Environmental variation mediates the evolution of anticipatory parental effects

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Abstract

Environments vary over time and if this variation is predictable, environments that are

similar across generations should favour evolution of anticipatory parental effects to

benefit offspring. In contrast, the absence of correlation between parental and

offspring environments should select against parental effects. However, experimental

evidence is scarce. We investigated the evolution of maternal effects using

experimental evolution. Populations of the nematode *Caenorhabditis remanei*,

adapted to 20°C, were exposed to a novel temperature (25°C) for 30 generations with

either positive or zero correlation between parent and offspring temperature. We

found that populations evolving in environments with positive correlations had a

positive maternal effect, since they required maternal exposure to 25°C to achieve

maximum reproduction and fitness in 25°C. In contrast, populations evolving under

zero correlation had lost this positive maternal effect. This shows that parental effects

can aid population viability in warming environments. Correspondingly, ill-fitting

parental effects can be rapidly lost.

Key words: *Caenorhabditis*, Environmental heterogeneity, Maternal effects,

Reproduction, Temperature, Transgenerational plasticity

Introduction

The role of environmental variation in the adaptive expression of phenotypes has

gathered considerable interest (Chevin *et al.* 2010; Hollander *et al.* 2015; Donelson *et*

al. 2017; Lind & Spagopoulou 2018). Not only is heterogeneity common, it is also

predicted to change evolutionary outcomes. While stable environments generally

should select for genetic specialisation, environmental heterogeneity can select for

environmental input on this process. Variable environments is expected to select for

phenotypic plasticity (if environmental cues are reliable) or bet-hedging (if the cues

are not reliable) (Moran 1992; Simons 2011). However, a developing organism may

not be able to acquire and/or interpret the environmental cues itself; therefore, the

parental environment can also function as a developmental cue (Marshall & Uller

2007; Leimar & McNamara 2015). Consequently, recent theory maintains that high

environmental correlation between generations can select for adaptive parental effects

and/or epigenetic inheritance of an environmentally induced phenotype (Lachmann &

Jablonka 1996; Bonduriansky & Day 2009; Rivoire & Leibler 2014; Kuijper & Hoyle

2015; Leimar & McNamara 2015; Uller *et al.* 2015; Proulx & Teotónio 2017),

mechanisms collectively referred to as transgenerational plasticity (Donelson *et al.*

2017). The sign of the environmental correlation is expected to result in a similar sign

of the parental effect, so that parents can prepare their offspring for the same

environment as they are themselves experiencing (positive correlation), the opposite

environment (negative correlation) or does not influence the phenotype of their

offspring (zero correlation between parent and offspring environments). The

exception is constant (perfectly correlated) environments, where genetic specialisation

is predicted to evolve (Leimar & McNamara 2015) together with a negative trans-

generational effect in order to reduce phenotypic variance (Hoyle & Ezard 2012;

Kuijper & Hoyle 2015).

However, while the theory is well developed, the empirical evidence is mixed. In its

most basic form, the theory of anticipatory parental effects predicts that offspring

should have higher performance if parental and offspring environments are matching

(Marshall & Uller 2007; Burgess & Marshall 2014). There are some striking

examples of this (e.g. Gustafsson *et al.* 2005; Galloway & Etterson 2007; Jensen *et al.*

2014; Kishimoto *et al.* 2017; Ryu *et al.* 2018; Toyota *et al.* 2019), but across studies

there is only weak support for this prediction in natural systems, and the effects are

small compared to the direct effects of offspring environment (reviewed in Uller *et al.*

2013). Since environmental predictability across generations is seldom quantified, one

reason for the scarcity of clear examples of adaptive parental effects is that

environments may not often be correlated across generations, and, therefore, provide

little opportunity for selection for such anticipatory effects (Burgess & Marshall

2014). Even if natural environments are correlated, the evidence for stronger parental

effects in more stable environments is mixed, and varies between traits (Walsh *et al.*

2016). Thus, there is a call for studies estimating parent and offspring fitness when the

environmental predictability between generations are well known, and maternal

effects are expected to be under selection (Burgess & Marshall 2014).

