Title: Reduced seed viability and reductions in plant size provide evidence for costs of

glyphosate resistance in an agricultural weed

Short title: Fitness cost of herbicide resistance

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Summary

• Although fitness costs of adaptation to herbicide are widely expected, they are not

universally detected, calling into question their generality. This lack of identified

costs could be due to experimental constraints such as a limited focus on

measures of plant fecundity and low replication.

• Here we examine the potential for a cost of herbicide resistance on seed

germination, root growth, and above-ground growth using 43 naturally occurring

populations of *Ipomoea purpurea* that vary in their resistance to RoundUp®, the

most commonly used herbicide worldwide.

• We show substantial costs in all three traits. Highly resistant populations had

lower germination rates, slower root growth and slower above-ground growth. A

visual exploration of the data indicated that the type of cost may differ and even

trade off among populations.

• We place our findings into a broad context using a simple model to show that the

strength of the germination cost could act to slow the rate of resistance evolution

in this species. Our results demonstrate that costs of adaptation may be present at

stages other than the production of progeny. Additionally, the cumulative effect of

costs at multiple life cycle stages can result in severe consequences to fitness

when adapting to novel environments.

Keywords: cost of herbicide resistance, fitness cost, glyphosate, *Ipomoea purpurea*,

trade-offs

Introduction

The study of adaptation to xenobiotics (chemicals such as pesticides and herbicides) can

provide unique opportunities to examine the evolutionary process: we know when

selection began, the strength of selection, and often its frequency. Researchers have

uncovered broad insights about the process of adaptation from such studies—for

example, information on the nature of genetic variation underlying resistance traits

(Preston & Powles, 2002; Hartley *et al.*, 2006), the effect size of alleles (Macnair, 1991;

Daborn *et al.*, 2002), and the repeatability of evolution (Andreev *et al.*, 1999; Menchari *et*

al., 2006; Délye *et al.*, 2010). Many investigations of herbicide and pesticide resistance

have also examined the central idea that novel traits incur costs—e.g., the expectation of

lower fitness of resistant (R) compared to susceptible (S) genotypes in the absence of the

xenobiotic (reviewed in Vila-Aiub *et al.*, 2011). Fitness costs of resistance are

hypothesized to arise from increased investment to defense/resistance and a subsequent

reduction in allocation to other traits (Coley *et al.*, 1985; Herms & Mattson, 1992), and/or

from mutations that decrease the efficiency of proteins on their native substrate (Fisher,

1958; Lande, 1983). Costs are universally expected since genetic variation is often found

to underlie resistance traits in nature (Baucom & Mauricio, 2004; Menchari *et al.*, 2006;

Délye *et al.*, 2010; Kuester *et al.*, *2015*); if there were no costs, resistance would increase

to fixation and every individual would be highly defended (Rausher & Simms, 1989).

Despite our expectations, however, reviews of the literature consistently show that

costs are not ubiquitous across herbicide resistant weeds (Bergelson & Purrington, 1996;

Gemmill & Read, 1998; Vila-Aiub *et al.*, 2009b). Three main ideas have been proposed

to explain their absence. First, there are a diverse number of potential mechanisms

responsible for resistance, and such mechanisms vary according to the particular

herbicide and weed species in question (Powles & Yu, 2010; Vogwill *et al.*, 2012). It is

possible that some mechanisms—such as those that incur resistance to a range of

different herbicides—are more likely to impose costs compared to others—i.e., a single

gene nucleotide substitution that does not alter the efficiency of translated proteins (e.g.

Yu *et al.*, 2007; Yu *et al.*, 2010). Second, costs may not be detected if the genetic

background is not properly controlled (Bergelson & Purrington, 1996; Vila-Aiub *et al.*,

2009b; Vila-Aiub *et al.*, 2011). Control of the genetic background, either by performing

crosses between R and S plants to homogenize the genetic background (Baucom &

Mauricio, 2004; Menchari *et al.*, 2008; Giacomini *et al.*, 2014) or ensuring replication of

multiple genetic backgrounds (Cousens *et al.*, 1997; Strauss *et al.*, 2002) increases the

likelihood that a cost will be detected (Bergelson & Purrington, 1996). Third, researchers

often examine only a portion of the life cycle (i.e., seed production or fecundity) and do

so in artificial and/or non-competitive conditions (Vila-Aiub *et al.*, 2009b; Vila-Aiub *et*

al., 2011). Studies that examine a range of traits are more likely to identify potential

growth and/or fitness differences between R and S genotypes compared to those that

focus solely on measures of fecundity (Vila-Aiub *et al.*, 2009b).

Despite recommendations to control/increase the number of genetic backgrounds

(Bergelson & Purrington, 1996), and to examine multiple life history stages when

determining if resistance incurs a cost (Primack & Kang, 1989; Vila-Aiub *et al.*, 2009b),

a recent review shows that only 25% of herbicide resistance studies control for

background effects and only 7-10% of cost studies examine multiple stages of the life

cycle (Vila-Aiub *et al.*, 2009b). Fewer still examine the potential for fitness costs using a

large number of populations sampled from a species' range, an approach suggested

almost 20 years ago (Cousens *et al.*, 1997; Strauss *et al.*, 2002). Just as the mechanism of

resistance can vary among species, populations of the same weed have been shown to

harbor different mechanisms of resistance to the same herbicide (Christopher *et al.*, 1991;

Christopher *et al.*, 1992; Christopher *et al.*, 1994; Preston & Powles, 1998; Yu *et al.*,

2008; Délye *et al.*, 2010), thus increasing the likelihood that costs may likewise vary

among populations. It is also possible, though rarely tested, that fitness costs have been

ameliorated in some herbicide resistant populations relative to other populations due to

the evolution of modifier loci (Darmency *et al.*, 2015). The above hypotheses for the lack

of costs are all interrelated: because resistance could be due to a variety of mechanisms

(Délye *et al.*, 2013a), costs may be apparent at only certain life history stages, expressed

in particular environments (Vila-Aiub *et al.*, 2009b), or apparent in some populations but

not others. Thus, there remain crucial gaps in our understanding of where in the life

history of a plant costs might be apparent, and further, how ubiquitous costs may be

across a species' range (Vila-Aiub *et al.*, 2011; Neve *et al.*, 2014).

The common morning glory, *Ipomoea purpurea*, a noxious weed of US

agriculture (Webster & MacDonald, 2001), provides an excellent system to examine the

strength and type of costs that may be present in natural populations. This species

exhibits variability in resistance to glyphosate (Baucom & Mauricio, 2008; Kuester *et al.*,

2015), which is the main ingredient in the herbicide RoundUp®. RoundUp® is currently

the most widely used herbicide in agriculture (Fernandez-Cornejo *et al.*, 2014), and of the

approximately 30 resistant weeds that have been examined (Heap, 2015), only a third are

reported to express fitness costs (Ismail *et al.*, 2002; Pedersen *et al.*, 2007; Brabham *et*

al., 2011; Giacomini *et al.,* 2014; Shrestha *et al.,* 2014; Vila-Aiub *et al.,* 2014; Glettner &

Stoltenberg, 2015; Goh *et al.,* 2015). *I. purpurea* has long been considered to exhibit low-

level resistance to glyphosate (considered 'tolerance' in Culpepper (2006)), and

previously we have shown that this low-level resistance has an additive genetic basis and

is under positive selection in the presence of the herbicide (Baucom & Mauricio, 2008).

