

# Can increased prenatal exposure to thyroid hormones alter physiology and behaviour in the long-term? Insights from an experimental study in Japanese quails

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1	Can increased prenatal exposure to thyroid hormones alter physiology
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4	
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# 32 Abstract

33 Maternal thyroid hormones (triiodothyronine T3 and thyroxine T4) are important regulators of embryonic development and gene expression. While maternal thyroid 34 35 disorders are known to cause developmental issues in humans, variation in maternal thyroid hormones in healthy mothers have also been found to correlate with infant 36 37 and child phenotypes. This suggests a potential impact of maternal thyroid hormones on offspring phenotype in an eco-evolutionary context. In chickens, prenatal thyroid 38 39 hormone treatment has been shown to influence embryonic gene expression and postnatal treatment to influence imprinting and learning. However, the potential long-40 41 term effects of maternal thyroid hormones on physiology and behaviour are unclear. 42 This study aims to investigate the long-term effects of maternal thyroid hormones on 43 behaviour, plasma thyroid hormone levels and brain gene expression using the 44 Japanese quail (*Coturnix japonica*) as a model. Egg hormone levels were elevated by 45 injecting unincubated eggs with either saline (control), T3, T4 or a mixture of T3 and 46 T4. Social motivation, boldness and fearfulness to predators were tested shortly after 47 hatching and as adults. Plasma thyroid hormone levels and pallial expression of thyroid hormone receptor A, type 2 deiodinase, and nuclear receptor coactivator 1 were 48 49 measured in adulthood. We found no evidence that elevated thyroid hormone levels in 50 eggs affected behaviour, plasma hormone levels, or gene expression. This is the first 51 study examining the potential long-term effects of elevated maternal thyroid 52 hormones within the natural range. Although we found no evidence of long-term 53 effects, other traits may still be affected and remain to be studied.

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55 Keywords: maternal hormones, maternal effects, endocrinology, phenotypic plasticity,

56 bird

# 57 Introduction

58 Maternal effects refer to the effect of the maternal phenotype or environment on 59 the phenotype of the offspring. The effect can be mediated by hormones passed from the 60 mother to the offspring, the maternal hormones. As the levels of maternal hormones are 61 influenced by the maternal environment, they can transmit information about the maternal 62 environment to the offspring. This allows the development of the offspring phenotype to be 63 controlled by environmental factors (Marshall and Uller, 2007; Mousseau and Fox, 1998). 64 Such hormone-mediated maternal effects can be observed in a wide range of traits, 65 including physiology and behaviour (Groothuis et al., 2019, 2005; Ruuskanen and Hsu, 66 2018), and can have long-lasting effects into adulthood with potential fitness consequences 67 (Groothuis et al., 2005; Schoech et al., 2011; von Engelhardt and Groothuis, 2011).

68 Compared to the more extensively studied androgens and glucocorticoids (Podmokła 69 et al., 2018), the effects of maternally transferred thyroid hormones (THs) have been less 70 explored (Darras, 2019; Ruuskanen and Hsu, 2018). THs (triiodothyronine, T3, and thyroxine, 71 T4) play important roles in metabolism and development, where they regulate the 72 expression of numerous genes (Bernal, 2017; Darras, 2019; Moog et al., 2017). TH action is 73 regulated by several components of the thyroid system. TH receptors (THRA and THRB) and 74 coactivators control the transcription of TH-responsive genes by forming a complex with the 75 active form, T3 (Brent, 2012). The intracellular availability of T3 is regulated by several 76 transmembrane TH transporters (e.g. MCT8, OATP1c1) that differentially transport either T4 77 or T3 (Groeneweg et al., 2020). Inside the cell, T3 availability can be further regulated by 78 deiodinase enzymes that either convert T4 to T3 (deiodinase 2, D2) or convert both to 79 inactive forms (deiodinase 3, D3) (Gereben et al., 2008). The expression of these 80 components (TH receptors, transporters and deiodinases) begins very early in embryonic

development, allowing the embryo to regulate TH levels in an organ and cell-specific manner (Darras, 2019; Darras et al., 2011; Flamant and Samarut, 1998; Ruuskanen et al., 2022; Van Herck et al., 2012). However, the development of the thyroid gland occurs much later in development (Forhead and Fowden, 2014; McNichols and McNabb, 1988; Porazzi et al., 2009). Thus, in almost all vertebrates studied so far, proper embryonic development depends on maternally transferred THs (Brown et al., 2014; Darras, 2019; Moog et al., 2017; Patel et al., 2011).

