed set may be due to either maternal investment or

pollen quantity. Heterozygosity had no impact on sexual fecundity (χ^2=1.24, df=1,

P=0.26; Table S8).

Herbivores targeted hybrid asexuals

33% of the experiment (1379 of 4080 plants) experienced leaf herbivory. The removal of

29 high-herbivory outliers resulted in a dataset of 1350 individuals (809 asexuals, 541

sexuals). Asexual herbivory was higher than sexual, with an average of 5.45%±0.002se

leaf area damaged per asexual individual (estimated marginal mean 2.14%±0.048se,

95%CI 1.72 – 2.66%). By comparison, mean sexual leaf damage across years was

3.16%±0.002se (estimated marginal mean 1.23%±0.05se, 95%CI 0.98 – 1.55%), a

significant difference (F=22.55, df=1, 48.29, P=9.33e-05; Table 2A, Figure 2). Herbivory

varied by garden (F=6.40, df=3, 62.99, P=2.99×10^{-3}; Table 2A, Figure S7), consistent

with spatial variation in herbivore communities. Herbivory was highest in hybrid

asexuals (Figure 2). Group had a significant effect on herbivory (F=18.08, df=2, 42.39,

$P=1.26\times10^{-5}$; Table 2B), with both hybrid and non-hybrid asexuals experiencing higher

leaf herbivory than sexuals (Figure 2). Plant height was significant in both models (Table

2).

Herbivory negatively impacted survival to subsequent years

What are the effects of herbivory on fitness, and do these effects differ between sexuals

and asexuals? We next predicted two separate fitness components, fecundity and survival

to a second year, using models incorporating a main effect of log-transformed herbivory,

and its interaction with both group and garden.

Over 90% of experimental fitness occurred in the first year of life. In both years,

more asexuals than sexuals survived to a second year. In experimental year 1, 8.2% of

plants survived for a second year (N=163: 108 asexuals, 55 sexuals). Similarly, 5.9% of

plants in year 2 survived for a second year (N=123: 104 asexuals, 19 sexuals). Roughly

1% of plants reproduced in their second year (Year 1: 1.4%, N=28: 24 asexuals, 4

sexuals; Year 2: 1.3%, N=26: 25 asexuals, 1 sexual). Nevertheless, survival to subsequent

years could substantially increase lifetime fitness.

In temperate montane environments, herbivory and seed set co-occur during the

short growing season, and may thus influence one another. We found no effect of

herbivory on fecundity ($\chi^2=0.28$, df=1, $P=0.598$; Table S9), but herbivory decreased

survival to a second year ($\chi^2=21.26$, df=1, $P=8.02\times10^{-6}$; Table S10). Group also affected

subsequent survival ($\chi^2=32.88$, df=2, $P=5.79\times10^{-7}$; Table S10), with more hybrid

asexuals than non-hybrids or sexuals surviving (Year 1: 82 hybrid asexuals, 17 non-

hybrid asexuals, 41 sexuals; Year 2: 75 hybrid asexuals, 28 non-hybrid asexuals, 17

sexuals). In all groups, herbivory and probability of survival were negatively correlated

(Figure 3). Although the interaction between group and herbivory was non-significant,

the negative relationship was strongest in hybrid asexuals (Figure 3), consistent with

herbivory limiting hybrid asexual fitness.

<u>Discussion</u>

Here we show that both abiotic and biotic factors play key roles in the maintenance of sex

in nature. While viability selection favors both inter- and intraspecific asexuals,

implicating shared causative traits such as asexuality or heterozygosity, biotic

interactions are especially influenced by hybridization. Together, these results suggest

that, although survival of adverse abiotic conditions is facilitated by asexuality or

associated heterozygosity, hybridization plays a singular role in the interaction between

asexuality and herbivory. The precise targets of selection are currently unknown, but our

results suggest that the costs of sex depends on ones' sexual partners, and may be

mediated by interactions with biological antagonists.

The complex ecology of sex

Despite extensive theoretical work and recurrent calls for field-based studies

(Agrawal 2006, 2009), few researchers have examined the ecology of sex. Identifying

ecologically-relevant traits associated with sex, and the selective processes acting in

natural environments, could elucidate the phenotypes and ultimately the molecular

mechanisms underlying the cost of sex in wild populations.

Although numerous selective agents are likely important to the evolution of sex

(Neiman *et al.* 2018), we have little understanding of how these biotic and abiotic

selection pressures maintain sex in the wild. Previous work shows that sex may permit

survival during abiotic environmental shifts (Stross & Hill 1965) or enable avoidance of

fitness-reducing biotic interactions (Lively 1987). We found that strong viability

selection early in life results in higher lifetime fitness for asexual lineages, but increased

insect herbivory may provide an upper limit on asexual fitness. That different agents act

at different life stages suggests that long-term studies are key to understanding the

evolution of sex.