More recently, a replicated dose-response experiment showed that some populations of *I.*

purpurea exhibit ~100% survival after application of the field dose of RoundUp® (*i.e.,*

resistance), whereas other populations exhibit high susceptibility (Kuester *et al., 2015*).

Strikingly, the fecundity of individuals from a single population artificially selected for

increased resistance was not significantly lower than susceptible individuals in the

absence of the herbicide, suggesting there may not be a fitness cost associated with

resistance in this species. However, there was some indication that progeny quality may

be lower in resistant individuals – resistant lines exhibited a trend for reduced seed

viability compared to susceptible lines (Debban *et al.*, *In press*). These results suggest

that reductions in progeny quality, rather than seed quantity, might be the mechanism of

fitness costs associated with glyphosate resistance in this species.

Here we determine if there is a cost to resistance in progeny quality measured at

multiple stages of the life cycle and the prevalence of this cost across populations by

examining germination, early root growth and above-ground growth across 43

populations of *I. purpurea*. We specifically ask the following: (i) is germination

correlated to resistance across this species' range in the US?, (ii) do resistant populations

exhibit smaller size at early life history stages (i.e., early germinant, young plant)?, and

(iii) do resistant populations exhibit the same type of fitness cost? Finally, we place our

findings of a strong fitness cost associated with germination (see Results) into a broad

context by using a simple model to examine the potential evolutionary trajectory of

resistance in this species.

Materials and Methods

Seed collection and control of maternal/environmental effects

Multiple fruits were collected from up to 79 individuals separated by at least 2 m from 43

populations located across the Midwest and Southeastern US (Table S1). These seeds

(hereafter field-collected seeds) were used in several experiments to determine resistance,

germination and early growth characteristics. To homogenize the effects of maternal

environment on seed quality, we chose a subset of the populations (N=18), grew them in

a common greenhouse for one generation and collected the autonomously self-pollinated

seeds from a similar growing and mating system environment (hereafter once-selfed

seeds).

Estimate of herbicide resistance

To determine glyphosate resistance across populations, a dose-response experiment was

conducted by planting a single field-collected seed from 10 randomly chosen maternal

lines from each population in six glyphosate treatments (including a non-herbicide

control treatment) in each of two greenhouse rooms. Full details of the dose-response

experiment are presented in Kuester *et al.* (*2015*) - for simplicity, we present resistance as

the percent survival per population at 1.70 kg a.i./ha of glyphosate, a rate which is

slightly higher than the suggested field rate of 1.54 kg a.i./ha. Individual seeds were

scarified, planted, allowed to grow until approximately 13 cm in height, and then treated

with the herbicide using a hand-held CO_2 pressurized sprayer (Spraying Systems Co.,

Wheaton, IL). Survival was scored three weeks after treatment application, and the

population estimate of resistance was determined as the proportion of individuals that

survived glyphosate.

Germination

To determine the effects of resistance on seed traits, we measured seed weight and

germination characteristics using field-collected seeds from each population (N=43). Up

to five (ave 4.6) seeds from 8-79 maternal lines per population (ave 38, total 1621, see

Table S1 for exact sample sizes per population) were haphazardly chosen for the

germination test. From this pool of seeds we randomly chose a subset of families per

population (8-49 maternal lines per population; Table S1) for which the selected seeds

were weighed (as a group) to determine the average seed weight. All of the selected seeds

were placed in a small petri dish (one dish per family) with water and allowed to

germinate in the lab under ambient light. Petri dishes were completely randomized across

lab benches. Germination was scored after 16 days, with successful germination

considered the emergence of a normal radicle. At this time, seeds that had not imbibed

water were scarified and germination was again scored after 1 week. We calculated the

percentage of seeds exhibiting normal germination, the percentage of seeds needing

scarification, the percentage of scarified seeds that germinated, and the percentage that

had abnormal germination. In addition, to compare results between this petri dish assay

and germination in soil, we examined germination data from one replicate of the dose-

response experiment mentioned earlier in which seeds were scarified and planted in

conetainers. Germination was scored after three weeks and used to calculate the

percentage of seeds that germinated.

We next repeated the germination assay experiment using seeds from maternal

lines that were selfed once in the greenhouse to examine the potential that germination

rates were influenced by the environment. Two sets of five seeds for up to 8 maternal

lines (randomly selected) for each of 18 populations were placed in petri dishes with

water. Germination was scored after 11 days. If seeds had not imbibed water they were

scarified and scored again in one week. We calculated the percentage of seeds with

normal germination prior to scarification, the percentage of seeds that germinated after

scarification, and the percentage that had abnormal germination.

Early root and above-ground growth

We carried out another assay to measure early root growth by placing scarified once-

selfed seeds in water and measured root growth after four days. We chose to first scarify

the seeds in this assay to standardize water absorption among individuals. Two sets of

five seeds for up to 8 maternal lines (randomly selected) for each of 18 populations were

scarified and placed in petri dishes with water. Germination was scored after 4 days and

petri dishes were scanned. For each germinated seed the root length was measured using

Image J (Abramoff *et al.*, 2004).

We next examined early growth traits of greenhouse-grown individuals to determine if

there was a relationship between resistance and plant size (i.e., are plants from resistant

populations smaller?). To do so we used measurements from plants from the dose-

response experiment prior to herbicide application. Approximately three weeks after

planting (and prior to spraying) we measured the height of the stem, the number of leaves

and length of the largest leaf on each individual planted per treatment per population

(total N=2908, Table S1 for exact sample sizes per population). Means for each trait per

population were obtained by calculating the LSMeans from models with replicate,

population, and planting tray within replicate as predictors using PROC GENMOD (SAS

v9.1), using the distribution resulting in normal residuals and the lowest AIC value

(normal for leaf number and leaf length; negative binomial for height).

Statistical analysis

To determine if there was a relationship between resistance and progeny quality, the

population mean of each trait averaged across maternal lines was used as the dependent

variable in individual regressions with the population level of resistance using PROC

REG in SAS (v9.1); residuals were examined to confirm normality. For the field-

collected seed germination assay the traits analyzed were average seed weight, percent

germination, percent abnormal germination, percent needing scarification and the percent

germination for scarified seeds. Visual examination of the regression between percent

germination of field-collected seeds and resistance suggested a different pattern between

resistant and susceptible populations. Therefore we applied separate regressions for

susceptible (defined as <50% survival) and resistant populations (>50% survival).

