1   How do fluctuating temperatures alter the cost of development?

Abstract

1. Quantifying how variable temperature regimes affect energy expenditure during
development is crucial for understanding how future thermal regimes may impact
early life survival and population persistence.

2. Developmental Cost Theory (DCT) suggests that there is an optimal temperature
(T_{opt}) that minimises energy expenditure during development (the "cost of
development"). Exposure to fluctuating temperatures around an average of T_{opt} are
anticipated to increase either development time or metabolic rate. As a result,
embryos will rapidly deplete yolk reserves, and consequently hatch at a smaller size
or with less residual yolk to support postnatal survival and growth.

3. Here, we studied total embryonic energy expenditure (development time and rate of
CO_2 production) and conversion of yolk into tissue in common wall lizards (*Podarcis*
muralis) under three incubation treatments anticipated, based on DCT, to increase the
cost of development: no variance (T_{opt} constant, 24 °C), low variance (22 – 24 °C),
and high variance (18 – 30 °C).

4. As predicted, we found that increasing variance around T_{opt} increased the cost of
development, despite reducing time to hatching. As a consequence, embryos on
average hatched with 59% lower residual yolk reserves under high variance versus
constant incubation temperatures.

5. Our results highlight how the relative temperature sensitivities of development time
and metabolic rate determine the cost of development, which in turn may predict the
ability of egg-laying ectotherms to persist in variable environments. We show that
DCT can provide a mechanistic framework for understanding the widespread, but
often seemingly idiosyncratic, effects of fluctuating incubation temperatures on
hatchling tissue and residual yolk mass.

Keywords: developmental cost theory**,** hatching, incubation, metabolic rate, temperature,
wall lizard, yolk

Introduction

The development of an embryo from a single-celled zygote to a nutritionally independent hatchling is a critical life stage that is likely to be an important bottleneck for survival under increasingly extreme and variable environments. In egg-laying species, embryonic development is fuelled by maternally-provisioned yolk (Blaxter, 1969; Deeming & Ferguson, 1991; Mousseau & Fox, 1998). Energy derived from yolk is used for crucial processes during prenatal development including cell division and differentiation, growth, and maintenance, while residual yolk at hatching can be internalised to support postnatal growth and survival (Marshall et al., 2020; Murakami et al., 1992; Troyer, 1983; Vidal et al., 2002). The total energy expenditure during the period of embryonic development, also known as the "cost of development", will be a product of 1) development time (time from fertilisation to hatching; D), and 2) rate of energy expenditure (metabolic rate; MR) (Pettersen et al., 2019). Any factor that increases either D or MR during prenatal development is anticipated to deplete yolk reserves which may in turn reduce residual yolk and/or the amount of tissue at hatching, with likely fitness consequences.

In ectotherms, temperature affects both developmental and metabolic rates (Pettersen et al., 2019). Developmental cost theory (DCT) posits that the thermal sensitivity of development time (D_T) relative to the thermal sensitivity of metabolic rate (MR_T) determines how the overall cost of development ($D \times MR$) scales with temperature (Marshall et al., 2020). If the slope of D_T and MR_T functions differ, as shown for many ectotherm species, there will be an optimal temperature that minimises energy consumption during development (T_{opt}), and thereby maximises residual yolk upon completion of development (Pettersen et al., 2019). At temperatures below T_{opt}, $D_T > MR_T$ and development time is extended to a greater extent than metabolic rate is reduced. Beyond T_{opt}, $D_T < MR_T$ and metabolic rate increases more than development time decreases with temperature. Using experimental data across seven phyla, DCT has been used to predict T_{opt} for 71 ectotherm species (Marshall et al., 2020). The effects of temperature on embryonic development is crucial for predicting the effect of changing temperature on population persistence, yet it remains poorly understood how well DCT predicts energy utilisation within species (Du et al., 2019).

Natural environments are thermally variable, and it is well-known that the relationship between performance and temperature is often nonlinear in ectotherms (Bernhardt et al., 2018; Denny, 2017). While DCT predicts that incubation at temperatures that deviate from T_{opt} should increase the cost of development, the extent to which fluctuating temperatures will exacerbate total energy expenditure, due to an increase in D, MR, or both,

has yet to be empirically tested. Depending on the shape of the cost-temperature function,

embryos experiencing variable temperatures may use less or more energy to complete

development than those under constant temperatures, even when variable and constant

temperatures have the same average temperature (Slein et al., 2023). For offspring, time spent

developing at temperatures that deviate from T_{opt} therefore necessitates that they will either

need to sacrifice energy reserves, hatching with lower residual yolk, or compromise the

conversion of yolk into tissue, by hatching at a smaller size. Elucidating the effect of

developmental temperatures that vary around a mean of T_{opt} on both the cost of development,

and embryonic phenotypes, is thus crucial for understanding how well DCT predicts energy

acquisition and allocation under fluctuating temperatures.

