Elevated exposure to prenatal thyroid hormones affects embryonic mortality but has no effects into adulthood

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Summary statement: Thyroid hormones are important hormones in all vertebrates, although overlooked in the context of maternal effect. We found short-term effects of prenatal THs but no evidence of programming effects.

Abstract

Maternal thyroid hormones (THs) are known to be crucial in embryonic development in humans, but their influence on other, especially wild, animals remains poorly understood. So far, the studies that experimentally investigated the consequences of maternal THs focused on short-term effects, while long-term organisational effects, as shown for other prenatal hormones, could also be expected. In this study, we aimed at investigating both the short- and long-term effects of prenatal THs in a bird species, the Japanese quail *Coturnix japonica*. We experimentally elevated yolk TH content (the prohormone T_4, and its active metabolite T_3, as well as a combination of both hormone) We analysed hatching success, embryonic development, offspring growth and oxidative stress as well as their potential organisational effects on reproduction, moult, and oxidative stress in adulthood. We found that eggs injected with both hormones had a higher hatching success compared with control eggs, suggesting conversion of T_4 into T_3 by the embryo. We detected no other short-term or organisational effects of yolk THs. Unfortunately, sex-specific responses could not be properly tested due to low sample sizes. These results suggest that yolk thyroid hormones are important in the embryonic stage of precocial birds, but may not have other short and long-term consequences, at least not in captivity. Research on maternal thyroid hormones will greatly benefit from studies investigating how embryos use and respond to this maternal signalling. Long-term studies on prenatal THs in other taxa in the wild are needed for a better understanding of this hormone-mediated maternal pathway.

Introduction

Maternal effects represent all the non-genetic influences of a mother on her offspring and have received increasing attention in evolutionary and behavioural ecology. Through maternal effects, mothers can influence the fitness of their progeny by adapting their phenotype to expected environmental conditions ("adaptive maternal effects" in Marshall and Uller, 2007; Mousseau and Fox, 1998), and this view is now also incorporated in the human disease literature (Gluckman et al., 2005). Maternal hormones transferred to the offspring can mediate important maternal effects. Historically, research on maternal hormones has mostly focused on steroid hormones (Groothuis et al., 2005; von Engelhardt and Groothuis, 2011). Recently, thyroid hormones (THs) also received attention in the context of hormone-mediated maternal effects (Ruuskanen and Hsu, 2018).

Thyroid hormones (THs) are metabolic hormones produced by the thyroid gland and are present in two main forms: the prohormone thyroxine (T_4) and the biologically active form triiodothyronine (T_3). THs play a crucial role in various aspects of an individual's life, e.g. development, metabolism and reproduction, across vertebrates, including humans (Krassas et al., 2010; Morreale de Escobar et al., 2004). In humans, physiological variation of maternal THs (i.e. no clinical symptoms in both mothers and foetuses) is found to be associated with infant birth weight and IQ in older children (Korevaar et al., 2016; Medici et al., 2013). In birds as well, THs play a role in brain development and neuronal turnover (reviewed in McNabb, 2007). THs control the endothermic heat production, and are therefore important in thermoregulation in juveniles and adults (McNabb and Darras, 2015).

THs can act, in concert with other hormonal axes, as mediators of life stage transitions across vertebrates (e.g., reviewed in Watanabe et al., 2016). The interaction between thyroid hormones and corticosteroids on amphibian metamorphosis is a well-known example of such effect on life stage transition (Kikuyama et al., 1993; Wada, 2008). THs are involved in gonadal development, and hyperthyroidism tends to fasten maturation (Holsberger and Cooke, 2005), and coordinate the transition between reproduction and moult (McNabb and Darras, 2015). Administration of exogenous THs is known to stop egg laying and induce moult in birds (Keshavarz and Quimby, 2002; Sekimoto et al., 1987). THs are also involved in photoperiodic control in seasonal breeding (Dardente et al., 2014). For example, thyroidectomised starlings transferred to long photoperiods became insensitive to future changes in photoperiod, and short photoperiod did not induce gonadal regression (Dawson, 1993).

While there has been recent research effort on the influence of maternal THs on offspring traits across vertebrate taxa, there are still substantial knowledge gaps. First, research on maternal thyroid hormones up to date has mainly investigated the short-term effects of prenatal THs on developing fish (Brown et al., 1988; Raine et al., 2004) and amphibians (Duarte-Guterman et al., 2010; Fini et al., 2012) and pre-fledging birds (Hsu et al., 2017; Hsu et al., 2019; Ruuskanen et al., 2016). So far, only a study on rock pigeons has looked at the influence of yolk THs on post-fledging survival and found no effect (Hsu et al., 2017). None of these studies in any taxa investigated the potential organisational effects of prenatal THs on life-history stage transitions in adult life.

Second, previous studies on prenatal THs in birds focused only on altricial species (great tits, Ruuskanen et al., 2016; rock pigeons, Hsu et al., 2017; collared flycatchers, Hsu et al., 2019). Embryonic development differs substantially between altricial and precocial species. In the latter, embryonic development is more advanced than in the former. In addition, precocial embryos start their endogenous production of TH around mid-incubation, considerably earlier than their altricial counterparts, in which endogenous TH production begins only after hatching (McNabb et al., 1998). While embryonic hormone production may limit the influence of maternal hormones, prenatal hormones have been shown to affect chick endogenous production and sensitivity (Pfannkuche et al., 2011). Overall, exposure to maternal hormones may be of different importance in these two developmental modes.

Third, previous research has studied the effects of T_3 only (Fini et al., 2012; Raine et al., 2004; Walpita et al., 2007) or a combination of T_3 and T_4 (Hsu et al., 2017; Hsu et al., 2019; Ruuskanen et al., 2016), where the effects of the two forms cannot be separated. Although T_3 is the biologically active form that binds to the receptors, both are deposited in eggs (Prati et al., 1992) and T_4 may be converted to T_3 via deiodinases from the mother or the developing embryo (Van Herck et al., 2015) or may still exert non-genomic actions (reviewed in Davis et al., 2016). Manipulating yolk T_4 and T_3 independently would help understanding the relative contribution of these two hormones.