Additionally, previous studies had indicated there was a geographic pattern of resistance

in this species (Kuester *et al.*, *2015*). To ensure that the above results were not an artifact

of geography, we included resistance as well as latitude and longitude of the population

in the above regressions. Similar to the field-collected seeds, the population means for the

once-selfed seeds were used in separate regressions between population-level resistance

and percent germination prior to scarification or percent germination after scarification.

To examine early plant size traits (root length of once-selfed seeds and height, leaf

number and leaf size of field-collected seeds), we used population means in a regression

with population-level resistance.

Finally, we performed a Principle Components Analysis (PCA) using the population

averages of several traits from the field-collected seeds, which included seed weight,

germination percentage, percentage of abnormally germinating seeds, percentage of

successfully germinating scarified seeds, early plant height, leaf number and leaf size to

determine if different populations had different suites of costs—i.e., a difference in

germination costs or differences in early plant size. This analysis was performed using

PROC FACTOR in SAS with a varimax rotation to obtain more easily interpretable axes.

Loadings and the proportion variance explained for each factor with an eigenvalue >1 can

be found in Table S3.

Results

Germination

We found a strong and significant negative correlation between resistance and the

percentage of field-collected seeds that germinated (b = -0.87, r^2 = 0.54, t_{41} = -6.94, P <

0.0001; Fig. 1a); increasing resistance by 1% decreased germination by 0.87%. When we

grouped populations into resistant (R) and susceptible (S) populations based on greater or

less than 50% survival after glyphosate application, respectively, we found that, on

average, R populations exhibit ~35% germination compared to ~76% germination among

S populations. Interestingly, the negative trend between germination and resistance is

present among R populations but not among S populations (Fig. 1b,c). Thus, the fitness

cost in this species is apparent only above a certain resistance threshold. We also

uncovered a negative relationship between germination and resistance when seeds from

these populations were planted in soil in the greenhouse (b = -0.18, r^2 = 0.31, t_{41} = -4.24,

P = 0.0001), showing that decreased germination associated with increased resistance is

consistent across experimental conditions.

In addition to a decline in germination, several other measures of seed quality also

declined with increasing resistance. We found a higher percentage of abnormally

germinating seeds (b = 0.88, r^2 = 0.59, t_{41} = 7.59, P < 0.0001) in that, instead of

exhibiting normal germination, a non-viable embryo would be ejected from the seed coat

with no further growth. Furthermore, some seeds simply did not imbibe water; we

scarified these seeds to determine if they were viable but potentially dormant.

Populations with greater resistance had more seeds that needed scarification (b = 0.16, r^2

= 0.14, t_{41} = 2.58, P = 0.01), of which fewer seeds that subsequently germinated (b = -

0.52, r^2 = 0.17, t_{38} = -2.76, P = 0.009). We also found that populations with higher

resistance produced lighter seeds (b = -0.006, r^2 = 0.11, t_{41} = -2.26, P = 0.03), indicating

that resistance influenced multiple measures of seed quality.

Using multiple regression analysis, we confirmed a geographic pattern in

resistance, namely a decline in resistance with increasing latitude (b = -0.033, t_{40} = -1.99,

P = 0.05) but not longitude (b = -0.02, t_{40} = -1.72, P = 0.09). However, resistance

remained significant for all traits after accounting for the effect of geography except for

the proportion of seeds needing scarification (Table S2). This suggests that the patterns

we find between seed quality and resistance are not entirely due to geography.

The negative relationship between resistance and germination was supported by

the results from the once-selfed seeds grown in a common environment for a generation.

Very few of the greenhouse grown seeds germinated prior to scarification (2.0%) and

there was no effect of resistance (b = -0.006, r^2 = 0.01, t_{17} = -0.324, P = 0.75). However,

after scarification there was a negative relationship between germination and resistance

(b = -0.13, r^2 = 0.37, t_{17} = -3.08, P = 0.007, Fig. 3a) as was found in the field-collected

seeds. As opposed to the field-collected seeds, there was a very low rate of abnormal

germination (~10%) and the level of abnormal germination showed no relationship with

resistance (b = 0.02, r^2 = 0.01, t_{16} = 0.318, P = 0.75).

Early root and aboveground growth

There was a significant negative correlation between root length and resistance (b = -

0.02, r^2 = 0.01, t_{17} = -2.63, P = 0.009; Fig. 3b), indicating that the more resistant

populations have either smaller or slower growing roots. Furthermore, plants from more

resistant populations had smaller above-ground structures on average than plants from

less resistant populations (height: b = -3.41, r^2 = 0.25, t_{41} = -3.66, P = 0.0007; leaf

number: b = -0.80, r^2 = 0.27, t_{41} = -3.89, P = 0.0004; largest leaf width: b = -1.43, r^2 =

0.45, t_{41} = -4.92, P < 0.0001). Specifically, plants from resistant populations (i.e.,

populations that exhibited >50% survival post-spray) were ~10% shorter, and had ~7%

fewer and ~13% smaller leaves.

Visualization of cost-related traits

We next examined germination and early growth traits from the original field-collected

seeds using a principle components analysis (PCA) to determine if there was variation

among populations in the expression of cost-related traits (full results Table S3). The first

3 factors of the PCA explained 77% of the variance, with the first factor loading with the

early growth traits while the second loaded with seed traits and the third with the

proportion of seeds that required scarification to successfully germinate. Populations with

higher resistance scored lower on factor 1 (b = -2.22, r^2 = 0.28, t_{41} = -4.03, P = 0.0002)

and factor 2 (b = -2.12, r^2 = 0.26, t_{41} = -3.79, P = 0.0005), but not on factor 3 (b = -1.02, r^2

= 0.06, t_{41} = -1.61, P = 0.12). Using the first two factors to plot the results, resistant

populations occur mostly in the lower left quadrant (smaller plants, lighter seeds, lower

germination and more abnormally germinating seeds) and have a wider spread than less

resistant populations (Fig. 3). Furthermore, while R populations scored lower on factors 1

and 2 in comparison to S populations, some R populations exhibited early growth traits

that were similar to S populations and yet scored very low on germination traits (e.g., pop

num 5) whereas other R populations exhibited similar germination traits compared to the

S populations, but were smaller in stature than S populations (e.g., pop num 51). Thus, it

appears that the type of cost may vary among populations sampled from North America.

Discussion

Here we show that glyphosate resistant populations of the common morning glory

exhibit costs of herbicide resistance, and, that the type of cost may vary among

populations. Our series of experiments uncovered three notable findings: First, we found

a strong negative linear correlation between germination and resistance indicating that

resistant populations have a lower germination rate than susceptible populations. We

uncovered similar trends using seeds generated from a common greenhouse environment

showing that this result is not due simply to field environmental and/or maternal effects.

