Environmental oscillations favor the evolution of adaptive transgenerational plasticity

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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.

Author Summary

Parental environments can alter progeny development, a phenomenon that has received renewed

focus as interest in epigenetic inheritance has surged. Mathematical models indicate that these effects

can evolve to adaptively match progeny phenotypes and environments when conditions change

predictably, but few studies have explored whether such conditions actually exist in nature. This

study reveals that patterns of precipitation and temperature variation in many regions of the US

should favor the adaptive evolution of parental effects, but the optimal extent of these effects varies

widely. Patterns of environmental fluctuations are heterogeneous across the landscape such that not

only will there be variation in optimal phenotypes across space, but that there will also be variation

for the optimal system of inheritance.

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 (1). 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 (2,3). Over the last three decades, it has become clear that *transgenerational plasticity* is also remarkably common (reviewed by (4-9), yet highly variable (10-12) and nearly absent in some cases (13). 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 (note that these mechanisms are not mutually exclusive; (14,15). 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, 8), and more generally parental x current environment interaction effects appear prevalent (10,11, 16, 17). 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 (18). Theoretically, these streams of information can be integrated in order to fine-tune the expression of a given trait for the selective environment that is most likely to be encountered. Whether organisms have in fact evolved to express adaptive transgenerational plasticity, and under what situations adaptive transgenerational plasticity is expected to evolve, has been the subject of intense research in recent years.

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 over time (19,20). For instance, within-

generation adaptive plasticity is expected to evolve when individuals encounter variable conditions

during their lives by either moving to different environments in a heterogeneous landscape, or by

conditions changing temporally and with some degree of predictability within an individual's

lifetime. Furthermore, there must be little to no cost of responding to environmental cues and

maintaining the molecular machinery to do so, and there must be genetic variation for individual

phenotypic responses to the immediate environment (see e.g., 21,22; reviewed by 23,24).

These same factors – predictable environmental variation and genotype-by-environment

interaction – also influence the evolution of transgenerational plasticity (18,25-30). 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. Such

intergenerational environmental correlations occur when the same environmental variable correlates

with itself across years, as is the case for temperature when hot years are followed by hot years (i.e., a

positive autocorrelation in temperature), or when several environmental factors change

simultaneously to indicate the changing of seasons (31). 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 (10, 32-35).

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. Thus, understanding how patterns of

variation and predictability favor or hinder different forms of phenotypic expression represents an

unexplored and potentially key factor in determining the populations that are most at risk under

future climate change scenarios. 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 environmental 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., 36-46; reviewed

by 6,7,14). For instance, when *Mimulus guttatus* plants experience herbivory, their offspring increase

production of defensive leaf trichomes (11,47,48). 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 (36).

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. We found 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, we constructed each model using raw

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

considering both raw data values and residuals in tandem, we are able to parse out the role of

transgenerational effects and within generation plasticity in periodically oscillating environments

(residual model), as well as those with periodic oscillations in addition to a linear change in mean

temperature (raw data model). In all other ways, the two variants of this model are identical.

Differences in the results of raw vs. residual models highlight 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., 49,50), and in some cases for

far longer (51,52). 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. In both of these models we calculate the fitness of genotypes as a product of the

difference between the individual's phenotype value and the environmental optimum for that year.

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 (54), 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.

Precipitation model

For a given generation T the expected precipitation (phenotype) is defined as $\dot{P}_T$, while the

actual precipitation is P_T. The two parameters considered in this model are m (maternal or parental

effect) and g (multigenerational persistence). The m value in this model alters the expected

precipitation (phenotype) for a given year $(\dot{P}_T)$ based on how deviant the precipitation was during a

preceding year or years $(P_{T-X} - \bar{P})$. In other words, m adjusts offspring phenotype based on how

different parental environment was from long term average environment. When there are

interannual autocorrelations, values of m that match the degree of interannual autocorrelation will

be adaptive.

