Vegetative Phase Change Causes Age-Dependent Changes in Phenotypic Plasticity

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Abstract

1. Phenotypic plasticity allows organisms to optimize traits for their environment. As

organisms age, they experience diverse environments that merit varying degrees of

phenotypic plasticity. Developmental transitions can control these age-dependent

changes in plasticity and as such, the timing of these transitions can determine when

plasticity changes in an organism.

2. Here we investigate how the transition from juvenile-to adult-vegetative development

known as vegetative phase change (VPC) contributes to age-dependent changes in

phenotypic plasticity using both natural accessions and mutant lines in the model plant

Arabidopsis thaliana. Further, we look at how the timing of this transition and the

concordant shifts in plasticity change across accessions and environments.

3. We found that the adult phase of vegetative development has greater plasticity than

the juvenile phase and confirmed that this difference in plasticity is caused by VPC

using mutant lines. Further, we found that the timing of VPC, and therefore the time

when increased plasticity is acquired, varies significantly across genotypes and

environments.

4. This genetic and environmental variation in the timing of VPC indicates the potential

for population-level adaptive evolution of VPC. The consistent age-dependent changes

in plasticity caused by VPC add further support to the hypothesis that VPC is adaptive.

Key words: age-dependent plasticity, developmental timing, ontogeny, phenotypic

plasticity, plant development, vegetative phase change

Introduction

Phenotypic plasticity, and its inverse – robustness -- are attributes of
development in all organisms. Whether plasticity or robustness in a given trait is most
beneficial for fitness depends on the degree of environmental variability, reliability in
environmental cues and the costs associated with phenotypic adjustment (Fischer *et al.*,
2014). Although there are many examples of adaptive plasticity, we still have much to
learn about what regulates plasticity in traits, including when plasticity changes, which
environmental cues bring about plasticity, and the degree of plasticity in a trait.

As organisms age, they transition through different developmental phases that
result in changing tradeoffs, while simultaneously experiencing diverse environments
during those phases. As such, they might benefit from varying degrees of phenotypic
plasticity across their lifespan. For example, cichlid fish display increased plasticity in
response to alarm cues to increase morphological defenses (i.e. body size and
coloration) when they are most vulnerable—shortly after birth and at the onset of
reproduction (Meuthen *et al.*, 2018). Age-dependent changes in plasticity associated
with whole plant development are often overlooked, though some have investigated the
distinct changes in plasticity associated with the age of a particular organ (e.g., changes
in photosynthetic traits associated with aging leaves (Niinemets, 2016)). However, a few
studies demonstrate variation in plasticity surrounding reproduction and at different
phases of vegetative growth. Two different studies found that *Arabidopsis thaliana* and
the aquatic plant *Sagittaria latifolia* have greater phenotypic plasticity in response to
changes in nutrients post-flowering compared to pre-flowering (Zhang & Lechowicz,
1994; Dorken & Barrett, 2004). Older but pre-flowering *Plantago lanceolata* individuals
show greater plasticity of chemical response to herbivory compared to younger
individuals (Barton, 2008). In some cases, the impacts of ontogeny on plasticity causes
individuals of *different species* at the same developmental stage to respond more alike
to environmental variation than *conspecifics* in different developmental phases (Parrish
& Bazzaz, 1985). Despite the significance of developmentally regulated changes in
plasticity and its impacts on fitness in animals (Hoverman & Relyea, 2007; Fischer *et*
al., 2014; Nilsson-Örtman *et al.*, 2015; Meuthen *et al.*, 2018; Sebestyén *et al.*, 2020) we
know relatively little about developmental changes in plasticity in plants.

In contrast, many studies examine plasticity in the timing of plant developmental

transitions, particularly flowering time and germination. Plasticity in the timing of

developmental transitions allows plants to optimize their life history for their

environment. For example, longer days and warmer temperatures accelerate flowering

in *A. thaliana* (Levy & Dean, 1998; Blázquez *et al.*, 2003) to promote flowering in the

spring and summer when conditions are favorable, and seeds that mature during short

photoperiodic days (i.e. autumn) are more sensitive to cold, delaying germination until

after winter (Munir *et al.*, 2001). Furthermore, Zhang & Lechowicz (1994) showed that

the timing of developmental transitions and the degree of plasticity in other traits during

developmental phases can be interconnected. Specifically, they found that in *A. thaliana*

a delay in flowering time was associated with greater plasticity in post-flowering traits

(Zhang & Lechowicz, 1994). Together this indicates that the timing of developmental

transitions can alter an organism's phenotypic plasticity both by dictating which

developmental phase the organism is in, and the degree of trait plasticity within that

phase (Fig. **1**).

All plants transition between distinct juvenile and adult phases during vegetative

development, which involves a wide-range of changes in physiology and morphology.

However, in contrast to reproduction and germination, almost nothing is known about

plasticity in the timing of vegetative phase change (VPC) or how it may impact the

degree of phenotypic plasticity during vegetative development.

VPC is regulated by a highly conserved microRNA, miR156, and its targets, the

Squamosa Promoter Binding-Like (SPL) transcription factors (Wu & Poethig, 2006;

Willmann & Poethig, 2007; Wu *et al.*, 2009; He *et al.*, 2018). As individuals transition

from a seedling to adult, changes in expression of the miR156/SPL module lead to

phase specific differences in leaf morphology, photosynthetic traits, growth strategies

and reproductive competence (Poethig, 1990; Bassiri *et al.*, 1992; Bongard-Pierce *et al.*,

1996; Telfer *et al.*, 1997; Wang *et al.*, 2011; Feng *et al.*, 2016; Leichty & Poethig, 2019;

Silva *et al.*, 2019; Lawrence *et al.*, 2020, 2021b,a). Specifically, as plants transition from

juvenile to adult, leaves become larger with decreased specific leaf area (SLA), and

often display increased photosynthetic rates per unit leaf area. These changes in leaf

morphology and physiology lead to a switch from a fast- to slow-growth strategy as

plants transition from producing low-cost juvenile leaves to expensive adult leaves with

long lifespans. Because VPC alters how plants function, the types of structures

produced, and the plant's ability to transition to reproduction, the timing of this transition

is likely to have significant consequences for fitness. Further, the model presented in

Fischer *et al.*, (2014) suggests that age-dependent changes in plasticity that align with

the timing of VPC are beneficial for fitness. An increase in plasticity during the adult

vegetative phase would provide time during the juvenile phase to collect enough

information about the environment before investing in any costly phenotypic adjustment

during the adult phase, and is therefore predicted to be an optimal strategy.