Egg-laying lizards provide a useful system to test DCT under fluctuating, ecologically

relevant temperatures since egg incubation temperatures are variable in the field, often

cycling through a consistent diel regime during incubation (Oufiero & Angilletta Jr., 2010b).

Many lizard species hatch with a significant amount of residual yolk that is internalized

before hatching – this energy resource can sustain offspring for several days post-hatching,

and is likely to be under strong selection when food availability is low or unpredictable

(Radder et al., 2004; Troyer, 1987; but see Radder et al. 2007). On the other hand, converting

a greater proportion of yolk reserves towards tissue production may be beneficial if larger

hatchling size confers greater survival, sprint speed, or reproductive output later in life

(Sinervo, 1990; Sorci & Clobert, 1991; Uller & Olsson, 2010). Past studies have measured

the effect of fluctuating versus constant incubation temperatures on hatchling phenotypes in

lizards, including effects on embryonic development (see recent review by Raynal et al.,

(2022)), and metabolic rate (Du & Shine, 2010; Hall & Warner, 2020; Oufiero & Angilletta

Jr., 2010a). Yet no studies have tested how deviations from the theoretical T_{opt} influence total

energy expenditure and the conversion of yolk into tissue during development. This makes it

difficult to assess patterns of developmental plasticity in relation to DCT, and their likely

fitness implications.

Here, we estimated total energy expenditure ($D \times MR$) and embryonic conversion of

yolk into tissue in common wall lizards (*Podarcis muralis*) to test DCT under ecologically

relevant, fluctuating incubation temperatures. First, we used existing measures of D_T and

MR_T to estimate the constant temperature that minimises the cost of development (T_{opt}) for *P.*

muralis. Second, we measured how developmental cost varies under incubation temperatures

that fluctuate around T_{opt} using three temperature regimes. All three temperature regimes had

the same average temperature (24 °C; optimal for this species in terms of energy expenditure,

Figure 1) but differed in the daily magnitude of fluctuation around this average. Third, we

measured hatchling wet mass, dry tissue, and residual yolk mass across the three incubation

temperatures regimes. Based on DCT we predicted that higher fluctuations around T_{opt} would

impose a higher energetic cost of development, necessitating embryos that developed under

fluctuating temperatures to either (*i*) compromise energy reserves at hatching and hatch at the

same mass but with less residual yolk, or (*ii*) compromise conversion of yolk to tissue and

hatch at a smaller size, than embryos developing at T_{opt}.

Materials and methods

Calculation of P. muralis embryo thermal optima (T_{opt})

All analyses were conducted in R version 4.0.3 (R Core Team, 2023). We used development

(incubation) time (*D*; (Pettersen et al., 2022; While et al., 2015) and metabolic rate (*MR*;

Pettersen et al., (2022)) data previously collected on Italian *P. muralis* populations, to

calculate the temperature that minimises the cost of development (*C*) according to

developmental cost theory (DCT), where:

[1]

Exponential models were fit using the "nls" function to determine parameter estimates for

MR and *D*, whereby:

[2]

[3]

where   and   represent the temperature ( ) dependencies of *MR* and *D* respectively, and   ,

 , and   are additional parameters that combined describe the decrease of development time,

and increase of metabolic rate, with increases in incubation temperature. Based on the

combined functions of [2] and [3], the temperature that minimises predicted *C* (referred to as

"T$_{opt}$") was calculated to be 23.9 °C (Figure 1; see Supporting Information for details).

Figure 1. Graphical representation of *P. muralis* embryonic **A)** Temperature dependence of

metabolic rate (ml CO_2 h^{-1}); MR_T based on previous work (Pettersen et al., 2022), and **B)**

Temperature dependence of development time (incubation time: number of days from

oviposition to hatching); D_T (Pettersen et al., 2022; While et al., 2015), used to calculate **C)**

Cost of development; C_T (total CO_2 production from oviposition until hatching). Grey

vertical lines represent the temperature that minimises the cost of development (T_{opt}, 24 °C)

and the stable temperature used for the "no variance" (NV) treatment. Light grey dashed lines

represent temperature range of "low variance" (LV: 22 – 26 °C) and black dotted lines

represent temperature range of "high variance" (HV: 18 – 30 °C) treatments used in this

study. Given that C_T is a convex function, embryos developing under variable temperatures

are predicted to have greater cost of development than embryos developing under constant 24

°C. **D)** Programmed incubator settings over a 24h cycle throughout development for the three

temperature treatments: NV (light grey line), LV (dashed grey line), and HV (dotted black

line).