In this study, we aimed at assessing the effects of maternal THs on development and life-history traits in a precocial bird species, the Japanese quail (*Coturnix japonica*). To do so, manipulated eggs received either an injection of T_4 or T_3 separately, a combination of both hormones, or a control injection of the vehicle saline solution. This design allowed us to explore the effects of T_4 and T_3 separately. The elevation in yolk THs remained within the natural range of this species, a crucial condition to obtain relevant results for an eco-evolutionary context. We measured traits known to be influenced by circulating and yolk THs:

hatching success, age at embryonic mortality, growth, transition between life-history stages (i.e., reproductive state and moult) and oxidative stress. First, we hypothesise that elevation of yolk THs in Japanese quails positively affects hatching success, as found in two studies on collared flycatchers and rock pigeons (Hsu et al., 2017; Hsu et al., 2019, but see Ruuskanen et al., 2016). Second, elevation of yolk THs is predicted to increase the proportion of well-developed embryos before hatching, as found in rock pigeons (Hsu et al., 2017). We therefore looked at the age at mortality in unhatched eggs. Third, we expect elevated yolk THs to affect chick growth (in body mass, tarsus and wing length) either positively (Hsu et al., 2019; Wilson and McNabb, 1997), negatively (Hsu et al., 2017), or in a sex-specific manner (Ruuskanen et al., 2016). Fourth, we predict that yolk THs will have organisational effects on life-history stage transitions; that is, age at sexual maturity and male gonadal regression (using cloacal gland size as a proxy), and moult when birds are exposed to short photoperiod. Based on the literature mentioned above we expect elevated yolk THs to advance the timing of puberty, gonadal regression and moult. The rate of moult should also be influenced, with birds receiving experimental TH elevation moulting faster. We also explored the effects of yolk THs on reproductive investment in females, another important fitness aspect. Finally, yolk THs may increase oxidative stress due to their stimulating effects on metabolism.

Material and Methods

Parental generation and egg collection

The parental generation was composed of adult Japanese quails provided by Finnish private local breeders that were kept in two acclimated rooms. Twenty-four breeding pairs were formed by pairing birds from different breeders. Individuals were identified using metal leg bands. The floor was covered with 3–5cm sawdust bedding. A hiding place, sand and calcium grit were provided. Each pair was housed in a 1 m² pen. The temperature was set to 20°C with a 16L:8D photoperiod (light from 06.00 to 22.00). Food (Poultry complete feed, “Kanan Paras Täysrehu”, Hankkija, Finland) was provided *ad libitum* and water was changed every day.

Pairs were monitored every morning to collect eggs. Eggs were individually marked (non-toxic marker), weighed and stored for 7 days in a climate-controlled chamber at 15°C and 50% relative humidity. On the last day of collection, an average of 6.6 eggs per pair (range = 4–8 eggs) were injected with a solution (see next section).

136 ***Preparation of the solution, injection procedure and incubation***

137 The preparation of hormone solution and the procedure of injection were based on previous
138 studies (Hsu et al., 2017; Ruuskanen et al., 2016). In brief, crystal T_4 (L-thyroxine, ≥ 98%
139 HPCL, CAS number 51-48-9, Sigma-Aldrich) and T_3 (3,3',5-triiodo-L-thyronine, > 95%
140 HPCL, CAS number 6893-02-3, Sigma-Aldrich) were first dissolved in 0.1M NaOH and then
141 diluted in 0.9% NaCl. The injection of thyroid hormones resulted in an increase of two
142 standard deviations (T_4 = 8.9 ng/egg; T_3 = 4.7 ng/egg), a recommended procedure for
143 hormone manipulation within the natural range (Hsu et al., 2017; Podmokła et al., 2018;
144 Ruuskanen et al., 2016). The control solution (CO) was a saline solution (0.9% NaCl). The
145 concentrations of the hormone solutions were based on previous measurements of 15 eggs
146 from the same flock (T_4 content per egg (SD) = 15.3 (4.4) ng, T_3 content per egg (SD) = 7.6
147 (2.3) ng).

148 Hormone injections were performed at room temperature in a laminar hood. Eggs
149 were put sideways, allowing yolks to float up to the middle position. Before injection, the
150 shell was disinfected with a cotton pad dipped in 70% EtOH. We used a 27G needle (BD
151 Microlance ™) to pierce the eggshell and then used a 0.3 ml syringe to deliver 50 µl of the
152 respective hormone solution or control. After injection, the hole was sealed with a sterile
153 plaster (OPSITE Flexigrid, Smith&Nephew).

154 In total, 158 eggs were injected and divided as follows over the treatments: T_3
155 treatment (N = 39); T_4 treatment (N = 39); T_3+T_4 treatment (N = 40); and control, CO (N =
156 40). To balance the genetic background of the parents and the effect of storage, each egg laid
157 by the same female was sequentially assigned to a different treatment and the order of
158 treatments was rotated among females. After injection, eggs were placed in an incubator at
159 37.8°C and 55% relative humidity. Until day 14 after starting incubation, eggs were
160 automatically tilted every hour by 90°. On day 14, tilting was halted and each egg was
161 transferred to an individual container to monitor which chick hatched from which egg. On day
162 16 after injection, (normal incubation time = 17 days), the temperature was set to 37.5°C and
163 the relative humidity to 70%. Eggs were checked for hatching every 4 hours from day 16
164 onwards. Four days after the first egg hatched, all unhatched eggs were stored in a freezer and
165 dissected to determine the presence of an embryo. The age of developed embryos was
166 assessed according to Ainsworth et al. (2010).