Direct experimental evidence for the evolution of positive anticipatory parental effects

of an environmentally induced parental phenotype is currently lacking. On the other

hand, if parents and offspring live in negatively correlated environments, the parental

phenotype should not be inherited, but the parents can still anticipate the offspring

environment. As such, the evolution of a negative parental effect could be adaptive, a

prediction that has recently received experimental support in a study by Dey *et al.*

(Dey *et al.* 2016), who found that populations of the nematode *Caenorhabditis*

elegans that evolved under strictly alternating hypoxia-normoxia conditions in every

generation evolved a negative maternal provisioning effect. This suggests an adaptive

benefit of maternal effects when the maternal environment is a strong cue for the

offspring environment (in this case a perfect negative correlation), as predicted by

theory (Uller *et al.* 2015; Proulx & Teotónio 2017). More generally, theory also

predicts that positive trans-generational correlations would result in the evolution of a

positive parental effect and, importantly, if the environmental state is uncorrelated

across generations, parental effects would be maladaptive and selected against

(Lachmann & Jablonka 1996; Kuijper & Hoyle 2015; Leimar & McNamara 2015;

Uller *et al.* 2015). The latter is considered a reason why adaptive parental effects are

generally weak (Uller *et al.* 2013). However, these scenarios have not been

investigated experimentally. Moreover, previous studies on evolution of parental

effects under environmental heterogeneity have investigated only the case of non-

overlapping generations (Dey *et al.* 2016). Most natural populations have however

overlapping generations and age structure, which can influence evolution in both

stable and heterogeneous environments, especially with respect to the evolution of life

history strategies (Cotto & Ronce 2014).

Taken together, environmental heterogeneity and environmental correlations over

generations predict the adaptive value of parental effects (Lachmann & Jablonka

1996; Kuijper & Hoyle 2015; Uller *et al.* 2015). We set out to test this using

experimental evolution in the dioecious free-living nematode *Caenorhabditis*

remanei, adapting to different temperature regimes. Genetically heterogeneous

populations, previously adapted to 20°C, were evolving for 30 generations in control

conditions or adapting to 25°C, in either constant 25°C, increased warming to 25°C or

a heterogeneous environment with fluctuating temperatures. We found positive

anticipatory maternal effects on reproduction in populations evolving in environments

that were positively correlated across generations. Moreover, we found the evolution

of a reduced maternal effect on reproduction in heterogeneous environments where

parent and offspring environments were not correlated during experimental evolution.

Material and methods

Experimental evolution

As founder population, we used the wild-type SP8 strain of *C. remanei*, obtained from

N. Timmermeyer at the Department of Biology, University of Tübingen, Germany.

This strain was created by crossing three wild-type isolates of *C. remanei* (SB146,

MY31, and PB206), harbour substantial standing genetic variation for life-history

traits (Chen & Maklakov 2012; Lind *et al.* 2017), and has been lab-adapted to 20°C

for 15 generations prior to setup of experimental evolution.

Experimental evolution was conducted for 30 generations in three climate cabinets;

one set to 20°C, one to 25°C and one with a slowly increasing temperature (see

below). Five selection regimes were used. *Control 20°C* was experiencing 20°C for

30 generations, and *Warm 25°C* was experiencing 25°C for 30 generations. *Increased*

warming started in 20°C, the cabinet temperature was then raised by 0.1°C every 2.13

day (rounded to whole days) to reach 25°C the last day of selection. *Slow temperature*

cycles spend their first 10 generations in 20°C, were then moved to the 25°C cabinet

for 10 generations, to finish the last 10 generations in the 20°C cabinet. Finally, the

Fast temperature cycles regime were moved between 20°C and 25°C every second

generation, thus experiencing 15 temperature shifts.

Generation time in 20°C and 25°C was defined as the average difference in age

between parents and offspring (Charlesworth 1994) and was calculated from the life-

table of age-specific reproduction and survival by solving the Euler-Lotka equation

(Stearns 1992; Charlesworth 1994) with trial data from the SP8 lines, and was 4.0

days in 20°C and 3.4 days in 25°C. This resulted 120 days of selection for *Control*

20°C, 114 days for *Slow temperature cycles*, 110 days for *Increased warming* and

Fast temperature cycles and 101 days for *Warm 25°C*. For the two temperature cycle

treatments, the worms spent shorter chronological time in 25°C than in 20°C, because

of the faster generation time in 25°C. This ensured equal exposure to the two

temperatures over biological time.