Many of the best-known asexual systems are polyploid (e.g., Fox *et al.* 1996;

Thompson & Whitton 2006), hybrid (e.g., Xu *et al.* 2015) or both (e.g., Lutes *et al.* 2010;

Gutekunst *et al.* 2018). Yet these biological features are rarely considered in either

studies of these systems or in broader discussions of the costs of sex. We explicitly

considered traits correlated with reproductive mode in assessing patterns of selection on

sexual reproduction, finding that viability is especially reduced in homozygous sexual

lineages, consistent with inbreeding depression acting as a cost of sex. Hybrid asexuals

were more prone to herbivory, suggesting that the costs of asexuality may be related to

the correlated trait of hybridity. Clearly, selective processes that influence the evolution

of sex are shaped by both the origin and composition of asexual lineages and patterns of

mating in sexuals.

Mating system influences the evolution and ecology of sex

The fundamental role of sex in all organisms is to reshuffle variation across the

genome through recombination and segregation. But the effects of these processes are

tempered by the efficacy of such shuffling. When sexual partners resemble one another

due to population structure, limited population genomic variation and/or inbreeding, the

traditional costs and benefits of sex can change dramatically (De Visser & Elena 2007).

For example, self-fertilization cannot unite beneficial mutations arising in different

lineages. Practically speaking, the costs and benefits of sex are always a function of mate

availability and identity.

Despite the widespread existence of asexual reproduction in flowering plants,

little is known about the costs of sex in angiosperms. Researchers studying the evolution

of sex in natural populations have largely focused on animal systems, where the existence

of separate sexes provide clear predictions about the relevant costs and benefits

(Lehtonen *et al.* 2012). Although hermaphroditic asexuals are still predicted to have a

1.5-fold fitness advantage, sex via self-fertilization muddles these predictions

(Charlesworth 1980), as do additional biological traits common in plants (Mogie 1996).

For example, both selfing and pseudogamous apomictic *Boechera* must produce

sufficient pollen for self-fertilization, reducing the applicability of any cost of males.

Additionally, the strong link between hybridization and asexuality in numerous vertebrate

and plant taxa (Asker & Jerling 1992; Kearney 2005) suggests that recurrent independent

hybridization events generate novel asexuals. In *Boechera*, both sexual and asexual

lineages have limited variation, but high diversity among asexual lineages may be

generated by multiple independent origins of asexuality. Such cases require consideration

of additional costs and benefits of sex absent from most models.

We hypothesize that inbreeding depression manifests as the primary cost of sex in

sexual *Boechera*. Our experiment showed fitness costs and benefits associated with sex

that follow predictions for inbreeding depression. Sexuals had reduced overwinter

survival (Figure 1), particularly in homozygous lineages (Figure S4), that varied both

spatially and temporally (Figure 1, Table 1, Table S4), consistent with inbreeding

depression (Armbruster & Reed 2005; Cheptou & Donohue 2010). Our experiment

contained few heterozygous sexuals, and was thus underpowered to detect a strong

relationship between heterozygosity and fitness. Nonetheless, our results suggest that the

main cost of sex in *Boechera* may be related to deleterious load, and that outcrossing may

be sufficient to increase survival in this system. This pattern, known as a heterozygosity-

fitness correlation or HFC, is widespread in wild systems (Szulkin *et al.* 2010). Despite

reduced survival in homozygous sexual lineages, we saw no evidence that survival

differed between asexuals and heterozygous sexuals (Table S5), consistent with the

hypothesis that the benefit of heterozygosity is the masking of genome-wide recessive

load.

Although inbreeding likely poses a main cost of sex in this system, other factors

may also be at play. First, asexual lineages likely arise frequently and those found in

nature have survived selection, potentially resulting in an overrepresentation of high-

fitness asexual genotypes in the field. Indeed, the majority of de novo *Boechera* hybrids

have lower fitness than their selfed progenitors (Rushworth & Mitchell-Olds 2020),

suggesting that fitness variation may be more common in the wild than our experiment

suggests. Additionally, the recurrent origins of asexual lineages (Sharbel & Mitchell-Olds

2001) suggest that asexuals comprise a range of ages. With deleterious mutations

accumulating over the lifespan of a given asexual lineage, older lineages may have

reduced fitness. Lastly, reproductive isolation that varies across populations may also

result in fitness that depends on interactions between parental genotypes. Each of these

scenarios is plausible and provides avenues for further research.

Self-fertilization also has clear benefits. Sex and recombination enable adaptation

to novel environments (Felsenstein 1974), and inbreeding may speed local adaptation. In

situations where environmental changes are substantial, fixation of beneficial recessive

alleles through self-fertilization results in rapid adaptation (Glémin & Ronfort 2012).