167 ***Rearing conditions of the experimental birds***

168 In total, 66 chicks hatched (N = 10 CO, 15 T_3, 20 T_4 and 21 T_3T_4). The overall hatching

success was rather low (ca. 40%), partly due to the injection procedure itself (Groothuis and von Engelhardt, 2005), although low hatching success in quails has also been reported in unmanipulated conditions previously (e.g. Okuliarová et al., 2007). Among the unhatched eggs, 33.7% (31 out of 92) had no developed embryos. Twelve hours after hatching, the chicks were marked by a unique combination of coloured rings and nail coding and transferred to two 1 m² cages (ca. 30 chicks/cage, sex and treatments mixed together). The chicks were provided with heating mats and lamps as extra heat sources for the first two weeks. The chicks were fed with sieved commercial poultry feed ("Punaheltta paras poikanen", Hankkija, Finland), and provided with Calcium and bathing sand. Two weeks after hatching, the chicks were separated in four 1 m² cages of about 16 individuals. Around 3 weeks after hatching, coloured rings were replaced by unique metal rings. On week 4 after hatching, birds were transferred to eight 1 m² pens (average of 7.1 birds/pen, range = 4–9), under the same conditions as the parents. Around the age of sexual maturity (ca. 6–8 weeks after hatching), the birds were separated by sex in twelve 1 m² pens (average of 4.8 birds/pen, range = 4–5).

Monitoring of growth and reproductive maturation

Body mass and wing length were measured twelve hours after hatching. Tarsus was not measured because it bends easily, resulting in inaccurate measures and potential harm for the young. From day 3 to day 15, these three traits were monitored every 3 days. From day 15 to day 78 (ca. 12 weeks), chicks were measured once a week. Body mass was recorded using a digital balance to the nearest 0.1 g. Wing and tarsus lengths were respectively measured with a ruler and a calliper to the nearest 0.5 mm and 0.1 mm. From week 6 to week 10, we monitored cloacal gland development and foam production in 28 males. Cloacal glands were measured every other day with a calliper to the nearest 0.1 mm as a proxy for testes development and sexual maturation (Biswas et al., 2007). Foam production (by gently squeezing the cloacal gland) was assessed at the same time and coded from 0 (no foam) to 3 (high production of foam), as a proxy of cloacal gland function (Cheng et al., 1989a; Cheng et al., 1989b). The same observer performed all measurements. We collected eggs produced by 10-week-old females over a 6-day period, and measured the short and long axes of the eggs with a calliper to the nearest 0.01 mm and record their mass to the nearest 0.1 g. We collected on average 5.7 eggs (range = 4–7) per female from 28 females.

Monitoring of cloacal gland regression and moult

In Japanese quails, exposure to short photoperiod and cold temperature triggers reproductive inhibition and postnuptial moulting (Tsuyoshi and Wada, 1992). Thyroid hormones are known to coordinate these two responses (see introduction). When the birds reached the age of ca. 7 months, we exposed birds to short photoperiod (8L:16D, i.e., light from 08.00 to 16.00) with a 12:12-h cycle of normal (20°C) and low (9°C) temperature (low temperature was effective from 18.00 to 06.00). Cloacal gland regression (as a proxy for testes regression) was monitored every other day for 2 weeks with a calliper by measuring the width and length to obtain the area of the gland to the nearest 0.1 mm^2 (N = 26 males). Primary moult was recorded from a single wing by giving a score to each primary from 0 (old feather) to 5 (new fully-grown feather) following Ginn and Melville (1983) (N = 54 males and females). The total score of moulting was obtained by adding the score of all feathers.

Oxidative status biomarker analyses

Two blood samples were drawn, when birds were 2 weeks (N = 51 chicks) and 4 months old (N = 49 adults), respectively. 200 µl of blood was collected from the brachial vein in heparinized capillaries and directly frozen in liquid nitrogen. Then, the samples were stored at -80°C until analyses. We measured various biomarkers of antioxidant status; the antioxidant glutathione (tGSH), the ratio of reduced and oxidised glutathione (GSH:GSSG) and activity of the antioxidant enzymes glutathione peroxidase (GPx), catalase (CAT) and superoxide dismutase (SOD) from the blood. It is important to measure multiple biomarkers of oxidative and antioxidant status for a broader understanding of the mechanism. Also, the interpretation of the results is more reliable if multiple markers show similar patterns. Of the measured biomarkers, the ratio of GSH:GSSG represents the overall oxidative state of cells and consequently, deviations in this ratio is often used as an indicator of oxidative stress (Halliwell and Gutteridge, 2015; Hoffman, 2002; Isaksson et al., 2005; Lilley et al., 2013; Rainio et al., 2013). GPx enzymes catalyse the glutathione cycle, whereas CAT and SOD directly regulate the level of reactive oxygen species (ROS) (Ercal et al., 2001; Halliwell and Gutteridge, 2015). The methodology for measuring each biomarker is described in detail in Rainio et al. (2015). All analyses were conducted blindly of the treatment. Briefly, the samples were analyzed using a microplate reader (EnVision, PerkinElmer-Wallac, Finland). All antioxidant and enzyme activities were measured in triplicate (intra-assay coefficient of variability [CV] < 10% in all cases) using 96-(CAT) or 384-well (GPx, SOD, tGSH and GSH:GSSG) microplates. Three control samples were used with each plate, to be able to

correct inter-assay precision with the ratio specific to the particular plate. Overall protein concentration (mg/ml) was measured according to the Bradford method (Bradford, 1976) using BioRad stock (BioRad, Finland) diluted with dH_2O (1:5) and BSA (bovine serum albumin, 1 mg/ml) (Sigma Chemicals, USA) as a standard. GPx-assay was conducted using Sigma CGP1 kit, CAT-assay using SigmaCAT100 kit and SOD-assay using Fluka 19160 SOD determination kit. Total GSH and the ratio of GSH:GSSG were measured with the ThioStar® glutathione detection reagent (Arbor Assays, USA) according to kit instructions, using reduced glutathione as a standard (Sigma Chemicals, USA).

Ethics

The study complied with Finnish regulation and was approved by the Finnish Animal Experiment Board (ESAVI/1018/04.10.07/2016).