Experimental evolution was conducted on 92 mm NGM-plates (Stiernagle 2006) and

to combat infections the agar and bacterial LB also contained the antibiotics

kanamycin and streptomycin, and the fungicide nystatin (Lionaki & Tavernarakis

2013; Lind *et al.* 2016). The plates were seeded with 2mL of the antibiotic resistant *E.*

coli strain OP50-1 (pUC4K) obtained from J. Ewbank at the Centre d'Immunologie

de Marseille- Luminy, France. To keep worm populations age-structured in

overlapping generations, the lines were always kept in experimental growth face by

cutting out a bit of agar containing 150 individuals of mixed ages and transferring this

to freshly seeded plates. Transfer was conducted when needed (every 1-2 day), always

before food was finished. Six independent replicate populations of each selection

treatment were set up, resulting in a total of 30 populations. All populations were

expanded for two generations and frozen after 30 generations.

Phenotypic assays

Before assays, worms were recover from freezing and grown 2 generations in

common garden, each generation synchronized by bleaching, a standard procedure

that kills all life-stages but eggs (Stiernagle 2006). The common garden temperature

was 20°C or 25°C (see below).

Fitness assays were performed to test for local adaptation to the selective regime and

the evolution of adaptive trans-generational effects. We therefore carried out three

assays, by varying parental temperature (the 2 generations of common garden after

defrosting the lines) and testing temperature for offspring. The *20°C - 20°C assay* had

both common garden and testing in 20°C. This is the environment the *Control 20°C*

lines have experienced. Likewise, in the *25°C - 25°C assay* both parents and testing

worms experience 25°C, which is the selective environment for *Warm 25°C* and very

close to the final environment for *Increased warming*. Finally, the *20°C - 25°C assay*

have 20°C as parental temperature, while the testing worms have their whole

development in 25°C. This assay represents strong temperature fluctuations between

generations, which is the selective environments for the *Fast temperature cycle* lines,

and by comparing this assay to the *25°C - 25°C assay* we can estimate the importance

of trans-generational effects on fitness when adapting to a novel environment.

The assays were initiated by synchronised egg-laying in the testing temperature by 40

females of each population. After 5h, females were killed by bleaching, and setup of

L4 larvae was initiated 39h later (in 25°C) or 50h later (in 20°C), due to temperature-

specific development time. The setup consisted of 8 testing females per plate, together

with the same number of background males from the SP8 line. Sex ratio was kept 1:1

throughout the experiment by adjusting the number of males to match the number of

females alive. Age-specific fecundity was measured by each day allowing the females

3h of egg-laying on an empty plate, where after the females were returned to a new

plate (together with the males) and the number of hatched offspring on the egg-laying

plate were counted 2 days later. The exact time the females were added to and

removed from each plate was noted, and the number of offspring was corrected by

exact number of minutes available for egg laying, and the number of females alive.

Thus, we did not collect individual level data on total reproduction, but daily

snapshot, in order to increase the number of individuals assayed and improve the

reproduction estimate of each population. Daily reproduction was collected until all

reproduction had ceased. Four replicate plates of each population was set up, and for

the *20°C - 20°C* and *20°C - 25°C* assays the replicates were evenly split between two

climate cabinets per temperature. However, for logistical reasons, the *20°C - 25°C*

assay was reduced. We excluded the *Slow temperature cycle* treatment from this

assay, and unfortunately we lost two *Warm 25°C* populations during common garden

(due to overcrowding and subsequent starving, which is known to induce epigenetic

effects (Rechavi *et al.* 2014) and therefore these populations were excluded), leaving

us with 4 replicate population of this treatment. This resulted in 30 replicate

populations and 960 female worms for the *20°C - 20°C* and *20°C - 25°C* assays, and

23 replicate populations and 736 female worms for the *25°C - 25°C assay.*

Statistical analyses

The age-specific reproduction data was analysed as rate-sensitive individual fitness

