Title: Local adaptation to an altitudinal gradient: the interplay between mean phenotypic trait

variation and phenotypic plasticity in *Mimulus laciniatus*

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Keywords: phenotypic plasticity, local adaptation, heterogeneity, elevation gradient, trait

variation, leaf lobing

ABSTRACT

Organisms can adapt to environmental heterogeneity through two mechanisms: (1) expression of

population genetic variation or (2) phenotypic plasticity. In this study we investigated whether

patterns of variation in both trait means and phenotypic plasticity along elevational and

latitudinal clines in a North American endemic plant, *Mimulus laciniatus*, were consistent with

local adaptation. We grew inbred lines of *M. laciniatus* from across the species' range in two

common gardens varying in day length to measure mean and plastic trait expression in several

traits previously shown to be involved in adaptation to *M. laciniatus's* rocky outcrop

microhabitat: flowering time, size-related traits, and leaf shape. We examined correlations

between the mean phenotype and phenotypic plasticity, and tested for a relationship between trait

variation and population elevation and latitude. We did not find a strong correlation between

mean and plastic trait expression at the individual genotype level suggesting that they operate

under independent genetic controls. We identified multiple traits that show patterns consistent

with local adaptation to elevation: critical photoperiod, flowering time, flower size, mean leaf

lobing, and leaf lobing plasticity. These trends occur along multiple geographically independent

altitudinal clines indicating that selection is a more likely cause of this pattern than gene flow

among nearby populations with similar trait values. We also found that population variation in

mean leaf lobing is associated with latitude. Our results indicate that both having more highly

lobed leaves and greater leaf shape plasticity may be adaptive at high elevation within *M.*

laciniatus. Our data strongly suggest that traits known to be under divergent selection between

M. laciniatus and close relative *Mimulus guttatus* are also under locally varying selection within

M. laciniatus.

INTRODUCTION

Environmental heterogeneity can come in multiple forms, such as spatial and temporal variation in resource availability (Cortés and Wheeler 2018). This heterogeneity is a driver for numerous ecological and evolutionary processes, such as community composition, species richness, and local adaptation (Stein, Gerstner, and Kreft 2014; Yang et al. 2015; Forester et al. 2016). Where environments vary, populations with enough genetic variation have the opportunity to adapt to their local conditions via natural selection and this is a common phenomenon across many taxa (Via and Conner 1995; Klich 2000; Hereford 2009; Nunez et al. 2021). Local adaptation occurs when an organism's trait expression optimizes fitness values for its local environment, and is achieved by low gene flow and strong selection between populations (Kawecki and Ebert 2004). Mean phenotypes, here defined as the average expression of a trait due to genetic differences in a common environment, have been the primary focus of studies in the local adaptation literature (Hereford 2009; Savolainen, Lascoux, and Merilä 2013). Phenotypic plasticity, or the ability of one genotype to express multiple phenotypes in response to the environment, has also been hypothesized to be locally adaptive, particularly in response to predictable environmental heterogeneity (Snell-Rood and Ehlman 2021). In order for phenotypic plasticity to respond to locally varying natural selection there must be genetic variation in plasticity, or a genotype by environment interaction (GxE) (Josephs 2018). Therefore, demonstrating that a plastic trait exhibits GxE is the first step in investigating whether phenotypic plasticity is locally adaptive. Although classic studies in *Impatiens campensis* have detected an adaptive role for plasticity (Donohue et al. 2000, 2001), this has been difficult to test in most systems (Thompson 1991; Via et al. 1995; Donohue et al. 2001; Scheiner 2013).

In a natural landscape, mean phenotypic expression and phenotypic plasticity both contribute to an individual's lifetime fitness, and therefore both have the potential to play a role in local adaptation. It can be difficult to identify when plasticity is locally adaptive in natural populations because the amount of plasticity in a trait must be measured using the same genotype in different environmental conditions. These constraints can be overcome with experimental designs using clonal replicates, or half-siblings in systems where inbreeding or cloning cannot be performed (Relyea 2002; Cooper et al. 2022). Plants are sessile and many of them can be inbred or replicated clonally making them ideal for investigating the relative contributions of mean phenotype and plastic trait expression to local adaptation.

Characterizing the plastic response of a species to environmentally relevant conditions is

important in understanding its life history (Wilczek et al. 2009). Sarkar (2004) postulated that, if

tested for, plasticity is bound to be present in different environments and yet there are still

unanswered questions about the variation in, mechanisms of, and significance of phenotypic

plasticity in most natural populations. While plasticity in a single trait has been examined in

numerous systems (Chevin and Hoffmann 2017), few studies have examined plasticity in

multiple traits across an organism's lifetime. Therefore we lack data about the degree of

correlation between plastic expression of multiple traits and how these correlations may

influence fitness (Nielsen and Papaj 2022). Although the genetic basis of phenotypic plasticity

has been mapped in traits like flowering time (Fournier-Level et al. 2022) and branching pattern

in *Arabidopsis* (de Jong et al. 2019), whether plastic variation in a given trait is controlled by

separate genetic loci from variation in the trait mean is still not well understood (but see

Fournier-Level et al. 2022). By comparing the correlation between plastic and mean phenotypic

trait expression in a population exhibiting GxE we can investigate whether plasticity has

independent genetic controls (Holeski, Chase-Alone, and Kelly 2010). This information is

foundational in determining how plasticity is expressed from genotype to phenotype.

Elevation and latitude exert spatially varying selection across populations. Due to their

sessile nature, plants experience strong natural selection based on the environment around them.

Local adaptation to elevation has been found in many plant systems such as in *Arabidopsis*

arenosa populations from four foothill-alpine clines (Wos et al. 2022) and in *M. guttatus* in

response to drought stress (Kooyers et al. 2015). Phenotypic plasticity is known to be involved in

local adaptation in *Populus fremontii* populations across Arizona, where plasticity in traits such

as bud set and plant height were found to vary along an elevational gradient (Cooper et al. 2022).

Local adaptation of phenotypic plasticity in a number of plant functional traits in *Phragmites*

australis was found to follow a latitudinal gradient (Ren et al. (2020).

