1   **Environmental oscillations favor the evolution of adaptive transgenerational plasticity**

6   Keywords: Transgenerational plasticity, epigenetics, local adaptation

9   Classification

10   Major: Biological Sciences

11   Minor: Evolution, Environmental Sciences

Abstract

Abstract

Effects of parental environment on offspring traits have been well known for decades. Interest in

this transgenerational form of phenotypic plasticity has recently surged due to advances in our

understanding of its mechanistic basis. Theoretical research has simultaneously advanced by

predicting the environmental conditions that should favor the adaptive evolution of

transgenerational plasticity. Yet whether such conditions actually exist in nature remains largely

unexplored. Here, using long-term climate data, we modeled optimal levels of transgenerational

plasticity for an organism with a one-year life cycle at a spatial resolution of $4km^2$ across the

continental US. Both annual temperature and precipitation levels were often autocorrelated, but the

strength and direction of these autocorrelations varied considerably across the continental US and

even among nearby sites. When present, such environmental autocorrelations render offspring

environments statistically predictable based on the parental environment, a key condition for the

adaptive evolution of transgenerational plasticity. Our optimality models confirmed this prediction:

high levels of transgenerational plasticity were favored at sites with strong environmental

autocorrelations, and little-to-no transgenerational plasticity was favored at sites with weak or non-

existent autocorrelations. These results suggest that transgenerational plasticity is highly variable in

nature, depending on site-specific patterns of environmental variation.

Introduction

The capacity for life to evolve adaptively results in part from patterns in environmental variation. Selection can produce adaptation only if the selective environment is reliably encountered over generations, or in other words, if selective environments are statistically predictable. Early models of evolution envisioned fitness landscapes that were static, such that populations adapt over the course of generations to one or another environment (Fisher 1930). While this form of adaptation optimizes phenotypes for homogenous environments, the more realistic scenario of environmental heterogeneity in both space and time limits the adaptive value of such constitutive genetic expression. Rather, in variable environments, the capacity to modify phenotypes in response to predictive environmental cues allows organisms to adaptively match their traits to the specific patch of habitat in which they find themselves.

This key mode of adaptation to environmental heterogeneity is termed adaptive phenotypic plasticity, a strategy in which individual organisms make functionally appropriate changes to their physiology, development, or behavior in response to specific environmental cues (Ghalambor et al. 2007; Nicotra et al. 2010). Over the last three decades, it has become clear that *transgenerational plasticity* is also remarkably common (reviewed by (Mousseau and Fox 1998; Uller 2008; Bonduriansky and Day 2009; Holeski et al. 2012; Conrath et al. 2015; Sultan 2015), yet highly variable (Herman and Sultan 2016; Colicchio 2017; Groot et al. 2017) and nearly absent in some cases (Ganguly et al. 2017). This form of plasticity (also known as parental effects) occurs when environments experienced by parents influence the traits of their offspring, without changing the DNA sequence. Such inherited environmental effects can be transmitted from parent to offspring (and to additional generations in some cases) by diverse mechanisms, including heritable epigenetic modifications (i.e., DNA methylation marks, histone modifications, and small RNAs) and the allocation of nutritive resources, hormones, mRNAs, and regulatory proteins to seeds or eggs (these mechanisms are not mutually exclusive; (Herman and Sultan 2011; Jablonka 2013). Additionally, rather than parental and current environment acting in an additive and independent fashion, some transgenerational effects only present themselves when offspring are exposed to particular environments, such as the same stress (e.g., priming, Conrath et al. 2015), and more generally parental x current environment interaction effects appear prevalent (Herman and Sultan 2016; Colicchio 2017; Sultan et al. 2009; Galloway et al. 2009). Thus in addition to genetic inheritance, organisms have at their disposal relevant information from the environments they have experienced and the environments experienced by previous generations (English et al. 2015).

One of the major findings of these studies has been that natural selection should favor

specific forms of within- and transgenerational plasticity depending on the precise patterns of

environmental variation experienced by a population (Shea et al. 2011; Sultan and Spencer 2002).

For instance, within-generation adaptive plasticity is expected to evolve when individuals encounter

variable conditions during their lives, there is little to no cost of responding to environmental cues,

and there is genetic variation for individual phenotypic responses to the environment (Tufto 2015;

reviewed by Scheiner 1993; Schlichting and Pigliucci 1998). These same factors also influence the

evolution of transgenerational plasticity (Lachmann and Jablonka 1996; Räsänen and Kruuk 2007;

Kuijper et al. 2014; Prizak et al. 2014; Leimar and McNamara 2015). First, environments must vary

across generations, either due to temporal change in environmental states at the same site, or by

offspring dispersal across spatially heterogeneous environments. Second, the environments

experienced by parents must predict the selective environments that offspring experience. This

scenario occurs when parental and offspring environments are correlated. Third, there must be

genetic variation for transgenerational plasticity, that is to say, genotype-by-parental environment

variation. This form of genetic variation appears to be common (Herman and Sultan 2016, Schmitt

et al. 1992; Sultan 1996; Stjernman and Little 2011; Alsdurf et al. 2013).

From a more practical perspective, the ability of natural and agricultural populations to

transmit environmental information transgenerationally will determine their ability to persist in the

face of a rapidly fluctuating and changing environments, and thus profoundly influence the health

and productivity of our planet as climate change proceeds. While anthropogenic climate change

differs greatly from natural climatic oscillations, both elevate the value of information from the prior

generation relative to the long-term average conditions at a site, and thus transgenerational plasticity

that has evolved in response to environmental oscillations may provide fitness benefits under

climate change. On the other hand, climate change is expected to increase environmental variation,

including more extreme weather events, potentially decreasing the relative fitness of organisms that

may have evolved transgenerational plasticity to cope with previously reliable patterns.

Empirical investigations in diverse plant and animal systems have confirmed that

transgenerational environmental effects can be adaptive when parent and progeny environments

match (i.e., under positive intergenerational environmental autocorrelations; see e.g., Bilichak et al.

2012; Herman et al. 2012; Rasmann et al. 2012; Slaughter et al. 2012; Verhoeven and van Gurp 2012;

Dantzer et al. 2013; Lopez Sanchez et al. 2016; Walsh et al. 2016; Wibowo et al. 2016). For instance,

when *Mimulus guttatus* plants experience herbivory, their offspring increase production of defensive

leaf trichomes (Holeski 2007; Colicchio et al. 2015; Colicchio 2017). Similarly, when the aquatic

crustacean *Daphnia cuculatta* is exposed to predator cues, it produces offspring with a defensive

'helmet' that protects against predation by midge larvae and cladocerans (Agrawal et al. 1999).

Despite this surge of interest, to date no study has examined long-term environmental data

for the presence of inter-generational environmental correlations that are expected to favor the

adaptive evolution of transgenerational plasticity. Although we tend to think about how the average

environmental conditions differ across a landscape, there is no reason to expect that the scale and

predictability of environmental variation is any less complex or ubiquitous across a landscape. Here

we identify *spatial variation* in the *temporal predictability* of environmental variation, and leverage this

variation to compare the adaptive value of within-generation plasticity, transgenerational plasticity,

and multigenerational epigenetic inheritance across different local regimes of climate variability. We

utilize 120 years of fine-scale (4km^2) climate data spanning the coterminous U.S. to test for auto- and

cross-correlations in temperature and precipitation levels across years, and find many significant

correlations that vary widely in both magnitude and direction across the US. We then construct

separate models for temperature and precipitation to determine the degree of transgenerational

plasticity that would maximize fitness in each of these sites across the U.S. with respect to the

observed environmental measurements. Additionally, by running each model using raw

environmental data and the residuals after factoring out the effects of directional climate change we

were able to inspect how climate change alters the benefits associated with transgenerational

plasticity. These models were constructed with summer annual plants in mind, but are generalizable

to other organisms with generation times of one year.

In our precipitation model, we examine transgenerational effects that persist for up to three

generations (Figure 1a), as multiple experimental studies have found that environmentally induced

epigenetic and phenotypic effects can persist for at least this long (e.g., Whittle et al. 2009;

Akkerman et al. 2016), and in some cases for far longer (Vastenhouw et al. 20026; Rechavi et al.