88 In mammals, maternal THs are transferred across the placenta, whereas in egg-laying 89 animals such as birds, maternal THs are deposited in the egg yolk. In birds, high natural 90 variation in egg TH levels has been observed, up to 5-fold in magnitude (Ruuskanen and Hsu, 91 2018). This variation may be due to environmental factors such as food availability or 92 temperature (Hsu et al., 2016; Ruuskanen et al., 2016c), and has been reported at both the 93 inter- and intra-individual levels (Hsu et al., 2016, 2019b; Ruuskanen et al., 2016a, 2016c). 94 As THs are important in development (Darras et al., 2009; Decuypere et al., 2005; McNabb, 95 2007; Wilson and McNabb, 1997), and as increasing yolk TH levels has been found to 96 increase embryonic TH levels (Van Herck et al., 2012), high variation in egg TH levels could 97 therefore be a source of offspring phenotypic variation.

To date, most of what is known about phenotypic differences due to maternal THs comes mainly from biomedical studies (Moog et al., 2017). However, studies using pathological TH concentrations do not provide relevant information on the ecology or evolution of maternal THs. Therefore, the effects of maternal THs within the natural range need to be studied to understand whether variation has effects on traits affecting offspring fitness. In addition, experiments manipulating maternal THs in mammals would inevitably interfere with maternal physiology, making it difficult to attribute results solely to maternal

105 THs. Egg-laying animals, such as birds, have therefore become common models for studying 106 the functions of maternal hormones since maternal THs deposited in eggs can be 107 manipulated directly and independently. So far, elevated TH levels in the egg within the 108 natural range have been found to affect hatching success, offspring growth, plasma T3 level 109 and telomere length (Hsu et al., 2023a, 2019a, 2017; Ruuskanen et al., 2016b; Sarraude et 110 al., 2020; Stier et al., 2020). Nevertheless, some of the reported effects appear to be 111 transient during the early post-hatching period (Hsu et al., 2019a) Whether the effects of 112 maternal THs can be long-lasting and potentially influence offspring fitness remains unclear.

113 To investigate the potential long-term effects of variation in maternal THs, we 114 experimentally elevated egg TH levels within the natural range in Japanese quail (Coturnix 115 japonica) and measured changes in behaviour, plasma TH levels and brain gene expression. 116 While maternal hyperthyroidism has been shown to affect behaviour, for example anxiety 117 (Laureano-Melo et al., 2019; Zhang et al., 2010, 2008), it is not known whether increasing 118 maternal THs within the natural range affects behaviour. Also, as postnatal THs have been 119 shown to influence filial imprinting and learning behaviour (Yamaguchi et al., 2012), it is 120 interesting to study whether maternal THs, acting earlier in development, may also have 121 behavioural effects. Potential behavioural effects were investigated in both chicks and 122 adults, allowing for a broader timescale, by examining boldness, social motivation and 123 fearfulness to predators. Due to the lack of studies with natural levels of elevated maternal 124 THs and inconsistent results from studies on maternal hyperthyroidism and offspring 125 anxiety (Laureano-Melo et al., 2019; Zhang et al., 2010, 2008), the direction of potential 126 behavioural changes is difficult to predict. In addition, plasma TH levels were measured in 127 adults, and changes in these would indicate long lasting effects on TH function. While 128 plasma TH levels have been reported to be affected by elevated egg TH levels in a sex-

129 dependent manner in the rock pigeons (Hsu et al., 2017), some more recent studies in wild 130 passerines found no clear effects of egg THs on nestling plasma THs (Hsu et al., 2023b, 131 2020). In terms of gene expression, previous studies have examined the effect of the natural 132 variation in maternal THs on brain gene expression only in bird embryos or young chicks 133 (Van Herck et al., 2015, 2012; Yamaguchi et al., 2012), while the potential long-term effects 134 of prenatal TH exposure remain unknown. Following our previous experiment manipulating 135 egg TH levels in Japanese quail (Sarraude et al., 2020), here we studied several genes 136 involved in TH function: THRA (thyroid hormone receptor A), NCOA1 (nuclear receptor 137 coactivator 1) and DIO2 (type 2 deiodinase) in the pallium of adults. Expression of these 138 genes has been shown to be influenced by THs in other species (Misiti et al., 1998; Walpita 139 et al., 2007).