Study system, field collection, and animal husbandry

The common wall lizard (*Podarcis muralis* Laurenti, 1768) is a small (50-70mm snout-to-

vent length) oviparous lacertid. Adult, gravid females (n = 62) and adult males (n = 38)

captured in the field during the prior reproductive season across three regions in central Italy

(for details, see Pettersen et al., (2022)), were reared in animal facilities at Lund University,

Sweden. During the reproductive season from April – June 2020, lizards were housed in a

temperature-controlled room (24 °C day/20 °C night) with a light cycle of 12 L:12 D. One to

two females and one male were housed in cages (590 × 390 × 415 mm) containing a sand

substrate, bricks and logs as shelter, a sand box for egg laying, and a water bowl. Each cage

contained a basking lamp (60 W) switched on for 8 h per day to create a thermal gradient (24

°C – 40 °C) and a UV light (EXO☐TERRA 10.0 UVB fluorescent tube) for 6 h per day.

Lizards were fed mealworms or crickets daily. Sand boxes were inspected daily for clutches

to ensure egg collection within 24 h post-oviposition.

Incubation treatments

Upon egg collection, we weighed the mother and eggs to the nearest 0.01g (Ohaus Scout

SKX Precision Balance). Both females in the cage were visually inspected for evidence of

having recently laid a clutch. Over the reproductive season, females laid between 1-5 clutches

(mean: 3.48 ± SD: 1.09) with clutch sizes ranging 1-10 eggs (mean: 4.6 ± SD: 1.8). For the

purpose of this experiment, we only sampled the 1st and 2nd clutches. We used a split clutch

design to allocate eggs across three incubation treatments: one constant at T_{opt}, "no variance"

(NV; 24 °C), and two fluctuating thermal regimes with a mean temperature of 24 °C and

either "low variance" (LV; ± 2°C, fluctuating between 22-26 °C) or "high variance" (HV; ±

6°C, fluctuating between 18-30 °C) to determine whether fluctuating temperatures that still

maintain a mean T_{opt} will alter the energetic cost of development as manifested at the

organismal level by smaller hatchling size, less residual yolk, or both. Incubators (Snijders

Labs Micro Clima-Series Economic Lux Chamber) were programmed on a 24h cycle: 9h at

the low temperature of the treatment range (LV:22 °C/HV:18 °C), followed by a 3h ramping

period to the upper treatment range (LV:26 °C/HV:30 °C) maintained for 9h and an

equivalent ramping down, or a continuous 24 °C for NV treatment (Figure 1d). Eggs were

incubated in small plastic containers filled two-thirds with moist vermiculite (5:1 vermiculite

to water ratio) and sealed with clingfilm throughout development (except when they were

temporarily removed for metabolic rate measurements, see details below) until hatching. The

level at which inferences were made is individual developing embryos, however the level at

which each treatment was applied is individual females (since some females contributed

more than one egg to a given treatment). The number of females used per treatment is

provided in Table 1, and the number of embryos measured for each treatment per female is

provided in Table S1.

Table 1. Replication Statement. for the scale of inferences made, the scale at which the

treatment was applied, and the number of replicated for each treatment (exact numbers per

measurement provided in Table S1).

Scale of inference	Scale at which treatment of interest was applied	Number of replicates at the appropriate scale
Individuals	Females	9-26

Measurement of embryonic metabolic rate

Pilot data from dissections on a subsample of first clutches was used to estimate days post

oviposition (dpo) for embryos when they were 75% through development (36-39 dpo).

Embryo metabolic rate (*MR*) was then measured at 75% of development for the temperatures

at which each embryo was exposed to within their treatment (NV:24 °C/LV: 22 °C, 24 °C, 26