2011). In our temperature model (Figure 1b), we also determine the degree of within-generation

plasticity that would maximize fitness, in response to both early and late-season temperatures. With

different genotypes producing their phenotypes based on different weightings of within generation

plasticity, transgenerational plasticity, multi-generation epigenetic inheritance, and genetic

inheritance, we find that across the US there is extraordinary variation in the optimal ratio of how

these various classes of information should inform an individual's phenotype. These findings imply

that just as landscape level variation in mean historical conditions selects for locally adapted

populations, landscape level variation in the predictability of environmental variation may select for

locally adapted patterns of plasticity within and across generations.

Methods

Descriptive statistics

Mean monthly temperature and precipitation at a 4km resolution from 1895-2014 (LT81m) were

downloaded from the PRISM climate group web server (53). In short, PRISM uses climate normal

from between 1981-2010 as a predictor grid, and then utilizes station networks with at least 20 years

of data to model monthly temperature and precipitation across the US. The emphasis on this

dataset is long-term consistency making it ideal for our purposes. Individual yearly values were

concatenated using the QGIS merge raster function (Quantum GIS geographic information system

2012), and exported in the .RData format for downstream analysis. For precipitation data, October

was chosen to represent the start of the "hydrologic" year in order to more accurately capture water

availability patterns during the growing season. For temperature data, mean daily maximum

temperature was calculated for March-May as a measure of early growing season temperature for a

given year, and July-September mean daily maximum temperature for late growing season

temperature. Custom R scripts were used to calculate descriptive statistics for precipitation and

temperature separately. Autocorrelations were calculated at lags between 1 and 12 years (i.e.,

environmental correlations were calculated between year X and year X+1, year X and year

X+2…year X and year X+12).

General modeling framework

Mathematical models were constructed in R for both precipitation and temperature patterns

to compare how individuals that utilize within-generation plasticity, transgenerational plasticity, and

genetic inheritance to varying degrees differ in their capacity to match their phenotype with the

environmentally optimal phenotype for a given year. In these models, there are hundreds

(precipitation models) or thousands (temperature models) of competing genotypes, each

representing unique points of parameter space for alleles that modify the extent to which

environment affects phenotype. Trait value is a measure of the expected environment (temperature

or precipitation) and is determined by a combination of the mean environment at a given site over

all years, and terms that modify this value based on recent environmental information. Each

genotype is in essence a climatologist, that utilizes genetic information (based on mean precipitation

over the 120 years at a site), transgenerational plasticity, or within generation plasticity (only in
temperature model) to come up with an expected environment that it will face. This expected
environmental value is equivalent to a phenotype, and the closer this phenotype is to the actual
environment experienced, the higher the fitness that genotype will have for a given generation.
While this framework is identical for precipitation and temperature modeling, inherent
differences in precipitation and temperature variables lead to us considering a different set of
parameters for each variable, allowing us to ask related but unique questions regarding
transgenerational inheritance. Precipitation can accumulate as snowpack, bodies of water, or soil
moisture, such that the cumulative precipitation over the course of the water year will determine to a
large extent the amount of water available to a plant. On the other hand, the effects of temperature
are much more immediate and transient, such that a particularly cold spring will not "keep the plant
cool" over the summer, in the way that a particularly wet spring could provide moisture during a
summer of drought. For this reason, we decided to extend our temperature models to compare
patterns across different segments of the growing season, and different forms of plasticity both
within and between a single generation. For precipitation we only considered annual hydrologic year
precipitation without breaking it down by seasons, but did consider the possibility of multi-
generation persistence of transgenerational plasticity.
For a given genotype, the absolute distance between the expected and actual environment at
a given time imposes a linear cost on fitness. If the expected and actual environment in a given
generation are identical, that genotype's fitness will equal 1, and as the difference increases, fitness
will drop down to a minimum value of 0.2. This minimum value was put in place to prevent a single
year with extremely low fitness from dominating final genotype frequencies, but was only used in
under 0.01% of fitness measurements. After every generation, there is a round of competition,
where the frequency of a given genotype will change based on the relative fitness of that genotype
compared to the mean phenotype for that year. This scenario is equivalent to an infinite population
size evolutionary model in which the relative frequency of genotypes changes due to selection and
reproduction once per year.
For each locale, mean annual precipitation (or temperature) across the 120 years ($\bar{P}$) is
calculated, and this statistic is used as the baseline phenotype of all genotypes in the raw data variant
of the model. Additionally, a linear regression of precipitation on year was fit for each site over the
120 years considered, and residuals were calculated for every year. In the residual variant of this
formula, the expected values are 0, and residuals are used as the precipitation values for each year. A

description of the models developed can be found in Appendix 1, and code can be found at

github.com/XXXX.

Results

Both mean annual precipitation and growing season temperatures vary immensely across the

US (Figure 2 and S1), but for the evolution of locally adaptive phenotypic plasticity, it is the patterns

of variation that are more relevant. The standard deviation of a site's annual precipitation and

growing season temperatures over the past 120 years also varied dramatically (Table 1 and S1), with

precipitation inter-annual standard deviation (IASD) varying from 40mm to 800m, spring

temperature IASD from 0.77C to 1.95C, and summer temperature IASD from 0.35C to 1.32C

(Figure 2 and S1 and Table S1). The southwest US generally had the highest precipitation IASD

relative to its mean precipitation, with IASD being nearly equal to the mean precipitation in some

regions (Figure 2). Interestingly, spring temperature IASD was significantly higher than summer

temperature IASD across the US. This pattern was largely driven by extremely high spring

temperature IASD over the north central US (Figure 2), and consistently low summer temperature

IASDs across much of the coastal US.

Directional climate change over the past 120 years was prevalent and variable across the US

(Figure 2). Mean annual precipitation has declined over much of the Sierra Nevada mountain range,

southern California, and other scattered regions over the last 120 years, while precipitation levels

have increased in the Midwestern and much of the northeastern US. Both spring and summer

temperatures have gotten substantially higher with the exception of the southeast, where spring

temperatures have decreased and summer temperatures have changed little (Figure 2). This

phenomenon has been noted numerous times (Knappenberger et al. 2001; Ellenburg et al. 2016) and

seems to be largely due to a switch from cropland to natural forest ecosystems across the

southeastern US during the past 120 years that has led to greater transpiration cooling. \

Although a variable environment is necessary for the evolution of adaptive phenotypic

plasticity, it is the patterns of this variation that influence which forms of plasticity will be favored.

We examined these patterns by calculating autocorrelations in annual temperature and precipitation

levels between successive years, and found that their magnitude and directions were highly variable

across the US (Figure 2 and S1, Table 1 and S1).

Averaged across all sites, the precipitation autocorrelation (AC) at lag-1 (i.e., the correlation

between the precipitation during one year and the next) was slightly positive (mean=0.04, Table 1,

Figure S1), and was reduced by half after taking linear changes in precipitation into account

(mean=0.02, Table 1, Figure S1). Spatially, we found that the southeastern gulf coast was the largest

region with negative lag-1 ACF (dry years tend to be followed by wet years), while the northeastern

US was the largest region of substantially positive lag-1 ACF (Figure 2). Somewhat surprisingly,

there were many more sites with moderately positive (62,693: lag-2 PACF > 0.2, vs. 21,671: lag-1

ACF >0.2) and negative (5,088 lag-2 PACF <-0.2 vs. 441 lag-1 ACF <-0.2) lag-2 partial

autocorrelation (PAC) than lag-1 ACF. This suggest that climatic oscillations impacting annual

precipitation tend to operate over more than two years in these regions, and that on a year to year

basis, variation is more stochastic (leading to lower absolute lag-1 ACF). The northeastern and

northwestern US show substantially positive precipitation lag-2 PACF; in these regions, multi-year

climatic oscillations may dictate over 20% of inter-annual precipitation variation (Figure 2).

Patterns of temperature autocorrelations extended over larger regions, and were often more

extreme than the patterns observed for precipitation autocorrelations (Figure 2). Lag-1 ACF for

spring and summer temperatures varied a great deal, with patterns of summer temperature

autocorrelation substantially more positive than those of spring (summer ACF-1 mean: 0.24, spring

mean: -0.01, Figure S1). In both cases, however, the western US tended to have more positive

autocorrelations than the rest of the country (with the exception of southern Florida; Figure 2). The

mean lag-2 PACF for spring temperature was negative (mean: -0.04, Figure 3) and more variable

than lag-1 ACF (sd=0.1 vs. 0.08), with much of the north-central US displaying lag-2 PACF of less

than -0.2 (Figure 2). The mean lag-2 ACF for summer temperatures was positive (mean: 0.09), but

substantially lower than the mean lag-1 ACF (mean: 0.24). At a lag of 3, both spring-temperature

ACF (mean: 0.05) and summer-temperature ACF (mean: 0.11) were positive (Figure 3), suggesting

that long-term climatic trends play a role in temperature variation during the past 120 years.