λ_{ind} (Brommer *et al.* 2002) as well as the total reproduction of each replicate plate,

adjusted to the number of females present each day. Individual fitness encompasses

the timing and number of offspring and is analogous to the intrinsic rate of population

growth (Stearns 1992) and was calculated by solving the Euler-Lotka equation for

each replicate plate. Individual fitness and total reproduction in 20°C was analysed in

separate mixed effect models with election treatment as fixed effect and population

and cabinet as random effects. Both response variables were log-transformed before

analysis. In 25°C, individual fitness and total reproduction was analysed with

selection treatment and parental temperature as crossed fixed effects, and replicate

population as random effect. Since the *25°C - 25°C assay* was conducted in only one

cabinet, and moreover the *Slow temperature cycle* treatment was not run, the random

effect of cabinet was excluded from the models, as was the *Slow temperature cycle*

treatment.

Results

For individual fitness (λ_{ind}) in 25°C, we found a significant selection regime ×

parental temperature interaction (Selection regime: $\chi^2 = 1.302$, df = 3, p = 0.729;

Parental temperature: $\chi^2 = 0.114$, df = 1, p = 0.736, Selection regime × Parental

temperature: $\chi^2 = 15.562$, df = 3, p = 0.001). This interaction was caused by

significantly opposite slope for *Fast temperature cycles* compared to *Increased*

warming and *Warm 25°C*, with highest fitness in 25°C for *Fast temperature cycles*

when parents were grown in 20°C, while highest fitness in 25°C for *Increased*

warming and *Warm 25°C* was achieved when their parents were also grown in 25°C

(Figure 1A).

A significant selection regime × parental temperature interaction was also found for

total reproduction in 25°C (Selection regime: $\chi^2 = 0.995$, df = 3, p = 0.802; Parental

temperature: $\chi^2 = 41.373$, df = 1, p < 0.001, Selection regime × Parental temperature:

$\chi^2 = 10.747$, df = 3, p = 0.013), caused by a much weaker positive effect of parental

exposure to 25°C for *Fast temperature cycles* than for the other selection regimes

(Figure 1B). Finally, we found a fitness cost of adaptation, since all lines had reduced

individual fitness (λ_{ind}) in 20°C relative to the *Control 20°C* lines ($\chi^2 = 19.799$, df = 4,

p < 0.001, Figure 2). The cost was however not detected in total reproduction ($\chi^2 =$

2.238, df = 4, p = 0.692, Supplementary figure 1).

Discussion

The degree of environmental variation can influence the expression of phenotypes

(Moran 1992; Uller *et al.* 2015; Proulx & Teotónio 2017). In addition to genetic

specialisation, the phenotype can match the environment either by phenotypic

plasticity, where the offspring matches its development to the current environment, or

by anticipatory parental effects, where the offspring non-genetically inherit the

parents phenotype to match its environment, thus improving offspring performance if

parent and offspring environments are matching (Marshall & Uller 2007; Donelson *et*

al. 2017). While within-generation phenotypic plasticity is common (DeWitt &

Scheiner 2004), parental effects seem, in contrast, to be generally weak (Uller *et al.*

2013), despite some well known examples (Gustafsson *et al.* 2005; Galloway &

Etterson 2007; Jensen *et al.* 2014; Kishimoto *et al.* 2017; Ryu *et al.* 2018; Toyota *et*

al. 2019). Since trans-generational effects should evolve only if there is a strong

correlation between parent and offspring environments (Kuijper & Hoyle 2015;

Leimar & McNamara 2015; Uller *et al.* 2015), it is possible that environments

generally are not highly correlated between generations, thus explaining why such

anticipatory effects are uncommon (Uller *et al.* 2013). However, the environmental

predictability is seldom investigated in studies of parental effects (Burgess &

Marshall 2014). Therefore, we investigated whether the degree of temporal

environmental variation, as well as the correlation between parent and offspring

environment, influenced the evolution of maternal effects.