°C/HV:18 °C, 24 °C, 30 °C; Figure 1a). For LV and HV, measures were taken in increasing

order of temperatures, reflecting the daily ramping of incubation temperature. To determine

whether *MR* remained broadly consistent across development, we also measured NV

embryos at 25% (11-14 dpo) and 50% (23-26 dpo) through development (at a single 24 °C

only). Rate of CO_2 production, or $\dot{V}CO_2$, was calculated from measures of fractional CO_2

using closed-system, stop-flow respirometry as per methods described previously (Cordero et

al., 2017; Pettersen et al., 2022). In brief, individual eggs were placed into 60ml glass

chambers containing 50ml moist vermiculite. Each chamber was then connected to and

flushed with dry scrubbed pressurised air (using drierite desiccant; Hammond Drierite at a

flow rate of 200 ($\pm$20) ml min^{-1} (standard temperature and pressure, dry; STPD; registered

using a FM-8 mass flow meter from Sable Systems International, Las Vegas, NV, USA). The

sealed vials were then placed into incubators set at the measurement temperature. After 180

min, a baseline measurement of CO_2 was taken before re-connecting the chambers to the

airflow. Excurrent air (at approximately 200 ml min^{-1}, STPD) from the chambers was dried

using magnesium perchlorate ($Mg(ClO_4)_2$), after which we measured the total volume of CO_2

accumulated in the chambers using a FC-10 CO_2 analyser (Sable Systems International).

Every embryo MR was measured at each temperature within their respective treatment. $\dot{V}CO_2$

in ml h^{-1} was calculated using the *metabR* package (Noble, 2021), by integrating the area

under the $\%CO_2$ curve above the baseline.

Development time and hatchling traits

Eggs were checked daily for signs of hatching to determine development time; the number of

days from oviposition until hatching (*D*). Within 24 h of hatching, individuals were humanely

euthanised and measures of hatchling wet mass (to 0.01g), hatchling dry mass and yolk dry

mass (to 0.01 mg), snout-to-vent length (svl; mm), and total length (mm) were taken. All

hatchling condition measurements were carried out by the same author. Since hatchling wet

mass includes residual yolk, it is not an informative metric for quantifying total energy

allocation towards tissue synthesis. The ability to dissect residual yolk from the hatchling

allows for direct measurement of the allocation of initial yolk reserves towards tissue

production and residual yolk, under temperature regimes that are more or less costly in terms

of energy expenditure. Hence, hatchling tissue and yolk were placed into a drying oven at 60

°C for 3 days, then weighed using a microbalance (Sartorius Research 200D). Further drying

did not lead to any further reductions in sample weights (personal observation).

Hypothesis tests

To quantify how variation in incubation temperature around T_{opt} (24 °C) alters the overall

energy cost of development, and conversion of yolk to tissue during development, we fitted

linear mixed effects models for the following response variables: (1) metabolic rate (MR,

measured as $\dot{V}CO_2$), (2) development time (D, measured as time from oviposition until

hatching), (3) cost of development (C, as a function of temperature-dependent MR x total

time spent at each temperature until hatching), (4) wet mass at hatching, (5) dry mass at

hatching and (6) dry residual yolk mass at hatching. All models were fit using the "lmer"

function within the *lme4* package (Bates et al., 2015). Fixed predictor variables of incubation

'Treatment' (NV, LV, HV), 'Egg mass', 'Region' of parent origin (high, mid, low altitude),

as well as two-way interactions among these, and 'Clutch order' (1-2; for D and wet mass at

hatching), were tested. For MR, we first tested for differences among treatments at the

intermediate temperature (24 °C). We then ran separate models for each treatment since MR

was measured at different temperatures within each treatment, and 'Temperature' (and its

interaction with Region) were tested. Maternal identity was included as a random effect in all

models. To account for non-independent replication when eggs from the same female were

included within a treatment, we used the average value per clutch (Harrison et al., 2018). Egg

mass and metabolic rate were natural log transformed for all hypothesis tests, to satisfy the

assumption of heteroscedasticity, and to allow for meaningful interpretation of scaling

patterns (Glazier, 2021; Niklas & Hammond, 2014).

Model selection and parameter estimation

Candidate models (summarised in Table 2) were ranked using Akaike Information Criterion

(AIC) averaged for those with Δ conditional Akaike information criterion (AICc) <4 using

the R package *MuMin* v.1.43.17 (Barton, 2009). Post-hoc tests using the *emmeans* (Lenth,

2022) package were run to compare pairwise differences in fixed factors retained in the final

models.

Table 2. Summary of candidate models used in model selection for 1. Log_n Metabolic rate, 2.

Log_n Development time, 3. Log_n Cost of Development, 4. Hatchling wet mass, 5. Hatchling

tissue dry mass, and 6. Hatchling residual yolk dry mass. Fixed predictor variables were

Temperature (within each Treatment: LV = 22, 24, 26 °C, HV = 18, 24, 30 °C), Log_n Egg

mass, Region (High, Mid, Low) and Clutch order (1st, 2nd). Mother ID was included as a

random effect for all models. Clutch order is included in models for development time and

hatchling wet mass (only 2nd clutches were measured for MR, C, and hatchling dry mass and

residual yolk mass). Note for Response 1 (Log_n Metabolic rate), Temperature was not

included for models of NV treatment measured at a single temperature (24 °C).