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# 141 Materials and methods

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## 143 1.1 Materials

This study was conducted in accordance with the Finnish legislation and approved by the Finnish Animal Ethics Committee (Eläinkoelautakunta; ESAVI / 1018 / 04.10.07 / 2016). Japanese quail (*Coturnix japonica*), a commonly used laboratory model with a reference genome, was used. Twenty-four breeding pairs were housed in cages with food and water available ad libitum. The photoperiod was long (16L:8D) and the temperature stable at 20° C. Eggs were collected from the cages in the morning and stored (15° C, 50% relative humidity) for seven days before treatment.

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# 152 1.2 Treatment of eggs, rearing of chicks and collection of samples

153 We manipulated egg TH levels by injecting hormones directly into the eggs prior to 154 embryonic development at a dose that mimicked increased maternal transfer (see details in 155 Sarraude et al., 2020). Eggs from each breeding pair were randomly assigned to four 156 treatment groups: T3, T4, T3 + T4 and control (0.9% saline). Eggs laid by the same pair were 157 always evenly distributed among all treatment groups. The injected hormone solutions were 158 prepared from crystallised powder (T3: CAS number 6893-02-3, Sigma-Aldrich, St. Louis, 159 MO, USA and T4: CAS number 51-48-9, Sigma-Aldrich, St. Louis, MO, USA). The amount of 160 hormone injected was based on previous measurements in the parental generation (mean  $\pm$ 161 SD: T4, 15.3  $\pm$  4.43 ng / egg; T3, 7.58  $\pm$  2.32 ng / egg) and the aim was to increase hormone 162 contents in the yolk by two standard deviations (as recommended by Podmokla et al., 163 2018). The injection protocol has been described in detail in Sarraude et al. (2020). Briefly, 164 each egg was injected with 50  $\mu$ l of either the respective hormone solution (T3, T4 or T3+T4, 165 in 0.9% saline) or 0.9% saline (control, CO), delivered by using 27G insulin syringes (BD 166 Microlance<sup>TM</sup>). After sealing the hole with sterile plaster (OPSITE Flexigrid, Smith & 167 Nephew), the eggs were incubated artificially (37.8 °C and 55% relative humidity). A total of 168 158 eggs were treated (39 T3, 39 T4, 40 T3+T4 and 40 CO), of which 66 chicks hatched (15 169 T3, 20 T4, 21 T3 + T4 and 10 CO), with sex distributions of chicks surviving to adulthood 170 (females / males): 7/4 for T3, 10/8 for T4, 9/12 for T3 + T4 and 3/4 for CO. Sex was 171 determined at maturity by examining the presence of the cloacal gland. The overall hatching 172 rate (~40%) was consistent with previous studies in domestic quail (Okuliarová et al., 2007). 173 The rearing conditions of the chicks have been described in detail in a previous 174 publication (Sarraude et al., 2020). In short, chicks were reared in groups (4-16 individuals,

175 depending on age) with treatments and sexes mixed, until sexual maturity (ca. 6–8 weeks

after hatching in Japanese quails), after which the birds were separated by sex to smaller groups (4-5 individuals). Photoperiod was 16L:8D for the first 7 months and 8L:16 thereafter. At approximately 4.5 months of age, we collected blood samples from the brachial vein, separated plasma, and stored it at -80° C. Birds were euthanised by decapitation at 10 months of age. After decapitation, 40 mg of tissue was collected from the pallium in PBS buffer, snap frozen in liquid nitrogen and stored at -80° C.

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183 **1.3** Behavioural experiments

184 To assess the potential effects of maternal THs on behaviour, we conducted a series of 185 behavioural tests (open arena, emergence and tonic immobility tests) at different life 186 stages. We followed previously published protocols that have been used extensively with 187 many different model organisms. The tests were conducted in separate rooms where the 188 birds had no visual or auditory contact with other individuals. The exception was the open 189 arena test, where the subject was able to hear the vocalisations of another bird being tested 190 at the same time in another room. All experiments were recorded, and the videos were 191 analysed blindly of the treatment. All videos of the same test at both ages (chick/adult) 192 were analysed by one person, except for the open arena test, where one person analysed 193 chick videos and another person analysed adult videos.