One important environmental cue in plants that may drive local adaptation is

photoperiod. Photoperiod varies temporally both predictably and unpredictably during the

growing season for sessile alpine and subalpine organisms. Photoperiod varies predictably with

shorter days occurring in the winter and the longest days occurring at the summer solstice. In this

way, photoperiod is a reliable sensing cue, which is hypothesized to be important for the

development of phenotypic plasticity (Snell-Rood and Ehlman 2021). Elevation and latitude are

two environmental gradients that interact with photoperiod in a heterogeneous manner as a result

of annually varying temperatures and snowpack. High latitude populations will always

experience longer photoperiods during the spring and summer growing seasons than low latitude

populations. Experienced photoperiod varies less predictably in mountainous environments

(Figure 1a and 1b). High elevations accumulate more snowfall and have a later timing of

snowmelt than low elevations, which means that for plants growing at the same latitude low

elevation populations will germinate and develop under shorter photoperiods than those

occurring at higher altitudes. However over time the amount of snowpack varies significantly

meaning that the experienced photoperiod of a short-lived annual plant can fluctuate from year to

year in alpine environments (Bormann et al. 2018). Therefore, photoperiod may be an important

cue that young seedlings utilize in alpine environments to determine time of year and their

subsequent developmental trajectory.

Mimulus laciniatus, a member of the *M. guttatus* species complex*,* is a Sierra Nevadan

endemic plant that grows on rocky granite outcrops from 1000-3300 meters in elevation (Fenster

and Ritland 1994; DeMarche, Rice, and Sexton 2013). Depending upon elevation and the timing

of snow melt, *M. laciniatus* populations experience different day lengths during the growing

season, making this an interesting system to examine patterns of local adaptation in response to

photoperiod fluctuation. A previous study by Ferris and Willis (2018) found that flowering time,

plant height, and leaf shape were under divergent selection between *M. laciniatus* and its close

relative *M. guttatus*. Leaf shape, and in particular leaf dissection, has been found to vary clinally

across elevation and latitude within other plant species (Gurevitch 1988; Campitelli and

Stinchcombe 2013; Wong et al. 2020). In many species it is unclear whether this trait variation

and potential adaptation arises via genetic variation in mean phenotype expression, phenotypic

plasticity, or a combination of the two (but see Bright and Rausher (2008); Campitelli and

Stinchcombe (2013)).

In this study we investigated whether mean and plastic variation in leaf shape, flowering

time, and floral size among *M. laciniatus* populations are consistent with local adaptation to

elevation and latitude. We grew replicates of genotypes from twenty-one populations across *M.*

laciniatus' elevational and latitudinal range in long and short day common garden experiments.

We then tested whether patterns of mean and plastic trait expression were associated with

elevation or latitude in the population of origin. This allowed us to examine whether the same

traits that are under divergent selection between closely related sympatric species may also be

under locally varying selection within *M. laciniatus*.

MATERIALS AND METHODS

<u>Study system</u>

Mimulus laciniatus is a small, self-fertilizing annual plant, with low levels of gene flow

expected between most populations (Ferris, Sexton, and Willis 2014; Sexton et al. 2016). Unlike

M. guttatus which displays rounded leaves, *M. laciniatus* has deeply lobed, narrow (laciniate)

leaves as a defining characteristic of the species (Ferris et al. 2015). We created inbred lines from

genotypes collected from natural populations of *M. laciniatus* across its geographic range to

investigate phenotypic responses to photoperiod fluctuation. Due to its high rate of self-

fertilization (93-97%) *M. laciniatus* is already highly inbred when first collected from the field

(Ferris et al 2014). We self-fertilized each genotype for at least 5 additional generations in the

lab. Twenty-one populations of *M. laciniatus* from across the central and southern Sierra Nevada

mountain range (California, USA) were used in this experiment (Table 1, Figure 2). These

populations originate from across 1000-2500m in elevation and two degrees of latitude (~36°-

38°; Figure 2) and six distinct watersheds (Figure S1). Low levels of gene flow between

populations have been observed in previous studies of *M. laciniatus* (Ferris, Sexton, and Willis

2014; Sexton et al. 2016).

<u>Experimental design</u>

To compare the variation in mean phenotype and plastic expression of quantitative traits

among *M. laciniatus* populations from across the species range we grew genetic replicates in 1) a

common garden in a long day environment to measure mean phenotypic expression of traits at

UC Davis in 2017, 2) another long day common garden experiment at Tulane University in

2020, and 3) a common garden in short days at Tulane University in 2021 to compare to the

Tulane long day garden and measure plasticity. We repeated the long day garden at Tulane to

control for possible effects of chamber and location on our phenotypes when measuring

plasticity. By replicating our long day garden, we are also able to avoid the statistical artifact of

regression to the mean when testing for differences in patterns of variation in mean phenotype

vs. phenotypic plasticity (Gunderson and Revell 2022). To measure genetic variation in mean

expression in a single environment we grew plants at UC Davis in a Conviron walk-in chamber

under 15 hour days at 21°C. To measure phenotypic plasticity, we sequentially grew plants at

two different photoperiods (11 hour & 15 hour days) in a Conviron walk-in chamber at Tulane

University. These long (15 hour) and short (11 hour) photoperiods represent the longest and

shortest natural day lengths within *M. laciniatus'* native range.

161   We grew three to ten unique inbred lines from each of twenty-one populations for a total

of 120 inbred lines in each growth chamber experiment. Seventy-one of those inbred lines were

grown in both the mean phenotype experiment at UC Davis and plasticity grow-outs at Tulane.

Three to five replicates of each inbred line per grow-out gave us a total of 1015 plants used to

measure mean expression of traits and 677 plants used to measure plasticity in this study. We

planted seeds atop 3" pots filled with wetted Sun Gro Fafard 3B/Metro-Mix 830 soil. Pots were

stratified in dark Percival chambers set at 4°C for 10 days and misted every other day. After 10

days, we moved pots to a Conviron walk-in growth chamber set to 21°C. We kept pots

constantly wet on full flood benches and misted daily. After seedlings showed their first set of

true leaves, we flooded benches with water for 8 hours (during light hours) and kept them empty

for 16 hours daily.