Response	Model no.	Parameters
1	1	Temperature + 1
1	2	Temperature + Log_n Egg mass
1	3	Temperature + Log_n Egg mass + Region
1	4	Temperature + Log_n Egg mass + Region + Temperature x Log_n Egg mass
1	5	Temperature + Log_n Egg mass + Region + Temperature x Region
1	6	Temperature + Log_n Egg mass + Region + Log_n Egg mass x Region
2,3,4,5,6	7	Treatment + 1
2,3,4,5,6	8	Treatment + Log_n Egg mass
2,3,4,5,6	9	Treatment + Log_n Egg mass + Region
2,4	10	Treatment + Log_n Egg mass + Region + Clutch order
2,4	11	Treatment + Log_n Egg mass + Region + Clutch order + Treatment x Log_n Egg mass
2,4	12	Treatment + Log_n Egg mass + Region + Clutch order + Treatment x Region
2,4	13	Treatment + Log_n Egg mass + Region + Clutch order + Log_n Egg mass x Region
3,5,6	14	Treatment + Log_n Egg mass + Region + Treatment x Log_n Egg mass
3,5,6	15	Treatment + Log_n Egg mass + Region + Treatment x Region
3,5,6	16	Treatment + Log_n Egg mass + Region + Log_n Egg mass x Region

Results

(i) Metabolic rate

Metabolic rate (*MR*; CO_2) measured at 24 °C (T_{opt}) was significantly higher in the high

variance (HV; 18-30 °C, mean ± SD: $1.51 \cdot 10^{-2} \pm 2.75 \cdot 10^{-3}$ ml CO_2 h^{-1}) and low variance

(LV; 22-26 °C, mean ± SD: $1.38 \cdot 10^{-2} \pm 2.76 \cdot 10^{-3}$ ml CO_2 h^{-1}) treatments compared with the

no variance treatment (NV; 24 °C, mean ± SD: $1.26 \cdot 10^{-2} \pm 2.00 \cdot 10^{-3}$ ml CO_2 h^{-1}; $F_{2,168}$ =

23.49, p <0.0001). Within LV and HV treatments, *MR* increased with measurement

temperature (Table 3, Figure 2). We found a significant effect of $\log_n$ Egg mass on $\log_n$ *MR*

for the NV treatment, but this relationship was non-significant for LV and HV (Table 3). *MR*

measured at 24 °C increased approximately 1.6x across development, but this was only

significantly different between 25% (mean ± SD: $8.10 \cdot 10^{-3} \pm 2.29 \cdot 10^{-3}$ ml CO_2 h^{-1}) and 75%

(mean ± SD: $1.26 \cdot 10^{-2} \pm 2.00 \cdot 10^{-3}$ ml CO_2 h^{-1}) completion of development. There was no

observed interaction between development stage and egg mass, indicating consistent

metabolic scaling across early ontogeny (Table S3, Figure S1).

Figure 2. Response of Metabolic rate ($\log_n$ CO_2; ml CO_2 h^{-1}), to temperature across three

incubation treatments: **A)** constant, "no variance" (NV; measured at 24 °C), **B)** low variance

(LV; measured at 22 °C, 24 °C, 26 °C), and **C)** high variance (HV; measured at 18 °C, 24 °C,

30 °C). Data points show raw data and lines are fitted to individual responses with a single

line () fitted using a linear model. Orange data points are measures of metabolic rate at T_{opt}

(24 °C). Lighter and darker colours represent deviations below and above T_{opt}, respectively.

(ii) Development time

Embryos incubated at HV (18-30 ℃) hatched, on average, five days earlier (HV mean ± SD:

1142.54 ± 54.23 hours) than embryos incubated in LV (HV mean ± SD: 1250.17 ± 46.94

hours) and NV (HV mean ± SD: 1262.48 ± 46.94 hours) treatments, while no significant

differences in development time between LV and NV were observed (Table 3, Figure 3a).

We observed no significant effect of egg mass, region, clutch order (or interactions of these)

on development time (Table S2).