Modeling Results

Optimal levels of transgenerational plasticity: precipitation

As expected, the dramatic variability of precipitation autocorrelations across the US leads to

a great deal of variation in the optimal levels of plasticity in our evolutionary models (Figure 3, Table

S1). In the raw variant of the model, optimal parental effect values were positive in 314,118 cases

(65%), zero in 32,352 (7%), and negative in 135,161 (28%), compared to 55%, 7%, and 38%

respectively in the residual variant. The most common "parental effect" value (m , see Appendix 1)

in the precipitation model was 0.1 (22.5% and 21.8% of sites in the raw and residual models,

respectively, Figure 3a.). This level of parental effect indicates that 90% of phenotypic variance is

dictated by the long-term average (genetic effects), and 10% by the difference between the parental

environment and the long-term average environment. The second most common optimal value of

m was 0.2 (19.15% in the raw model, 18.6% in residual model), followed by -0.1 (14.3% in the raw

model, 17.2% in the residual, Figure 3a). A value of -0.1 means that 10% of phenotypic variation

stems from transgenerational plasticity and 90% from genetic variation, but that the precipitation

experienced by parents will influence the progeny in the opposite direction. Optimal values of m

ranged from -0.6 to 0.9, and were approximately normally distributed, except a dearth of sites for

which the optimal allele value was 0 due to the lack of multigenerational environmental effects in

this situation.

The multigenerational persistence(g, Appendix 1) of transgenerational effects was also

found to vary greatly across the US with the two most common values being 1 (40.7% raw, 39.5%

residual) and 0 (18.6% raw, 16.8% residual) (Figure 3b). Here, a value of 1 indicates that the

precipitation one, two, and three years prior all contribute equally to the expected precipitation at a

given site. A g value of 0 indicates that only the previous year's precipitation is predictive of the

current precipitation level. The remaining 40.7% of sites (in the raw variant) have intermediate

optimal values of g, suggesting that in these locations the precipitation of each of the past three

years is informative, but information from the immediately preceding year is of the highest value.

Of the nine values considered between 0 and 1, all were represented by between 3.1% and 5.7% of

sites, with a value of 0.8 being the most frequent strategy, and 0.1 being the least frequent (Figure 4).

Interestingly, full multigenerational persistence (g =1) was more frequently optimal at sites with

negative transgenerational effect values compared to those with positive values (44.9% vs. 38.9%,

respectively), where intermediate multigenerational persistence was more common (Figure S2).

Additionally, we found that the adaptive value of multigenerational persistence increases as the

strength of the transgenerational effect increases, such that little-to-no multigenerational persistence

is optimal when the transgenerational effect is close to zero, and strong multigenerational effects are

optimal when the transgenerational effect is extreme (Figure S2).

Spatial variation for optimal precipitation plasticity values largely paralleled the spatial

distribution of inter-annual precipitation autocorrelation patterns (compare Figure 2a to Figure 4a).

At the broadest level, the northern latitudes show the highest optimal transgenerational precipitation

plasticity values (Figure 4a), but not necessarily multi-generation persistence of transgenerational

effects (Figure 4b). Optimal transgenerational plasticity values were on average 0.057 lower in the

residual variant of the model compared to the raw variant, with the vast majority of sites having

equal values (56%), decreasing by 0.1 (24%), decreasing by 0.2 (8.5%), or increasing by 0.1 (4.3%).

The northeast US and the Yellowstone National Park region, where precipitation increased most

(Figure 2), also saw the greatest proportion of their optimal transgenerational plasticity values

diminished after factoring out linear climate change (Figure 4c).

Optimal levels of transgenerational plasticity: Temperature

Purely positive transgenerational effects ($m_{EE} \geq 0$, $m_{EL} \geq 0, m_{LL} \geq 0, m_{LE} \geq 0$) of

temperature were optimal in 70.2% of sites (338,327 out of 481,631) in the raw version of the model

and 55.7% of sites (268,307) in the residual variant. Conversely, only 1.4% of sites (7,018) in the raw

model and 3.1% (14,777) in the residual version included only negative transgenerational plasticity

values. Only 0.4% (raw model) or 0.9% (residual model) of sites totally lacked transgenerational

plasticity (either positive or negative) as part of the optimal strategy. The optimal strategies in the

remaining sites (28% raw model, 40% residual model) comprised a mixture of positive and negative

transgenerational plasticity values. Positive within-generation plasticity was favored in 79.7% of sites

(383,667), compared to only 0.03% of sites (157) in which negative within-generation plasticity (w,

Appendix 1) was favored, and 20.3% of sites (97,807) in which no within-generation plasticity was

favored (Figure 3g). Optimal levels of within-generation plasticity were generally positive and

relatively weak across the US; 72% (346,603/481,631) of sites had an optimal w value of 0.1 (Table

S1).

The most common optimal form of transgenerational plasticity to temperature in both the

raw and residual models was the effect of late growing season temperature on the next generation's

late growing season phenotype (m_{LL}, Figure 3f, Table 2a). Effects of late season temperature on the

next generation's early season phenotype (m_{LE}) were the most variable, with a substantial number of

sites having negative transgenerational plasticity values (63,881 raw, 100,267 residual) and many

others having moderate (98,524 raw, 66,625 residual) and major (26,967 raw, 10,080 residual)

positive values (Figure 3e). Effects of early growing season temperature on the next generation's

early season phenotype (m_{EE}) were slightly more likely to be positive in the raw version of the

model (98,089 vs. 89,547), and more likely to be negative in the residual variant (68,441 vs. 132,352).

In both cases however, no m_{EE} plasticity was the most common optimal strategy. Finally, the effect

of early season temperature on the next generation's late season phenotype (m_{EL}) was most often

minor (0.1) in both raw (251,691) and residual (229,301) variants. When considering the combined

plasticity value profile of a site, we find the most common combination of plasticity values is, EE:

none (0), EL: minor (0.1), LE: minor (0.1), LL: moderate (0.3), WP: minor (0.1) (Table 2b).

Summing the four transgenerational plasticity alleles together we find the southwest US has the

highest optimal values of transgenerational plasticity, while the Great Lakes region has the lowest

optimal values (Figure 4d). Texas and Florida tend to have the greatest variability for

transgenerational plasticity values, with many areas within these regions having both positive and

negative transgenerational values leading to intermediate total transgenerational plasticity (Figures 4e

and 4h). Optimal total transgenerational plasticity values were on average 0.146 (+/- 0.21) higher in

the raw relative to the residual models (although the most common change was 0). In the

southwestern US, where temperature increased the most over the past 120 years (Figure 2), the

difference between the raw and residual model was the greatest (Figure 4g).

Variation in different classes of temperature autocorrelations between seasons explains a

large portion of the variation in the optimal transgenerational response to temperature at a given

site. For example, the autocorrelation between early season growing temperature and the next year's

late season growing temperature is the factor that explains the largest amount of variation in optimal

levels of m_{EL} (Table 3). We assessed potential tradeoffs between different forms of

transgenerational plasticity to temperature by first calculating the residuals of a particular plasticity

term after accounting for the effects of environmental autocorrelations, then testing the effect of the

other four plasticity terms on these residuals. There was a highly significant negative association

between m_{LE} and m_{EE} plasticity, and between m_{LL} and m_{EL} plasticity (Figure S3a). As higher levels

of late-late transgenerational plasticity were favored, the optimal levels of EL plasticity also

decreased across all environmental autocorrelation values. These associations suggest that, for a

given life history stage in this model, there are tradeoffs between using transgenerational information

from the previous generation's early vs. late season temperature (Table S2). For example, there are

many sites where no plasticity, m_{EE} plasticity, and m_{LE} plasticity all have higher fitness than

individuals exhibiting both m_{EE} and m_{LE} plasticity (Figure S3b).