<u>Phenotypic measurements</u>

To examine ecologically relevant phenotypic variation in *M. laciniatus* across populations,

inbred lines, and photoperiods, we measured traits that have previously been shown to be under

selection in *M. laciniatus'* native habitat: flowering time (days from germination to first flower),

leaf lobing, and plant height, as well as mating system-related traits. The specific mating system

traits measured were: corolla length, corolla width, longest stamen length, shortest stamen

length, pistil length, and pseudo-cleistogamy (denoted as either pseudo-cleistogamous for a

closed flower or chasmogamous for an open flower). *Mimulus laciniatu*s is pseudo-

cleistogamous rather than truly cleistogamous because while the size and shape of the outer

corolla changes, the internal structure of the plant does not (Lord 1981). Table S1 provides a

chart of all traits measured. Critical photoperiod was also measured and calculated as the

proportion of individuals per genotype that reached flowering under the 11h photoperiod. All

plants flowered under 15h. For plants that flowered, we collected all phenotypic data on the day

of first flower. For plants that did not flower under 11 hours, but instead maintained a vegetative

state, we collected leaf lobing and plant height data two months after germination. We collected

leaf lobing data by harvesting the most lobed leaf from the second node of each plant. We taped

leaves to a sheet of white paper and scanned them for digital convex hull analysis of leaf lobing

in ImageJ (Ferris et al. 2015). The convex hull is the shape that is created by connecting the

outermost points of a leaf. The lobing index provided by convex hull analysis is calculated by

dividing the difference between the area of the true leaf and the area of the convex hull by the

larger area of the convex hull (Figure S2). The lobing index is a metric of leaf lobing from 0 to 1,

with a higher number equaling higher lobing (Ferris et al 2015). We measured plant height in

millimeters (mm) to the apical meristem of each plant using a ruler, and all floral traits were

measured in mm with calipers after removing the corolla from each flower using forceps. We

calculated herkogamy (stigma-anther separation) for each plant by subtracting the longest stamen

length from the pistil length. Herkogamy acts as a metric of self-fertilization, where a smaller

difference means a higher likelihood of self-fertilization (Karron et al. 1997).

<u>Statistical analyses</u>

To understand whether the plastic and mean values of the same phenotype were genetically

correlated, and therefore potentially subject to correlational selection, we performed a correlation

analysis. We input all of our measured mean and plastic trait values into the R package *corrplot*

which created a correlation matrix (Wei and Simko 2021). This allowed us to test, for example,

the degree to which genotypes that expressed a highly lobed mean leaf shape in a long-day

environment also displayed a large plastic change in leaf shape in response to different

photoperiods.

To test for significant phenotypic plasticity and GxE in all *M. laciniatus* traits across

photoperiods, we used a two-way ANOVA including genotype, photoperiod environment, and

genotype by environment interaction as fixed effects (Lorts and Lasky 2020; Zhang and

Lechowicz 1994). We calculated plasticity strength for each genotype by taking the difference

between 15 and 11h mean phenotypic values of each genotype in each photoperiod (Valladares,

Sanchez-Gomez, and Zavala 2006).

To examine patterns of trait variation consistent with local adaptation we tested for a

relationship between mean trait expression and population elevation and latitude. We used an

ANOVA with mean phenotype as the dependent variable for all traits measured. Elevation and

latitude were fixed effects in our models while genotype was nested within population as a
random effect. We used this same method to examine the relationship between the strength of
trait plasticity and population elevation and latitude.
Reaction norm and environmental cline figures were created in R with the package
ggplot2 (Wickham 2016). Maps were created in R using the package *ggmap* as well as QGIS
(Kahle and Wickham 2013; QGIS Development Team 2023).
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RESULTS
<u>Weak correlations between trait means and plasticity</u>
We performed a correlation analysis to investigate whether the mean phenotype and plasticity of
the same trait were genetically correlated and possibly subject to correlational selection. We
found that at the individual genotype level mean trait expression was not tightly associated with
the strength of plasticity in that same trait (Figure 3). Several mean traits were strongly
correlated ($r > +/-0.5$) with one another. Leaf area, plant height and flowering time were all
strongly correlated with each other as were corolla length, corolla width and stamen length.
Mean leaf shape was most strongly correlated with mean leaf size ($r = 0.59$) and more weakly
correlated with other mean traits. All floral plasticity measurements were tightly correlated with
each other: stamen length (long and short), pistil length, corolla length and corolla width ($r > +/-$
0.5). Due to the strong correlation between floral size traits we focus on corolla width as a
representative floral trait for the rest of the results. For the majority of traits measured there was
not a strong correlation between mean phenotype and plastic expression of the same phenotype
($r < +/- 0.5$). For example, our results show that mean leaf shape expression of a genotype was
only weakly correlated ($r = 0.11$) with the magnitude of leaf shape plasticity in that same
genotype (Figure 3).
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<u>Genotypes from across the species' range respond plastically to photoperiod</u>
To test whether *M. laciniatus* exhibited plasticity in response to different photoperiods, we grew
replicate genotypes in long and short day controlled environments. Significant phenotypic
plasticity, measured by a significant effect of photoperiod environment in our models, was found
in all traits that we studied (Figure 4, Table 2). Plant height showed the highest proportion of
genotypes that exhibited significant plastic response to photoperiod change indicating that height

in *M. laciniatus* is an environmentally labile trait. Across all genotypes the degree of plastic

change in height was on average an increase in 45 mm between short and long days. Other

phenotypes were less plastic at the species level in response to photoperiod such as leaf lobing,

which on average only increased on our leaf lobing index by 0.03 between short and long days.