We found that the presence of environmental variation mediated the evolution of

maternal effects on reproduction and individual fitness (λ_{ind}) in *C. remanei* nematode

worms adapting to a novel and stressful warm temperature (25°C) for 30 generations

(see predictions and findings summarized in Supplementary table 1). For all

populations evolving in stable or slowly increasing temperature (*Control 20°C*, *Warm*

25°C, *Increased warming*), a strong positive maternal effect on reproductive output

resulted in an increased offspring production in 25°C when the parents were also

cultured in 25°C and not in 20°C. Since these populations have evolved in

environments with high and positive environmental correlations over generations,

trans-generational effects are adaptive and predicted by theory (Mousseau & Fox

1998; Kuijper & Hoyle 2015; Leimar & McNamara 2015; Uller *et al.* 2015). This

result is also in agreement with a recent study by Dey *et al.* (2016) who found the

evolution of an anticipatory negative maternal effect in *C. elegans* evolving in

perfectly negatively correlated environments. Our finding of positive anticipatory

maternal effects in positively correlated environments highlights the importance of

experimental evolution studies with known environmental correlations to study the

evolution of adaptive trans-generational effects.

In contrast to environments with high positive correlations, the *Fast temperature*

cycles populations evolved in a fluctuating environment where the temperature

changed every second generation. Thus, the next generation would with equal

likelihood be exposed to the same or a different temperature as the parents, resulting

in zero correlation between parent and offspring environments. In this environment,

trans-generational effects are not considered adaptive (Mousseau & Fox 1998; Uller

et al. 2015), and, in agreement with the theory, we found a loss of the positive

maternal effect, since the reproductive output in 25°C of these lines was independent

of the environment their parents. The adaptive value of these differences in parental

effects is illustrated in the individual fitness (λ_{ind}) of the different selection treatments.

Fast temperature cycles populations had highest λ_{ind} in 25°C only when the parents

were cultured for two generations in 20°C, a situation mimicking the fluctuating

environments they were exposed to during evolution. Although adaptive, this should

be defined as negative maternal on λ_{ind}, which is not predicted by theory. Thus,

maternal effects on reproduction and λ_{ind} does not follow the same pattern for the *Fast*

temperature cycles, but importantly, none of the measures show the positive maternal

effects present in lines from more positively correlated environments (figure 1). In

contrast, populations adapting to more or less stable warm temperatures (*Warm 25°C*

and *Increased warming*) improved λ_{ind} when parents were also cultured in 25°C,

which mimics stable temperatures over generations and a positive maternal effect on

λ_{ind}. Whether the negative maternal effect on fitness in the *Fast temperature cycles* is

a result of the fact that the assays with parents and offspring in 25°C represent a

stability not experienced evolutionary by the *Fast temperature cycles* populations

(two generations common garden for the parents and one generation offspring testing

equals to three generations in the same environment), or is a general response when

parent and offspring environment is not matching is unknown.

In addition, we also found a fitness cost of adaptation, since all populations evolving

in new environments had lower λ_{ind} in the original environment (20°C), compared to

the *Control 20°C* populations. Moreover, although *Control 20°C* showed a positive

maternal effect on reproduction, parental exposure to 25°C did not improve their

individual fitness, suggesting that an evolutionary history in 25°C is needed for

maximum fitness in this temperature.

While the positive maternal effect present in both stable (*Control 20°C*, *Warm 25°C*)

and slowly increasing (*Increased warming*) temperature regimes is anticipated when

there is positive autocorrelation between parent and offspring environments

(Lachmann & Jablonka 1996; Uller *et al.* 2015), perfectly stable environments are

actually predicted to select for negative trans-generational effects (Hoyle & Ezard

2012; Kuijper & Hoyle 2015). When a population is well adapted to a stable

optimum, a negative maternal effect reduces phenotypic variance between

generations. However, we find no support for this prediction, since both treatments in

stable environments (*Control 20°C*, *Warm 25°C*) showed a positive maternal effect,

even after 30 generations in stable conditions. It is however possible that these lines

still show transient dynamics, since a positive trans-generational effect is predicted to

evolve as a transient response when experiencing a novel environment (Kuijper &

Hoyle 2015), in a similar way to how phenotypic plasticity is predicted to aid

adaptation to new environments (Price *et al.* 2003; Lande 2009; Chevin *et al.* 2010;