Statistical analysis

Data were analysed with the software R version 3.5.3 (R core team, 2019). In this study, two different statistical approaches were used: null-hypothesis testing with Generalised Linear Mixed Models (GLMMs) and Linear Mixed Models (LMMs), and multimodel inference with Generalised Additive Mixed Models (GAMMs). GAMMs were used to analyse the data on body and cloacal gland growth to account for its non-linear pattern (see *Growth*). In this analysis, we preferred multimodel inference as GAMMs generate many candidate models that cannot be directly compared (e.g., by the Kenward-Roger approach). Instead, candidate models were ranked based on their Akaike Information Criterion (AIC) values. Models with a $\Delta AIC \leq 2$ from the top-ranked model were retained in the set of best models. Akaike weights of all models were calculated following (Burnham and Anderson, 2002), and evidence ratios of the top-ranked models were calculated as the weight of a model divided by the weight of the null model (Burnham et al., 2011). To estimate the effect of the predictors, we computed the 95% confidence intervals from the best models using the *nlme* package (Pinheiro et al., 2018). GLMMs and LMMs were fitted using the R package *lme4* (Bates et al., 2015), and GAMMs were fitted using the package *mgcv* (Wood, 2017). P-values for GLMMs were obtained by parametric bootstrapping with 1,000 simulations and p-values for LMMs were calculated by model comparison using Kenward-Roger approximation, using the package *pbkrtest* in both cases (Halekoh and Højsgaard, 2014). Post-hoc Tukey analyses were conducted with the package *multcomp* (Hothorn et al., 2008). Due to our experiment design, eggs injected with both hormones received a higher absolute amount of hormones than eggs

injected with T_4 or T_3 only. Therefore, we also tested a potential dose-dependent effect of the treatment on the response variables when treatment groups showed significant differences. Model residuals were checked visually for normality and homoscedasticity. Covariates and interactions were removed when non-significant ($\alpha = 0.05$).

Hatching success

To analyse hatching success, each egg was given a binary score: 0 for unhatched egg and 1 for hatched egg. A series of GLMMs were fitted with a binomial error distribution (logit link) and mother identity as a random intercept. The first model included the 4-level treatment (treatments: CO, T_3, T_4 and T_3T_4) as the predictor, while in a second model treatment was converted into an ordered variable, following the increasing levels (i.e. CO, T_3, T_4 and T_3T_4). The second model was meant to test for a potential dose-dependent effect as the eggs received an increasing amount of total THs as a potential source of T_3, the most potent hormone. Egg mass might affect hatchability and was therefore added as a covariate in both models. The potential effect of storage duration on hatchability (Reis et al., 1997) was accounted for by including laying order as a covariate in both models.

Duration of embryonic period, age at embryonic mortality and mass at hatching

Duration of embryonic period and mass at hatching were modelled with LMMs. Treatment, sex of the individuals and egg mass were included as fixed factors. Laying order was added as a covariate to account for potential effects of storage duration on hatching time and on chick weight (Reis et al., 1997). Mother identity was included as a random intercept. In the model for mass at hatching, duration of embryonic period was further added as a covariate.

The data for embryonic age had a skewed distribution and residuals were not normally distributed and heterogenous, which violated LMM assumptions on residual distribution. We therefore performed a simple Kruskal-Wallis test.

Growth

As growth curves typically reach an asymptote, we fitted non-linear GAMMs to these curves. Growth in body mass, tarsus and wing length were analysed in separate GAMMs. Growth was analysed until week 10 after hatching as all birds appeared to have reached their maximum body mass and tarsus and wing length. The data are composed of repeated measurements of the same individuals over time; therefore, we first corrected for temporal autocorrelation between the measurements using an ARMA(1,1) model for the residuals (Zuur et al., 2009). Second, as mothers produced several eggs, the models included nested random

effects, with measured individuals nested into mother identity, allowing for random intercepts. GAMMs allow modelling the vertical shift of the curves (i.e., changes in intercepts) and their shape. Treatment and sex were included as predictors. A smoothing function for the age of the birds was included to model the changes in the growth curves, and was allowed to vary by sex or treatment only, or none of these predictors. The interaction between sex and treatment was not analysed due to low statistical power. Additive effect of treatment and sex was tested for the intercept but could not be computed for curve shape. All combinations of the relevant predictors were tested for both shape parameters (i.e., intercept and curve shape).

Reproductive maturation, regression and investment

Due to low sample sizes in sex-specific responses, we could not perform robust statistical analyses. We therefore present these analyses and results in the supplementary material and only briefly discuss them.

Oxidative stress

A principal component analysis (PCA) was first performed on measured antioxidant markers (SOD, CAT, GPx, tGSH and GST), to reduce the number of metrics for subsequent analyses. The first and the second principal components (PCs) explained together 60.2% of the variance (Table 1). PC1 and PC2 were then used as dependent variables in separate LMMs. LMMs included the treatment, sex and age of individuals (2 weeks and 4 months old) as fixed factors and the 2-way interactions between treatment and sex, and treatment and age. Mother and individual identities, to account for repeated measures, were added as random intercepts. Malondialdehyde (MDA) is a marker of oxidative damage, which is a different measure from antioxidant activity, and was therefore analysed in a separate LMM using the same parameters as for PC1 and PC2, adding the batch of the assay as an additional random intercept. The marker of cell oxidative status (GSH:GSSG ratio) was analysed with the same model used for PC1 and PC2.

Moult

Two parameters of moult were analysed in separate LMMs: the timing of moult (i.e., the moult score after one week of short photoperiod), and the rate of moult (i.e., how fast birds moulted). Both models included treatment and sex as fixed factors, and mother identity as a random intercept. The rate of moult was tested by fitting an interaction between treatment and age. This model also included the main effect of age and individual identity, nested within mother identity, as a random intercept to account for repeated measures. Estimated marginal

means and standard errors (EMMs ± SE) were derived from the model using the package *emmeans* (Lenth, 2019).

Results

Effects of prenatal THs on hatching success and age of embryo mortality

There was a significant effect of elevated prenatal THs on hatching success (GLMM, p = 0.05, Fig. 1). Tukey post-hoc analysis revealed that hatching success in the T_3T_4 group was significantly higher than in the CO group (Estimate±SE = 1.24±0.50, Tukey z = 2.46, p = 0.05). The other groups (T_3 and T_4) were not different from the control group (all $z < 2.22$ and $p > 0.09$). The data suggested a dose-dependent effect that we tested by changing the treatment factor to an ordered variable, as the eggs received an increasing amount of TH from T_3 to T_3T_4 injections ($T_3 < T_4 < T_3T_4$). We found a linear positive dose-dependent effect of yolk TH elevation (GLMM, Estimate±SE = 0.96±0.36, z = 2.69, p = 0.007), but no quadratic or cubic effects ($p > 0.53$).