The timing of VPC is responsive to environmental cues, specifically light, and

defoliation (Yang *et al.*, 2011; Leichty & Poethig, 2019; Xu *et al.*, 2021). While the exact

effects on the timing of VPC are unknown, expression of the miR156/SPL module is

also altered by temperature, and drought (Kong *et al.*, 2010; Lee *et al.*, 2010; May *et al.*,

2013; Arshad *et al.*, 2018). Furthermore, the observation that juvenile and adult phases

of alfalfa exhibit different tolerance to drought (Arshad *et al.*, 2017) suggests genotypic

or species differences in the timing of this transition could be adaptive. However, the

validity of this hypothesis is difficult to assess because the range of variation in the

timing of VPC across species or even multiple genotypes within a species has only

been investigated in a few cases.

Intimate relationships between development and plasticity lead us to examine

variation in the timing of VPC and explore the effects of this transition on plasticity of

vegetative traits. In this study we used ten natural accessions of *A. thaliana* to

investigate how much genetic variation and plasticity exists in the timing of VPC

(plasticity in developmental timing, Fig. **1**) as well as how VPC impacts leaf trait

plasticity (age/phase-dependent plasticity, Fig. **1**)*.* We also examined eight transgenic

lines with varying expression levels of the miR156/SPL module, both to confirm the role

of this developmental transition in altering juvenile and adult trait plasticity and to further

investigate how alterations in the timing of VPC impact plant growth under varying

environments, independent of genetic variation at other loci.

Materials and Methods

Plant growth and materials

Ten natural accessions of *Arabidopsis thaliana* (Table **S1**)—selected based on a

preliminary screen for variation in the timing of VPC—were obtained from the

Arabidopsis Biological Resource Center (Ohio State University, Columbus, OH, USA).

All mutant lines used in this study were in a Col genetic background. The *mir156a-2*,

mir156c-1, *mir157c-1*, *35S::MIM156*, *35S::MIR156a*, and *spl9-4 spl13-1* lines have been

previously described (Wu & Poethig, 2006; Franco-Zorrilla *et al.*, 2007; Yang *et al.*,

2013; Xu *et al.*, 2016; He *et al.*, 2018). The *MIR156A* genomic line developed in the

Poethig lab is homozygous for a T-DNA insertion containing a 5.5 kb fragment spanning

the region between the genes upstream and downstream of *MIR156A* in a

pCAMBIA3301 backbone. These mutant and transgenic lines allowed us to evaluate

developmentally juvenile and adult leaves at every leaf position within a single genetic

background, thus distinguishing the effects of VPC from other factors such as length of

exposure to the environment, and size or age of the plant, which could influence

phenotypic plasticity.

Seeds were planted in 96 well flats in Fafard-2 growing mix supplemented with

Peters 20-10-20 fertilizer. Beneficial nematodes (*Steinernema feltiae*, BioLogic, Willow

Hill, PA), Marathon 1% granular insecticide and diatomaceous earth were added to the

growing mix to control insects. Plants were placed in 4°C for 5 days before being grown

in their respective treatment conditions (Table **S2**) in Percival growth chambers. Short

day (10 hrs. light/14 hrs. dark, at 120 or 90 µmol m^{-2} s^{-1}) light conditions were used to

prevent flowering so adult vegetative phenotypes were not impacted by the reproductive

transition. Plants were grown under 13.5W full spectrum LED lights. A reduction in light

intensity for the low light treatment was achieved by increasing the distance between

the lights and plants. Plants in all treatments were watered every other day with either

700 mL per flat until seeds germinated and throughout the 28-day growth period for

control, heat and low light treatments or 350 mL per flat for drought treatment. Soil

moisture was recorded with a Delta T HH2 Moisture Meter (Meter Group, Pullman, WA,

USA) at a depth of approximately 1.5 cm twice a week both before and after watering at

a minimum of two locations within the growth chamber. Temperature, light level and

average soil moisture conditions for each treatment are reported in Table **S2**

Plant Phenotyping

Plants were harvested at the soil surface 28 days following transfer to growth chambers.

Leaves were removed from the shoot and placed flat between two transparency sheets

in the order they were initiated (leaf one is the first leaf to be produced) and scanned

using a flatbed scanner (CanoScan LiDE220) at 300 dpi. All shoot material was then

transferred to coin envelopes and dried at 60°C until constant mass.

Whole plant growth phenotypes were measured to determine the influence of the

timing of VPC on plant productivity in the tested environments. Whole shoot mass from

dried plants was recorded and shoot area was analyzed from scanned images using

FIJI (Fiji software). The number of leaves initiated was counted as the total number of

fully expanded leaves and leaf primordia present at the end of the 28-day growing

period. This value was used to calculate the average leaf initiation rate across the

growing period by dividing the total number of leaves initiated by 28 days.

Leaf morphological traits were analyzed from scanned images using FIJI.

Specifically, for leaf morphological measures, base leaf angle was measured as the

angle between the edges of the leaf on either side of the petiole. Petioles were then

removed and the wand tool was used to select each leaf individually from binary images

to measure area, perimeter, length, width and circularity and the presence or absence

of serrations was noted. These traits describe both leaf shape and size to indicate

phenotypic change across development and environment while also being reliably and

easily measured from leaf scans.

Determination of Developmental Phase

Vegetative phase change in *Arabidopsis thaliana* is marked by changes in leaf

morphology that include leaf base angle, length/width ratio, circularity, the presence and

absence of serrations, and the presence or absence of abaxial trichomes (Telfer *et al.*,

1997; Tsukaya *et al.*, 2000; He *et al.*, 2018). While the presence of abaxial trichomes is

the most commonly used marker for the onset of the adult phase (Yang *et al.*, 2013; Xu

et al., 2016; He *et al.*, 2018), we chose to use a combination of leaf morphological traits

more easily measured from leaf scans to determine juvenile and adult phases. This

method allows us to avoid the influence of any phase change-independent effects on

trichome development (i.e. photoperiodic and UV-B effects on trichome regulators

gibberelic acid biosynthesis and GLABRA3, respectively) that could arise in plants

grown under different environments (Chien & Sussex, 1996; Olszewski *et al.*, 2002; Yan

et al., 2012). Further, by using a combination of traits we can get a more holistic

representation of when vegetative phase change occurs, as the onset of different adult

traits are often not perfectly synchronized.