Self-fertilization will also reduce the potential for reshuffling of beneficial allelic

combinations, as high homozygosity limits opportunities for functional disruption

(Hartfield *et al.* 2017). Our results showed that sexual lineages experienced less insect

herbivory than asexuals (Figure 2). Local adaptation underlies variation in chemical

defense both among populations of *Boechera stricta* (Prasad *et al.* 2012) and between

species (Windsor *et al.* 2005). We thus hypothesize that interactions between divergent

alleles in outcrossed asexuals disrupt chemical defenses, consistent with outbreeding

depression.

Herbivory as an ecological cost of asexuality

Classic literature posits multiple costs of asexuality, including the accumulation of

deleterious mutations and the extinction of asexual clones with the highest mutation load

(Müller 1964; Felsenstein 1974). While asexual *Boechera* do show elevated putatively

deleterious mutations (Lovell *et al.* 2017), the heterozygous state of these mutations, the

higher fitness of asexuals observed here, and the replenishment of asexuals by novel

hybridization events, suggests that these intrinsic genetic costs have relatively little to do

with the costs of asexuality in *Boechera*.

Rather, we found evidence for a short-term ecological cost associated with

asexuality in this system – a 73% increase in asexual insect herbivory (Figure 2),

consistent with predictions from theoretical literature (Stearns 1987). This pattern may be

underlain by asexuality, heterozygosity, hybridization, or some combination of all three.

In functionally asexual *Oenothera*, generalist herbivory is elevated across multiple

species (Johnson *et al.* 2009), supporting a role for asexuality in the macroevolution of

plant defense. Microevolutionary patterns suggest the same; viral infection is elevated in

asexual and common sexual grass genotypes (Kelley 1994). Although elevated in both

asexual groups, insect herbivory is highest in hybrid asexuals (Figure 2). Ample research

also shows that hybridization strongly alters plant defense and herbivore preference

(Strauss 1994; Fritz 1999; Whitham *et al.* 1999; Orians 2000), and hybrid taxa may

produce greater or lesser quantities of chemical compounds, and/or novel ineffective

compounds (Orians 2000). Disentangling the influence of asexuality and hybridization on

defense offers a fruitful avenue for future research.

While the costs of herbivory were insufficient to overcome the other benefits of

asexuality in our two-year study, these costs could manifest in other traits or at later life

stages (Marquis 1992). We found that herbivory decreased survival to subsequent years

(Figure 3), consistent with the negative correlation between fitness and herbivory

documented in *Boechera stricta* (Prasad *et al.* 2012). Similar patterns have been

identified in animals, suggesting that downstream consequences of parasitism or

herbivory are broadly applicable. One recent study of Soay sheep showed that fecundity

and parasite load collectively reduce overwinter survival (Leivesley *et al.* 2019).

Additionally, we found that the relationship between second-year survival and herbivory

is strongest in hybrid asexuals, suggesting that hybridization may limit the spread of

asexuality via herbivory. However, more hybrids survive to a second year of life overall.

Further work, including herbivore exposure experiments, are needed to understand the

interplay between fitness and herbivory in this system.

Collectively, the work presented here shows that sexual/asexual dynamics in wild

plant populations are largely shaped by environmental complexity and correlated traits,

including mating system. Asexual *Boechera* have consistently higher fitness and higher

herbivory damage than self-fertilizing sexual lineages, and survival is lowest in

homozygous sexuals. Our results highlight the importance of both the biotic and abiotic

environment in the evolution of sex, and suggest that inbreeding and outbreeding

depression underlie the ecological costs and benefits of sex in this system. Collectively,

this work shows that the complex natural environment and the choice of a sexual partner

work in tandem to shape patterns of reproductive variation in the wild.

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Tables and Figures

Table 1. **Reproductive system (sexual vs. asexual) and hybridity predict lifetime**

fitness via spatially- and temporally-variable selection. Results for the main effect of

group and two interactions (group and garden, and group and year) for zero-inflated

negative binomial GLMMs of lifetime fitness. Left, the zero-inflation portion models

structural zeros or plants that did not survive to reproduce. Center, the conditional portion

models seed set in plants that survived (fecundity). Right, significance estimates for the

overall model, incorporating both portions. Group indicates sexuals, "sex" vs. asexual

hybrids, "hyb" vs. asexual non-hybrids, the reference category. Estimates come from

conditional models, while test statistics (χ^2 deviance, degrees of freedom, and P values)

come from likelihood ratio tests for each overall effect. Significant P-values are in bold.