However, a significant GxE interaction was detected in all traits studied indicating there is a

great deal of genetic variation for phenotypic plasticity in this species (Table 2). In addition,

some phenotypes had more similar patterns of plastic change than others, with over 40% of all

genotypes growing taller, flowing faster, and with larger flowers in long days. This indicates that

the direction of plasticity is mostly fixed across the geographic range of *M. laciniatus* in height,

flowering time, and flower size. Genetic variation in leaf shape plasticity however was not as

consistent in direction or degree across populations making it particularly interesting for follow

up analyses. The detection of GxE in plant height, flowering time, floral size, and leaf shape

makes these traits the focus of our remaining analyses testing for patterns of local adaptation

within *M. laciniatus.*

<u>Mean leaf shape varies clinally with both elevation and latitude</u>

To examine phenotypic patterns consistent with local adaptation in *M. laciniatus* we first tested

for associations between variation in mean phenotype and elevation or latitude of origin. We

found that mean phenotype expression was significantly associated with origin of population

elevation in both morphological and life history traits of *M. laciniatus* (Table 3). Critical

photoperiod, the photoperiod threshold required to reach flowering, was positively associated

with population elevation of origin ($R^2 = 0.2$, p = 0.003, Figure 5). Critical photoperiod is longer

for high elevation populations, the majority of which did not flower in our 11 hour day treatment

but did flower under 15 hour days. On the other hand, many genotypes from low elevation

populations were able to flower under short days. Mean leaf shape also shows a significant

positive association with population elevation, with leaves becoming more lobed at higher

elevations ($R^2 = 0.36$, p = 0.003, Figure 5). Two mean traits were marginally associated with

population elevation in our first long day experiment: flowering time and corolla width ($R^2 =$

0.11 and 0.12, p = 0.056 and 0.058, Figure 5). In these traits populations from higher elevations

flowered later and had larger flowers than low elevation populations.

Mean leaf lobing was the only trait which had a relationship with latitude ($R^2 = 0.36$, p =

0.0663; Figure 6). Even across the short latitudinal range of *M. laciniatus* we found that high

latitude populations have more deeply lobed leaves than low latitudes. All other traits had no

relationship with latitude (Figure 6, Table 4). Mean leaf lobing was therefore the only trait to be

significantly associated with variation in both elevation and latitude across the species' range.

<u>Phenotypic plasticity in *M. laciniatus* varies only with elevation</u>

We next set out to detect patterns consistent with local adaptation in phenotypic plasticity. Using

a mixed model approach, we found that the strength of phenotypic plasticity in both life history

and morphology was significantly associated with population elevation across the geographic

range of *M. laciniatus* (Table 5). The degree of plastic change in leaf shape in response to our

photoperiod treatments was greater in high than in low elevation populations ($R^2 = 0.26$, p =

0.0663, Figure 7). High elevation populations became significantly more lobed in long days.

Low elevation *M. laciniatus* populations had either very little change in leaf shape between

treatments or the amount of lobing actually decreased slightly in response to long days.

Therefore, both mean leaf lobing and leaf lobing plasticity increase in higher elevation

populations. Plasticity in a number of floral size traits showed marginally significant trends (0.1

> p > 0.05) with elevation. The amount of phenotypic plasticity increased in higher elevation

populations for long and short stamen length, pistil length, and pseudocleistogamy (Figure 7).

Variation in flowering time and corolla size plasticity was not associated with elevation even

though the mean values of these traits increased in higher elevation populations. Plant height,

leaf size, and herkogamy plasticity also did not associate with altitude (Figure 7). When

plasticity was analyzed for a relationship with latitude, no trait showed any clinal association

(Figure S3, Table S2).

DISCUSSION

In this study we investigated whether quantitative traits that are involved in adaptation to *M.*

laciniatus's harsh granite outcrop environment - flowering time, leaf shape and flower size

(Ferris and Willis 2018) - also appear to be under selection in response to environmental

variation across the species' range. To test for patterns consistent with local adaptation to

elevation and latitude we grew replicates of genotypes from across *M. laciniatus*'s geographic

range under both long and short day lengths. Plasticity and GxE were detected in response to

photoperiod in all traits that we measured. For local adaptation of plasticity to evolve there must

be GxE in the traits' expression and genetic variation in plasticity associated with variation in the

environment. We found that patterns of genetic variation in the mean phenotypic expression of

flowering time, flower size, and critical photoperiod were associated with elevation making them

interesting traits to investigate in future studies about adaptation across *M. laciniatus's*

geographic range. In our study both mean and plastic leaf shape expression were significantly

associated with elevation, strongly suggesting that these traits are involved in local adaptation

within *M. laciniatus*. Furthermore, we found that leaf lobing expression in long days was

positively associated with latitude as well as elevation indicating that increased lobing may be

adaptive in colder environments.

<u>Mean phenotype and plastic trait expression seem genetically independent</u>

In our study of range-wide phenotypic variation in *M. lacinaitus* mean trait expression had

minimal influence on the degree of plasticity in the same trait (Figure 3). Notably in leaf lobing,

which both mean and plastic variation are similarly associated with elevation there was little

correlation between how lobed a leaf was in a long day environment and how plastic that

genotype's leaf shape expression was between day lengths. The independence of expression in

these two traits implies that there is likely a separate genetic basis for mean and plastic

expression of leaf lobing. Independence between mean and plastic trait expression has been

theorized in a handful of studies (Ehrenreich and Pfennig 2016; Johansson et al. 2013), and

identified in others (Lind et al. 2011; Snell-Rood et al. 2011; Tétard-Jones, Kertesz, and Preziosi

2011). While Snell-Rood et al. (2011) focus on an extreme form of plasticity, polyphenism, their

study aimed to investigate their developmental divergence hypothesis first discussed in the

literature with regards to plasticity by West-Eberhard (1989). Snell-Rood et al. (2011) found

divergence in patterns of gene expression between different morphs in two species of horned

beetle, *Onthophagus taurus* and *Onthophagus nigriventris*. Plasticity-biased genes were found to

be more divergent than those with shared patterns of expression between beetle morphs. Tétard-

Jones et al. (2011) also found independent plasticity QTL located separately from main effect

QTL in barley growth, *Hordeum vulgare*, adding more evidence that plasticity is controlled by

separate genetic loci than mean trait expression loci.

In a trait that is up- or down-regulated by common developmental pathways, expression

of phenotypic plasticity is not linked to specific environmental cues and is more likely to be

linked with mean trait expression (Snell-Rood et al. 2011). However, traits may also be

controlled by environment-specific gene regulation, which adjusts expression in these common

developmental pathways and can contribute to phenotypic plasticity. Given the lack of

correlation between mean and plastic expression in *M. laciniatus* perhaps additional genes are

triggered to cause differences in leaf shape in different environments. These adjustments would

reduce pleiotropic constraints and in this way could facilitate rapid adaptation to different

environments (West-Eberhard 1989, 2005).