Leaf stage was determined from leaf scans using a principal component analysis

of summarizing phase specific traits (leaf base angle, the length:width ratio of the

lamina, the presence vs. absence of serrations, and the circularity of the lamina), and

comparing the PC1 value obtained by this approach to the PC1 value for plants over-

expressing miR156 (35S::MIR156a, "OX"), in which all leaves are juvenile (Fig. **S1**).

Leaves with a PC1 value greater than or equal to the value for the miR156-over

expressing line were considered juvenile leaves, and leaves with a PC1 value less than

this number were considered adult leaves. Specifically, the minimum PC1 value of all

OX leaf samples for each treatment was identified. All leaves with PC1 values below

this threshold were considered adult leaves. PC1 threshold values used here were 0 for

control, 0.3 for drought, -2 for heat and 0 for low light conditions. Setting independent

thresholds for each environment using leaves produced across all leaf positions

eliminates the influence of any non-phase-change specific changes in PC1 values (i.e.

plant size, age, or environment) in the determination of juvenile and adult leaves.

Data Analysis and Statistics

All statistical analyses were performed in JMP® Pro v. 15.0.0 (SAS Institute Inc., Cary,

NC). Differences in the time of transition among genotypes were compared by two-way

ANOVA where genotype and environmental treatment were the main effects. Because

treatment had a significant effect ($p < 0.05$) Tukey's HSD was used to determine which

treatments were significantly different from each other. Within each genotype, ANOVA

was used to determine if environmental treatment had a significant effect on the time of

transition measures, total leaves initiated and leaf initiation rate. For genotypes where

treatment had a significant effect ($p < 0.05$), Tukey's HSD was used to determine which

treatments were significantly different from each other. We used one-way ANOVA with

natural accession genotypes as the main effect to estimate broad-sense heritability (R^2

from the ANOVA).

To understand how the timing of VPC impacts plant growth and productivity

across environments, we looked at relationships between the time of transition and

whole plant growth phenotypes of shoot mass, shoot area, and leaves initiated in each

of the four environments. Significant effects of the timing of VPC on these traits was

determined using least square linear regression analysis for both accessions and

mutant genotypes. Further, we investigated how the timing of VPC impacts plasticity in

these traits using phenotypic plasticity index and the change in each trait between

control and each of the three environmental stress treatments. Phenotypic plasticity

index was calculated as the absolute difference between the maximum and minimum

mean values among all four growth conditions divided by the maximum mean value,

following Valladares *et al.*, (2000) and Valladares *et al.*, (2006). While the plasticity

index describes the overall plasticity in a trait across all environments, the signed

change in trait describes plasticity induced by a specific environment and was

calculated as the difference between the mean control value and the mean treatment

value. Linear regression was used to evaluate whether there was a significant impact of

the timing of VPC on plasticity in plant growth and productivity traits using the mean

number of juvenile leaves produced or number of days initiating juvenile leaves for each

genotype.

To determine if vegetative phase change contributes to age-dependent changes

in phenotypic plasticity, we tested whether developmental phase (i.e. juvenile or adult)

impacts plasticity of leaf morphology using phenotypic plasticity index and change in

trait described above. A student's T-test was used to evaluate whether differences

between juvenile and adult plasticity was significant. Tests were conducted between

phenotypic plasticity index for juvenile and adult leaves of all accessions or all leaf

positions 1-8 for genotypes in the Col-0 background (i.e. miR156/SPL mutants and Col-

0 accession). For evaluating changes in trait data, tests compared the change in trait

between control and treatment phenotypes of juvenile and adult leaves for all

accessions or all leaf positions 1-8 for genotypes in the Col-0 background for each of

the three treatment growth conditions.

To briefly investigate whether there was any relationship between plasticity in the

timing of vegetative phase change and an accession's climate of origin, we used the

geoclimatic variables from the WorldClim2 data set (Fick & Hijmans, 2017), associated

with georeferences for each of the 10 accessions published as part of the 1001

genomes project (Alonso-Blanco *et al.*, 2016). We used linear regression to determine

any significant relationships between the bioclimatic variables and the phenotypic

plasticity index for the time of transition calculated as described above using the number

of juvenile leaves.

Results

The timing of vegetative phase change varies among genotypes and environments

Leaf identity was determined using a principal component analysis of phase

specific leaf morphology traits and the timing of VPC was measured in two ways; by the

number of juvenile leaves produced and the number of days the plant was initiating

juvenile leaves calculated using leaf initiation rate. Both juvenile leaf number and days

initiating juvenile leaves significantly differed among genotypes and was altered by

abiotic environment (ANOVA: Genotype $p < 0.0001$, Treatment $p < 0.0001$, G x T $p <$

0.0001) (Table 1). Differences in the amount of time plants spent in the juvenile phase

were largely due to the effect of environmental conditions on the rate of leaf production,

as evident from the differences in the total number of leaves produced by 28 days under

these conditions. Among the accessions examined here, VPC occurred as early as leaf

3 or day 6, and as late as leaf 8 or day 18 in "control" conditions. Transgenic genotypes

with altered miR156 or *SPL* gene expression in the Col-0 background had VPC that

occurred from leaf 1 or day 1 to leaf 25 or the full 28-day period of the experiment (i.e.

these plants never transitioned to adult phase). Broad-sense heritability of the timing of
VPC was high, ranging between 0.82 in control and 0.64 in low light environments when
measured by the number of juvenile leaves and 0.61 in drought and 0.46 in control
environments when measured by days initiating juvenile leaves (Table S3).