$$\dot{P}_T = \bar{P} + \frac{m(P_{T-1} - \bar{P}) + mg(P_{T-2} - \bar{P}) + mg^2(P_{T-3} - \bar{P})}{1 + g + g^2}$$

When $m = 0$ the expected precipitation will always be equivelant to the mean precipitation

at a site, when $m = 1$ and $g = 0$, the expected precipitation will always equal the precipitation

experienced the prior year. In the simplest case when $g = 0$ only the prior year's information alters

the expected precipitation. The g value in this model weighs how much the precipitation

experienced two and three generations back should change expected precipitation values relative to

the prior year's precipitation. At a value of $g = 1$ the previous three years will all contribute equally

to the expected precipitation. As g decreases, the relative information from two generations back

decreases linearly, and the relative information contribution from three generations back decreases

quadratically. At $g = 0.5$, the grandparental precipitation will contribute half as much information

as the parental generation, and the great-grandparental generation will contribute one quarter the

information of the parental generation. In other words, the g parameter represents how much the

transgenerational information diminishes each year after the first. The cumulative effect of the

previous generations environment is scaled by a factor of $(1 + g + g^2)$ in order to standardize the

total contribution of transgenerational effects for a given value of m independent of g. That is to

say, m affects the total amount of transgenerational information transmitted between generations,

and g effects the distribution of the information disseminated from each of the prior three years.

A test of a random subset of 2000 sites led us to restrict the values of m we investigated to

between -0.6 and 0.9 at increments of 0.1 (from a possible range of -1 to 1). Values less than -0.6

and greater than 0.9 were not optimal in any of the 2000 sites and were therefore excluded for

computational reasons. The full range of g values (i.e., 0 to 1) was retained. We considered genotypes

for every possible combination of m and g values selected at intervals of 0.1 (for a total of 16×11

= 176 genotypes). For each genotype we calculated expected phenotype for every year between

1897-2015. Next, for each genotype and each year, we calculated the difference between the

expected phenotype and the actual phenotype for that year.

273
$$w_{MT} = 1 - \frac{\left| \dot{P}_T - P_T \right|}{\bar{P}}$$

Geometric mean fitness over the 118 years was calculated for each of the 176 genotypes.

The values of m and g for which fitness was maximized was recorded for each of the 481,631 sites

and plotted in qGIS. This approach identifies the most fit genotype at a given site over the

generations, or more explicitly, the genotype that would be at the highest frequency in the year 2015.

In order to compare evolutionary dynamics in more detail, we constructed fitness

landscapes and compared the relative frequency of competing genotypes over time for twelve sites

representing a wide range of geographic and parameter space [may need to say more about the

parameter space here]. For these sites, the mean fitness across the 176 genotypes was calculated for

the first year (1897). The fitness of each genotype in that year was then divided by the mean fitness

to produce the relative fitness of each genotype. This relative fitness was then used as the relative

frequency of each genotype for the following year. For each year following the first, the absolute

fitness of each genotype was divided by a weighted mean of the fitness of all individuals for that year

($w_X^{Rel} = w_X^{Abs} / \sum_{i=1}^{154} w_i^{Abs} * Freq(i)$) to calculate the relative fitness of each genotype for each

generation. This value was then multiplied by the frequency of that genotype in that generation to

get the following generation's frequency. The final generation frequency was used for the

construction of fitness surface heatmaps.

Temperature model

The general framework of the temperature model is similar, but rather than allowing for the

persistence of transgenerational effects across multiple years, it considers early and late season

transgenerational effects, as well as within-generation plasticity. We ran the same model on raw data

and residuals, calculated as in the precipitation model. Annual plants were used as the motivation for

this model, with a growing season from March-September. The growing season was broken down

into three parts: early growing season (March, April, May), transitional growing season (June), and

late growing season (July, August, September). Within this model there are two bouts of selection,

one after the early growing season, and one after the late season. While early season phenotype can

only be modified by the previous year's temperature (because organisms are newly born in the spring

in this scenario, and therefore cannot respond themselves to temperatures), late growing season

phenotype can be modified by the previous year's temperature, and/or the temperature earlier in the

growing season. Additionally, this model allows for both early and late season phenotype to be

modified by both the previous season's early and late temperatures independently. In other words,

early season phenotype can be modified by the previous year's early season temperature, or the

previous year's late season temperature; late season phenotype can be modified by these same two

factors as well as the current year's early season temperature (i.e., by within-generation plasticity).