Fitness Landscapes

In the previous analyses we used restricted parameter space to identify optimal site-specific

combinations of plasticity values across the entire contiguous U.S., but further insight can be gained

by comparing fitness landscapes across the full parameter space at individual sites. In the

precipitation model we found that, among sites where fitness optima are located near zero

transgenerational effects, a vertical fitness ridge formed that was centered near parental effect values

of zero. This result is due to transgenerational persistence levels (y-axis) having a minimal impact on

phenotype when parental effects are marginal. As absolute optimal parental effect values increased,

however, the fitness landscape shifted from a ridge to a peak, with certain values of

transgenerational persistence imparting extreme fitness advantages over others (Figure 5). Site B

(North Central Minnesota) exemplifies a unique and unexpected outcome of this model: under

certain conditions, there can be multiple local fitness maxima with divergent levels of

transgenerational plasticity (Figure 5). Two fitness maxima exist at this site, one in which the optimal

strategy comprises slightly negative parental effect values with no multigenerational persistence, and

a second in which the optimal strategy comprises slightly to moderately positive parental effect

values with high levels of multigenerational persistence. This situation occurs when two conditions

hold: the lag-1 autocorrelation is in a different direction than the average of the lag-2 and lag-3

autocorrelations, and the absolute value of the lag-1 autocorrelation is less than the average of the

lag-2 and lag-3 autocorrelations. In these cases, plasticity in the opposite direction of the previous

generation, or in the same direction as the average environment of the past three generations, would

be adaptive. This scenario occurs in approximately 90k out of the 480k sites, but only in 30k sites are

lag-2 and lag-3 average values greater than 0.1 and therefore likely to show up as bimodal peaks in

our model. Over the course of the 120 year simulation, the frequency of plasticity genotypes

fluctuated somewhat dramatically, suggesting that the interactions between climatic oscillations and

other effects might have favored positive plasticity values over some decades, and negative in the

next (Figure S4)

Discussion

Although transgenerational environmental effects on phenotypic expression have been

recognized for decades (Falconer 1981; Roach and Wulff 1987), interest in these effects has surged

recently due to increased appreciation for the potential role of transgenerational plasticity in

adaptation (Donelson et al. 2018). Despite this renewed interest, a critical question has remained

unanswered: do natural patterns of environmental variation favor the adaptive evolution of

transgenerational plasticity? Our analysis of 120 years of climatic data from the continental U.S.

revealed that such patterns are indeed widespread. Specifically, we analyzed how inter-annual

variation in precipitation and temperature impacts the optimal mode of adaptation for clonally

reproducing organisms with a life cycle meant to mimic that of an annual plant. When there are

correlations between the parental and offspring environments, neither traditional genetic selection

nor within-generation plasticity take full advantage of the available information inherent in the

environment. Instead, under such correlations selection should favor the genetic evolution of

mechanisms that transmit plastic responses from one generation to the next. Absent such

correlations, the information provided by the parental environment may not be relevant to

offspring, and indeed may prove to be maladaptive (reviewed by Herman et al. 2014).

Our modeling results revealed that the vast majority of sites in the contiguous US

experienced autocorrelations in precipitation and temperature that should favor the evolution of

adaptive transgenerational plasticity. As predicted by other models, the predictability of an

environmental variable based on its autocorrelation or its association with other environmental

variables is a major factor driving the optimal level of plasticity (e.g., Groot et al. 2017; English et al.

2015; Sultan and Spencer 2002; Scheiner 2016). Furthermore, we find that the strength and direction

of autocorrelations in precipitation and temperature varied substantially across the U.S., and

consequently, the optimal levels of plasticity were also highly variable. Although both precipitation

and temperature interact to shape the moisture availability and rates of transpiration for a plant, here

we consider these terms separately. Future studies considering temperature and precipitation in

tandem should shed new light into the transmission of more complex environmental information.

Precipitation

Local adaptation to variable water regimes has been a major focus of plant evolutionary

ecology for many years. This research shows that plants have evolved a wide range of physiological,

phenological, and morphological adaptations to handle site-specific patterns of water availability

(Kooyers 2015). These adaptive phenotypes may be expressed constitutively or may be induced by

an environmental cue that predicts a change in water availability later in the life of the organism.

Increasingly, experimental studies show that the parental soil moisture regime can also adaptively

influence the development of progeny (e.g., Alsdurf et al. 2013; Alsdurf et al. 2015), providing a

third route by which plants can fine tune the phenotypes of their offspring to local soil-moisture

levels. For instance, in Massachusetts genotypes of the annual plant *Polygonum persicaria*, offspring of

drought-stressed parents make more extensive root systems and deploy them faster in response to

drought as compared to offspring of well-watered parents. This drought-induced change in growth

and development can be inherited for at least two generations, resulting in increased survival of

grandoffspring under severe drought stress (Sultan et al 2009; Herman et al. 2012). Furthermore,

these epigenetic effects of drought are genetically variable in *P. persicaria*: some genotypes strongly

increase root length and biomass in response to parental drought, while other genotypes do so only

moderately or not at all (Herman and Sultan 2011).

Our analysis revealed substantial and spatially variable interannual autocorrelations in

precipitation, indicating that precipitation levels in one year are often predictive of precipitation

levels up to three years later. For example, across the coterminous U.S., lag-1 interannual

precipitation autocorrelations varied from moderately negative (-0.27) to strongly positive (0.69),

including some values near zero. In turn, the optimal direction and strength of transgenerational

effects of precipitation also varied. Positive parental effects, wherein individuals are developmentally

predisposed to perform better in environments that match their parents' environment, were optimal

across more than twice as many regions (65% of sites) as negative transgenerational effects (28% of

sites), wherein individuals perform better in a different environment than their parents. Relatively

strong parental effect values of 0.3 or higher were optimal in nearly 30% of sites. By contrast,

complete absence of parental effects was favored in only 7% of sites.

Multigenerational persistence values of 0 (18.7% of sites) and 1 (40.7% of sites) were most

common, representing strategies in which transgenerational effects lasted only a single generation or

persisted fully to the third generation, respectively. The remaining persistence values were somewhat

evenly distributed between 0 and 1 and represent strategies in which environmental information gets

passed through three generations, but the environment of more recent years is weighted more

heavily.

The optimal level of transgenerational effects varied on multiple scales. On the largest scale,

we found that the western and northern US experience conditions that select for the highest levels

of transgenerational plasticity (Figure 4a). There was a striking contrast between the northeast,

where positive transgenerational plasticity was generally optimal, and the southeast, where negative

transgenerational plasticity predominated. These large-scale differences in plasticity levels stemmed

from substantial differences in precipitation autocorrelations between these regions. There was also

considerable variation in optimal levels of transgenerational plasticity on much finer scales. In some

cases, optimal levels of transgenerational plasticity were highly divergent between adjacent sites (e.g.,

in central Texas and Minnesota). Furthermore, some sites had bimodal fitness landscapes, in which

genotypes that express either positive or negative transgenerational effects were favored over

genotypes that do not express plasticity (Figure 5b). For example, after a wetter-than-average year

but two drier than average years prior to that, a genotype that prepares its offspring for another

wetter-than-average year (i.e., a positive transgenerational effect), or one that prepares its offspring

for a drier-than-average year (i.e., a negative transgenerational effect), will have higher fitness than a

genotype that does not express transgenerational plasticity. To our knowledge, this represents the

first demonstration that patterns of environmental variation can favor both positive and negative

transgenerational effects in the same site.

Temperature

Temperature is vitally important to plant function and fitness, as it impacts the rate of

physiological reactions, cues developmental transitions, and in extremes can cause stress and

mortality. Plants adapt to variable temperature regimes in a host of ways, including the production

of heat shock proteins and cold-response factors, and the development of morphologies that

mitigate the experience of temperature extremes. Experimental studies have identified adaptive

plastic responses to temperature changes in plants, both within and across generations. For example,

ambient temperature in Arabidopsis has been shown to influence the expression and splicing of

hundreds of genes, leading to changes in underlying histone methylation (Pajoro et al. 2017), and

significantly shifts flowering time (Donohue 2009) and many other phenotypes (Adams et al. 2016)

in genotype-specific ways. Additionally, recent work has demonstrated the lasting impact of

temperature regime on *Arabidopsis thaliana* plants for generations to come (Whittle et al. 2009; Suter

and Widmer 2013a; Suter and Widmer 2013b; Groot et al. 2017).   In order for these responses to

adaptively match phenotypes with natural environments, there must be substantial correlations in

temperature within and between growing seasons.

We found significant autocorrelations in temperature, both within and between years. Within

a single growing season, temperatures early and late in the growing season tended to be positively

correlated across the U.S. Furthermore, we found that the temperatures of the late growing season

months (July, August, September) were generally strongly autocorrelated between successive years.

Interannual correlations between the temperatures of the early growing season months (February,

March, April) were often much lower. Surprisingly, early growing season temperatures were

generally more predictive of the following late seasons temperature than they were of the following

early seasons temperature.