Both measures of the timing of VPC were significantly altered by abiotic
environment, and this response differed among genotypes as indicated by a significant
($p < 0.05$) ANOVA interaction term (genotype x treatment). Among the accessions, heat
increased the number of juvenile leaves by an average of 1.3 from control and
decreased the rate of leaf initiation by 38%, thus increasing the average number of days
plants were in the juvenile phase by 11.15 ($p < 0.05$ Tukey's HSD). Low light intensity
decreased the average number of juvenile leaves, but decreased the rate of leaf
initiation by 49%, which was sufficient to increase the number of days plants were in the
juvenile phase by 6.39. Drought caused no significant difference in the number of
juvenile leaves or the rate of leaf initiation for most genotypes, and thus had no effect on
the days plants were in the juvenile phase (Table 1). All three environmental treatments
subjected plants to some degree of stress as indicated by losses of biomass averaging
8.4, 7.2 and 3 mg in low light, heat and drought respectively. It should be noted that it is
unclear why we observed a decrease in the number of juvenile leaves under low light
conditions, given that previous studies have shown that low light intensity increases
juvenile leaf number (Leichty & Poethig, 2019; Xu *et al.*, 2021).

Plasticity in the number of juvenile leaves and the rate of leaf initiation varied
among genotypes in response to abiotic environments, causing variation in the amount
of time plants spent in the juvenile phase. This led to variation in the order genotypes
transitioned to the adult phase between environments (Fig. **S2**). For example, Strand
(ST) is the 10th genotype to transition in control, 7th in drought, 17th in heat and 13th in
low light when measured by days initiating juvenile leaves. In the MIM156 target
mimicry line (MI) and 35S:*MIR156a* (OX) mutants, where miR156 abundance is highly
constrained (functionally non-existent or in excess respectively), there was no plasticity
in the days spent in the juvenile phase, although the rate of leaf initiation was
significantly affected by heat and light intensity in these genotypes. MI plants produce
only adult leaves and OX plants produce only juvenile leaves in all environments (Table

1), indicating that variation in miR156 abundance is necessary for the days spent in the

juvenile phase to be altered in response to the tested environments.

Plasticity in the number of juvenile leaves in the natural accessions across all

four environments, measured by phenotypic plasticity index, was significantly related to

the temperature of the driest quarter (R^2 = 0.493, $p < 0.05$) and precipitation of the

warmest quarter (R^2=0.502, $p < 0.05$) for each accession's climate-of-origin (Fig. **S3**).

Specifically, increased plasticity in the timing of VPC was related to higher temperatures

during the driest quarter and lower precipitation during the warmest quarter.

The timing of vegetative phase change correlates with plant productivity and

performance among miR156/SPL mutants in the Col-0 background but not across

natural accessions.

To understand how changes in the timing of VPC impacts plant productivity and

performance in different environments, we explored the relationships between the

number of juvenile leaves or days spent in the juvenile phase and the whole plant

growth phenotypes of shoot mass, shoot area, and number of leaves initiated during a

28-day period. Among miR156/SPL mutant genotypes, later phase change (i.e. an

increase in the number of juvenile leaves or longer time spent initiating juvenile leaves)

was significantly associated with increases in all three whole plant phenotypes across

most environments (the sole exception being total shoot area vs. days initiating juvenile

leaves in heat) (Fig. **2, S4**). The effect of these genotypes on the total number of leaves

produced is consistent with previous studies showing that variation in the expression of

the mir156/SPL module has a significant effect on the rate of leaf initiation (Schwab *et*

al., 2005; Wang *et al.*, 2008; Feng *et al.*, 2016; Lawrence *et al.*, 2021a).

By contrast, natural genetic variation in the number of juvenile leaves in

accessions showed mostly non-significant ($p > 0.05$) or weak ($R^2 < 0.1$) relationships

with whole plant phenotypes within each environment (Fig. **2**). An exception to this was

that a more rapid rate of leaf initiation was associated with a larger number of juvenile

leaves in all environments, similar to what was observed in the mutants (Fig. **2**).

Interestingly, some growth traits had contrasting relationships with the number of days

plants were initiating juvenile leaves in natural accessions compared to mutant

genotypes. Specifically, shoot mass, and shoot area in control and heat environments,

and leaves initiated in the control treatment, were negatively related to the time spent in

the juvenile phase in natural accessions (Fig. **S4**). That is, natural accessions that

transitioned earlier had greater mass and leaf area in control and heat growth

treatments while mutants that transitioned later were bigger. This suggests natural

variation in the rate of leaf initiation (the major driver of time spent in the juvenile phase)

could be regulated by more than just miR156 abundance or that the impacts of VPC on

growth rate vary across genetic backgrounds. However more data are needed to

understand why the relationships between the timing of VPC and plant growth differ

when the number of juvenile leaves or time spent in the juvenile phase it is altered

within a single genetic background (i.e. the isogenic mutants) compared to natural

accessions. For example, if greater miR156 abundance leads to increases in vegetative

growth (as suggested by our data and other studies (Fu *et al.*, 2012; Wang & Wang,

2015; Zheng *et al.*, 2016)) but natural variation in the threshold needed to suppress *SPL*

expression exists, then genotypes with comparable levels of miR156 could undergo

VPC at different times.

The adult vegetative phase has greater phenotypic plasticity than the juvenile phase

Plasticity of leaf morphology was greater across the four growth environments for adult

leaves than for juvenile leaves. This greater plasticity was true for both accessions and

for miR156/SPL mutants (Fig. **3a,b**). Specifically, adult leaves had significantly greater

plasticity in leaf base angle, area, perimeter, length/width ratio, and circularity and the

average of those traits regardless of leaf position in the Col-0 lines, and significantly

greater plasticity in leaf base angle, length/width ratio, and circularity and the average of

all measured traits in the accessions (Student's T, $p < 0.05$)(Fig. **3a,b**). The consistency

with which adult leaves display greater plasticity across genotypes with different

numbers of juvenile leaves (i.e. independent of both exposure time and resources,

which has also been associated with increased phenotypic plasticity (Kleunen &

Fischer, 2005; Fischer *et al.*, 2014)) confirms the effect of this developmental program

on age-dependent changes in plasticity. The greater morphological plasticity of adult

leaves compared to juvenile leaves can be visualized for all accessions in Figure **4**.