Second, we found that individuals from resistant populations were smaller than

individuals from susceptible populations, indicating that resistance may incur a cost in

terms of plant growth. Interestingly, we also found evidence that the two types of cost

may differ among populations—using PCA, we show that some R populations produce

normally sized plants, but score low on germination rate, and *vice versa*—some R

populations germinate at a rate similar to S populations, but tend to have smaller

individuals than S populations. Below, we detail how these results add further strength to

the suggestion that a variety of life stages and populations sampled across the species'

range should be assessed when testing the hypothesis that resistance incurs a fitness cost

(Délye *et al.*, 2013a). We then place our results into a broader context using a simple

model to examine how the cost identified in this work may influence the evolutionary

trajectory of resistance in this species.

Fitness costs: seed germination and early plant size

It is difficult to ascertain the frequency of germination costs associated with

herbicide resistance in nature since many studies of resistant weeds focus on seed

quantity rather than seed quality. There is some indication that germination may be

affected in other glyphosate resistant species–Dinelli *et al.* (2013) found reduced

germination of glyphosate resistant *Ambrosia trifida* populations, while Ismail *et al.*

(2002) found greater germination of resistant biotypes of goosegrass (*Eleusine indica)*.

Costs may be specific to the species or the type of mutation conferring resistance

(O'Donovan *et al.*, 1999; Vila-Aiub *et al.*, 2005; Délye *et al.*, 2013b). For example, only

one of two different resistance mutations in ACCase resistant *Lolium rigidum* had more

stringent germination requirements (germinated poorly in the dark and required

fluctuating temperatures to break seed dormancy) than the susceptible genotype (Vila-

Aiub *et al.*, 2005). Similarly, Délye *et al.* (2013b) found differential effects on

germination among resistance mutations to ACCase in *Alopecurus myosuroides*. Both of

these studies report that the resistance mutation led to delayed germination; such a delay

in germination may affect fitness, especially in agricultural settings where germinating

too early can lead to removal by pre-sowing practices and germinating too late can lead

to intensified competition with already established plants (Weaver & Cavers, 1979;

Barrett, 1983; Mortimer, 1997; Forcella *et al.*, 2000; Owen *et al.*, 2014).

While delays in germination may explain some of the results we present, in

general, the low germination we uncovered in the R populations was due to non-viable

embryos (53% of R seeds vs 18% of S seeds) rather than later germination. In the normal

progression of seed germination in this species, the radicle first emerges from the seed

coat and the cotyledons rapidly follow. In comparison, the non-viable seeds swell, split in

half, and eject the entire embryo from the seed coat. Many of these seeds appear fully

provisioned, and, although they weigh less than seeds from S populations, the differences

in weight are not drastic, nor are there visible indications that would suggest that seeds

from R populations would be of such low quality.

The reduced growth of R compared to S populations that we uncovered could lead

to decreased competitive ability and subsequent lower fitness in the presence of

competition, as has been found in other herbicide resistant species (Weaver & Warwick,

1982; Ahrens & Stoller, 1983; Holt, 1988; Alcocer-Ruthling *et al.*, 1992; Williams *et al.*,

1995; Vila-Aiub *et al.*, 2005; Tardif *et al.*, 2006; Vila-Aiub *et al.*, 2009a). However,

similar to studies of germination differences, results from studies comparing growth of

resistant and susceptible genotypes suggest this cost is species, mutation and environment

specific. For example, *Lolium rigidum* has evolved herbicide resistance via a variety of

mutations ranging from target site (Christopher *et al.*, 1992; Yu *et al.*, 2008) to non-target

site (Christopher *et al.*, 1991; Christopher *et al.*, 1994; Preston & Powles, 1998). Target

site mutations in the acetohydroxyacid synthase gene result in little cost in growth (Yu *et*

al., 2010). On the other hand, herbicide resistance mediated by the cytochrome P450

complex resulted in reduced biomass and decreased competitive ability (Vila-Aiub *et al.*,

2009a).

An alternative explanation for the decline in germination and growth we identify

using the field-collected seeds is that some other co-varying population characteristic

such as soil fertility, spraying regime, herbivore levels, or the many other biotic and

abiotic factors that can influence seed development differed among R and S populations

(Roach & Wulff, 1987; Fenner, 1991; Schmitt *et al.*, 1992; Platenkamp & Shaw, 1993;

Galloway, 2001). However, several lines of evidence suggest the relationship between

resistance and germination that we uncovered represents a true fitness cost rather than

simply an effect of the environment (e.g., driven by maternal effects, latitude, spray

environment). First, we identified a very strong linear correlation between resistance and

germination. If the fitness cost was due simply to glyphosate exposure, with resistant

populations exhibiting abnormal seed development after surviving glyphosate

application, we would expect to see a binary distribution of seed quality of populations

that had been sprayed and those that had not, rather than a linear trend with resistance.

Second, the negative correlation between germination and resistance is present only

among resistant populations—strongly suggesting that as populations become more

resistant, they exhibit a correspondingly higher fitness penalty when in the absence of the

herbicide. Third, the negative relationship is maintained after a generation in a common

growing/mating system environment. Finally, our results parallel those from the recent

experiment that specifically controlled for genetic background and environmental effects

using *I. purpurea* plants from a single population (Debban *et al.*, *In press*). Individuals

from this population were artificially selected for increased or decreased resistance for

three generations under controlled greenhouse conditions, and, similar to results

presented here, the increased resistance lines had a larger percentage of "bad" seeds that

ejected the embryo. That the results from one population utilizing a controlled genetic

background are mirrored across many populations collected from the landscape provides

strong evidence that a fitness cost of glyphosate resistance is present in this weed species.

Interestingly, by visually examining the germination and growth traits in a PCA,

we find variation in the type of cost trait among R populations. While some R

populations fell into the "poor germination" axis, other R populations fell into the "poor

growth" axis compared to S populations. There are at least three possible reasons for this

difference: different resistance genes, different compensatory mutations or different

genetic backgrounds. First, the gene(s) involved in resistance may vary among

populations leading to different costs. Independent origins of resistance to herbicide have

been found in other species (Délye *et al.*, 2010) and these different mutations often incur

different fitness costs (Vila-Aiub *et al.*, 2005; Délye *et al.*, 2013b). Second, the resistance

gene(s) may be the same amongst populations but each population may have different

compensatory mutations that lead to different costs (Darmency *et al.*, 2015). Third, the

resistance gene(s) may behave differently in different genetic backgrounds (Paris *et al.*,

2008). These distinctions are important because they would differentially affect the

evolutionary trajectory of herbicide resistance. For example, if populations differ in the

gene(s) involved, each population may have a very different set of costs, benefits and

evolutionary trajectories, which would need to be incorporated in models.