The open arena test measured social motivation and boldness and it followed a setup developed by Forkman et al. (2007). The test was performed on both sexes at three days and approximately 4.5 months of age. In the test, the bird was placed in a circular illuminated arena and the movement duration and the number of escape attempts (attempt to jump/fly over the wall) were measured from videos for 10 minutes (from the time the researcher left the room). Only 10 of the 55 adults attempted to escape, so adult escape

attempts were not analysed. Subjects who moved less and made fewer escape attempts
 were interpreted as less bold and less socially motivated.

202 The emergence test measured social motivation and boldness. The test was 203 performed on both sexes, twice on chicks (6-9d old) and once on adults (4.5 months old). 204 The test followed a setup developed by Forkman et al. (2007), in which a bird was placed in 205 a dark chamber connected to another, bright chamber. One minute after the subject was 206 placed in the dark chamber, a door connecting the chambers was removed. After removing the door, the researcher left the observation room and the time it took for the subject to 207 208 completely exit the dark chamber was measured. Subjects who left the dark chamber more 209 quickly were interpreted as bolder and more socially motivated. The test lasted five 210 minutes. Whether the subject remained in the dark chamber (0) or left it (1) during the test 211 was also recorded.

212 The tonic immobility test (Mills and Faure, 1991) measured fearfulness to predators. 213 The test was performed on both sexes at 12 days and approximately 4.5 months of age. To 214 induce immobility, the bird was placed on its back in the centre of a V-shaped test apparatus 215 and restrained by hand for 15 seconds, after which the hand was removed, and the time 216 taken for the bird to resume a standing posture was measured. If the subject resumed the 217 standing posture in less than 10 seconds, this was recorded as a failed induction and the 218 induction procedure was repeated. Four adult subjects who failed five times were excluded. 219 If the subject remained stationary for more than 10 seconds, then the time to recovery was 220 measured. The longer a subject took to recover, the more fearful it was interpreted. The 221 test lasted 10 minutes. Five subjects (1 chick and 4 adults) did not recover for 10 minutes 222 and were excluded from the analysis (approximately 4% of all tests).

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# 224 1.4 Plasma thyroid hormone levels

To test whether elevated maternal THs affect plasma TH levels later in life (4.5 months of age), we extracted THs from plasma samples and measured them using a previously validated LC-MS / MS protocol (Ruuskanen et al., 2018). The sample size was 48 individuals (T3: 10, T4: 16, T3T4: 15, and 7 in the control). Hormone concentrations are expressed in pmol/ml plasma.

230

# 231 1.5 Gene expression

232 To test for potential long-term changes in the expression of TH related genes (DIO2, 233 THRA, and NCOA1), we measured gene expression from pallium samples (10 months old). 234 RNA was extracted from the homogenised pallium samples using the NucleoSpin RNA Plus 235 kit (MACHEREY-NAGEL). The sample size was 31 (7 in control and 8 per treatment group). 236 Extraction was done from 50  $\mu$ l of the homogenised brain sample according to the 237 manufacturer's instructions, except that the gDNA removal column was centrifuged longer 238 (1 min, 11000 g) and the RNA was eluted once with 40  $\mu$ l of RNase-free water. RNA 239 concentration and purity were quantified by optical density, and all samples not meeting 240 our quality criteria (i.e. RNA concentration > 30 ng/ $\mu$ l and 260/280 > 1.80) were excluded 241 from further analysis. RNA integrity was checked using an E-Gel 2% electrophoresis system 242 (Invitrogen), and the ribosomal RNA 18S vs. 28S band intensity and was deemed 243 satisfactory. Samples were stored at -80° C for 2 weeks prior to cDNA synthesis. cDNA 244 synthesis was performed using the SensiFAST ™ cDNA Synthesis Kit (Bioline) according to 245 the manufacturer's instructions. The reaction was performed using 500 ng of RNA as the

starting material. A control (noRT) was prepared without reverse transcriptase enzyme. The

cDNA was diluted with water to a concentration of 1.2 ng/ $\mu$ l and stored at -20° C.