Changes in leaf morphological traits between leaves grown in control and

treatment conditions was greater for adult leaves than juvenile leaves in response to

heat and low light (Fig. **3c,d**). Across accessions, significantly greater changes in leaf

area, perimeter and L/W ratio were observed in adult leaves grown in heat and low light

conditions, and in base angle and circularity for leaves grown in heat (Student's T, $p <$

0.05). Changes between leaves at positions 1-8 in the Col-0 lines were significantly

greater among adult leaves in L/W ratio and circularity in response to both heat and low

light, in base angle in response to heat, and in area and perimeter in response to low

light (Student's T, $p < 0.05$). Interestingly, the drought treatment—which caused no

overall change in the number of juvenile leaves or the amount of time spent in the

juvenile phase, and was the least stressful of the three treatments as indicated by

biomass loss—lead to no significant differences in the change of morphological traits

between juvenile and adult leaves (i.e. juvenile and adult leaf morphology responded

similarly to drought). Because miR156 expression has previously been shown to

respond to drought (Katiyar *et al.*, 2015; Liu *et al.*, 2019), it is possible that plasticity in

VPC would occur in a more severe drought treatment.

Plasticity of whole plant growth phenotypes is not related to the timing of vegetative

phase change

Despite differences in plasticity between the juvenile and adult phases of development,

the number of juvenile leaves did not impact plasticity of whole plant growth phenotypes

(shoot area, shoot mass or leaves initiated) in the accessions across any of the tested

environments (Fig. **5**). However, in miR156/SPL mutants, we found that an increase in

the number of juvenile leaves resulted in a greater reduction in these phenotypes when

grown in heat and low light conditions ($p < 0.05$) (Fig. **5b**). Even within the mutant lines,

overall plasticity measured as the phenotypic plasticity index considering all four growth

conditions was not related to the number of juvenile leaves (Fig. **5a**).

Discussion

Morphological and physiological plasticity and the timing of development are key

ways that organisms cope with fluctuating and unpredictable environments. Plasticity is

an important element of development, both as it pertains to the timing of transitions as

well as its role in dictating the degree of plasticity of an organism. Vegetative phase

change (VPC) is a highly conserved developmental transition in plants, but the role of

VPC in environmental adaptation is not known. Our study shows how the timing of VPC

responds to environmental factors, and demonstrates that the transition alters the

degree of phenotypic plasticity in response to environment. The existence of natural

genetic variation in VPC and associated plasticity suggests the potential for selection,

though the VPC-environment-fitness map remains to be dissected.

We found diverse natural accessions consistently showed increases in plasticity

with vegetative phase change, that is, the juvenile phase was more robust than the

adult phase. This consistency suggests this change in plasticity is adaptive. This

increase in plasticity might be favored if it is advantageous to delay high levels of

plasticity until after a plant has accumulated a significant amount of information about

the environment, but while it still has sufficient time to benefit from a phenotypic

adjustment (Fischer *et al.*, 2014). Deviations from this pattern, such as in the OX line

which remains in the robust juvenile phase, suggests that there are fitness costs to this

strategy, as these OX plants had the greatest reductions in mass, area and leaf

initiation in response to environmental stress (Fig **5b**). In addition, Fischer *et al.*, (2014)

predicted the initial more robust phenotype would improve fitness if adapted to the most

likely environment encountered during that lifestage. Previous studies show the

morphology and physiology of the juvenile phase is well suited to the low light

environments often encountered by juvenile leaves emerging in a shaded understory

and quickly overtopped by newly initiated leaves (Lawrence *et al.*, 2020, 2021b; Xu *et*

al., 2021), providing further support for the hypothesis that VPC modulates phenotypes

in an adaptive way.

Changes in phenotypic plasticity modulated by VPC were consistent regardless

of genetic backgrounds or when the transition occurs. Adult leaves had greater plasticity

whether they were produced at node one (i.e. in the MI line) or node ten. This confirms

that VPC, and even more so, expression levels of miR156, is responsible for these

changes in plasticity, rather than simply the length of exposure to environmental cues.

High levels of miR156, which are associated with a more robust juvenile phase, likely

represent an example of microRNA buffering. MicroRNAs buffer by silencing changes in

target gene expression (i.e. in response to environmental cues) when target genes are

below a certain threshold. Once transcription of the target genes exceeds this threshold,

protein output becomes sensitive to any transcriptional changes (Posadas & Carthew,

2014). There are numerous examples of microRNAs contributing to developmental

robustness, such as miR164, which buffers its targets and causes robust plant organ

development (Sieber *et al.*, 2007). The abundance of miR156 early in the juvenile phase

far exceeds the level necessary to repress SPL genes, and is therefore likely to

contribute to the more robust phenotypes through this buffering mechanism (He *et al.*,

2018). Early studies of VPC noted stable phenotypes that were unresponsive to

environmental and hormonal treatments in the first two leaves of *Arabidopsis* plants,

where miR156 abundance is highest (Telfer *et al.*, 1997). An example of miR156

buffering occurs through the microRNA's role in plant responses to the stress-related

phytohormone abscisic acid (ABA). High miR156 levels, as found during the juvenile

phase, suppress plant ABA responses through repression of its target SPL genes,

which when expressed during the adult phase, interact directly with the transcription

factor ABA-INSENSITIVE5 (ABI5) to facilitate ABA signaling (Dong *et al.*, 2021).

Because phenotypic plasticity increases between juvenile and adult phases

across genotypes, genetic and environmental variation in the timing of VPC determines

when increased plasticity is acquired. We found natural variation in the timing of this

transition and significant genotype x environment interactions, indicating the time of

VPC has the potential to respond to selection. Determining the timing of VPC is not as

straightforward as other developmental transitions because it is not marked by any

single morphological change, but by multiple traits that do not always appear

simultaneously. Because each phase-specific trait can be differentially sensitive to

environmental cues, using combinations of traits (as we did) is likely necessary when

multiple environments are tested. Despite this increased complexity, understanding how

the timing of VPC responds to environments could provide important insights into plant

adaptation and acclimation. Here, we add further evidence that VPC impacts plant

growth and productivity as increased levels of miR156 leads to greater vegetative

growth across environments, consistent with observations in other species (Fu *et al.*,

2012; Rubinelli *et al.*, 2013; Wang & Wang, 2015; Zheng *et al.*, 2016), while bringing

forth new questions about whether these relationships remain true among plants of

different genotypes (Fig **2**). Within the few *Arabidopsis* accessions examined here, we

found significant relationships between plasticity in the timing of VPC and multiple

climate-of- origin variables (Fig. **S3**), suggesting further exploration of population

differences in VPC. Further, trait plasticity can influence a plant's vulnerability to

changing climates. For example, plasticity in the leaf traits of 12 perennial plant species

was an important determinant of susceptibility to climate change scenarios in a

Mongolian steppe (Liancourt *et al.*, 2015). Subsequently, climate can alter selection on

plasticity. For instance, end-of-season drought conditions selected for increased

plasticity of water use efficiency in *A. thaliana* (Kenney *et al.*, 2014) whereas plasticity of

flowering time in response to temperature was maladaptive (Stinchcombe *et al.*, 2004).