It is currently unknown if the costs identified here are pleiotropic or due to

linkage to the resistance gene. The most restrictive definition of a cost requires that the

decrease in fitness is due to the resistance allele itself - either the actual allele or through

it acting pleiotropically (Bergelson & Purrington, 1996). Given that we do not know the

identity of the loci involved in either resistance or the abnormal germination and reduced

growth, we cannot entirely rule out physical linkage between resistance genes and cost

genes, in which case the "cost" could quickly become unlinked over generations

(Lewontin, 1974; Hartl & Clark, 1989). For some species, easily identifiable mutations in

the enzyme targeted by the herbicide can be linked to resistance, i.e. target site resistance

(TSR). However, preliminary work suggests that glyphosate resistance in *I. purpurea* is

due to non-target site mechanism (NTSR; Leslie and Baucom, *unpublished data*), and as

such elucidating the genetic basis of both resistance and the cost will be a non-trivial

endeavor. Furthermore, it is rare that genes underlying costs are identified; most

documented cases of the genes involved in the cost of resistance is when TSR mutations

lead to poor performance of the enzyme on its natural substrate (Vila-Aiub *et al.*, 2009b).

As far as we are aware, no study has identified the genes involved in the cost of

resistance when the mechanism of resistance is NTSR. One intriguing possibility for this

species stems from a previous study that compared transcript expression levels of

artificially selected lines of resistant and susceptible *I. purpurea* plants following

herbicide application (Leslie & Baucom, 2014). One of the differences between the

replicated R and S lines was a lower expression of pectin methylesterase (PME) in the

resistant plants. This enzyme has been shown to play a role in breaking seed dormancy

(Ren & Kermode, 2000) and stem elongation (Pilling *et al.*, 2000). Thus, the decreased

expression of PME in resistant plants may explain both the reduced germination and

growth in populations with higher resistance.

How might costs influence the evolutionary trajectory of resistance in this species?

Recent work in this system has shown that populations of the common morning

glory sampled from 2012 exhibit higher levels of resistance compared to the same

populations sampled in 2003 (Kuester et al, *In* Review). Interestingly, however, the

difference in resistance between sampling years was only slight, i.e., 62% survival at 1.7

kg ai/ha in 2012 vs 57% survival in 2003 samples. It is very likely that the fitness cost we

identified here is responsible for maintaining resistance between sampling years. In light

of this finding, we elected to model the influence of the germination cost on the evolution

of resistance in this species, making simplifying assumptions about the genetic basis of

resistance and pattern of mating in this species. We used a single locus Hardy-Weinberg

model of a panmictic population with the resistance allele dominant to the susceptible

allele. We examined the frequency of a resistance allele as a function of the fitness cost

and the proportion of the population not under selection (*i.e.* not sprayed). We ran the

model using one of three costs – no cost, a moderate cost, or a high cost to demonstrate

the differing evolutionary trajectories given different costs. On one extreme, we show

that the frequency of a resistance allele without a cost increases relative to the proportion

of the population under selection (Fig. 4a). On the other extreme, if a resistance mutation

leads to a strong cost—similar to that of the germination cost identified here—the

frequency of the resistance allele can decrease when enough of the population is free

from selection, i.e., not sprayed with herbicide (Fig. 4b). Populations or species with high

costs will have a slower rate of increase of the resistant allele compared to those that

exhibit a moderate cost (Fig. 4c) (such as found when examining the negative

repercussions of slower growth on competitiveness of resistant plants (Vila-Aiub *et al.*,

2009a)).

478   This simple model can also demonstrate the importance of accurate estimations of

costs when making management decisions. For example, we can use a "back of the

envelope" calculation to examine the threshold proportion of selection-free land that will

result in a decrease in resistance for a given cost of resistance. We estimated the threshold

proportion of non-treated land as $(1-C_R/(C_R+C_S))$, or one minus the cost of resistance

divided by the sum of the cost of susceptibility (i.e. herbicide efficiency) and the cost of

resistance. From the cost obtained in this study using the field-collected seeds (and

assuming complete control of susceptible individuals when spraying), this proportion is

0.54, *i.e.* more than 54% of acreage would need to be non-treated to decrease the

frequency of the resistant allele. With no fitness cost, 100% of acreage would need to be

non-treated. Clearly, estimates of fitness costs are necessary to decide which management

decisions will be both feasible and effective. Additionally, if populations differ in their

costs, these management decisions will likely differ by population. Although the

calculation we present here is consistent with expectations of a simple model of allele

frequency change (Jasieniuk & Maxwell, 1994), we caution that our estimates do not

include important dynamics such as the influence of population subdivision, selfing rates,

and the contribution of migrants from the seed bank (among other biologically relevant

forces).

In summary, we found fitness costs in seed quality across replicated populations

of the common morning glory and demonstrate that these costs can play a substantial role

in herbicide resistance evolution. Although most studies use seed quantity as a proxy for

fitness, our results highlight that reductions in progeny quality are an equally, if not more,

important cost of adaptation. Given that fitness costs are thought to arise from a variety of

mechanisms (allocation of resources, ecological costs, etc.), our results suggest that a

high priority should be placed on the examination of multiple stages of the life cycle

when assessing potential costs and not just seed quantity. Furthermore, because the

strength of this cost could maintain the efficacy of a globally important herbicide, this

work illustrates the utility and importance of integrating evolutionary principles into

management scenarios (Gould, 1995).

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Figure Legends

Fig 1. Regression between survival at 1.7 kg a.i./ha glyphosate and germination rate of

field-collected seeds across all populations (a), in susceptible populations (b) and

resistant populations (c).

Fig 2. Regression between survival at 1.7 kg a.i./ha glyphosate and germination rate (a)

and root growth (b) of once-selfed seeds generated in the greenhouse.

Fig 3. Scatter plot of PCA results showing average factor 1 (higher values indicate larger

plants) and factor 2 (higher values indicate heavier seeds, higher germination, and fewer

abnormally germinating seeds) values for field-collected seeds with circle color

indicating survival at 1.7 kg/ha glyphosate.

Fig 4. A simulation of resistance allele frequencies over time given different selection

regimes. Without a cost to resistance (a) the frequency of resistance increases if under

selection. When costs are high (b; C_R=0.86, from field-collected germination rate) some

selection regimes lead to a decrease in resistance frequency. At moderate levels of costs

(c), such as reproductive output under competitive environments found in Vila-Auib *et al.*

(2009a, C_R=0.11) resistance can decrease under a small subset of conditions. We used a

simple single-locus, 2 allele Hardy-Weinberg population genetics model, with resistance

being completely dominant to susceptibility. The net strength of selection for resistance

was the weighted average of the strength of selection in sprayed (0) and non-sprayed

(cost of resistance = C_R) areas weighted by the proportion of land in each. Similarly the

net strength of selection for susceptibility was the weighted average of the strength of

selection in sprayed (efficacy of spraying = C_S = 1) and non-sprayed areas (0) weighted

by the proportion of land in each.

 Supporting information

542 Table S1. Population characteristics including population number, long name, state,

543 crop type, latitude and longitude and sample sizes for experiments with field-

544 collected seeds.