Mimulus laciniatus exhibits range-wide plasticity and GxE in response to photoperiod

Phenotypic plasticity is expected to evolve when future environmental conditions will vary, and

presently sensed cues are reliable in predicting those future conditions (Snell-Rood and Ehlman

2021). We therefore may expect that plasticity evolved in response to *M. laciniatus*' annual

variation in climate and photoperiod, and was subsequently selected against in populations which

have more stable, temporally homogeneous environments from year to year. We identified

significant plasticity and GxE in all traits measured in different photoperiods in our set of range-

wide *M. laciniatus* genotypes. Identifying GxE means there is genetic variation in plastic trait

expression, and therefore selection can act on these different reaction norms potentially giving

rise to local adaption.

Some traits such as plant height and flowering time appear to show a strong plastic

response across genotypes, but less variation in strength and direction of plasticity among

genotypes. Plant height and flowering time are strongly phenotypically correlated in our

experiment (r = 0.72) which makes sense because these two phenotypes have been shown to be

controlled by the same light sensing genetic pathway in other plants (Pigliucci and Schlichting

1996; Mockler et al. 2003). This is well documented in the *Arabidopsis* system, in which much is

known about genetic controls linking flowering and plant height (Alvarez et al. 1992; Salehi et

al. 2005). In *A. thaliana* the *terminal flower* and *FLOWERING LOCUS C* loci act

antagonistically to genetic pathways that regulate a vegetative-to-reproductive switch in resource

allocation. Given the likely genetic correlation this more universal plastic response in flowering

and height across *M. laciniatus* populations may be due to genetic constraint (Auld, Agrawal,

and Relyea 2010). Previous reciprocal transplant experiments have demonstrated that early

flowering time is under strong selection in *M. laciniatus*' harsh rocky outcrop environments

(Tataru et al 2023; Ferris and Willis 2018). Given the ephemeral nature of water availability in

the shallow rocky soils of this species' habitat and the significant variation in the amount and

timing of snow melt from year to year it seems likely that an ability to respond plastically in the

timing of flowering is adaptive across *M. laciniatus'* range. Therefore, plasticity in crucial life

history traits like flowering time and size could be universally, rather than locally, adaptive in

our short-lived rocky outcrop specialist.

<u>Patterns of local adaptation to elevation in *M. laciniatus* present in both mean and plastic traits</u>

Reciprocal transplant experiments have found selection for early flowering time, more highly

lobed leaves, and larger flowers in *M. laciniatus'*s harsh granite outcrop habitat (Ferris and

Willis 2018; Tataru, Wheeler, and Ferris 2023). However, it is not known whether these same

traits are involved in local adaptation across *M. laciniatus'*s range. One environmental variable

that is likely to drive population level adaptation in this system is variation in elevation across

the Sierra Nevada, CA. Using a long day common garden we found that variation in critical

photoperiod, mean flowering time, flower size, and leaf lobing as well as plasticity in leaf lobing

was associated with population elevation. Previous studies have found low levels of gene flow

between populations of *M. laciniatus* (Ferris, Sexton, and Willis 2014; Sexton et al. 2016),

therefore associations between phenotypic variation and elevation that are repeated across

multiple mountain clines may be more likely caused by natural selection than by gene flow

alone.

In our sample of *M. laciniatus* populations critical photoperiod increases with elevation.

This pattern could be indicative of local adaptation to flower under shorter day lengths that lower

elevation populations experience during their early spring growing season. These results are

supported by Friedman & Willis (2013) who previously found that flowering in higher elevation

M. laciniatus populations requires a longer photoperiod. However, Leinonen et al. (2020) found

that with a more limited population sample mid-elevation populations required the longest

photoperiod to reach flowering. In addition, Leinonen et al. (2020) found no significant

association of population elevation with other *M. laciniatus* phenological traits in response to

photoperiod. These previous studies examined only a few populations of *M. laciniatus* (<6

populations) compared to our increased sampling of the species range (21 populations total).

Mimulus guttatus is also known to have locally adapted to photoperiod, one such example being

the geothermally adapted *M. guttatus* of Yellowstone National Park, WY. These plants have

adapted to flower under short winter days when geothermal vents make the surrounding

environment wet and warm, suitable for plant growth and reproduction (Lekberg et al. 2012).

Photoperiodism has been documented in many plant species as a crucial cue for seasonal change

(Davis 2002; Mockler et al. 2003; MacQueen et al. 2021). High elevation populations in other

species have been found to be locally adapted to longer photoperiods similar to our findings,

such as in *Populus* and red spruce tree species (Pauley and Perry 1954; Butnor et al. 2019) as

well as more herbaceous alpine plant species such as *Arabidopsis arenosa* (Wos et al. 2022).

High-elevation adaptation to longer photoperiod has also been identified in non-plant species,

such as pitcher plant mosquitoes and burying beetles (Bradshaw and Lounibos 1977; Tsai et al.

2020). In turn, photoperiodism can inform how and when resources are directed during a plant's

life span (i.e. to growth or reproduction). In our study while short photoperiod leaves of *M.*

laciniatus are less lobed, they are larger on average than long photoperiod leaves. For genotypes

that didn't flower in a short photoperiod, the resources were instead being shunted into

production of vegetative tissue, which could be a resource conservation measure (Freschet et al.

2010; Reich 2014). A trait that allows for sensing time of year, such as photoperiodism would be

beneficial for *M. laciniatus*, as it provides information to these plants on how much time there is

left to grow in the season before the occurrence of seasonal drought and senescence in their

ephemeral rocky environment.

Patterns of flowering time and corolla width are also consistent with local adaptation to

elevation in *M. laciniatus* as higher elevation populations flower later and have wider corollas.