(iii) Cost of development

The total CO_2 production to complete development, calculated from the temperature-specific

$\dot{V}CO_2$ (*MR*) at 75% of development and total development time (*D*) increased with

incubation temperature variation (Table 3). Based on the final best-fitting model, the 'cost of

development' for an average size egg (0.28 g) was lowest under NV treatment (mean ± SD:

15.87 ± 2.54 ml CO_2), intermediate for LV (mean ± SD: 17.19 ± 3.24 ml CO_2), and highest

for HV (mean ± SD: 17.84 ± 2.96 ml CO_2), however differences between LV and HV were

marginally non-significant (Figure 3b). The cost of development did not differ across regions

(Tables S2 and S3).

(iv) Wet mass at hatching

Hatchling wet mass (tissue and residual yolk; g) was significantly higher in the NV (mean ±

SD: 0.36 ± 0.05g) and LV (mean ± SD: 0.35 ± 0.04g) treatments, compared with HV (mean ±

SD: 0.34 ± 0.05g; Table 3, Figure 3c). Wet mass at hatching was significantly positively

correlated with initial egg mass, however there was no significant difference in hatchling wet

mass between regions or clutch order (Table S2). For an average size egg (0.28g) we

calculated that embryos incubated at NV hatched 6% and 2% heavier than those under HV

and LV, respectively.

(v) Dry hatchling and yolk mass

Analysis of the dry mass of hatchlings and residual yolk revealed that hatchling dry mass was

consistent across the incubation temperature treatments (Table 3 and S3, Figure 3d). We

found that on average, embryos incubated at constant 24 °C (NV; mean ± SD: $1.41 \cdot 10^{-3}$ ± $1.09 \cdot 10^{-3}$ g) hatched with 20% and 59% more yolk reserves, compared with the LV (mean ± SD: $1.14 \cdot 10^{-3}$ ± $8.38 \cdot 10^{-4}$ g) and HV (mean ± SD: $1.02 \cdot 10^{-3}$ ± $9.56 \cdot 10^{-4}$ g) respectively. However, there were no significant difference in dry yolk mass between LV and HV (Table 3, Figure 3e). While larger eggs resulted in hatchlings with higher tissue dry mass, there was no effect of egg mass on residual yolk dry mass at hatching (Table S2).

Table 3. Summary of significance tests and marginal means of fixed effects from best-fitting linear mixed effects models describing the relationship between 1. Metabolic rate, 2. Development time, 3. Cost of development, 4. Wet mass, 5. Dry mass, 6. Dry residual yolk mass, with predictor variables: Temperature (within treatment), Treatment (NV = no variance, LV = low variance, HV = high variance), Log_n Egg mass, Region, and Clutch order, and their interactions as shown in Table 2. Final model parameters were included based on model selection (AICc) shown in Table S2. Significant estimates shown in bold: *p < 0.05, **p < 0.01, ***p < 0.001. Letters represent significant pairwise differences across levels (a > b > c). Note that responses for 1, 2, and 3 are back transformed from model output log_n estimates.

	χ^2	df	p	emmean (±SE)
1a. Log_n Metabolic rate (NV)				
Intercept	200.80	1	**<0.001*****	
Log_n Egg mass	10.21	1	**<0.01****	
1b. Log_n Metabolic rate (LV)				
Intercept	132.30	1	**<0.001*****	
Temperature	216.54	2	**<0.001*****	22 °C: $1.19 \cdot 10^{-2}$ ($3.41 \cdot 10^{-4}$) [c] 24 °C: $1.38 \cdot 10^{-2}$ ($3.96 \cdot 10^{-4}$) [b] 26 °C: $1.58 \cdot 10^{-2}$ ($4.51 \cdot 10^{-4}$) [a]
Log_n Egg mass	3.15	1	0.08	
1c. Log_n Metabolic rate (HV)				
Intercept	256.09	1	**<0.001*****	
Temperature	1530.51	2	**<0.001*****	18 °C: $9.75 \cdot 10^{-3}$ ($1.99 \cdot 10^{-4}$) [c] 24 °C: $1.47 \cdot 10^{-2}$ ($3.71 \cdot 10^{-4}$) [b] 30 °C: $2.17 \cdot 10^{-2}$ ($5.49 \cdot 10^{-4}$) [a]
Log_n Egg mass	0.01	1	0.91	
2. Log_n Development time				
Intercept	2003063.25	1	**<0.001*****	
Treatment	711.43	2	**<0.001*****	NV: 1257.65 (1.00) [a] LV: 1251.38 (1.00) [a] HV: 1132.29 (1.00) [b]
3. Log_n Cost of development				
Intercept	274.05	1	**<0.001*****	
Treatment	26.16	2	**<0.001*****	NV: 15.64 (1.02) [b] LV: 16.95 (1.02) [a] HV: 17.64 (1.02) [a]