Dissection of the unhatched eggs showed that age of embryo mortality did not differ between the treatments (Kruskal-Wallis $\chi^2 = 7.22$, df = 3, p = 0.07). Finally, the manipulation of yolk THs did not affect the duration of embryonic period (LMM, $F_{3,42.0} = 0.57$, p = 0.64, Fig. S1). Sex of the embryo or egg mass (LMM sex, $F_{1,49.7} = 2.63$, p = 0.11; LMM egg mass, $F_{1,19.3} = 0.01$, p = 0.92) were also not associated with the duration of the embryonic period.

Effects of prenatal THs on growth

Mass at hatching was not influenced by the elevation of prenatal THs (LMM, $F_{3,35.0} = 0.81$, p = 0.50, Fig. S2). Mass at hatching was positively correlated with egg mass (LMM, Estimate±SE = 0.72±0.10 g, $F_{1,24.1} = 46.9$, $p < 0.001$), while duration of embryonic period was negatively correlated with mass at hatching (LMM, Estimate±SE = -0.008±0.003 g, $F_{1,46.7} = 4.49$, p = 0.04).

Regarding body mass growth, the top-ranked model showed that the curve shape and the intercept differ according to sex (Table 2). After 10 weeks, females had a larger body mass than males (mean±SE females = 214.4±5.7 g, males = 172.4±4.5 g, Fig. 2), which was supported by the 95% CIs (Table 3). Based on model selection we conclude that the treatment had no effect on body mass growth (Table 2).

For wing length, the top-ranked model ($\Delta AIC \leq 2$) included sex in the intercept, while treatment was not included in the best supported model (Table S1). The 95% CIs (Table 3)

confirmed that males had a lower wing length than females (Fig. S3).

Concerning tarsus length, the models within $\Delta AIC \leq 2$ included no predictors for the curve shape but included treatment for the intercept (Table S2). The 95% CIs of the parameter estimates from these models suggested that there was a slight negative effect of T_3T_4 treatment on tarsus growth (Table 3, Fig. S4). However, as the estimates were close to 0 (Table 3) and evidence ratios showed that the model with treatment as a predictor was only 3.5 times more supported than the null model (Table S2), we conclude that the effect of THs on tarsus length is likely to be very small. Likewise, the second model for tarsus length included sex as a predictor for the intercept, but its 95% CIs overlapped with 0 (Table 3). We therefore conclude that sex had no effect on tarsus growth.

Effects of prenatal THs on postnuptial moult

As expected, birds started to moult soon after being exposed to short photoperiod, with an average increase of moult score by 6 per week (SE = 0.2, $F_{1,254.0} = 827.4$, $p < 0.001$, Fig. 3). The first moult score (assessed one week after switching to short photoperiod) was not affected by the treatment (LMM, $F_{3,42.7} = 0.36$, $p = 0.78$), but was influenced by sex, with females having a higher score than male (EMMs ± SE: female = 21.4 ± 1.6, male = 7.2 ± 1.7; LMM $F_{1,45.3} = 41.9$, $p < 0.001$). Yolk TH elevation did not affect the rate of moult (LMM interaction treatment × time, $F_{3,251.0} = 0.59$, $p = 0.62$, Fig. 3).

Effects of prenatal THs on oxidative stress

The elevation of yolk THs had no effect on PC1 or PC2 of antioxidants at either 2 weeks ("chicks") or 4 months ("adults") old (LMM on PC1, $F_{3,40.3} = 2.40$, $p = 0.08$; LMM on PC2, $F_{3,42.2} = 0.92$, $p = 0.44$, treatment × age, $F < 0.91$, $p > 0.44$). The age of the birds had a highly significant effect on PC1, with chicks generally having higher antioxidant capacities (CAT, GST and tGSH) than adults (LMM, Estimate±SE = -1.34±0.19, $F_{1,49.2} = 52.1$, $p < 0.0001$). All the other predictors had no effect on either PC1 or PC2 (all $F < 2.93$ and all $p > 0.09$).

The marker of oxidative damage, MDA, was affected by the elevation of yolk THs (LMM, $F_{3,43.6} = 3.08$, $p = 0.04$, Fig. 4). Tukey post-hoc analysis showed that the T4 group had higher MDA values than the T3 group (Estimate±SE = 0.01±0.004, Tukey contrast $p = 0.01$), but none of the groups differed from the control (Tukey p-values > 0.19). However, this result became non-significant when removing the outlier in the T_4 group (LMM, $F_{3,43.1} = 2.68$, $p = 0.06$). MDA levels were not affected by the age or the sex of individuals (LMM age, $F_{1,54.4} = 0.30$, $p = 0.59$; LMM sex, $F_{1,42.0} = 1.47$, $p = 0.23$).

The marker of cell oxidative balance, GSH:GSSG, was not influenced by the yolk THs nor by the sex of the birds (LMM treatment, $F_{3,33.0} = 0.85$, p = 0.48; LMM sex, $F_{1,40.6} = 0.57$, p = 0.45). However, chicks had a higher GSH:GSSG ratio than adults (LMM, Estimate±SE = 0.17±0.04, $F_{1,50.0} = 18.3$, p < 0.0001).

Discussion

The aim of this experimental study was to investigate the short-term and organisational effects of maternal thyroid hormones (THs) in a precocial species, the Japanese quail, by experimental elevation of THs in eggs. Our study is the first to investigate the effects of yolk T_3 and T_4 separately, within the natural range of the study model. In addition we studied both short and long term effects on embryonic development, growth, life stage transitions and oxidative stress. We only detected a positive effect of yolk THs on hatching success. All other response variables studied were not affected by elevated prenatal THs.