Thus, this model includes five plasticity parameters, reflecting the effect of 1) the previous

generation's early season temperature on current early season phenotype (m_{EE}), 2) the previous

generation's late season temperature on current early season phenotype (m_{LE}), 3) the previous

generation's early season temperature on the current late season phenotype(m_{EL}), 4) the previous

generation's late season temperature on current late season phenotype (m_{LL}), and 5) the current

early season temperature on the current late season phenotype (b). Note that the mean and variance

of temperature data are not correlated, unlike precipitation data. For this reason, fitness is scaled by

a constant 20 rather than the mean temperature of a given site. Results were exceedingly similar

when fitness was scaled by constants of 10 and 100; 20 was chosen as it is closer to the mean annual

temperature across the US.

$$\dot{P}_{TE} = \overline{P_E} + m_{EE}(P_{ET-1} - \overline{P_E}) + m_{LE}(P_{LT-1} - \overline{P_L})$$

$$w_E = 1 - \frac{|\dot{P}_{TE} - P_{TE}|}{20}$$

$$\dot{P}_{TL} = \overline{P_L} + m_{EL}(P_{ET-1} - \overline{P_E}) + m_{LL}(P_{LT-1} - \overline{P_L}) + b(P_{ET} - \overline{P_E})$$

$$w_L = 1 - \frac{|\dot{P}_{TL} - P_{TL}|}{20}$$

$$w_F = w_E * w_L$$

The geometric mean of genotype fitness was taken over the 119 years for each of the

genotypes. For each of the five plasticity (transgenerational and within generational) terms in this

model we considered possible values of -0.2 (negative plasticity), 0 (no plasticity), 0.1 (minor

plasticity), 0.3 (moderate plasticity), and 0.5 (major plasticity). While this resolution is somewhat

coarse, focusing on this subset of values allowed us to consider this model for the full factorial

combination of parameter space. In total, fitness was calculated for 3125 genotypes, across each of

119 years, at all 481631 locations, for a total of 179 billion measures of fitness. The values of

plasticity at which fitness was maximized were recorded from each grid point across the US. Finally,

for the same 12 focal sites used in the precipitation model, we calculated the changes in genotype

frequency across 119 years using the same methodology as above. All code for these simulations,

and other modeling results are available at (github.com/XXXX).

Results

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

US (Figure 2 and S1). Mean annual temperature and precipitation are both primary forces driving

local adaptation to the diversity of climate regimes across the US, 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 or 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 (55,56) 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.

Patterns of climatic oscillation varied substantially across the US over the last 120 years, suggesting

the potential for climatically driven variation in locally adaptive plasticity (Figure 2 and S1, Table 1

and S1). We examined these patterns by calculating autocorrelations in annual temperature and

precipitation between successive years. The climate (temperature or precipitation) of a given year

and that of the prior year(s) were often significantly correlated across the US. However, the

magnitude and direction of correlations varied depending on the region and climatic parameter.

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 was higher mean and standard deviation for lag-2 partial ACF than lag-1 ACF (Figure S1).

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

ACF >0.2) and significantly negative (5,088 lag-2 PACF <-0.2 vs. 441 lag-1 ACF <-0.2) lag-2 partial autocorrelation (PAC) than lag-1 ACF. These correlations 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 the variation in inter-annual precipitation variation (Figure 2). By contrast, across the southeastern US, negative lag-1 ACF and lag-2 PACF suggest a negative feedback dynamic in which multiple dry years tend to be followed by a wet year, and vice versa (Figure 2).