Modulating age-dependent changes in plasticity might contribute to any role VPC has in

adaptation. Additional studies are needed to understand what, if any, type of selection is

on VPC and how the timing of vegetative phase change contributes to adaptation and

acclimation in plants.

The effects of VPC on plasticity found here in *A. thaliana* are likely to be

conserved across species. VPC is regulated by miR156 and its sister microRNAs

across all land plants, allowing for microRNA buffering to create a consistently more

robust juvenile phase. For example, across 40 species of *Passiflora,* juvenile leaf

morphology is conserved despite the highly variable morphology among adult leaves

(Chitwood & Otoni, 2017), similar to what we observed here between the various

genotypes (Fig. **3, 4**). This further suggests the age-dependent changes in plasticity due

to VPC could lead to varying degrees of both intra and interspecific trait variation

through time among populations. It seems likely many published studies capture

patterns of variation driven by VPC within and among species, however are unaware of

this transition (i.e. (Mason *et al.*, 2013; Garbowski *et al.*, 2021). Intraspecific variation

can play an important role in community- and ecosystem-level processes (Bolnick *et al.*,

2011; Madritch *et al.*, 2014; Turner *et al.*, 2020; Westerband *et al.*, 2021), though the

functional basis of this variation is usually not understood. To better understand the

implications of VPC on these large scale processes, further characterization of the

timing of VPC across plant species (of which there is currently few; Maize, (Poethig,

1988; Bongard-Pierce *et al.*, 1996); *Arabidopsis thaliana*, (Telfer *et al.*, 1997);

Eucalyptus globulus, (Jordan *et al.*, 1999); *Nicotiana tabacum*, (Feng *et al.*, 2016);

Sorghum, (Hashimoto *et al.*, 2019); Acacia, (Leichty & Poethig, 2019); *Passiflora edulis*,

(Silva *et al.*, 2019); Poplar, (Lawrence *et al.*, 2021a) and Rice, (Asai *et al.*, 2002)) and

variation among genotypes within species (even fewer; *Eucalyptus globulus*, (Jordan *et*

al., 1999); Maize, (Poethig, 1988; Foerster *et al.*, 2015); and *A. thaliana*, (Doody *et al.*,

2021)) is necessary.

VPC may contribute to plant adaptation by setting up periods of low and high

phenotypic plasticity when it is most beneficial. Here we demonstrate that the timing of

vegetative phase change varies across genotypes, interacts with the environment, and

alters the plasticity of vegetative traits. While more work is needed to fully understand

the functional, ecological, and evolutionary significance of VPC, its determination as a

modulator of age-dependent changes in plasticity provides new insights for

understanding plant environmental interactions.

Acknowledgments

We thank Erin Doody for providing preliminary data regarding the timing of vegetative

phase change in *Arabidopsis thaliana* accessions. This research was funded by NSF

grant DGE-1845298 awarded to EHL, NSF DEB-1927009 and NIH 1R35GM138300-01

grants awarded to JRL, and NIH GM51893 awarded to RSP.

Author Contributions

EHL, RSP and JRL contributed to research planning and design, EHL performed the

experiments and statistical analysis, EHL wrote the initial version of the manuscript and

all authors revised and provided comments.

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Tables

Table 1. Timing of vegetative phase change, measured as number of juvenile leaves produced or the number of days initiating juvenile leaves, and leaf initiation traits during 28-days of growth for natural accessions and miR156/SPL mutants grown in different environments. Values represent the mean ± standard error. Different lowercase letters indicate treatment groups that are significantly different ($p < 0.05$) based on Tukey's HSD, following a significant ANOVA result ($p < 0.05$). Groups significantly different from control are bolded. N represents the number of plants sampled.