Effects of prenatal THs on hatching success and embryonic development

We found that hatching success increased when the eggs received an injection of both T_4 and T_3. Previous similar studies reported comparable effects of yolk THs in rock pigeons (Hsu et al., 2017) and in collared flycatchers (Hsu et al., 2019). In these studies, injections consisted of a mixture of both T_3 and T_4. Our results point towards a dose-dependent effect of yolk THs, as found previously of androgen hormones (e.g., Muriel et al., 2015). Importantly, given that only T_3 binds to receptors, these results also suggest that embryos must express deiodinase enzymes to convert T_4 to T_3, and/or yolk may contain maternally derived deiodinase mRNA. Indeed, precocial embryos start to produce endogenous T_3 and deiodinase expression has previously been characterised in chicken embryos (Darras et al., 2009; Van Herck et al., 2012). In contrast with our study, a similar study in great tits detected no increased hatching success due to the injection of THs (Ruuskanen et al., 2016). The dissimilarities between the studies may come from inter-specific differences in terms of utilisation of yolk THs by the embryos or from context-dependent effects (e.g. due to other egg components). Further comparative and mechanistic studies could help understanding the dynamic of yolk THs during incubation.

Increased yolk THs did not improve age of embryo mortality. Similar to our study, Ruuskanen et al. (2016) did not find any difference in the timing of mortality in great tit embryos. Conversely, the study on rock pigeons found that yolk THs increased the proportion of well-developed embryos (Hsu et al., 2017). Similarly to our result on hatching success,

yolk THs' effects on embryonic development may differ in a species-specific manner.

Effects of prenatal THs on growth

We found no influence of yolk THs on growth, contrary to our expectations based on the recent literature. Other comparable studies found either a positive (Hsu et al., 2019), a negative (Hsu et al., 2017) or a sex-specific effect (Ruuskanen et al., 2016) of yolk THs on growth. This notable difference may be due to the captive conditions experienced by the Japanese quails in our study, with unrestricted access to food and water. Although the pigeon study also provided ad libitum food, parents still needed to process food before feeding their nestlings in the form of crop milk, whereas precocial quails have no such limitation. In addition, the Japanese quail has been domesticated for many generations, and probably selected for rapid growth for economic reasons. Whole-genome sequencing in chickens showed that domestication induced a strong positive selection on genes associated with growth (Rubin et al., 2010). Interestingly, that study also found a strong selection for a locus associated with thyroid stimulating hormone (TSH) receptor. TSH controls most of the TH production by the thyroid gland (McNabb and Darras, 2015), and this artificial selection may overshadow the effects of natural variations of prenatal THs on growth. Besides, the low number of individuals in the control and T_3 groups (7 and 11, respectively) may have limited statistical power to detect differences between the treatments. Repeating the study with a larger sample size may allow us to ascertain the effects of yolk THs on growth in precocial study models. Research on the influences of prenatal THs on growth will also benefit from experimental studies on wild precocial species.

Effects of prenatal THs on postnuptial moult

Short photoperiod in combination with cold temperature triggered primary moult, as expected. However, we detected no effect of yolk THs on the timing or speed of moult. Thyroid hormones are important in moult and feather growth (reviewed in Dawson, 2015). For example, thyroidectomised birds fail to moult after being exposed to long photoperiods (Dawson, 2015). In addition, thyroidectomised nestling starlings failed to grow normal adult plumage and grown feathers presented an abnormal structure (Dawson et al., 1994). By removing the thyroid gland, these two studies implemented extreme pharmacological protocols that differ drastically from our injection of physiological doses. In addition, our experimental design, increasing TH exposure (vs decreased TH exposure in the above-mentioned studies), may have different consequences. For example, there may be a threshold

456 above which any, additional hormones may not affect moult.

457 Overall, our results show no support for the hypothesis of programming effect of
458 prenatal THs on life stage transitions. Yet, due to small sample sizes in sex-specific analyses
459 (i.e., male gonadal maturation and regression, and female reproductive investment), there
460 remains a relatively high uncertainty about the potential programming effects of prenatal THs.
461 Replicate studies with larger samples sizes and different study models will reduce this
462 uncertainty.

463 ***Effects of prenatal THs on oxidative stress***

464 In contrast to our predictions, elevated yolk THs did not affect oxidative status during chick or
465 adult phase. We found no changes in antioxidant activities in relation to yolk THs and no
466 imbalance in the oxidative cell status. Nevertheless, T_4 birds had a higher level of oxidative
467 damage on lipids than T_3 birds, but this was a weak effect driven by one outlier. The lack of
468 effects on chick oxidative status among the treatment groups could be explained by the
469 absence of treatment effects on growth, given that high growth rates usually result in higher
470 oxidative stress and damage (e.g. Alonso-Alvarez et al., 2007). In turn, the lack of treatment
471 effects on adult oxidative status may suggest no organisational effects of prenatal THs on
472 adult metabolism. A recent study in an altricial species also found no influence of yolk THs on
473 nestling oxidative stress (Hsu et al., 2019). Our study shows for the first time that prenatal
474 THs have no influence on adult oxidative stress either. The previous study focused on a
475 limited set of biomarkers: one antioxidant enzyme, oxidative damage on lipids and oxidative
476 balance. In the present study, we measured 7 biomarkers, thus providing broader support to
477 the absence of effects of prenatal THs on oxidative stress.

478 ***Conclusion***

479 To our knowledge, this study is the first one to experimentally investigate the consequences of
480 natural variations of maternal THs not only early but also in adult physiology and postnuptial
481 moult in any vertebrate. Furthermore, this study explored for the first time the effects of
482 maternal T_3 and T_4 separately. We found no evidence for differential effects of maternal T_4
483 and T_3, while a dose-dependent effect on hatching success suggests that T_4 is converted into
484 T_3, the biologically active form during embryonic development. Contrary to similar studies on
485 wild altricial species, we found no influence of maternal THs on growth. Further research on
486 embryos utilisation of maternal THs may help understand the differences observed between
487 precocial and altricial species. Studies in other vertebrates are urgently needed to understand

488 the potential for long-term organising effects of maternal THs.