In our framework, temperature both acts as a selective agent, by which fitness is reduced

according to the deviation between the actual temperature and a genotype's phenotypic temperature

optimum, and second as a source of information, that a plant can use to alter its phenotype later in

the growing season or its offspring's phenotype in the following year. As expected, we find that, at a

given site, the strength of the correlation between the average temperature during the season in

which information is gathered and the average temperature during the season when selection occurs

is highly predictive of both the type and degree of plasticity that will be favored. For example,

warmer than average springs were very often followed by hotter than average summers, and this

information yielded benefits via within-generation responses to temperature in many sites. The

optimal strategy in more than 99% of sites across the U.S. contained some form of transgenerational

plasticity, suggesting that environmental oscillations provide valuable information that allows

transgenerational plasticity to improve the match between phenotypes and temperature regimes.

The most common form of transgenerational plasticity in this model was late-growing

season temperature impacting the following generation's phenotype late in the growing season,

which matches our expectations based on the patterns of temperature autocorrelation. Interestingly,

patterns of environmental oscillations lead to favorable strategies in which the current late-season

phenotype was more strongly impacted by the previous late-season temperature than it was by the

current generation early season temperature. Indeed, this pattern was found in over half of the

regions considered (270k/480k). Although intuition suggests that more recent information is of

higher value, this result suggests that parental environments can be more predictive of offspring

selective environments than environmental cues early in the offspring generation. This result stems

from the cyclic nature of seasonal environments (Auge et al. 2017). Since autocorrelations between

consecutive early growing seasons were generally low, it is not surprising that effects of early

growing season temperatures on phenotypes in the following early growing season was the least

common form of plasticity and was in the negative direction more often than the positive. Other

forms of transgenerational plasticity were present at intermediate levels and varied across the US.

The west coast of the US and southern Florida experienced the highest optimal

transgenerational plasticity values. Because these regions are due east of large bodies of water, their

climates are heavily influenced by maritime airflow including the prevailing westerlies, loop current,

and Coriolis affect (Lorentz 1966). As water has a substantially higher heat capacity than either rock

or soil, the location of these land masses downstream of maritime air may predispose them to

temperature autocorrelations between years. Additionally, we found highly variable correlations

between late growing season temperature and the following generation's early growing season

temperature. This result is intriguing because the temperature experienced during seed maturation

interacts with the post-dispersal temperature to strongly influence the dormancy and germination

behavior of seeds, with cascading effects throughout the life cycles of annual plants (Donohue 2009;

Burghardt et al. 2016). Consequently, site-specific correlations between maternal late-season

temperature and the early-season temperature in the next generation may select for divergent, site-

specific effects of maternal temperature on germination. Intriguingly, parental effects of temperature

on germination and flowering time are highly genetically variable in *Arabidopsis thaliana* (Burghardt et

al. 2016; Kerdaffrec and Nordburg 2017; Groot et al. 2017). In *Plantago lanceolata*, such genotype-by-

maternal temperature effects persist throughout the offspring life cycle to generate variation in

offspring reproduction in the field (Lacey and Herr 200). Our results suggest that such genetic

variation for maternal effects may derive in part from variable selection imposed by differences

among sites in intergenerational temperature correlations (see also Groot et al. 2017).

Common Themes and Future Directions

Although our precipitation and temperature models yielded distinct insights into the

dynamics of each of these factors, common themes emerged in both sets of analyses. For example,

we found higher levels of inter-annual autocorrelation, and therefore more prominent

transgenerational effects, at northern latitudes and along coastal regions within both models. Studies

that compare patterns of transgenerational plasticity across such large geographic regions will be

necessary to determine whether underlying differences in environmental patterns do in fact drive

differences in transgenerational plasticity. Another common finding of both the temperature and

precipitation models is that transgenerational effects are expected to provide greater benefits in

changing climates relative to purely oscillating climates, in which linear climate change has been

removed (i.e., the residual models). These results suggest that transgenerational effects may have an

important role in adaptation to human-induced climate change, and that rapid climate change should

select for more transgenerationally plastic individuals. However, there is an important caveat. In our

models we assume that genotypes are uniform in their mean phenotype, and do not allow for

mutations that could lead to genetic adaptation to changing conditions. The potential for

transgenerational plasticity to either promote or hinder genetic adaptation has been explored (Day

and Bonduriansky 2011), but our models do not address this issue. In the absence of genetic

evolution, it follows that if there is a linear trend towards hotter or drier years in addition to climatic

oscillations (as in the raw model variants), then there is more transgenerational information relative

to a situation in which only climatic oscillations are occurring (residual model variants). Theory

indicates that these dynamics become much more complex when local genetic adaptation to

changing conditions is allowed to occur along with plastic responses (Groot et al. 2017). For

instance, in some scenarios transgenerational effects can increase fitness in the short term, while

reducing it in the long term (Hoyle and Ezard 2012). In future studies, it will be critical to investigate

the influence of standing genetic variation for transgenerational plasticity on evolutionary

trajectories, and to examine the availability of novel mutations that alter the dynamics of

transgenerational effects.

Temperature and precipitation autocorrelations likely stem in part from the same broad-scale

climatic oscillations, such as the El Niño Southern Oscillation (Yang et al. 2018), the Quasi-biennial

oscillation (Baldwin et al. 2001), and the Pacific Decadal Oscillation (Mantua and Hare 2002;

Newman et al. 2016). Aside from these climatic oscillations, autocorrelations will arise due simply to

"red" or "pink" noise in which rare, large events and common, small events have equal power in

explaining variation (Szendro et al. 2001). It has been demonstrated that even without clear

underlying phenomena explaining variation, pink-noise is often the model that best explains patterns

of ecological and abiotic time-series variation (Halley 1996). These oscillations and general patterns

of red noise will interact with each other to varying degrees across different regions of the US,

leading to variable levels of autocorrelation at all lags for both precipitation and temperature.

Furthermore, because temperature and precipitation interact to alter moisture availability, it

is likely that organisms do not process temperature and precipitation information independently, but

rather use them in tandem along with other sources of information to fine tune phenotypes for the

most likely future environment. For instance, temperature influences water availability by influencing

rates of evaporation and transpiration. Interactions between temperature and water availability also

shape the collection of herbivores, pathogens, and competitors present in a given locality.

Understanding how these environmental factors jointly influence the expression of transgenerational

plasticity is an important goal for future research. A key element of this research direction is to study

environmental (auto)correlations at fine scales in the context of dispersal distances. It is possible that

transgenerational plasticity may be a more common mode of adaptation for organisms with short

dispersal distances, in which parents and offspring are more likely to grow and develop in similar

microsites. Finally, differences in life history strategies and generation times will alter the time-scales

and types of environmental autocorrelations relevant to transgenerational plasticity. Future studies

drawing inspiration from other species will provide insight into how these variables might impact

the evolution of transgenerational effects.

Conclusion

In summary, we demonstrate that patterns of climatic variation in nature may favor the

adaptive evolution of transgenerational plasticity in organisms with approximately annual generation

times, such as annual plants. Our models indicate that differing patterns of climatic oscillations

across the US lead to strikingly different optimal patterns of within- and transgenerational plasticity.

Thus, for a given species, one may expect that environmental variation across its range not only

selects for different locally adapted mean trait values, but also different classes and magnitudes of

plasticity. Perhaps the most meaningful result of this study is that the climatic patterns across

relatively small geographic regions vary so dramatically that the optimal value of transgenerational

plasticity ranges from extremely high to non-existent. It should therefore be expected that although

many species, environmental variables, or phenotypes of interest may show no evidence of

transgenerational plasticity, such results may be due to their specific situation rather than a

fundamental biological limitation. This applies equally strongly to the other side of the coin: because

a single population or species expresses strong transgenerational plasticity does not mean that this

process is a key driver of evolutionary processes. Rather, variation in transgenerational plasticity

should be expected, just as genetic variation is ubiquitous in natural populations. Transgenerational

plasticity is best considered in the specific ecological and evolutionary context of the study organism,

and broad generalizations about the role of these effects in evolution should be avoided until

considerably more field data are in hand. The results described here provide a source of testable

predictions for geographical variation in this mode of adaptation.

634 **References**

1. Adams, W.W., J.J. Stewart, C.M. Cohu, O. Muller, & B. Demmig-Adams. 2016. Habitat temperature and precipitation of Arabidopsis thaliana ecotypes determine the response of

foliar vasculature, photosynthesis, and transpiration to growth temperature. Front. Plant Sci. 7:1026. doi. 10.3389/fpls.2016.01026.