Coulson *et al.* 2017). It could possibly be argued that the positive trans-generational

effect is non-adaptive, caused by the lines being maladapted in 20°C and therefore

producing low-quality offspring in this temperature. However, the fact that the

Control 20°C lines, who have highest fitness in 20°C and who have not experienced

25°C for at least 45 generations show positive maternal effects on reproduction of

parental exposure to 25°C argues against a non-adaptive explanation and instead

reinforces the view that all lines from stable and slowly changing environment has an

adaptive positive maternal effect. In agreement with this, exposing a laboratory strain

of *C. elegans* to 25°C (Schott *et al.* 2014) or to starvation (Rechavi *et al.* 2014)

results in heritable changes in gene expression, directed by inherited small RNA

(Rechavi *et al.* 2014; Schott *et al.* 2014; Houri-Zeevi & Rechavi 2017), which further

suggests that transgenerational plasticity such as maternal effects and epigenetic

inheritance is important in *Caenorhabditis* nematodes.

We found that the environmental correlation between generations is driving the

evolution of anticipatory maternal effects, and the experimental design assured low

within-generation heterogeneity. While stable and slowly changing environments

select for positive anticipatory maternal effects, environments that fluctuate with no

correlation between generations select against parental influence on offspring

phenotype. This is the first empirical study that investigates the evolutionary loss of

anticipatory maternal effect, which follow theoretical predictions (Kuijper & Hoyle

2015; Uller *et al.* 2015) and suggest that one reason for the weak effects of parent

environment on offspring phenotype in natural systems (Uller *et al.* 2013) could be

that natural environments are not always temporally correlated across generations.

While most examples of positive parental effects comes from systems with short life-

cycles such as nematodes (Kishimoto *et al.* 2017), daphniids (Gustafsson *et al.* 2005;

Toyota *et al.* 2019), annelids (Jensen *et al.* 2014) and herbs (Galloway & Etterson

2007), it is also been found in fish where the generation time span years (Ryu *et al.*

2018). However, when investigating maternal effects in *Daphnia* from natural

populations with different degree of variation in predation intensity, Walsh *et al.*

(2016) found some support for stronger positive maternal effects in population from

more stable environments, but the effect differed between traits and no effect was

found on reproduction. Nevertheless, these types of studies where the environmental

predictability is known are vital for our understanding of the selection pressures

resulting in the presence or absence of adaptive maternal effects in natural populations

(Burgess & Marshall 2014), where deconstruction of environmental predictability to

seasonality and environmental colour noise (following Marshall & Burgess 2015) is a

very promising approach.

On-going climate change is not only resulting in warmer temperatures, but also

increased temperature variation, which can impact both the ecology and evolution of

populations and species (Vasseur *et al.* 2014; Vázquez *et al.* 2017), and trans-

generational acclimation can be an important response to deal with a warming climate

(Donelson *et al.* 2017; Ryu *et al.* 2018). We show that environmental heterogeneity

drives the evolution of maternal effects, and support the theoretical predictions

(Lachmann & Jablonka 1996; Uller *et al.* 2015) that the correlation between parent

and offspring environment is the driver of the evolution of transgenerational

plasticity.

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Competing interests

The authors report no competing interests.

Materials & Correspondence

Requests for material and correspondence should be addressed to MIL.

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Figure 1. The effect of parental temperature on fitness and reproduction in 25°C.

Individual fitness λ_{ind} (A) and total reproduction (B) in 25°C when parent were raised

for 2 generations in either 20°C or 25°C. Symbols represent selection treatments

(mean ± SE). *Control 20°C* and *Warm 25°C* have spent 30 generations in 20°C or

25°C, respectively. *Increased warming* has been subjected to slowly increased

temperatures, starting in 20°C and reaching 25°C at generation 30. *Fast temperature*

cycles have spent two generations in 20°C, followed by two generations in 25°C, this

cycle has then been repeated for 30 generations.

Figure 2. Fitness in original environment. Individual fitness λ_{ind} in 20°C when

parents are also grown for two generations in 20°C. Symbols represent selection

treatments (mean ± SE). Grey symbolises treatments groups that differ significantly in

fitness.