Full results from each separate model are in Supplementary Information.

term	condition	df	Zero-inflation				Conditional				Overall model		
			coef	se	χ^2	P	coef	se	χ^2	P	df	χ^2	P
Group	sex	2	0.339	0.468	12.22	**0.01**	-0.329	0.254	1.12	0.57	4	13.33	**0.039**
	hyb		-0.528	0.487			-0.357	0.250					
Group* Garden	sex*CAP	8	0.129	0.342	42.82	**6.66×10⁻⁶**	0.048	0.221	23.70	**0.01**	16	66.70	**3.00×10⁻⁷**
	hyb*CAP		0.977	0.349			0.485	0.208					
	sex*CUM		-0.596	0.495			0.377	0.319					
	hyb*CUM		0.376	0.517			0.682	0.322					
	sex*MIL		-0.278	0.394			0.052	0.233					
	hyb*MIL		1.063	0.402			0.435	0.228					
	sex*SIL		-1.167	0.661			1.843	0.481					
	hyb*SIL		-1.746	0.663			1.895	0.463					
Group* Year	sex*2	2	1.103	0.378	23.56	**5.36×10⁻⁵**	0.594	0.194	13.32	**5.16×10⁻³**	4	36.87	**1.53×10⁻⁶**
	hyb*2		-0.402	0.407			0.071	0.193					

Table 2. **Herbivory varies by reproductive system, hybridity, and garden.** (A) Model

results for reproductive system ("RS", sexual vs. asexual) LMM and (B) group (sexual

"sex" vs. asexual hybrid "hyb" vs. asexual non-hybrid, the reference category) LMM.

Significant P-values are in bold. Estimates apply to the conditional models, while test

statistics (F statistics, degrees of freedom, and P values) come from significance tests for

each overall effect.

A.

term	condition	estimate	se	df	*F*	*P*
	intercept	-1.633	0.071	-	-	-
RS	sexual	-0.299	0.079	1, 48.29	22.55	**9.33×10⁻⁵**
Garden	CAP	-0.002	0.074	3, 62.99	6.40	**2.99×10⁻³**
	MIL	0.340	0.095			
	SIL	-0.002	0.103			
RS*Garden	sexual*CAP	0.049	0.080	3, 1280.95	1.74	0.372
	sexual*MIL	0.059	0.115			
	sexual*SIL	0.234	0.105			
Year	2	-0.045	0.076	1, 67.59	0.38	1
RS*Year	sexual*2	-0.013	0.084	1, 1175.63	0.02	0.876
Plant height	-	0.045	0.016	1, 1318.25	7.52	**0.019**

765 B.

term	condition	estimate	se	df	F	P
	intercept	-1.719	0.095	-	-	-
Group	sex	-0.210	0.010	2, 42.39	18.08	**1.26×10⁻⁵**
	hyb	0.141	0.102			
Garden	CAP	-0.035	0.098	3, 62.98	6.37	**2.99×10⁻³**
	MIL	0.332	0.122			
	SIL	0.239	0.149			
Group*Garden	CAP*sex	0.083	0.102	6, 1273.93	1.99	0.065
	CAP*hybrid	0.060	0.102			
	MIL*sex	0.067	0.139			
	MIL*hybrid	0.016	0.132			
	SIL*sex	-0.005	0.150			
	SIL*hybrid	-0.324	0.152			
Year	2	-0.087	0.099	1, 66.98	0.30	1
Group*Year	sex*2	0.023	0.105	2, 1056.46	0.45	0.77
	hybrid*2	0.092	0.105			
Plant height	-	0.042	0.016	1, 1320.42	6.96	**0.019**

Figure 1. **Asexuals survival is higher than sexual.** Estimated marginal means for

survival show substantial temporal (A, group by year interaction) and spatial variation (B,

group by garden interaction). (A) Survival is reduced in sexuals in both experimental

years. (B) In all sites, hybrid asexuals have higher over-winter survival than sexuals; in

three of five sites, non-hybrid asexual survival is similarly high. Bars show 95%

confidence intervals. SIL was only included in year one, but fitness was predicted at this

site.

Figure 2. **Herbivory is elevated in hybrid asexuals.** Estimated marginal means for

herbivory from LMMs show the main effects of group (sexual vs. non-hybrid asexual vs.

hybrid asexual, left) and reproductive system (sexual vs. asexual, right). Bars show 95%

confidence intervals.

Figure 3. **Plants with less herbivory are more likely to survive to the next year.**

Herbivory has a significant impact on survival to a second year, suggesting that the

detrimental consequences of herbivory may not be observed in a single growing season.

Although the correlation between year 2 survival and herbivory differs qualitatively by

group (sexual vs. non-hybrid asexual vs. hybrid asexual), the interaction is not

significant. Estimated marginal means are shown averaged across five different values of

log herbivory. Points are jittered for ease of reading. Bars show 95% confidence

intervals.

Figure 1

804   **Figure 2**

Figure 3