2. Agrawal, A., C. Laforsch, and R. Tollrian. 1999. Transgenerational induction of defences in animals and plants. Nature 401:60-63.

3. Akkerman, K., A. Sattirin, J. K. Kelly, and A. G. Scoville. 2016. Transgenerational plasticity is sex-dependent and persistent in yellow monkeyflower (Mimulus guttatus). Environmental Epigenetics. doi: 10.1093/eep/dvw003

4. Alsdurf, J. C. Anderson, D.H. Siemens. 2015. Epigenetics of drought-induced trans-generational plasticity: consequences for range limit development. AoB Plants 8: plv146. doi: 10.1093/aobpla/plv146.

5. Alsdurf, J. D., T. J. Ripley, S. L. Matzner, and D. H. Siemens. 2013. Drought-induced trans-generational tradeoff between stress tolerance and defence: consequences for range limits? AoB Plants 5:plt038.

6. Auge, G. A., L. D. Leverett, B. R. Edwards, and K. Donohue. 2017. Adjusting phenotypes via within- and across-generational plasticity. New Phytol. 216:343-349.

7. Baldwin, M.P., L.J. Gray, T.J. Dunkerton, K. Hamilton, P.H. Haynes, W.J. Randel, J.R. Holton, M.J. Alexander, I. Hitora, T. Horinouchi, D.B.A. Jones, J.S. Kinnersley, C. Marquardt, K. Sato, M. Takahashi. 2001. The quasi-biennial oscillation. Rev. Geophys. 39:179-229. doi: 10.1029/1999rg000073.

8. Bilichak, A., Y. Ilnystkyy, J. Hollunder, and I. Kovalchuk. 2012. The Progeny of Arabidopsis thaliana Plants Exposed to Salt Exhibit Changes in DNA Methylation, Histone Modifications and Gene Expression. PLoS ONE 7:e30515. doi:10.1371/journal.pone.0030515.

9. Bonduriansky, R. and T. Day. 2009. Nongenetic inheritance and its evolutionary implications. Annu. Rev. Ecol., Evol. Syst. 40:103-125.

10. Burghardt, L.T., B.R. Edwards, K. Donohue. 2016. Multiple paths to similar germination behavior in Arabidopsis thaliana. New Phyt. 209:1301-1312. doi: 10.1111/nph.13685.

11. Colicchio, J. 2017. Transgenerational effects alter plant defence and resistance in nature. J. Evol. Biol. 30:664-680. doi: 10.1111/jeb.13042.

12. Colicchio, J., F. Miura, J. Kelly, T. Ito, L. Hileman. DNA methylation and gene expression in Mimulus guttatus. 2015. BMC Genomics 16:507. doi: 10.1186/s12864-015-1668-0.

13. Conrath, U., G. J. Beckers, C. J. Langenbach, and M. R. Jaskiewicz. 2015. Priming for enhanced defense. Annu Rev Phytopathol 53:97-119.

14. Daly, C., W. P. Gibson, G. H. Taylor, G. L. Johnson, and P. Pasteris. 2002. A knowledge-based approach to the statistical mapping of climate. Climate Research 22:99-113.

15. Dantzer, B., B. Dantzer, A. E. M. Newman, A. E. M. Newman, R. Boonstra, R. Boonstra, R. Palme, R. Palme, S. Boutin, S. Boutin, M. M. Humphries, M. M. Humphries, A. G. McAdam, and A. G. McAdam. 2013. Density Triggers Maternal Hormones That Increase Adaptive Offspring Growth in a Wild Mammal. Science. 340:1215-1217. doi: 10.1126/science.1235765.

16. Day & Bonduriansky. 2011. A unified approach to the evolutionary consequences of genetic and non-genetic inheritance. Am Nat. 178: E18-E36. doi: 10.1086/660911.

17. Donelson, J.M., S. Salinas, P.L. Munday, L.N.S. Shama. 2018. Transgenerational plasticity and climate change experiments: where do we go from here? Glob. Change. Biol. 24:13-34. doi: 10.1111/gcb.13903.

18. Donohue, K. 2009. Completing the cycle: maternal effects as the missing link in plant life histories. Phil. Trans. R. Soc. B. 364:1059-1074. doi: 10.1098/rstb.2008.0291.

19. Ellenburg, W.L., R.T. McNider, J.F. Cruise, J.R. Christy. 2016. Towards an understanding of the Twentieth-Century cooling trend in the southeastern United States: biogeophysical impacts of land-use change. Earth Interactions 20. doi: 10.1175/ei-d-15-0038.1.

20. English, S., I. Pen, N. Shea, and T. Uller. 2015. The information value of non-genetic inheritance in plants and animals. PloS One 10:e0116996. doi:10.1371/journal.pone.0116996.

21. Falconer, D. S. 1981. Introduction to Quantitative Genetics. London: Longman.

22. Fisher, R.A. 1930. The Genetical Theory of Natural Selection. Clarendon, Oxford.

23. Galloway, L., J. Etterson, & McGlothlin. 2009. Contribution of direct and maternal genetic effects to life-history evolution. New Phytol. 183:826-838. doi: 10.1111/j.1469-8137.2009.02939.x

24. Ganguly, D.R., P.A. Crisp, S.R. Eichten, B.J. Pogson. 2017. The Arabidopsis DNA Methylome is Stable Under Transgenerational Drought Stress. Plant Phys. 175:1893-1912. doi: 10.1104/pp.17.00744.

25. Ghalambor, C.K., J.K. McKay, S.P. Carroll, & D.N. Reznick. 2007. Adaptive versus non-adaptive phenotypic plasticity and the potential for contemporary adaptation in new environments. Funct. Ecol. 21:394-407.

26. Groot, M.,P., Kubisch A., Ouborg, N.J., Pagel, J., Schmid, K.J., Verger, P., Lampei, C.. Transgenerational effects of mild heat in Arabidopsis thaliana show strong genotype specificity that is explained by climate at origin. New Phyt. doi: 10.1111/nph.14642.

27. Halley, J. 1996. Ecology, evolution, and 1/f noise. Trends Ecol. Evol. 11:33-37.

28. Herman, J. J. and S. E. Sultan. 2011. Adaptive transgenerational plasticity in plants: case studies, mechanisms, and implications for natural populations. Front. Plant Sci. 2:1-10. doi: 10.3389/fpls.2011.00102 .

29. Herman, J. J. and S. E. Sultan. 2016. DNA methylation mediates genetic variation for adaptive transgenerational plasticity. Proc. R. Soc B. 283: 20160988. Doi: 10.1098/rspb.2016.0988.

30. Herman, J. J., S. E. Sultan, T. Horgan-Kobelski, and C. Riggs. 2012. Adaptive transgenerational plasticity in an annual plant: Grandparental and parental drought stress enhance performance of seedlings in dry soil. Integr. Comp. Biol. 52:77-88.

31. Herman, J.J., H.G. Spencer, K. Donohue, & S.E. Sultan. 2014. How stable 'should' epigenetic modifications be? Insights from adaptive plasticity and bet hedging. Evolution 68:632-643. doi: 10.1111/evo.12324.

32. Holeski, L. M., G. Jander, and A. A. Agrawal. 2012. Transgenerational defense induction and epigenetic inheritance in plants. Trends Ecol. Evol. 27:618-626.

33. Holeski, L.M. 2007. Within and between generation phenotypic plasticity in trichome density of Mimulus guttatus. J. Evol. Biol. 20:2092-2100. doi: 10.1111/j.1420-9101.2007.01434.x.

34. Hoyle, R. B. and T. H. Ezard. 2012. The benefits of maternal effects in novel and in stable environments. J R Soc Interface 9:2403-2413.

35. Jablonka, E. 2013. Epigenetic inheritance and plasticity: The responsive germline. Prog. Biophys. Mol. Biol. 111:99-107.

36. Kerdaffrec, E. & M. Nordborg. 2017. The maternal environment interacts with genetic variation in regulating seed dormancy in Swedish Arabidopsis thaliana. PLoS ONE 12: e0190242. doi: 10.1371/journal.pone.0190242.

37. Knappenberger, P.C., P.J. Michaels, R.E. Davis. 2001. Nature of observed temperature changes across the United States during the 20th century. Climate Res. 17:45-53.