Log_n Egg mass	7.40	1	**<0.01****	
4. Hatchling wet mass				
Intercept	816.09	1	**<0.001*****	
Treatment	45.03	2	**<0.001*****	NV: $3.49 \cdot 10^{-1}$ ($3.93 \cdot 10^{-3}$) [a] LV: $3.43 \cdot 10^{-1}$ ($3.92 \cdot 10^{-3}$) [a] HV: $3.33 \cdot 10^{-1}$ ($3.93 \cdot 10^{-3}$) [b]
Log_n Egg mass	273.23	1	**<0.001*****	
5. Hatchling tissue dry mass				
Intercept	281.35	1	**<0.001*****	NV: $6.76 \cdot 10^{-2}$ ($9.21 \cdot 10^{-4}$)
Treatment	3.54	2	0.17	LV: $6.88 \cdot 10^{-2}$ ($9.16 \cdot 10^{-4}$)
Log_n Egg mass	99.46	1	**<0.001*****	HV: $6.74 \cdot 10^{-2}$ ($9.21 \cdot 10^{-4}$)
6. Hatchling residual yolk dry mass				
Intercept	62.65	1	**<0.001*****	
Treatment	9.35	2	**<0.01****	NV: $1.41 \cdot 10^{-3}$ ($1.30 \cdot 10^{-4}$) [a] LV: $1.22 \cdot 10^{-3}$ ($1.29 \cdot 10^{-4}$) [ab] HV: $1.03 \cdot 10^{-2}$ ($1.30 \cdot 10^{-4}$) [b]

Figure 3. Tukey's boxplots for response variables: a) Development time, b) Cost of development, c) Hatchling wet mass (tissue and yolk mass combined), d) Hatchling tissue dry mass, and e) Hatchling residual yolk dry mass, to incubation temperature treatment (constant, "no variance" (NV; 24 °C; light shading), low variance (LV; 22-26 °C) and high variance (HV; 18-30 °C; dark shading)). Data points show raw data. Letters represent significant pairwise differences across levels (a > b > c).

Discussion

Many regions of the globe are experiencing increasingly variable temperatures. Here, we show evidence that thermal variability may exacerbate the cost of development for egg-laying ectotherms. Our results demonstrate that, in wall lizards, increasing variation around the optimal incubation temperature (T_{opt}) increases the cost of development such that hatchlings utilise more residual yolk and consequently begin life outside the egg with fewer energetic reserves. The extent to which these effects of fluctuating temperatures around T_{opt} will impact fitness is likely to be context dependent. Where resources in the postnatal environment are abundant then the consequences of the costs of development on fitness may be offset. However, where the postnatal environment is characterised by low or unpredictable resource availability then the consequences of the costs of reproduction may be severe. Indeed, environments that have greater fluctuating thermal and hydric regimes on a seasonal basis, often show equally strong seasonality in food availability (Gilbert & Miles, 2016). Hence, thermally variable environments may present a twofold cost for developing embryos: a greater cost of development and therefore greater immediate reliance on external sources of food as well as a reduced availability of those external resources upon hatching.

As predicted, embryo energy expenditure (cost of development) was higher under fluctuating versus stable incubation treatments. This expectation was based on the convex shape of the temperature-cost function – which is a product of both metabolic rate (*MR*) and development time (*D*). Based on data from constant incubation temperatures, we anticipated that embryos exposed to thermal variance would show higher developmental energy costs, relative to embryos incubated under constant conditions, potentially due to extended development time, increased metabolic rates, or both. While we observed that exposure to high fluctuating temperatures increased overall metabolic rates, it also caused embryos to hatch sooner than those under constant or low fluctuating temperatures. For *P. muralis* experiencing thermal variability during development, exposure to temperatures above T_{opt} *decreased* development time to a lesser extent than it *increased* metabolic rates, resulting in overall increased energy expenditure with increasing time spent at temperatures further away from the optimum.

Due to the nonlinear relationship between development time and temperature, we expected a positive net effect of fluctuating temperatures on development time yet observed a decrease in *D* under fluctuating temperatures. Faster development under fluctuating temperatures has also been observed in other reptile species (Du & Shine, 2010; Fischer et al., 2011; Georges et al., 2005; Hall & Warner, 2020; Lin et al., 2008; Lu et al., 2009; Shine,

2004; Shine & Harlow, 1996) and may be due to physiological plasticity (acclimatisation),

whereby embryos respond to shifts in temperature and compensate *D* accordingly.