List of symbols and abbreviations

- CAT: catalase
- CO: control treatment
- GP: glutathione peroxidase
- tGSH: oxidised glutathione
- GSSG: reduced glutathione
- GST: Glutathione S-transferase
- MDA: malonaldehyde
- RMR: resting metabolic rate
- SOD: super-oxide dismutase
- T_3: triiodothyronine
- T_4: thyroxine
- THs: thyroid hormones

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Competing interests

We declare no competing interests.

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Figure legends

Figure 1: Percentage of hatching success according to yolk TH manipulation treatments: CO (N = 40), T_3 (N = 39), T_4 (N = 39), T_3T_4 (N = 40). CO = control, T_4 (thyroxine) = injection of T_4, T_3 (triiodothyronine) = injection of T_3, T_3T_4 = injection of T_3 and T_4.

Figure 2: Growth curves in body mass of Japanese quails hatching from eggs treated with either T_3, T_4, a combination of both hormones, or a control solution. See Fig. 1 for a description of the treatments. Each line represents an individual bird, while thick coloured lines represent mean values. A: Growth curve according to yolk TH manipulation. N = 7 CO, 11 T_3, 18 T_4 and 21 T_3T_4. B: Growth curve according to sex. N = 29 females and 28 males.

Figure 3: Primary moult score in 7-month old Japanese quails according to yolk TH manipulation treatments: CO (N = 7), T_3 (N = 11), T_4 (N = 16), T_3T_4 (N = 20). See Fig. 1 for a description of the treatments. Measures were taken once a week after switching from long photoperiod (16L:8D) to short photoperiod (8L:16D, switch = time point 0 on x-axis). Each line represents an individual bird, while thick coloured lines represent group mean values.

Figure 4: MDA concentration according to yolk TH manipulation treatments, samples from two ages pooled: CO (N = 7 individuals), T_3 (N = 11), T_4 (N = 17), T_3T_4 (N = 20). See Fig. 1 for a description of the treatments.