Patterns of temperature autocorrelations extended over larger regions, and were often substantially more extreme, compared to the more complex 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).

Spring and summer temperatures displayed significant autocorrelations at lags 2 and 3. 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 vast variability of climatic autocorrelations lead to a great deal of variation in the optimal levels of plasticity in the precipitation evolutionary model (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) 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. For example, if the mean annual precipitation at a

site is 1000mm (dictating the genetic value), and the previous year's precipitation was 1100mm, then

the phenotype of the current generation would be 1010mm (1000*0.9 + (1,100-1000)*0.1) in the

case of an allele value of 0.1, and 990mm in the case of an m value of -0.1 (assuming there are no

multigenerational environmental effects). 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) 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 0.057 lower in the residual

variant of the precipitation model compared to the raw data 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)

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; e.g., 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 transgenerational plasticity 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

503   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. Fitness landscapes provide a snapshot of the final frequency at the end of the 120 years

of the modeling run, which is the product of differences in fitness within each generation and

subsequent changes in frequency (Figure 6). 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 6).

530   Fitness landscapes for the temperature model are substantially more complex due to the

larger number of terms, but do suggest that patterns of environmental variation can favor tradeoffs

between multiple forms of plasticity. As the effect of parental early season temperature on offspring

late season phenotypes (m_{EL}) increased, optimal m_{LL} values decreased (Figure S4) and vice versa.

The same is true for m_{EE} and m_{LE}. At a given stage in the offspring's life cycle there is only so

much information value that can be gleaned from the environments of previous generations; this

total value ($m_{EL} + m_{LL}$) is one factor that varies across the US in this model. The optimal balance

between the two forms of transgenerational plasticity that affect a given life history stage (m_{EL} :

m_{LL}) is the other parameter that varies, and this balance is what controls the degree to which the

early vs. late season parental conditions will modify offspring development at a given stage.

Discussion

Although transgenerational environmental effects on phenotypic expression have been

recognized for decades (57,58), interest in these effects has surged recently due to increased

appreciation for the potential role of transgenerational plasticity in adaptation (59). 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 60).

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., 12,18,21,61). 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

The availability of water is one of the environmental factors that most directly impacts

terrestrial plant life. 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 (62). 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., 35, 63), providing a third route by which plants can fine

tune their 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

(16, 38). 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 (10).

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 compared to that of their

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

but the most common transgenerational effect values were lower.

Intriguingly, multigenerational persistence values of 0 (18.7% of sites) and 1 (40.7% of sites)

were the most common, representing strategies in which transgenerational effects lasted only a

single generation or persisted fully through 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. By contrast, complete absence of parental effects was

favored in only 7% of sites.

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. Our modeling results suggest that genetic variation for

transgenerational precipitation effects may stem in part from fine-scale spatial variation in

precipitation autocorrelations. We predict that such variation in precipitation autocorrelations

translates into variable selection for transgenerationally plastic responses to precipitation, resulting in

the maintenance of genetic variation for transgenerational plasticity within species.

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 (64), and significantly shifts

flowering time (31) and many other phenotypes (65) in genotype-specific ways. Additionally, recent

work has demonstrated the lasting impact of temperature regime on *Arabidopsis thaliana* plants for

generations to come (12,49,66,67).   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. These results suggest that early spring temperatures may be a harbinger of

temperatures to come later in the same growing season and the latter half of the following growing

season but provide little information regarding the following year's early season temperature.

In our framework, temperature plays two roles related to fitness at both the early and late

growing season time points. First, temperature 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. Second, temperature is a source of information that a plant can use to alter its phenotype

later in the growing season, and to alter the phenotype of its offspring in the following year. As a

result, five classes of plasticity are available to plants in this model in order to best prepare

themselves and their progeny for the most likely temperature conditions in the future. These

plasticity classes include within-generation plasticity and four classes of transgenerational plasticity

representing all permutations of early and late season temperature effects across the parent and

offspring generations.

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 generally

provide valuable information that allows transgenerational plasticity to improve the match between

phenotypes and temperature regimes.