38. Kooyers, N.J. 2015. The evolution of drought escape and avoidance in natural herbaceous populations. Plant Sci. 234:155-162. doi: 10.1016/j.plantsci.2015.02.012.
39. Kuijper, B. and R. B. Hoyle. 2015. When to rely on maternal effects and when on phenotypic plasticity? Evolution 69:950-968.
40. Kuijper, B., R. A. Johnstone, and S. Townley. 2014. The evolution of multivariate maternal effects. PLoS Comput. Biol. 10:e1003550.
41. Lacey, E. & D. Herr. 2000. Parental effects in Plantago lanceolata L. III. Measuring parental temperature effects in the field. Evolution 54:1207-1217.
42. Lachmann, M. and E. Jablonka. 1996. The inheritance of phenotypes: an adaptation to fluctuating environments. J. of Theor. Biol. 181:1-9.
43. Leimar, O. and J. M. McNamara. 2015. The evolution of transgenerational integration of information in heterogeneous environments. Am. Nat. 185:E55-E69. Doi: 10.1086/679575.
44. Lopez Sanchez, A., J. H. Stassen, L. Furci, L. M. Smith, and J. Ton. 2016. The role of DNA (de)methylation in immune responsiveness of Arabidopsis. Plant J. doi: 10.1111/tpj.13252.
45. Lorentz, E. The circulation of the atmosphere. 1966. 2017Amer. Scientist 54:402-420.

46. Mantua, N. & S. Hare. 2002. The Pacific Decadal Oscillation. J. Oceanography 58:35-44.
47. Mousseau, T. A. and C. W. Fox. 1998. Maternal effects as adaptations. Oxford University Press, New York.
48. Newman, M., M.A. Alexander, T.R. Ault, K.M. Cobb, C. Deser, E. Di Lorenzo, N.J. Mantua, A.J. Miller, S. Minobe, H. Nakamura, N. Schneider, D.J. Vimont, A.S. Phillips, J.D. Scott & C.A. Smith. 2016. The Pacific Decadal Oscillation, revisited. 2016. J. Climate 29:4399-4427.
49. Nicotra, A. B., O. K. Atkin, S. P. Bonser, A. M. Davidson, E. J. Finnegan, U. Mathesius, P. Poot, M. D. Purugganan, C. L. Richards, F. Valladares, and M. v. Kleunen. 2010. Plant phenotypic plasticity in a changing climate. Trends Plant Sci. 15:684-692.
50. Pajoro, A., E. Severing, G.C. Angenent, & R.G. H. Immink. 2017. Histone H3 lysine 36 methylation affects temperature-induced alternative splicing and flowering in plants. Genome Biology. 18:102. doi: 10.1186/s13059-017-1235-x.
51. Prizak, R., T. H. Ezard, and R. B. Hoyle. 2014. Fitness consequences of maternal and grandmaternal effects. Ecol. and Evol. 4:3139-3145.
52. Quantum GIS geographic information system. 2012. Open source geospatial foundation project. Free Software Foundation, India.
53. Räsänen, K. and L. Kruuk. 2007. Maternal effects and evolution at ecological timescales. Funct. Ecol. 21:408-421.
54. Rasmann, S., M. De Vos, C. L. Casteel, D. Tian, R. Halitschke, J. Y. Sun, A. A. Agrawal, G. W. Felton, and G. Jander. 2012. Herbivory in the Previous Generation Primes Plants for Enhanced Insect Resistance. Plant Physiol. 158:854-863.
55. Rechavi, O., G. Minevich, and O. Hobert. 2011. Transgenerational inheritance of an acquired small RNA-based antiviral response in C. elegans. Cell 147:1248-1256.
56. Roach, D., and Wulff, R. 1987. Mater- nal effects in plants. Annu. Rev. Ecol. Syst. 18, 209–235.
57. Scheiner, S. 2016. Habitat Choice and Temporal Variation Alter the Balance between Adaptation by Genetic Differentiation, a Jack-of- All-Trades Strategy, and Phenotypic Plasticity. Am. Nat. 187:633-646. doi: 10.1086/685812.

58. Scheiner, S. M. 1993. Genetics and Evolution of Phenotypic Plasticity. Annu. Rev. Ecol. Syst. 24:35-68.

59. Schlichting, C. and M. Pigliucci. 1998. Phenotypic evolution: a reaction norm perspective. Sinauer, Sunderland, Massachusetts.

60. Schmitt, J., J. Niles, and R. Wulff. 1992. Norms of reaction of seed traits to maternal environments in Plantago lanceolata Am. Nat. 139:451-466.

61. Shea, N., I. Pen, and T. Uller. 2011. Three epigenetic information channels and their different roles in evolution. J. Evol. Biol. 24:1178-1187.

62. Slaughter, A., X. Daniel, V. Flors, E. Luna, B. Hohn, and B. Mauch-Mani. 2012. Descendants of Primed Arabidopsis Plants Exhibit Resistance to Biotic Stress. Plant Physiol. 158:835-843.

63. Stjernman, M. and T. J. Little. 2011. Genetic variation for maternal effects on parasite susceptibility. J. Evol. Biol. 24:2357-2363.

64. Sultan, S. E. 1996. Phenotypic plasticity for offspring traits in Polygonum persicaria. Ecology 77:1791-1807.

65. Sultan, S. E. 2015. Organism and Environment: Ecological Development, Niche Construction, and Adaptation. Oxford University Press, New York, NY, United States of America.

66. Sultan, S. E. and H. G. Spencer. 2002. Metapopulation structure favors plasticity over local adaptation. Am. Nat. 160:271-283.

67. Sultan, S. E., K. Barton, and A. M. Wilczek. 2009. Contrasting patterns of transgenerational plasticity in ecologically distinct congeners. Ecology 90:1831-1839.

68. Suter L, Widmer A. 2013a. Environmental heat and salt stress induce transgenerational phenotypic changes in Arabidopsis thaliana. PLoS ONE 8: e60364. doi:10.1371/journal.pone.0060364.

69. Suter L, Widmer A. 2013b. Phenotypic effects of salt and heat stress over three generations in Arabidopsis thaliana. PLoS ONE 8: e80819. doi:10.1371/journal.pone.0080819.

70. Szendro, P. G. Vincze, A. Szasz. Pink-noise behaviour of biosystems. 2001. European Biophys. J. 30:227-231.

71. Tufto, J. 2015. Genetic evolution, plasticity, and bet-hedging as adaptive responses to temporally autocorrelated fluctuating selection: A quantitative genetic model. Evolution 69:2034-2049.

72. Uller, T. 2008. Developmental plasticity and the evolution of parental effects. Trends Ecol. Evol. 23:432-438.

73. Uller, T., S. English, and I. Pen. 2015. When is incomplete epigenetic resetting in germ cells favoured by natural selection? Proc. R. Soc. B. 282:20150682. Doi: 10.1098/rspb.2015.0682.

74. Vastenhouw, N. L., K. Brunschwig, K. L. Okihara, F. Muller, M. Tijsterman, and R. H. Plasterk. 2006. Gene expression: long-term gene silencing by RNAi. Nature 442:882.

75. Verhoeven, K. J. F. and T. P. van Gurp. 2012. Transgenerational effects of stress exposure on offspring phenotypes in apomictic dandelion. PLoS ONE 7:e38605.

76. Walsh, M. R., T. Castoe, J. Holmes, M. Packer, K. Biles, M. Walsh, S. B. Munch, and D. M. Post. 2016. Local adaptation in transgenerational responses to predators. Proc. R. Soc. B. 283. 20152271. doi: 10.1098/rspb.2015.2271.

77. Whittle, C., S. Otto, M. Johnston, and J. Krochko. 2009. Adaptive epigenetic memory of ancestral temperature regime in Arabidopsis thaliana. Botany 87:650-657.

78. Wibowo, A., C. Becker, G. Marconi, J. Durr, J. Price, J. Hagmann, R. Papareddy, H. Putra, J. Kageyama, J. Becker, D. Weigel, and J. Gutierrez-Marcos. 2016. Hyperosmotic stress memory in Arabidopsis is mediated by distinct epigenetically labile sites in the genome and is restricted in the male germline by DNA glycosylase activity. Elife 5:e13546. doi: 10.7554/eLife.13546.

79. Yang, S., Z. Li, J. Yu, X. Hu, W. Dong, & S. He. 2018. El Ñino Southern Oscillation and its Impact in the Changing Climate. National Science Review doi:10.1090/nsr/nwy046.

662 Table 1: Summary statistics of climatic patterns relevant to the evolution of within and

663 transgenerational plasticity. Mean (s.d.).