Alternatively, faster development under fluctuating temperatures may be a result of the

thermal inertia of embryo body temperature cooling after exposure to the upper

environmental temperature, which may also explain why higher metabolic rates are

maintained in the thermal variance treatments when embryos were measured at 24 °C.

Investigating the complicated responses of thermal history on embryo physiology, and how

this may alter predicted implications of environmental change, is an important avenue of

future research (Denny, 2019; Du & Shine, 2010; Pottier et al., 2022).

While DCT predicts that fluctuating temperatures will commonly impose energetic costs to

developing embryos, what has been less clear, is *how* this increased cost will be paid for. In

P. muralis, the increased cost of development under fluctuating temperatures did not affect

the conversion of yolk into tissue but did result in lower energy reserves at hatching. This

suggests that wall lizard embryos may prioritise tissue production over hatchling energy

reserves when costs of development are high. There is evidence that internalised residual

yolk reserves at hatching act as an insurance mechanism that can sustain hatchlings in the

event of low foraging success (Radder et al., 2004; Troyer, 1983, 1987). On average,

embryos incubated under high (± 6 °C) and low (± 2 °C) temperature variability around T_{opt}

(24 °C) hatched with 59% and 20% lower yolk reserves, respectively, compared to incubation

at constant T_{opt}. This suggests that hatching at sufficient size, regardless of the energy needed

to attain it, may be of greater importance for post-hatching fitness components such as

survival and growth than retaining greater energy reserves. The importance of preserving

yolk relative to synthesising tissue is likely to be context dependent, yet measures of the

consequences of this trade-off for survival in the field are lacking.

Incubation temperature produces strong and pervasive effects on post-hatching survival in

ectotherms, that have been well documented in the literature, particularly in lizards (Noble,

Stenhouse, Riley, et al., 2018; While et al., 2018). Mechanistic explanations have been put

forward to explain these effects, yet few studies have considered the fitness consequences of

energetic cost of development (Cordero et al., 2017; Ji & Braña, 1999; Oufiero & Angilletta

Jr., 2010b), and measures of both the temperature-dependence of embryonic metabolic rates

and developmental time are rare. DCT provides a framework for predicting how the mean

and variation of incubation temperature will affect the cost of development, and through this,

fitness. Based on the shape of the cost function (as a product of non-invasive measures of

MR_T and D_T), deviations in incubation temperature from T_{opt}, including fluctuations around a

mean of T_{opt}, will increase developmental cost. However, exactly how much an increase in a

degree fluctuation of incubation temperature will cost in terms of energy expenditure depends

on the average temperature (as illustrated in Figure 1). This observation may help to explain

the highly variable responses of D and MR under fluctuating incubation temperatures that

have been reported in ectotherms (Bozinovic et al., 2011; Niehaus et al., 2012; Slein et al.,

2023).

While we recognise the diversity of temperature effects on embryogenesis, we

propose that many of the gross effects on size, mass, and energy consumption that have been

observed in ectotherms (Noble, Stenhouse, & Schwanz, 2018; While et al., 2018) will be well

explained by the temperature-dependence of metabolic rate relative to development time.

Combining these measures, while also collecting data on the consequences of these costs in

the wild, represents an important step towards understanding the mechanisms underlying the

effects of developmental temperature on survival.

Both metabolic and developmental rates can vary between populations, reflecting either local

adaptation (Conover & Schultz, 1995; Demarco, 1992; Pettersen, 2020) or plasticity (Cordero

et al., 2017; Gangloff et al., 2019). In wall lizards, developmental rate shows counter-gradient

variation, whereby embryos from cooler, high altitude and latitude populations develop faster

than those from warm climates when placed under common garden conditions (Pettersen et

al., 2022; While et al., 2015). In our study, we found that population differences had a

marginal influence on the optimal temperature from an energy-efficiency perspective –

embryos from different climatic regimes were found to generate similar estimates of the cost

of development and did not differ in their residual yolk at hatching. Under the assumption

that the cost of development may vary across a lineage (Marshall et al. 2020), we can predict

that populations that occupy the hot or cold extremes of the lineage's distribution will

therefore pay a higher cost. This raises the possibility that DCT may be able to predict

species ranges based on mean and variation in nest temperatures. Complementary to relying

on estimates of embryo thermal tolerance to inform species distributions (e.g., Kearney and

Porter (2009)), it is plausible that measures of fitness quantified through the cost of

development, can provide accurate predictions required to assess the impact of increasing

thermal variance on the response of populations under global change.

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