Natural Accessions

Genotype	Treatment	Number of Juvenile Leaves	Days Initiating Juvenile Leaves	Total Leaves Initiated	Leaf Initiation Rate	N
SH	Control	2 ± 0.45 ab	5.6 ± 1.39 b	10.8 ± 0.7 a	0.39 ± 0.03 a	6
	Drought	1 ± 0.52 b	2.4 ± 1.23 b	11.2 ± 0.48 a	0.4 ± 0.02 a	6
	Heat	3.6 ± 0.29 a	**15.8 ± 1.56 a**	**6.4 ± 0.29 b**	**0.23 ± 0.01 b**	9
	Low Light	1 ± 0.63 b	5.6 ± 3.54 b	**5.4 ± 0.24 b**	**0.19 ± 0.01 b**	5
AT	Control	3.3 ± 0.31 ab	8.4 ± 1.5 b	12.1 ± 1.04 a	0.43 ± 0.04 a	8
	Drought	2.5 ± 0.34 b	5 ± 0.73 b	14 ± 0.37 a	0.5 ± 0.01 a	6
	Heat	4.1 ± 0.13 a	**16.6 ± 1.81 a**	**7.4 ± 0.6 b**	**0.26 ± 0.02 b**	8
	Low Light	2.4 ± 0.34 b	10 ± 1.46 b	**6.8 ± 0.27 b**	**0.24 ± 0.01 b**	12
BC	Control	4.1 ± 0.35 b	7.9 ± 0.85 b	14.9 ± 0.58 a	0.53 ± 0.02 a	8
	Drought	3.4 ± 0.6 b	7.1 ± 1.22 b	13.6 ± 1.03 a	0.49 ± 0.04 a	5
	Heat	**5.8 ± 0.36 a**	**17.6 ± 1.42 a**	**9.6 ± 0.73 b**	**0.34 ± 0.03 b**	10
	Low Light	2.6 ± 0.53 b	11.2 ± 2.13 b	**6.3 ± 0.29 c**	**0.22 ± 0.01 c**	7
ST	Control	4.4 ± 0.37 b	11.4 ± 1.74 b	11.6 ± 0.81 a	0.41 ± 0.03 a	7
	Drought	3.8 ± 0.17 b	8.7 ± 0.46 b	12.3 ± 0.21 a	0.44 ± 0.01 a	6
	Heat	**6.4 ± 0.5 a**	**28 ± 0 a**	**6.4 ± 0.5 b**	**0.23 ± 0.02 b**	8
	Low Light	4.5 ± 0.29 b	**22.7 ± 3.12 a**	**5.8 ± 0.48 b**	**0.21 ± 0.02 b**	4
TU	Control	5.6 ± 0.38 ab	11.2 ± 1.19 c	14.5 ± 0.71 a	0.52 ± 0.03 a	8
	Drought	6 ± 0.52 ab	13.9 ± 2.91 bc	13.2 ± 1.14 a	0.47 ± 0.04 a	6
	Heat	6.2 ± 0.22 a	**22.4 ± 1.52 a**	**8.1 ± 0.65 b**	**0.29 ± 0.02 b**	9
	Low Light	4.8 ± 0.36 b	**19.4 ± 1.47 ab**	**6.9 ± 0.29 b**	**0.25 ± 0.01 b**	13
CO	Control	5.9 ± 0.14 a	11.5 ± 0.22 b	14.3 ± 0.47 a	0.51 ± 0.02 a	7
	Drought	5 ± 0.32 ab	10.2 ± 0.72 b	13.8 ± 0.8 a	0.49 ± 0.03 a	5
	Heat	5.5 ± 0.36 a	**18.1 ± 1.37 a**	**8.9 ± 0.71 b**	**0.32 ± 0.03 b**	12
	Low Light	**4.1 ± 0.15 b**	**16.2 ± 1.02 a**	**7.2 ± 0.28 b**	**0.26 ± 0.01 b**	15
VE	Control	6 ± 0 b	13.6 ± 1.08 b	13 ± 1.18 a	0.46 ± 0.04 a	8
	Drought	5.8 ± 0.48 b	12.6 ± 1.12 b	13.2 ± 0.91 a	0.47 ± 0.03 a	6
	Heat	**9 ± 0.8 a**	**25.7 ± 1.18 a**	10 ± 1.07 ab	0.36 ± 0.04 ab	8
	Low Light	5.6 ± 0.69 b	**23.2 ± 2.43 a**	**6.7 ± 0.36 b**	**0.24 ± 0.01 b**	7
CS	Control	6.3 ± 0.16 ab	12.3 ± 1.08 b	14.9 ± 1.01 a	0.53 ± 0.04 a	8
	Drought	5.6 ± 0.24 bc	12.6 ± 0.95 b	12.8 ± 1.24 a	0.46 ± 0.04 a	5
	Heat	7.4 ± 0.46 a	**24.3 ± 1.76 a**	**8.8 ± 0.62 b**	**0.31 ± 0.02 b**	8
	Low Light	**4.3 ± 0.42 c**	**23.2 ± 2.18 a**	**5.5 ± 0.72 b**	**0.2 ± 0.03 b**	6
VO	Control	6.7 ± 0.21	17 ± 2.15 b	11.7 ± 1.12 b	0.42 ± 0.04 b	6
	Drought	6.2 ± 0.17	12.1 ± 1.02 b	**14.7 ± 0.95 a**	**0.52 ± 0.03 a**	6
	Heat	7.3 ± 0.41	**25.1 ± 1.05 a**	**8.4 ± 0.68 c**	**0.3 ± 0.02 c**	11
	Low Light	6.1 ± 0.34	**23.4 ± 1.22 a**	**7.4 ± 0.22 c**	**0.26 ± 0.01 c**	16
BO	Control	7.3 ± 0.18 b	12.3 ± 0.69 c	16.1 ± 1.08 a	0.58 ± 0.04 a	8
	Drought	7 ± 0.37 b	11 ± 0.71 c	18 ± 0.73 a	0.64 ± 0.03 a	6
	Heat	**8.6 ± 0.29 a**	**27.3 ± 0.43 a**	**8.9 ± 0.32 b**	**0.32 ± 0.01 b**	15
	Low Light	6.3 ± 0.25 b	**16.8 ± 1.02 b**	**10.5 ± 0.65 b**	**0.38 ± 0.02 b**	4

