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 – 26 °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 (Figure 1). 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. The extent to which embryos experiencing variable

temperatures expend more energy to complete development compared to those developing

under constant temperatures, will depend on the shape of the cost-temperature function, 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., 2010a).

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., 2010b). 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*) from three locally adapted

altitude regions to test DCT under ecologically relevant, fluctuating incubation temperatures

(Pettersen et al., 2023). 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 (low: 24 ± 2 °C, high:

24 ± 6 °C). 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., (2023; While et al., 2015) and metabolic rate (*MR*;

Pettersen et al., (2023)) 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] $$C = MR \cdot D$$

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

MR and *D*, whereby:

[2] $$MR(T) = f \cdot e^{a \cdot T}$$

[3] $$D(T) = -d \cdot e^{(-b \cdot T)} + c$$

where a and b represent the temperature (*T*) dependencies of *MR* and *D* respectively, and f,

d, and c 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).

To estimate the expected increase in *C* under temperatures that fluctuate evenly around T_{opt}

over a "low variance" ($T_{opt} \pm 2$ °C) interval, we find the average value of function [8] (see

Supporting Information) over 22 – 26 °C given by

[4] $$\frac{1}{4} \int_{22}^{26} (293.927 - 22.514\, T^2)\, dT$$

where development under low variance is predicted to cost 15.1 ml CO_2, equivalent to a 4.1%

increase in *C* from T_{opt}. The expected increase in *C* when incubation temperature fluctuates

evenly around T_{opt} over a "high variance" ($T_{opt} \pm 6$ °C) interval, the average value of function

[8] (see Supporting Information) over 18 – 30 °C is given by

[5] $$\frac{1}{12} \int_{18}^{30} (293.927 - 22.514\, T^2)\, dT$$

where development under high variance is predicted to cost 20.0 ml CO_2, resulting in a 38%

increase in *C* compared with development at T_{opt}.

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

metabolic rate (ml $CO_2\,h^{-1}$ ± 95% confidence interval; CI); MR_T based on previous work

(Pettersen et al., 2022), and **B)** Temperature dependence of development time (incubation

time: number of hours from oviposition to hatching ± 95% CI); 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:

low-, mid-, and high-altitude (for details, see Pettersen et al., (2023)), 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. Fluctuating

temperature regimes within this range will be encountered in the wild because of daily

variation in soil temperatures (see also While et al. 2015 and Supporting Information in

Pettersen et al., (2023)). 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 (NV: 38 ± 1 dpo,
LV: 37 ± 2 dpo, HV: 35 ± 1 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. Embryos were kept in measurement vials with the lid
removed for 60 min prior to sealing for metabolic rate measurements, which ensured they
were in thermal equilibrium with the measurement temperature when the experiment started.
Hence, metabolic rates were unlikely to depend on the direction of the ramping profile (as
shown previously in Nord and Folkow _). 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 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 (±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, cost of development, and hatchling trait measurements

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

hours from oviposition until hatching (*D*). Since hatching success was high across all

treatments (NV: 100%, LV: 99.4%, HV: 98.7%) we did not analyse survival. The cost of

development (total ml CO_2 produced) by each individual was calculated as $\dot{V}CO_2$ at each

temperature (measured within each treatment) multiplied by the time in hours spent at each

temperature. 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 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 (AKP 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 (including predictor variables Treatment, Log_n Egg mass, Region, and

their interactions, 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: $1.51\cdot10^{-2} \pm 2.75\cdot10^{-3}$ ml CO$_2$ h^{-1}) and low variance

(LV; 22-26 °C, mean ± SD: $1.38\cdot10^{-2} \pm 2.76\cdot10^{-3}$ ml CO$_2$ h^{-1}) treatments compared with the

no variance treatment (NV; 24 °C, mean ± SD: $1.26\cdot10^{-2} \pm 2.00\cdot10^{-3}$ 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: $8.10\cdot10^{-3} \pm 2.29\cdot10^{-3}$ ml CO$_2$ h^{-1}) and 75%

(mean ± SD: $1.26\cdot10^{-2} \pm 2.00\cdot10^{-3}$ 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 °C) 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). This reflected an increase in cost of 8% and 12%

above NV for LV and HV, respectively. however no significant differences between LV and

HV were found ($p = 0.07$, 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: $1.41 \cdot 10^{-3} \pm$

$1.09 \cdot 10^{-3}$g) hatched with 20% and 59% more yolk reserves, compared with the LV (mean ±

SD: $1.14 \cdot 10^{-3} \pm 8.38 \cdot 10^{-4}$g) and HV (mean ± SD: $1.02 \cdot 10^{-3} \pm 9.56 \cdot 10^{-4}$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 model output (fixed and random effects) and 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.