The idea of later flowering being adaptive at high elevations is contradictory to what is typically

expected in alpine and high latitude plants because of the shortened growing season in these

environments. In most systems it is thought that populations at high elevation should flower

faster than those at low because if individuals flower too late in the season they may be killed by

an early frost before reproducing (Stock, Campitelli, and Stinchcombe 2014). However, *M.*

laciniatus's unique rocky outcrop environment combined with the Mediterranean climate of

California produce a different expectation in this system that is tied to water availability rather

than the timing of first frost. In the Sierra Nevada, CA the majority of seasonal soil moisture is

due to snow melt with little rain occurring during the growing season. Snowpack levels are lower

at low elevations meaning water resources will run out faster shortening the growing season due

to drought at low vs high mountain sites. Early flowering time allows plants to escape from the

onset of seasonal drought while bigger, later flowering plants usually occur in more continuously

moist environments (Heschel and Riginos 2005; Hall and Willis 2006; Franks, Sim, and Weis

2007; Kenney et al. 2014; Ferris and Willis 2018). Therefore plants adapted to low elevations

should flower faster in order to reproduce successfully compared to those at high elevations (Fox

1990). Dickman et al. (2019) and Leinonen et al. (2020) have also found that high elevation *M.*

laciniatus flower later than low elevation populations. Flower size has been shown to increase in

plant populations at high elevation due to the increased attractiveness of large flowers in an

environment where pollinators are scarce (Fabbro and Körner 2004; Maad, Armbruster, and

Fenster 2013). However, *M. laciniatus* is highly self-fertilizing so this explanation seems less

relevant in our study system. Increased flowering time and larger flowers at high elevations

where there is greater water availability may be adaptive in *M. laciniatus* because later flowering

plants can become larger and more fecund. A large body of work exists in local adaptation of

flowering time phenology and flower size, indicating that these traits often experience selection

in nature in order to optimize fitness (Lowry and Willis 2010; Wang et al. 2014; Dickman et al.

2019; Leinonen et al. 2020).

452   Leaf shape, size, thickness, and number have all been studied previously in various plant

species and identified as locally adaptive to harsh marginal habitats such as high light or extreme

hot or cold temperatures (Nagy 1997; Dudley 1996; Parker, Rodriguez, and Loik 2003;

Campitelli and Stinchcombe 2013; Ferris et al. 2015). In our study *Mimulus laciniatus*

populations from higher elevations had more highly lobed and highly plastic leaves than

populations from lower elevations (Figure 5, Figure 7). We also saw a positive trend between

leaf lobing and latitude which suggests that it is adaptive to have more highly lobed leaves in

northern populations. Lobed leaves have a thinner boundary layer which means they equilibrate

more efficiently to air temperature through convection and therefore should not drop below

ambient temperature on clear nights when leaves radiate heat to the sky (Givnish 1979). This

could prevent lobed leaves from incurring frost damage when ambient temperature is near, but

still above freezing (Schuepp 1993; Nobel 2009). Patterns consistent with increased cold

tolerance of lobed leaves have been found in other plant systems. For example, along the east

coast of North America there is a latitudinal cline in leaf shape in the Ivyleaf morning glory,

Ipomoea hederaceae, with northern populations fixed for the lobed morph while those in the

south have entire leaves (Campitelli and Stinchcombe 2013). When Campitelli, Gorton and

colleagues (2013) tested for differences in leaf temperature they did find that lobed *Ipomoea*

leaves stayed warmer at night than round leaves, although no significant difference in was found

in freeze tolerance itself. Preliminary work on leaf temperature in *Mimulus* found that the lobed

leaves of *M. laciniatus* stayed warmer throughout a 24-hour period than the round leaves of *M.*

guttatus in common garden experiment in the field (K. Ferris unpublished data). A study using

Solanum lycopersicum found that shade-avoidant plants had more complex leaf structure in both

lobing and serration, indicating that high light can also drive a more complex leaf shape

(Chitwood et al. 2015). The patterns we found of greater leaf lobing in both high elevation and

high latitude populations of *M. laciniatus* are consistent with leaf lobing being adaptive in colder

environments (Yapp 1912; Nobel 2009).

<u>Conclusion</u>

Our study investigated how mean and plastic trait variation can contribute to local adaptation in

heterogeneous environments. The evolution of leaf shape, critical photoperiod, and leaf shape

plasticity across 21 *M. laciniatus* populations spanning multiple watersheds is consistent with a

signature of local adaptation (James et al. 2023). The results of this study raise a number of

questions for future research. First, are these patterns of mean and plastic trait expression truly

due to natural selection? What genes underlie the plastic responses observed due to photoperiod?

Are the genes that control for a plastic response in changing photoperiod the same that control

mean trait expression in a long day environment, and does this hold true for all traits studied?

What is the function of leaf lobing in the field setting? While functions are hypothesized above,

there is currently a lack of data to validate these claims. In addition, by further studying the

strength of selection on plastic traits it would be possible to investigate costs associated with

plasticity in mismatched environments. Our data supports the idea that highly lobed leaf shape is

locally adaptive in high elevation and northern populations of *M. laciniatus*. In adaptation

between *Mimulus* species we see evidence that ephemeral water availability in *M. laciniatus's*

dry rocky habitat may be driving the evolution of lobed leaf shape (Tataru et al 2023; Ferris and

Willis 2018). However high elevation highly lobed populations of *M. laciniatus* tend to have

higher water availability than low elevation populations with less lobing, implying that the

selective force causing local adaptation in leaf shape within this species may be different than the

force driving divergent evolution between species. Future studies using genomic data in

combination with experimental manipulation in the field will be able to further investigate these

questions.

ACKNOWLEDGEMENTS

We thank Natalie Gonzalez, Aditi Mahesh, and Cecilia Hammond for help with planting and

data collection. We also thank Johanna Schmitt and Graham Coop for all of their support and

feedback on the portion of this project completed at University of California Davis, and Mirielle

Caton-Darby for assistance collecting the mean phenotypic data presented in this paper. This

research was supported by the National Institute of General Medical Sciences of the National

Institute of Health (NIH) under Award Number R35GM138224. The content is solely the

responsibility of the authors and does not necessarily represent the official views of the NIH.

This work was also supported by a Center for Population Biology Postdoctoral Fellowship at the

University of California Davis and a Tulane University Ecology and Evolutionary Biology

Graduate Student Grant.

For research conducted at UC Davis, we should take a moment to acknowledge the land

on which we gathered. For thousands of years, this land has been the home of Patwin people.