	Mean	IASD	RACF-1	ACF-1	ACF-2	ACF-3
Precipitation	763 (443)	145 (76)	0.02 (0.08)	0.04 (0.09)	0.05 (0.13)	0.01 (0.09)
Spring Temp	10.4 (5.5)	1.2 (0.2)	-0.04 (0.07)	-0.01 (0.08)	-0.04 (0.1)	0.05 (0.09)
Summer Temp	21.2 (4.2)	0.9 (0.2)	0.17 (0.1)	0.24 (0.12)	0.09 (0.1)	0.11 (0.09)

689   Table 2: Most common, second most common, and mean optimal plasticity values across all sites in

690   the US.

Climate Term		Raw		Residual	
		#1 / #2	Mean (s.d.)	#1 / #2	Mean (s.d.)_
Precipitation	M	0.1/0.2	0.094 (0.22)	0.1/0.2	0.036 (0.21)
	G	1/0	0.63 (0.40)	1/0	0.64 (0.40)
Temperature	m_{EE}	0 / 0.1	-0.016 (0.098)	0 / -0.2	-0.04 (0.11)
	m_{EL}	0.1 / 0	0.057 (0.073)	0.1 / 0	0.045 (0.073)
	m_{LE}	0.1 /0	0.096 (0.178)	0 / 0.1	0.042 (0.17)
	m_{LL}	0.3 / 0.1	0.204 (0.133)	0.1 / 0.3	0.148 (0.117)
	w	0.1 / 0	0.095 (0.072)	0.1 / 0	0.092 (0.072)

710　Table 3: Correlations between the temperature correlation between seasons and the optimal

711　transgenerational plasticity values.

	m_{EE}	m_{EL}	m_{LE}	m_{LL}
EE ACF	0.757139	-0.131583	-0.049501	-0.156768
EL ACF	-0.036229	0.5968488	-0.069245	-0.153818
LE ACF	0.1256068	-0.036048	1.3390581	-0.067451
LL ACF	0.0135772	-0.017808	-0.120773	0.9757164
Within ACF	-0.067861	-0.160476	0.0595802	0.0701295

Figure 1: Schematic depicting the ecological motivations (summer annual plants) and theoretical

underpinnings for the evolutionary modeling of plasticity traits (A and C), and the types of

environmental fluctuations that may influence their evolution. (A) Temperature plasticity model. In

the abbreviations, **E** denotes the Early growing season (spring) and **L** denotes the late growing

season (summer). The first letter represents the relevant season during the parental generation and

the second letter represents the relevant season in the offspring generation (e.g., EL denotes effects

of parental early growing season temperature on offspring phenotypes late in the growing season).

Within generation developmental changes in response to early season environment (W) are also

considered in this model. Additionally, the longterm average environmental conditions at a specific

area determine the genetic baseline phenotype of an individual (G). (B) On the left we see an

example of an environment with high within season autocorrelations for temperature (hot springs

tend to be followed by hot summers), but low inter-annual autocorrelations (a hot year does not

tend to be followed by another hot year) that selects for within generation plasticity but not

transgenerational plasticity. On the right, a situation where spring and summer temperatures are not

correlated with each other, but we do find that environmental oscillations lead to a string of warmer

than average springs and cooler than average summers, in this situation transgenerational plasticity

(EE and LL) but not within generation plasticity is expected to be optimal. (C) Precipitation

plasticity model. The amount of precipitation experienced by an individual can lead to

transgenerational effects in the next generation (T), as well as persist for two (TG) or three (TGG)

generations. (D) On the left, relatively gradual decadal oscillations give value to transgenerational

effects that persist for multiple seasons (T, TG, and TGG). On the right shorter period climatic

oscillations may favor parental effects (T), but not multi-generation effects (TG or TGG).

Figure 2: Maps depicting natural climatic variation across the conterminous US.

Figure 3: Distributions of optimal plasticity (A and B) and temperature (C-G) values across all 4km x

4km sites in the US. Histograms of optimal (A) precipitation transgenerational plasticity value (T),

(B) precipitation multi-generation persistence (G), (C) temperature spring (early season) -> spring

(early season) transgenerational plasticity (m_{EE}) , (D) temperature spring (early season) -> summer

(late season) transgenerational plasticity (m_{EL}), (E) temperature summer (late season) -> spring(early

season) transgenerational plasticity (m_{LE}), (F) temperature summer (late season) -> summer (late

season) transgenerational plasticity (m_{LL}), and within generation temperature plasticity (W)

Figure 4: Maps coded to show patterns of variability for optimal precipitation (A-C) and temperature

(D-H) plasticity values across the US. (A) Optimal transgenerational plasticity values for the one-

generation transmission of precipitation level information. (B) Optimal grandparental

transgenerational plasticity values coded blue (green) or red (orange) based on the direction of effect

(positive or negative). White regions have an optimal multi-generation persistence (G) of 0, while

red and blue both have optimal multigeneration persistence of 1, intermediate values (0>G>1) in

orange and green. (C) The difference between optimal transgenerational plasticity values in the raw

vs. residual variant of the mode. Higher values suggest that the primary value associated with

transgenerational plasticity over the past 120 years has been associated with allowing individuals to

keep up with linearly changing precipitation patterns. (D) Optimal total levels of transgenerational

temperature plasticity ((M_{EE} + M_{EL} + M_{LE} + M_{LL})/2). (E) Optimal transgenerational plasticity of

most extreme positive transgenerational plasticity allele. (F) Optimal transgenerational plasticity of

lowest transgenerational plasticity allele. Regions in orange have at least one form of

transgenerational plasticity for which negative transgenerational effects increase fitness. (G) The

difference between optimal transgenerational plasticity values in the raw vs. residual variant of the

mode. Higher values suggest that the primary value associated with transgenerational plasticity over

the past 120 years has been associated with allowing individuals to keep up with increasing

temperature. (H) Optimal within generation plasticity (W) values.

Figure 5: Fitness landscapes of transgenerational precipitation alleles for twelve sites across the US.

Sites with low optimal parental effects (D and I) have only very subtle fitness differences associated

with chanes in the multigeneration persistence (Y-axis) due to the minor role in any form of

transgenerational effect on fitness in these cases. More defined fitness peaks tend to occur in areas

where more substantial transgenerational effects are optimal (E, G, J, K, L). In some rare cases,

bimodal fitness landscapes arise (B) where lines with either positive (with high persistence) or

negative (with low persistence) transgenerational persistence have higher fitness than lines with no

transgenerational inheritance.

Figure S1: Distribution of climatic summary statistics across the 481k 4x4km grids in the US.

Dotted lines at 0 for autocorrelation histograms.

Figure S2: Mosaic plot showing the frequency of specific combinations of optimal transgenerational

effects and multi-generational persistence values. More subtle transgenerational effects more

frequently only have a single generation of persistence (G=0), while more extreme transgenerational

effects tend to coincide with full transgenerational persistence (G=1) where each of the prior three

years contribute equally. Additionally, positive transgenerational effects were more likely to have

intermediate levels of persistence than negative transgenerational effects.

Figure S3: Figures demonstrating tradeoffs between classes of plasticity. (A) Tradeoffs between

early-late (m12) and late-late (m22) transgenerational plasticity in relation to the autocorrelations in

summer temperature. Generally, as inter-annual summer autocorrelations increase, so too does the

frequency of minor and moderate early-late transgenerational plasticity. However, we also find that

areas that favor higher levels of late-late transgenerational plasticity tend to favor lower levels of

early-late plasticity compared to other sites with similar levels of temperature autocorrelation. (B)

Within a single site there are many examples of localities where either early-late or late-late

transgenerational plasticity lead to fitness increases relative to clones with no plasticity, but

individuals expressing both forms of plasticity have the lowest fitness of all.

Figure S4: Changes in the frequency of ten clonal types in a 120 year simulated competition

experiment for four different sites across the US. The top three final frequency clones are denoted

with stars.

Precipitation Plasticity Model

A — Number of Sites vs Parental Effect

B — Multigenerational Persistence

Temperature Plasticity Model

C — Spring -> Spring Transgenerational

D — Spring -> Summer Transgenerational

E — Summer -> Spring Transgenerational

F — Summer -> Summer Transgenerational

G — Within-Generation

A   4   Precipitation   D   Temperature

B   C   Plasticity Allele Value
-.3   .6

E   G

F   H

Table S1

Click here to access/download
Other (Video, Excel, large data files)
TableS1.xlsx

Table S2

Click here to access/download
Other (Video, Excel, large data files)
TableS2.xlsx