Population number	State	Crop	Latitude	Longitude	N_{fam} germ	N_{seeds} germ	N early growth
1	TN	Corn	35.775237	-85.903419	8	31	10
2	NC	Corn	34.595714	-77.927484	79	387	60
4	NC	Corn	34.556672	-79.125602	39	178	79
5	SC	Soy	33.859875	-79.909072	34	146	27
8	SC	Corn	34.297195	-79.991259	40	180	73
9	NC	Soy	34.924044	-77.796171	40	189	46
10	NC	Soy	34.983161	-78.039309	28	123	24
11	NC	Corn	34.527135	-78.756704	38	178	81
12	SC	Cotton	34.145812	-79.865313	42	202	79
14	NC	Soy	35.424763	-77.917121	40	192	76
15	SC	Soy	34.104209	-79.073735	41	193	81
16	SC	Alfalfa	34.10535	-79.183234	41	203	78
17	SC	Soy	34.159155	-79.272908	27	135	82
18	SC	Corn	34.156593	-79.27027	39	193	79
19	NC	Corn	34.508193	-78.70899	35	138	70
20	TN	Corn	35.830692	-85.777871	38	169	13
21	TN	Soy	35.369816	-77.877314	40	200	70
22	NC	Corn	36.1436	-78.053422	17	78	69
23	TN	Corn	35.067905	-86.62955	40	195	77
26	TN	Soy	35.533413	-85.951902	50	250	86
28	SC	Corn	34.097917	-80.377715	35	172	80
29	NC	Corn	34.705135	-78.738897	30	122	75
30	TN	Corn	35.31105	-85.945003	42	206	82
31	TN	Corn	35.608482	-85.846379	31	141	69
32	TN	Corn	35.099356	-86.225509	15	56	33
33	OH	Corn	39.858763	-83.669821	52	260	79
34	OH	Corn	39.44316	-83.910189	48	238	82
35	IN	Corn	39.853945	-85.770156	28	120	80

36	IN	Corn	40.565608	-85.503826	33	146	85
37	OH	Corn	39.583755	-83.758264	49	243	79
38	OH	Soy	39.515447	-83.407431	46	230	88
39	IN	Corn	39.988984	-85.742262	38	188	75
40	OH	Soy	41.284684	-83.847252	50	237	84
41	VA	Corn	38.636343	-78.472921	41	200	78
42	VA	Corn	38.373523	-78.662516	49	230	82
43	VA	Soy	36.886448	-78.553156	24	89	28
44	VA	Soy	38.285415	-78.797088	20	98	75
45	VA	Soy	36.847945	-78.595042	42	209	76
46	TN	Soy	35.536019	-86.17985	34	100	17
47	SC	Soy	34.282132	-79.746597	41	201	83
48	TN	Corn	35.31653	-87.35373	31	151	74
51	TN	Soy	35.533413	-85.951902	24	118	56
52	OH	Corn	41.284684	-83.847252	62	310	88

547   Table S2. Regression results between resistance and seed quality traits, accounting for

548   geography (latitude and longitude) for field-collected seeds.

Trait	d.f.	Survival at 1.7 kg a.i. /ha		Latitude		Longitude	
		b	t	*b*	t	*b*	t
Seed weight	39	-0.00506	-2.24*	0.00068	2.75**	-0.00033	-1.95
Percent germination	39	-0.8484	-7.65***	0.03227	2.64**	-0.01885	-2.3*
Percent abnormally germinating seeds	39	0.67661	5.88***	-0.02331	-1.84	0.024	2.82**
Percent needing scarification	39	0.09051	1.53	-0.01955	-3.0**	-0.01346	-3.07**
Percent of scarified that germinated	36	-0.49897	-2.44*	0.01504	0.66	-0.0022	-0.15
Height	39	-3.23167	-3.49**	-0.07628	-0.75	0.14928	2.18*
Leaf number	39	-0.71189	-3.36**	0.0067	0.29	0.03242	2.07*
Largest leaf size	39	-1.28809	-4.21**	0.0274	0.81	0.03655	1.61

552   Table S3. PCA loadings and variance explained.

Trait	Factor1	Factor2	Factor3
Seed weight	0.09506	0.8414	-0.16627
Percent germination	0.29276	0.82272	0.37541
Percent viable dormant seeds	0.0339	0.12801	0.92794
Percent abnormally germinating seeds	-0.15651	-0.81953	-0.42013
Height	0.94253	0.04143	0.05526
Leaf number	0.86728	0.13535	0.06511
Largest leaf size	0.74853	0.28353	0.00642
Prop. Variance explained	0.3316	0.3105	0.1734

References

Abramoff MD, Magelhaes PJ, Ram SJ. 2004. Image Processing with ImageJ. *Biophotonics International* **11**(7): 36-42.

Ahrens WH, Stoller E. 1983. Competition, growth rate, and CO_2 fixation in triazine-susceptible and-resistant smooth pigweed (*Amaranthus hybridus*). *Weed Science* **31**(4): 438-444.

Alcocer-Ruthling M, Thill DC, Shafii B. 1992. Differential competitiveness of sulfonylurea resistant and susceptible prickly lettuce (*Lactuca serriola*). *Weed Technology* **6**(2): 303-309.

Andreev D, Kreitman M, Phillips TW, Beeman RW. 1999. Multiple origins of cyclodiene insecticide resistance in *Tribolium castaneum* (Coleoptera: Tenebrionidae). *Journal of Molecular Evolution* **48**(5): 615-624.

Barrett SH. 1983. Crop mimicry in weeds. *Economic Botany* **37**(3): 255-282.

Baucom RS, Mauricio R. 2004. Fitness costs and benefits of novel herbicide tolerance in a noxious weed. *Proceedings of the National Academy of Sciences of the United States of America* **101**(36): 13386-13390.

Baucom RS, Mauricio R. 2008. Constraints on the evolution of tolerance to herbicide in the common morning glory: resistance and tolerance are mutually exclusive. *Evolution* **62**(11): 2842-2854.

Bergelson J, Purrington CB. 1996. Surveying patterns in the cost of resistance in plants. *American Naturalist* **148**(3): 536-558.

Brabham CB, Gerber CK, Johnson WG. 2011. Fate of glyphosate-resistant giant ragweed (*Ambrosia trifida)* in the presence and absence of glyphosate. *Weed Science* **59**(4): 506-511.

Christopher JT, Powles SB, Holtum JA. 1992. Resistance to acetolactate synthase-inhibiting herbicides in annual ryegrass (*Lolium rigidum*) involves at least two mechanisms. *Plant Physiology* **100**(4): 1909-1913.

Christopher JT, Powles SB, Liljegren DR, Holtum JA. 1991. Cross-resistance to herbicides in annual ryegrass (*Lolium rigidum*) II. chlorsulfuron resistance involves a wheat-like detoxification system. *Plant Physiology* **95**(4): 1036-1043.

Christopher JT, Preston C, Powles SB. 1994. Malathion antagonizes metabolism-based chlorsulfuron resistance in *Lolium rigidum*. *Pesticide Biochemistry and Physiology* **49**(3): 172-182.