Mutants

Genotype	Treatment	Number of Juvenile Leaves	Days Initiating Juvenile Leaves	Total Leaves Initiated	Leaf Initiation Rate	N
MI	Control	0 ± 0	0 ± 0	6 ± 1 a	0.21 ± 0.04 a	4
	Drought	0 ± 0	0 ± 0	4.9 ± 0.48 a	0.17 ± 0.02 a	8
	Heat	0 ± 0	0 ± 0	**2.3 ± 0.31 b**	**0.08 ± 0.01 b**	8
	Low Light	0 ± 0	0 ± 0	**2.1 ± 0.11 b**	**0.08 ± 0 b**	9
ACC	Control	1.4 ± 0.38 ab	5.7 ± 1.65 ab	7.3 ± 0.45 a	0.26 ± 0.02 a	8
	Drought	1.8 ± 0.56 a	7.3 ± 2.62 ab	7.3 ± 0.37 a	0.26 ± 0.01 a	8
	Heat	2.2 ± 0.15 a	10.5 ± 0.96 a	6.1 ± 0.31 a	0.22 ± 0.01 a	14
	Low Light	0.5 ± 0.26 b	3.3 ± 1.72 b	**4.8 ± 0.22 b**	**0.17 ± 0.01 b**	12
SAC	Control	2.3 ± 0.16 ab	7 ± 0.55 b	9.1 ± 0.55 ab	0.33 ± 0.02 ab	8
	Drought	2.3 ± 0.33 ab	6.2 ± 1.25 b	11 ± 0.58 a	0.39 ± 0.02 a	6
	Heat	3.5 ± 0.65 a	**16.7 ± 5.28 a**	7 ± 1.29 bc	0.25 ± 0.05 bc	4
	Low Light	1.8 ± 0.17 b	10.7 ± 1.12 ab	**4.8 ± 0.17 c**	**0.17 ± 0.01 c**	6
M6C	Control	4 ± 0.32 a	9.4 ± 1.2	12.4 ± 1.21 a	0.44 ± 0.04 a	5
	Drought	3.9 ± 0.35 a	9.6 ± 1.11	11.5 ± 0.57 ab	0.41 ± 0.02 a	8
	Heat	4.6 ± 0.29 a	13.4 ± 1.16	**9.8 ± 0.55 b**	**0.35 ± 0.02 b**	9
	Low Light	**2.2 ± 0.24 b**	10.1 ± 1.08	**6 ± 0.21 c**	**0.21 ± 0.01 c**	12
M7C	Control	4.4 ± 0.18 b	10.8 ± 1.14 b	11.9 ± 0.79 a	0.42 ± 0.03 a	8
	Drought	4 ± 0.19 b	9.3 ± 0.82 b	12.4 ± 0.56 a	0.44 ± 0.02 a	8
	Heat	**5.4 ± 0.34 a**	**17.9 ± 1.82 a**	**9 ± 0.79 b**	**0.32 ± 0.03 b**	10
	Low Light	**3.1 ± 0.3 c**	13.6 ± 1.51 ab	**6.6 ± 0.32 b**	**0.24 ± 0.01 b**	8
MA	Control	7 ± 0.37 a	12.7 ± 1.69 b	16.2 ± 1.17 a	0.58 ± 0.04 a	6
	Drought	7.4 ± 0.26 a	15.6 ± 2.02 b	14.3 ± 1.18 a	0.51 ± 0.04 a	8
	Heat	7.3 ± 0.53 a	**22.3 ± 1.53 a**	**9.3 ± 0.62 b**	**0.33 ± 0.02 b**	8
	Low Light	**4.7 ± 0.49 b**	16.7 ± 1.38 ab	**7.8 ± 0.48 b**	**0.28 ± 0.02 b**	6
NT	Control	9.7 ± 0.18 a	15.3 ± 0.61 b	18 ± 0.76 a	0.64 ± 0.03 a	7
	Drought	8.9 ± 0.13 a	16.7 ± 0.93 b	15.3 ± 0.88 a	0.54 ± 0.03 a	8
	Heat	9.2 ± 0.37 a	**23.4 ± 1.92 a**	**11.4 ± 1.21 b**	**0.41 ± 0.04 b**	5
	Low Light	**6.6 ± 0.69 b**	**23.5 ± 2.39 a**	**7.9 ± 0.4 c**	**0.28 ± 0.01 c**	7
OX	Control	24.8 ± 2.04 a	28 ± 0	24.8 ± 2.04 a	0.89 ± 0.07 a	6
	Drought	22.9 ± 1.27 a	28 ± 0	22.9 ± 1.27 a	0.82 ± 0.05 a	8
	Heat	**14.6 ± 2.05 b**	28 ± 0	**14.6 ± 2.05 b**	**0.52 ± 0.07 b**	9
	Low Light	**12.3 ± 0.4 b**	28 ± 0	**12.3 ± 0.4 b**	**0.44 ± 0.01 b**	10

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Figures

Figure 1. Illustration of two sources of plasticity involving vegetative phase change,
phase-dependent changes in plasticity that describe how the developmental phase of
an organ (juvenile or adult) impacts the degree of phenotypic plasticity of that organ,
and plasticity in developmental timing describing how the timing of vegetative phase
change can be accelerated or delayed in response to environment. In the above
example, we find greater plasticity in leaf morphology for adult (red) than juvenile (blue)
leaves between environment 1 and 2 as adult leaves change in both shape and size
whereas juvenile leaves only change in size. The timing of vegetative phase change is
delayed in environment 2 compared to environment 1 as indicated by the longer time
arrow and an increased number of juvenile leaves produced.

Figure 2. Relationships between the timing of vegetative phase change measured as
number of juvenile leaves, and whole plant growth phenotypes of shoot mass, shoot
area, and leaves initiated for accessions (purple) and miR156/SPL mutants (orange) in
all four environments (clockwise from top left: control, drought, heat and low light). R^2
and *p*-values for linear regression of accessions (A) and mutants (M) noted at the top of
each graph.

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Figure 3. Phenotypic plasticity of juvenile and adult leaf morphology. Phenotypic
plasticity index (a, b) and the change in trait between control and treatment conditions
(c, d) for leaf base angle, leaf area, leaf perimeter, length/width (L/W) ratio, and leaf
circularity for adult (red) and juvenile (blue) leaves of each accession (a, c) or leaf
positions 1-8 in plants of the Col-0 genetic background (b, d). Mutations in miR156/SPL
pathway cause both juvenile and adult leaves to be produced at each leaf position.
Significant differences in phenotypic plasticity index between adult and juvenile leaves
across all accessions or leaf positions for each trait denoted by asterisks (* $p < 0.05$, **
$p < 0.01$, *** $p < 0.001$). Significant differences in the change in traits between control
and each treatment between adult and juvenile leaves for all accessions or leaf
positions determined by Student's t denoted by asterisks (* $p < 0.05$, ** $p < 0.01$, *** $p <$
0.001).

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784 **Figure 3.** Representative leaf images of juvenile (left) and adult (right) leaves of each
785 accession from control, drought, heat and low light environments. Leaves are of one of
786 the first two positions of their respective developmental phase. Position of adult leaves
787 shown in parenthesis.

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Figure 4. Phenotypic plasticity in the whole plant growth traits of shoot mass, shoot
area, and leaves initiated. Phenotypic plasticity index (a) and the change in trait
between control and treatment conditions (b) for accessions (left) and miR156/SPL
mutants (right). Bar colors indicate time of vegetative phase transition based on number
of juvenile leaves produced in control conditions. Significant relationships between
number of juvenile leaves and change in trait between treatments were only present
among mutant genotypes and are denoted by asterisks (* $p < 0.05$, ** $p < 0.01$).
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