Today, there are three federally recognized Patwin tribes: Cachil DeHe Band of Wintun Indians

of the Colusa Indian Community, Kletsel Dehe Wintun Nation, and Yocha Dehe Wintun Nation.

The Patwin people have remained committed to the stewardship of this land over many

centuries. It has been cherished and protected, as elders have instructed the young through

generations. We are honored and grateful to be here today on their traditional lands.

For research conducted at Tulane University, we acknowledge and pay tribute to the

original inhabitants of this land. The city of New Orleans is a continuation of an indigenous trade

hub on the Mississippi River, known for thousands of years as Bulbancha. Native peoples have

lived on this land since time immemorial, and the resilient voices of Native Americans remain an

inseparable part of our local culture. With gratitude and honor, we acknowledge the indigenous

nations that have lived and continue to thrive here.

DATA AVAILABILITY

Data and code are available from the Dryad Digital Repository.

https://datadryad.org/stash/share/xyNBULQagWnkmI3oAbWKPEIJcZPvxR-1xg3zxwGEnKI

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830
Figure 1. a) Early in the growing season, only lower elevation populations have warmed up

enough to experience complete snowpack melt. These populations begin to grow earlier in the

spring, and therefore experience a shorter photoperiod. b) By early to mid summer, all snowpack

has melted from higher elevation sites, therefore high elevation populations grow during longer

photoperiods. c) Snowpack level changes from year to year, and the pace of warming may be

slower or faster. This creates fluctuation in experienced photoperiod over time as well as across

elevation. Snow depth data collected at Dana Meadows in Yosemite National Park (California

Data Exchange Center 2023). Created with BioRender.com.

840
841 Figure 2. A map of the Sierra Nevada mountain range with each population mapped to its
842 original collection location. Populations are colored based on latitude.
843

844
845 Figure 3. Correlation matrix of mean trait expression and plastic trait expression. Plastic trait
846 expression is denoted by (p). Blank squares had an insignificant correlation (p > 0.05) when
847 tested with a t-test.
848

850   Figure 4. Reaction norms for a selection of focal traits. Trait values of *M. laciniatus* inbred lines

851   were measured in controlled growth chamber conditions under two different photoperiods: 11

852   hour and 15 hour days. Reaction norms faceted by population can be found in Figure S2. See

853   Table 2 for the significance of genotype by environment interactions for each trait.

858   Figure 5. Trait mean expression per genotype plotted against elevation of origin for each

859   population. See Table 3 for the statistical analysis of each clinal association.

Figure 6. Trait mean expression per genotype plotted against latitude of origin for each

population. See Table 4 for a statistical analysis of the clinal association of each trait with

elevation.

866
867 Figure 7. Mean trait plasticity per genotype plotted against elevation of origin for each

868 population. See Table 5 for statistical analysis of the association between trait plasticity and

869 latitude.

 TABLES
875 Table 1. Geographic data for each *M. laciniatus* population used in the study.

Population name	GPS coordinates (Decimal Degrees)	Location description	Elevation (m)
BCL	37.2049, -119.246	Big Creek, Fresno County, CA	1493.45
BML	37.090509, -119.3150	Mill Run Lane, Shaver Lake, CA	1696.74
CECL	37.300143, -119.4835	Central Camp Rd, Madera County, CA	1274.67
CRV	37.812585, -119.50447	South of Olmstead Point parking area, Yosemite National Park, CA	2577.44
DNK	37.0511, -119.13102	Dinky Creek Rd, Sierra National Forest, CA	1851.57
GB	37.0708, -119.23165	Grand Bluff off Hwy 168, Sierra National Forest, CA	1679.37
HEH	37.89314, -119.84853	Evergreen Rd, Yosemite National Park, CA	1396.83
HHR	37.957206, -119.7860	Hetch Hetchy Reservoir trail, Yosemite National Park, CA	1207.56
HUL	37.14372, -119.09606	Hwy 168 before south turn off to Huntington Lake Rd, Fresno County, CA	2170.68
LIB	36.36958, -118.48498	Generals Hwy east of Little Baldy, Sequoia National Park, CA	2196.28
OPN	37.8107, -119.4852	Olmstead Point, Yosemite National Park, CA	2529.72
PETL	37.03346, -119.22122	Peterson Rd, Fresno County, CA	1256.63
QAL	36.7203, -118.02306	By Quail Flat off Generals Hwy, Tulare County, CA	1986.99
SBL	37.02384, -119.2455	Sandy Bluffs, Hwy 168, Sierra National Forest, CA	991.77
SHL	37.08682, -119.18388	Hwy 168, Shaver Lake, CA	1594.33
ST	37.76637, -119.54167	Snow Trail near falls, Yosemite National Park, CA	1869.95
TRT	37.42990, -119.42321	Hwy 41 below Turtleback Dome, Yosemite National Park, CA	1480.65
WCL	37.20227, -119.34225	Willow Creek by Bass Lake Rd, Fresno County, CA	1100.27
WLF	37.50492, -119.35631	Tioga Rd east of White Wolf, Yosemite National Park, CA	2395.31
WUV	36.681, -118.88028	Along road to Wuksachi Village, Sequoia National Park, CA	2129.638
YCN	37.8264, -119.5957	Yosemite Creek on Tioga Rd, Yosemite National Park, CA	2346.85

878 Table 2. Significance of genotype (G), environment (E), and genotype by environment (GxE)
879 interactions for each phenotype. A two-way ANOVA was used to generate these statistical
880 comparisons: df = degrees of freedom, F = F-value, p = p-value.