Coley PD, Bryant JP, Chapin III FS. 1985. Resource availability and plant antiherbivore defense. *Science* **230**: 895-899.

Cousens R, Gill G, Speijers EJ. 1997. Comment: number of sample populations required to determine the effects of herbicide resistance on plant growth and fitness. *Weed Research* **37**(1): 1-4.

Culpepper AS. 2006. Glyphosate-induced weed shifts. *Weed Technology* **20**(2): 277-281.

Daborn P, Yen J, Bogwitz M, Le Goff G, Feil E, Jeffers S, Tijet N, Perry T, Heckel D, Batterham P. 2002. A single P450 allele associated with insecticide resistance in *Drosophila*. *Science* **297**(5590): 2253-2256.

Darmency H, Menchari Y, Le Corre V, Délye C. 2015. Fitness cost due to herbicide
resistance may trigger genetic background evolution. *Evolution* **69**(1): 271-
278.
Debban C, Matthews S, Elliot L, Pieper K, Wilson A, Baucom RS. *In press.* An
examination of fitness costs of glyphosate resistance in the common morning
glory, *Ipomoea purpurea. Ecology and Evolution.*
Délye C, Jasieniuk M, Le Corre V. 2013a. Deciphering the evolution of herbicide
resistance in weeds. *Trends in Genetics* **29**(11): 649-658.
Délye C, Menchari Y, Michel S, Cadet É, Le Corre V. 2013b. A new insight into
arable weed adaptive evolution: mutations endowing herbicide resistance
also affect germination dynamics and seedling emergence. *Annals of Botany*
111(4): 681-691.
Délye C, Michel S, Bérard A, Chauvel B, Brunel D, Guillemin JP, Dessaint F, Le
Corre V. 2010. Geographical variation in resistance to acetyl‐coenzyme A
carboxylase‐inhibiting herbicides across the range of the arable weed
Alopecurus myosuroides (black‐grass). *New Phytologist* **186**(4): 1005-1017.
Dinelli G, Marotti I, Catizone P, Bosi S, Tanveer A, Abbas R, Pavlovic D. 2013.
Germination ecology of *Ambrosia artemisiifolia* L. and *Ambrosia trifida* L.
biotypes suspected of glyphosate resistance. *Open Life Sciences* **8**(3): 286-
296.
Fenner M. 1991. The effects of the parent environment on seed germinability. *Seed*
Science Research **1**(02): 75-84.
Fernandez-Cornejo J, Nehring R, Osteen C, Wechsler S, Martin A, Vialou A.
2014. Pesticide use in U.S. agriculture: 21 selected crops, 1960-2008, EIB-
124: U.S. Department of Agriculture, Economic Research Service.
Fisher RA. 1958. *The genetical theory of natural selection.* New York: Dover
Publishing Inc.
Forcella F, Arnold RLB, Sanchez R, Ghersa CM. 2000. Modeling seedling
emergence. *Field Crops Research* **67**(2): 123-139.
Galloway LF. 2001. The effect of maternal and paternal environments on seed
characters in the herbaceous plant *Campanula americana* (Campanulaceae).
American Journal of Botany **88**(5): 832-840.
Gemmill AW, Read AF. 1998. Counting the cost of disease resistance. *Trends in*
Ecology & Evolution **13**(1): 8-9.
Giacomini D, Westra P, Ward SM. 2014. Impact of genetic background in fitness
cost studies: an example from glyphosate-resistant *Palmer amaranth. Weed*
Science **62**(1): 29-37.
Glettner CE, Stoltenberg DE. 2015. Noncompetitive growth and fecundity of
Wisconsin giant ragweed resistant to glyphosate. *Weed Science* **63**(1): 273-
281.
Goh SS, Vila‐Aiub MM, Busi R, Powles SB. 2015. Glyphosate resistance in
Echinochloa colona: phenotypic characterisation and quantification of
selection intensity. *Pest management science*.
Gould F. 1995. Comparisons between resistance management strategies for insects
and weeds. *Weed Technology*: 830-839.

Hartl DL, Clark AG. 1989. *Principles of population genetics*. Sunderland, Mass.:
Sinauer associates.
Hartley C, Newcomb R, Russell R, Yong C, Stevens J, Yeates D, La Salle J,
Oakeshott J. 2006. Amplification of DNA from preserved specimens shows
blowflies were preadapted for the rapid evolution of insecticide resistance.
Proceedings of the National Academy of Sciences **103**(23): 8757-8762.
Heap I 2015. The International survey of herbicide resistant weeds.
Herms DA, Mattson WJ. 1992. The dilemma of plants: to grow or defend. *Quarterly*
Review of Biology **67**(3): 283-335.
Holt J. 1988. Reduced growth, competitiveness, and photosynthetic efficiency of
triazine-resistant *Senecio vulgaris* from California. *Journal of Applied Ecology*
25(1): 307-318.
Ismail B, Chuah T, Salmijah S, Teng Y, Schumacher R. 2002. Germination and
seedling emergence of glyphosate‐resistant and susceptible biotypes of
goosegrass (*Eleusine indica* [L.] Gaertn.). *Weed Biology and Management* **2**(4):
177-185.
Jasieniuk M, Maxwell BD. 1994. Populations genetics and the evolution of
herbicide resistance in weeds. *Phytoprotection* **75**(4): 25-35.
Kuester A, Chang SM, Baucom RS. *2015*. The geographic mosaic of herbicide
resistance evolution in the common morning glory, *Ipomoea purpurea*:
Evidence for resistance hotspots and low genetic differentiation across the
landscape. *Evolutionary Applications*.
Lande R. 1983. The response to selection on major and minor mutations affecting a
metrical trait. *Heredity* **50**(1): 47-65.
Leslie T, Baucom RS. 2014. De novo assembly and annotation of the transcriptome
of the agricultural weed *Ipomoea purpurea* uncovers gene expression
changes associated with herbicide resistance. *G3: Genes| Genomes| Genetics*
4(10): 2035-2047.
Lewontin RC. 1974. *The genetic basis of evolutionary change*. New York: Columbia
University Press.
Macnair M. 1991. Why the evolution of resistance to anthropogenic toxins
normally involves major gene changes: the limits to natural selection.
Genetica **84**(3): 213-219.
Menchari Y, Camilleri C, Michel S, Brunel D, Dessaint F, Le Corre V, Délye C.
2006. Weed response to herbicides: regional‐scale distribution of herbicide
resistance alleles in the grass weed *Alopecurus myosuroides. New Phytologist*
171(4): 861-874.
Menchari Y, Chauvel B, Darmency H, Délye C. 2008. Fitness costs associated with
three mutant acetyl‐coenzyme A carboxylase alleles endowing herbicide
resistance in black‐grass *Alopecurus myosuroides. Journal of Applied Ecology*
45(3): 939-947.
Mortimer AM. 1997. Phenological adaptation in weeds—an evolutionary response
to the use of herbicides? *Pesticide science* **51**(3): 299-304.