Figures

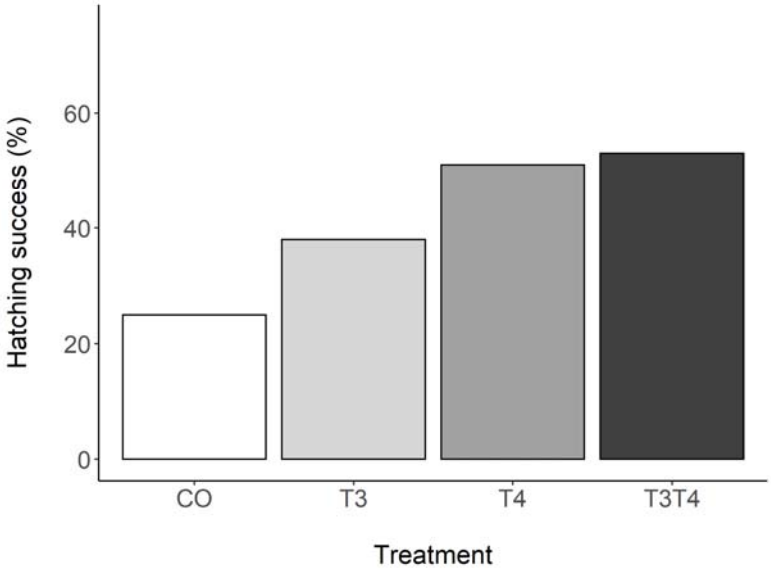

Figure 1

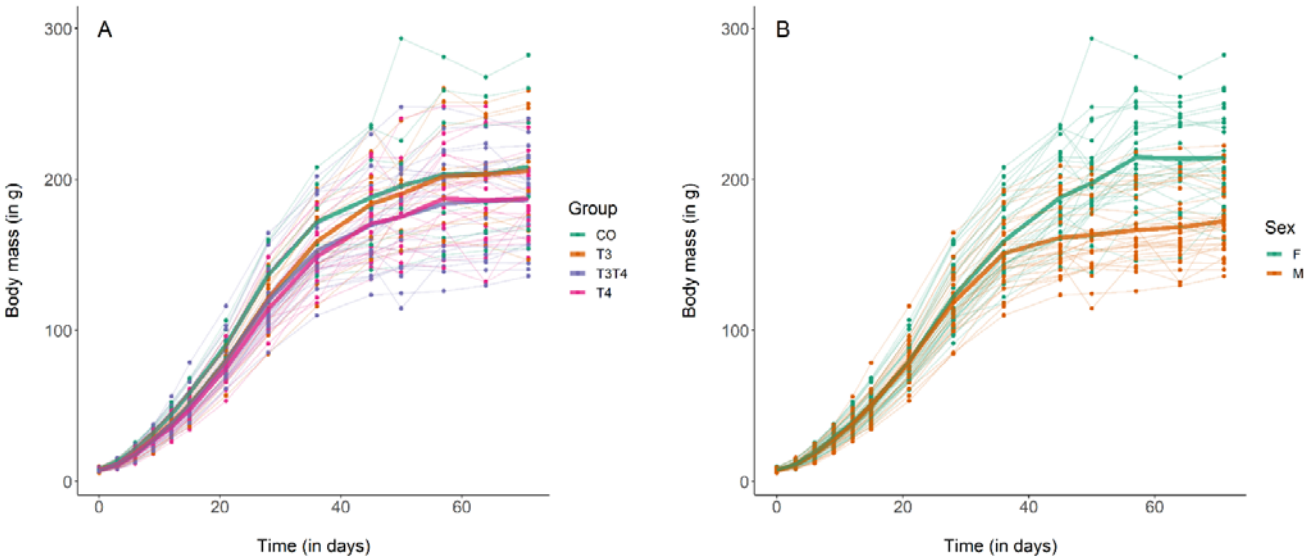

Figure 2

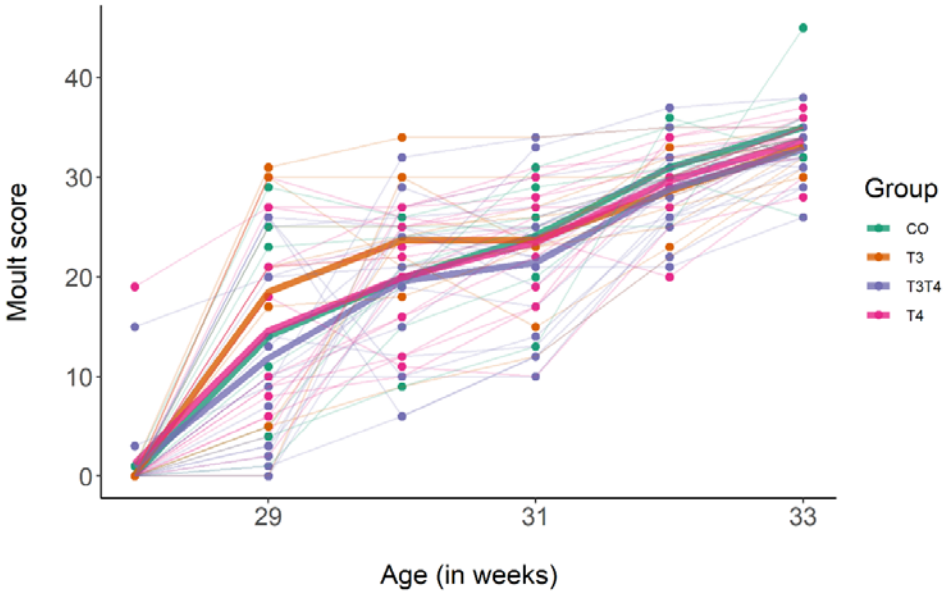

Figure 3

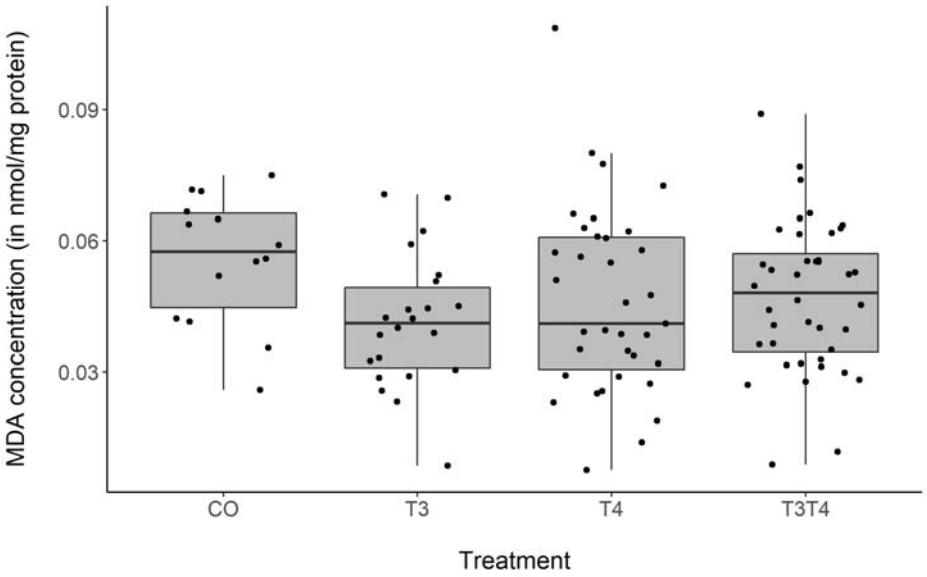

Figure 4

Tables

Table 1: Loadings of the different antioxidant biomarkers on the principal components 1 and 2.

Factor loadings	PC1 (34.0%)	PC2 (26.2%)
CAT	-0.49	0.14
SOD	0.20	-0.71
GST	-0.65	-0.10
GP	0.04	-0.63
tGSH	-0.60	-0.26

Table 2: Results of the Generalised Additive Mixed Models (GAMMs) on body mass growth, with sex and treatment fitted either as intercept, curve shape or both (all combinations tested). A total of 12 GAMMs were fitted and ranked based on their AIC, from the lowest to the highest. Weight: Akaike's weights.

Model	Intercept	Curve shape	ΔAIC	df	Weight
1	Sex	Sex	0.0	11	0.8430
8	Treatment + sex	Sex	3.5	14	0.1497
3	-	Sex	9.9	10	0.0061
2	Treatment	Sex	13.2	13	0.0012
11	Sex	-	77.6	9	<0.001
9	Treatment + sex	-	81.6	12	<0.001
12	-	-	91.2	8	<0.001
10	Treatment	-	95.0	11	<0.001
5	Sex	Treatment	147.9	15	<0.001
7	Treatment + sex	Treatment	151.7	18	<0.001
6	-	Treatment	161.2	14	<0.001
4	Treatment	Treatment	165.5	17	<0.001

Table 3: 95% confidence intervals of the predictors in the top-ranked models according to AIC values (see Tables 2, S1 and S2). Predictors in bold have confidence intervals that do not overlap with 0. For the intercept, the reference groups are female and CO for the predictors sex and treatment, respectively.

Curve parameter	**Predictors**	**Lower limit**	**Estimate**	**Upper limit**
(A) Body mass (Model 1)				
Intercept	**Sex (M)**	-19.7	-12.6	-5.5
Curve shape	**Sex (F)**	9.9	20.0	30.0
Curve shape	**Sex (M)**	14.3	24.5	34.7
(B) Wing length (Model 11)				
Intercept	**Sex (M)**	-2.3	-1.2	-0.1
Curve shape	**Age**	26.4	28.7	31.0
(C) Tarsus length (Model 10)				
Intercept	Treatment (T_3)	-0.8	0.02	0.8
Intercept	**Treatment (T_3T_4)**	-1.5	-0.8	-0.1
Intercept	Treatment (T_4)	-1.3	-0.6	0.2
Curve shape	**Age**	10.5	11.1	11.8
Tarsus length (Model 9)				
Intercept	Treatment (T_3)	-0.9	-0.07	0.7
Intercept	**Treatment (T_3T_4)**	-1.5	-0.8	-0.1
Intercept	Treatment (T_4)	-1.4	-0.6	0.1
Intercept	Sex (M)	-0.8	-0.3	0.3
Curve shape	**Age**	10.5	11.1	11.7