	Estimate	χ^2	df	p	emmean (±SE)
1a. Log_n Metabolic rate (NV)					
Intercept	-3.58	200.80	1	**<0.001***	
Log_n Egg mass	0.63	10.21	1	**<0.01**	
Mother ID (random)	0.09	0.92	1	0.34	
1b. Log_n Metabolic rate (LV)					
Intercept	-3.84	132.30	1	**<0.001***	
Temperature	24 °C: 0.15 26 °C: 0.28	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	0.46	3.15	1	0.08	
Mother ID (random)	0.18	132.66	1	**<0.001***	
1c. Log_n Metabolic rate (HV)					
Intercept	-4.60	256.09	1	**<0.001***	
Temperature	24 °C: 0.42 30 °C: 0.80	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	0.03	0.01	1	0.91	
Mother ID (random)	0.14	82.09	1	**<0.001***	
2. Log_n Development time					
Intercept	7.04	2003063.25	1	**<0.001***	

Treatment	NV: 0.10 LV: 0.10	711.43	2	**<0.001*****	NV: 1257.65 (1.00) [a] LV: 1251.38 (1.00) [a] HV: 1132.29 (1.00) [b]
Mother ID (random)	0.03	127.98	1	**<0.001*****	
3. Log_n Cost of development					
Intercept	3.43	274.05	1	**<0.001*****	
Treatment	NV: -0.12 LV: -0.04	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	0.44	7.40	1	**<0.01****	
Mother ID (random)	0.09	23.15	1	**<0.001*****	
4. Hatchling wet mass					
Intercept	0.79	816.09	1	**<0.001*****	
Treatment	NV: 0.02 LV: 0.02	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	0.36	273.23	1	**<0.001*****	
Mother ID (random)	0.02	42.92	1	**<0.001*****	
5. Hatchling tissue dry mass					
Intercept	0.16	281.35	1	**<0.001*****	
Treatment	NV: $1.40 \cdot 10^{-7}$ LV: $4.05 \cdot 10^{-4}$	3.54	2	0.17	NV: $6.76 \cdot 10^{-2}$ ($9.21 \cdot 10^{-4}$) [a] LV: $6.88 \cdot 10^{-2}$ ($9.16 \cdot 10^{-4}$) [a] HV: $6.74 \cdot 10^{-2}$ ($9.21 \cdot 10^{-4}$) [a]
Log_n Egg mass	0.08	99.46	1	**<0.001*****	
Mother ID (random)	$5.32 \cdot 10^{-3}$	46.26	1	**<0.001*****	
6. Hatchling residual yolk dry mass					
Intercept	$1.03 \cdot 10^{-3}$	62.65	1	**<0.001*****	
Treatment	NV: $3.76 \cdot 10^{-4}$ LV: $1.84 \cdot 10^{-4}$	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]
Mother ID (random)	$6.82 \cdot 10^{-4}$	51.13	1	**<0.001*****	

Figure 3. Tukey's boxplots for response variables: a) Metabolic rate (measured at 24 °C), b) Development time, c) Cost of development, d) Hatchling wet mass (tissue and yolk mass combined), e) Hatchling tissue dry mass, and f) 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) based on model output (note: raw data shown does not account for embryo mass included in all models).

Discussion

Many regions of the globe are experiencing altered patterns of temperature variability (Buckley & Huey, 2016; Wang & Dillon, 2014). 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. Despite exacerbating developmental cost, there may still be selection for shorter development times that potentially increase metabolic rate, however this relationship was not observed in our study. If faster development does incur greater overall energy cost, then it may only be selected for when resources in the postnatal environment are abundant such that any fitness consequences of high costs of development are offset. However, where the postnatal environment is characterised by low or unpredictable resource availability then the consequences of the costs of development 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. This may help to explain why our predicted increase in cost of development above NV (4% and 38% for LV and HV respectively) differed from the actual increase (8% and 12%). For *P. muralis* experiencing thermal variability during development, exposure to temperatures above and below T_{opt} *decreased* development time to a lesser extent than it *increased* overall 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., 2010a), 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; Macagno et al., 2018). 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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