The most common form of transgenerational plasticity observed in this model entailed

effects of late-growing season temperature on 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 (68). For instance,

cool temperatures in spring do not generally foreshadow similarly cool temperatures in late summer,

but warm summer temperatures in the previous generation do correlate with warm temperatures in

the following summer. 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 greatly across the US.

Although we identified general patterns in temperature autocorrelations, a key finding was

that these autocorrelations varied considerably across the U.S., leading to vastly different optimal

levels of within- and transgenerational plasticity among sites. 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 (69). 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

(31,70). 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 for species that set seed in late summer. Intriguingly, parental

effects of temperature on germination and flowering time are highly genetically variable in

Arabidopsis thaliana (12,70,71). In *Plantago lanceolata*, such genotype-by-maternal temperature effects

persist throughout the offspring life cycle to generate variation in offspring reproduction in the field

(72). 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 12).

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 raw 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). We also found that populations in regions undergoing the most

severe climate change are expected to benefit most by increasing transgenerational plasticity.

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 (73), but our models do not

address this issue. Instead, we set the baseline phenotype of an organism as the optimal phenotype

for the average conditions over the 120 years for which we have data, and then compared the fitness

of genotypes that vary in their expression of plasticity. 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 transgenerational plasticity (12). For instance, in some scenarios

transgenerational effects can increase fitness in the short term, while reducing it in the long term

(74). 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. Answering both of these questions in

the same system will be necessary to determine the likelihood that selection on transgenerational

plasticity is a viable route to adaptation in the face of changing conditions.

Temperature and precipitation autocorrelations and optimal plasticity values likely stem in

part from the same broad-scale climatic oscillations, such as the El Niño Southern Oscillation (75),

the Quasi-biennial oscillation (76), and the Pacific Decadal Oscillation (77,78). 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 (79). 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 (80). 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. Another critical goal of future research is to better

understand how within- and transgenerational plasticity interact to influence phenotypic expression.

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.

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

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)

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1002   Table 2: Most common, second most common, and mean optimal plasticity values across all sites in

1003   the US.

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		Raw		Residual	
Climate Term		#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)

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1023   Table 3: Correlations between the temperature correlation between seasons and the optimal

1024   transgenerational plasticity values.

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	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

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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).

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1069 Figure 2: Maps depicting natural climatic variation across the conterminous US.

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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)

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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.

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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.

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1107 Figure 6: Changes in the frequency of ten clonal types in a 120 year simulated competition

1108 experiment for four different sites across the US. The top three final frequency clones are denoted

1109 with stars.

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1111 Figure S1: Distribution of climatic summary statistics across the 481k 4x4km grids in the US.

1112 Dotted lines at 0 for autocorrelation histograms.

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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.

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1121   Figure S3: Multi-panel fitness landscape comparing the relative fitness of clonal types with different

1122   levels of the four transgenerational temperature plasticity terms.

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Figure 1

Figure 2

Figure 3

Precipitation Plasticity Model

A — Parental Effect (x-axis: -0.6 to 0.9); Number of Sites (y-axis: 0 to 100,000)

B — Multigenerational Persistence (x-axis: 0 to 1)

Temperature Plasticity Model

Number of Sites (y-axis: 0 to 400000)

C — Spring -> Spring Transgenerational (x-axis: -0.2, 0, 0.1, 0.3, 0.5)

D — Spring -> Summer Transgenerational (x-axis: -0.2, 0, 0.1, 0.3, 0.5)

E — Summer -> Spring Transgenerational (x-axis: -0.2, 0, 0.1, 0.3, 0.5)

F — Summer -> Summer Transgenerational (x-axis: -0.2, 0, 0.1, 0.3, 0.5)

G — Within-Generation (x-axis: -0.2, 0, 0.1, 0.3, 0.5)

Figure 4

Precipitation

Temperature

Plasticity Allele Value

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Figure 5

Figure 6

Figure S2

Figure S3

Figure S4