Neve P, Busi R, Renton M, Vila‐Aiub MM. 2014. Expanding the eco‐
evolutionary context of herbicide resistance research. *Pest management*
science **70**(9): 1385-1393.
O'Donovan J, Newman J, Blackshaw R, Harker K, Derksen D, Thomas A. 1999.
Growth, competitiveness, and seed germination of triallate/difenzoquat-
susceptible and-resistant wild oat populations. *Canadian Journal of Plant*
Science **79**(2): 303-312.
Owen MJ, Goggin DE, Powles SB. 2014. Intensive cropping systems select for
greater seed dormancy and increased herbicide resistance levels in *Lolium*
rigidum (annual ryegrass). *Pest management science.*
Paris M, Roux F, Berard A, Reboud X. 2008. The effects of the genetic background
on herbicide resistance fitness cost and its associated dominance in
Arabidopsis thaliana. Heredity **101**(6): 499-506.
Pedersen BP, Neve P, Andreasen C, Powles SB. 2007. Ecological fitness of a
glyphosate-resistant *Lolium rigidum* population: Growth and seed production
along a competition gradient. *Basic and Applied Ecology* **8**(3): 258-268.
Pilling J, Willmitzer L, Fisahn J. 2000. Expression of a *Petunia inflata* pectin
methyl esterase in *Solanum tuberosum* L. enhances stem elongation and
modifies cation distribution. *Planta* **210**(3): 391-399.
Platenkamp GA, Shaw RG. 1993. Environmental and genetic maternal effects on
seed characters in *Nemophila menziesii. Evolution* **47**(2): 540-555.
Powles SB, Yu Q. 2010. Evolution in action: plants resistant to herbicides. *Annual*
review of plant biology **61**: 317-347.
Preston C, Powles S. 2002. Evolution of herbicide resistance in weeds: initial
frequency of target site-based resistance to acetolactate synthase-inhibiting
herbicides in *Lolium rigidum. Heredity* **88**(1): 8-13.
Preston C, Powles SB. 1998. Amitrole inhibits diclofop metabolism and synergises
diclofop-methyl in a diclofop-methyl-resistant diotype of *Lolium rigidum.*
Pesticide Biochemistry and Physiology **62**(3): 179-189.
Primack RB, Kang H. 1989. Measuring fitness and natural selection in wild plant
populations. *Annual Review of Ecology and Systematics* **20**: 367-396.
Rausher MD, Simms EL. 1989. The evolution of resistance to herbivory in *Ipomoea*
purpurea. I. Attempts to detect selection. *Evolution* **43**(4): 563-572.
Ren C, Kermode AR. 2000. An increase in pectin methyl esterase activity
accompanies dormancy breakage and germination of yellow cedar seeds.
Plant Physiology **124**(1): 231-242.
Roach DA, Wulff RD. 1987. Maternal effects in plants. *Annual Review of Ecology and*
Systematics **18**: 209-235.
Schmitt J, Niles J, Wulff RD. 1992. Norms of reaction of seed traits to maternal
environments in *Plantago lanceolata. American Naturalist* **139**(3): 451-466.
Shrestha A, Steinhauer KM, Moretti ML, Hanson BD, Jasieniuk M, Hembree KJ,
Wright SD. 2014. Distribution of glyphosate-resistant and glyphosate-
susceptible hairy fleabane (*Conyza bonariensis*) in central California and their
phenological development. *Journal of pest science* **87**(1): 201-209.

Strauss SY, Rudgers JA, Lau JA, Irwin RE. 2002. Direct and ecological costs of
resistance to herbivory. *Trends in Ecology & Evolution* **17**(6): 278-285.
Tardif FJ, Rajcan I, Costea M. 2006. A mutation in the herbicide target site
acetohydroxyacid synthase produces morphological and structural
alterations and reduces fitness in *Amaranthus powellii. New Phytologist*
169(2): 251-264.
Vila-Aiub MM, Goh SS, Gaines TA, Han H, Busi R, Yu Q, Powles SB. 2014. No
fitness cost of glyphosate resistance endowed by massive EPSPS gene
amplification in *Amaranthus palmeri. Planta* **239**(4): 793-801.
Vila-Aiub MM, Neve P, Powles SB. 2009a. Evidence for an ecological cost of
enhanced herbicide metabolism in *Lolium rigidum. Journal of Ecology* **97**(4):
772-780.
Vila-Aiub MM, Neve P, Powles SB. 2009b. Fitness costs associated with evolved
herbicide resistance alleles in plants. *New Phytologist* **184**(4): 751-767.
Vila-Aiub MM, Neve P, Roux F. 2011. A unified approach to the estimation and
interpretation of resistance costs in plants. *Heredity* **107**(5): 386-394.
Vila-Aiub MM, Neve P, Steadman K, Powles S. 2005. Ecological fitness of a
multiple herbicide‐resistant *Lolium rigidum* population: dynamics of seed
germination and seedling emergence of resistant and susceptible
phenotypes. *Journal of Applied Ecology* **42**(2): 288-298.
Vogwill T, Lagator M, Colegrave N, Neve P. 2012. The experimental evolution of
herbicide resistance in *Chlamydomonas reinhardtii* results in a positive
correlation between fitness in the presence and absence of herbicides.
Journal of Evolutionary Biology **25**(10): 1955-1964.
Weaver S, Warwick S. 1982. Competitive relationships between atrazine resistant
and susceptible populations of *Amaranthus retroflexus* and *A. powellii* from
southern Ontario. *New Phytologist* **92**(1): 131-139.
Weaver SE, Cavers PB. 1979. The effects of date of emergence and emergence
order on seedling survival rates in *Rumex crispus* and *R. obtusifolius*.
Canadian Journal of Botany **57**(7): 730-738.
Webster TM, MacDonald GE. 2001. A survey of weeds in various crops in Georgia
1. *Weed Technology* **15**(4): 771-790.
Williams MM, Jordan N, Yerkes C. 1995. The fitness cost of triazine resistance in
jimsonweed (*Datura stramonium* L.). *American Midland Naturalist*: 131-137.
Yu Q, Collavo A, Zheng M-Q, Owen M, Sattin M, Powles SB. 2007. Diversity of
acetyl-coenzyme A carboxylase mutations in resistant *Lolium* populations:
evaluation using clethodim. *Plant Physiology* **145**(2): 547-558.
Yu Q, Han H, Powles SB. 2008. Mutations of the ALS gene endowing resistance to
ALS‐inhibiting herbicides in *Lolium rigidum* populations. *Pest management*
science **64**(12): 1229-1236.
Yu Q, Han H, Vila-Aiub MM, Powles SB. 2010. AHAS herbicide resistance
endowing mutations: effect on AHAS functionality and plant growth. *Journal*
of Experimental Botany **61**(14): 3925-3934.
773
774