Trait	G	E	GxE
Plant height	df = 112 F = 4.0948 p < 0.001	df = 1 F = 2251.7352 p < 0.001	df = 109 F = 3.4516 p < 0.001
Flowering time	df = 112 F = 8.47 p < 0.001	df = 1 F = 809.2933 p < 0.001	df = 88 F = 3.8501 p < 0.001
Leaf lobing	df = 112 F = 3.3609 p < 0.001	df = 1 F = 49.5744 p < 0.001	df = 108 F = 2.0924 p < 0.001
Leaf area	df = 112 F = 5.6028 p < 0.001	df = 1 F = 600.5746 p < 0.001	df = 108 F = 2.8207 p < 0.001
Corolla length	df = 112 F = 5.7080 p < 0.001	df = 1 F = 506.5164 p < 0.001	df = 88 F = 2.8753 P < 0.001
Corolla width	df = 112 F = 6.7879 p < 0.001	df = 1 F = 699.9515 p < 0.001	df = 88 F = 2.2367 p < 0.001
Short stamen length	df = 112 F = 4.433 p < 0.001	df = 1 F = 530.3004 p < 0.001	df = 88 F = 2.8457 P < 0.001
Long stamen length	df = 112 F = 4.9323 p < 0.001	df = 1 F = 477.1790 p < 0.001	df = 88 F = 2.4494 p < 0.001
Pistil length	df = 112 F = 4.5903 p < 0.001	df = 1 F = 409.9971 p < 0.001	df = 88 F = 1.8443 p < 0.001
Herkogamy	df = 112 F = 1.8216 p < 0.001	df = 1 F = 104.4350 p < 0.001	df = 88 F = 3.1339 p < 0.001
Cleistogamy	df = 112 F = 2.7126 p < 0.001	df = 1 F = 138.1892 p < 0.001	df = 87 F = 3.5217 p < 0.001

883 Table 3. Significance of association between mean trait expression and population elevation. An
884 ANOVA was used to generate these statistical comparisons. For all traits, numDF = 1, denDF =
885 18. *** = p < 0.01; ** = p < 0.05; * = p < 0.1.

Trait	F-value	p-value	R^2	Direction of correlation
Critical photoperiod	**13.44429**	**0.003*****	**0.2**	+
Leaf shape	**12.276596**	**0.003*****	**0.36**	+
Flowering time	4.179683	0.056*	0.11	-
Corolla width	4.086909	0.058*	0.12	+
Pistil length	2.57640	0.126	0.048	
Leaf area	2.4211552	0.137	0.11	
Corolla length	2.27840	0.149	0.094	
Herkogamy	1.8451199	0.191	0.015	
Plant height	0.397350	0.536	0.047	
Stamen length	0.34039	0.5668	0.039	

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888　　Table 4. Significance of association between mean trait expression and population latitude. An
889　　ANOVA was used to generate these statistical comparisons. For all traits, numDF = 1, denDF =
890　　13. *** = p < 0.01; ** = p < 0.05; * = p < 0.1.

Trait	F-value	p-value	R^2	Direction of correlation
Leaf lobing	4.017275	0.0663*	0.26	+
Corolla length	1.837361	0.1983	0.15	
Flowering time	1.753481	0.2083	0.0019	
Herkogamy	1.535635	0.2372	0.00018	
Corolla width	1.0265023	0.3295	0.12	
Anther length	0.7934602	0.3892	0.056	
Critical photoperiod	0.651162	0.4342	0.022	
Plant height	0.6417213	0.4375	0.0000056	
Pistil length	0.0470382	0.8317	0.016	
Leaf area	0.0344051	0.8557	0.017	

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893   Table 5. Significance of association between strength of plasticity and population elevation. An
894   ANOVA was used to generate these statistical comparisons. For all traits, numDF = 1, denDF =
895   14. *** = $p < 0.01$; ** = $p < 0.05$; * = $p < 0.1$.

Trait	F-value	p-value	R^2	Direction of correlation
Leaf shape plasticity	**8.790705**	**0.010*****	**0.14**	+
Short stamen length plasticity	4.068938	0.063*	0.14	+
Pseudocleistogamy plasticity	3.933972	0.0673*	0.087	+
Pistil length plasticity	3.836726	0.070*	0.13	+
Long stamen length plasticity	3.798262	0.072*	0.16	+
Corolla length plasticity	3.7866308	0.117	0.1	
Corolla width plasticity	1.746458	0.208	0.086	
Plant height plasticity	0.365843	0.555	0.0018	
Herkogamy plasticity	0.1446548	0.709	0.00000095	
Flowering time plasticity	0.115048	0.740	0.0049	
Leaf area plasticity	0.044490	0.836	0.0021	

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 SUPPORTING INFORMATION

899
900 Figure S1. A map of study populations and watershed boundaries in California.
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Figure S2. A visual depiction of the leaf lobing index used to measure leaf shape. The convex
hull is outlined in red. A greater difference between true leaf area and the convex hull means a
higher leaf lobing index. Leaf petioles were not included in leaf shape analysis.

Figure S2. Reaction norms for each trait, faceted by population.

Figure S3. Trait plasticity mean expression per genotype plotted against latitude of origin for

each population. See Table S2 for statistical analysis of clinal association.

Table S1: Each trait measured for mean and plastic phenotypes

Trait	Mean phenotype	Plastic
Corolla length	Y	Y
Corolla width	Y	Y
Critical photoperiod	Y	--
Flowering time	Y	Y
Herkogamy	Y	Y
Leaf area	Y	Y

Leaf lobing	Y	Y
Long stamen	Y	Y
Pistil length	Y	Y
Plant height	Y	Y
Pseudo-cleistogamy	--	Y
Short stamen	--	Y

914 Table S2. Significance of association between plastic trait expression and population latitude. An
915 ANOVA was used to generate these statistical comparisons. For all traits, numDF = 1, denDF =
916 13. *** = $p < 0.01$; ** = $p < 0.05$; * = $p < 0.1$.

Trait	F-value	p-value	R^2
Leaf area plasticity	0.3967746	0.5397	0.013
Flowering time plasticity	0.3773531	0.5496	0.02
Short stamen plasticity	0.293991	0.5968	0.0072
Long stamen plasticity	0.217931	0.6483	0.0026
Leaf lobing plasticity	0.173843	0.6835	0.00021
Corolla length plasticity	0.0889135	0.7703	0.0025
Plant height plasticity	0.0513789	0.8242	0.014
Pseudocleistogamy plasticity	0.030234	0.8646	0.0019
Pistil length plasticity	0.00739	0.9328	0.0000064
Herkogamy plasticity	0.00497844	0.9448	0.0034
Corolla width plasticity	0.0027088	0.9593	0.0023