248 GAPDH and PGK were used as control genes, using primers previously designed and 249 validated in Japanese quail (Vitorino Carvalho et al., 2019). Stability of expression between 250 experimental groups was further verified using geNorm software (geNorm M < 0.5, geNorm 251 V < 0.15), and thus the geometric mean of these two genes was used as our reference gene. 252 Primers for the target genes were designed using the Primer-BLAST tool (NCBI). The primer 253 pairs (Table 1) were designed to have at least one intron between them. The suitability of 254 the primers in gPCR was tested prior to the actual analysis using different primer 255 concentrations and different amounts of cDNA. The tests confirmed the sufficient 256 amplification efficiency of the primers (between 90 and 110% based on the standard curve) 257 and the formation of only one product of the expected size based on both the presence of a 258 single narrow peak in the melting curve, and the presence of a single band of the expected 259 size on the agarose gel.

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270 Table 1: qPCR primer set information and performance for gene expression assays. Cq refers to

271 qPCR quantification cycle, efficiency was evaluated using the LinReg method based on individual well

efficiency. Technical precision estimates are given for Cq as coefficient of variation CV (%) and for

- 273 expression (ratio of target to reference genes) as technical repeatability (*R*).
- 274

Gene	Primers	Cq ± SD	Efficiency	CV Cq	Repeatability expression
GAPDH	F: TCTCTGTTGTTGACCTGACCTG R: ATGGCTGTCACCATTGAAGTC	15.0±0.50	98.0%	0.59%	
PGK1	F: CAAGCTCACCCTGGACAAGT R: GGACGGCTGCCTTGATTCTT	21.1±0.86	85.5%	0.48%	
DIO2	F: CTGGAACAGCTTCCTCCTGG R: ATCCAGAAGACGGCACTTGG	23.6±0.83	92.9%	0.40%	0.91
NCOA1	F: CGACTCACGGAAGAGGAAGG R: GTGTCGATGTCCCCGATGTT	20.4±2.82	91.9%	0.38%	0.84
THRA	F: TGGTGAAGAGCAGCATGTCA R: AGCACCCATCGTACTTGCAG	21.4±0.82	96.0%	0.46%	0.82

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276 qPCR assays were performed on a Mic qPCR (Bio Molecular Systems) in a final volume of 12 277 μL containing 6 ng of cDNA, 800 nM of forward and reverse primers and SensiFAST SYBR Lo-278 ROX (Bioline). We ran gene-specific qPCR plates and each sample was analysed in duplicate. 279 Two plates were run for each gene and experimental groups and sex were always balanced 280 within each plate. We used a cDNA reference sample (*i.e.* ratio = 1), which was a pool of 10 281 different individuals on each plate. We used a two-step cycling with the following 282 conditions: 2 min at 95°C; then 45 cycles of 5s at 95°C followed by 20s at 60°C (fluorescence reading) for all reactions. The expression of each gene was calculated as  $(1+Ef_{Target})^{\Delta Cq(Target)}/Cq(Target)$ 283 geometric mean  $[(1+Ef_{GAPDH})^{\Delta Cq^{GAPDH}}; (1+Ef_{PGK1})^{\Delta Cq^{PGK1}}]$ , where Ef is the amplification's 284 285 efficiency and  $\Delta Cq$  is the difference between the Cq-values of the reference sample and the 286 sample of interest. Amplification in the no-RT controls never occurred until at least 8 cycles 287 after the lower Cq sample, and thus contamination by genomic DNA could not interfere with

our results. Technical precision and repeatability were generally good (Table 1), but a few samples with a CV% of duplicates (based on the final ratio) greater than 15% were excluded from the analysis (final sample size:  $N_{DIO2} = 27$ ,  $N_{NCOA1} = 28$ ,  $N_{THRa} = 26$ )

291

## 292 **1.6** Statistical analysis

293 All statistical analyses were performed with R software (version 3.6.2), using the packages 294 Ime4 version 1.1-23 (Bates et al., 2015) and emmeans version 1.6.0 (Lenth, 2021). Normality 295 and homoscedasticity of the residual distributions were checked visually. For the 296 behavioural analysis, recovery time (tonic immobility) and exit time (emergence test), were 297 transformed (natural logarithm) to ensure that the residuals were normally distributed. 298 Linear mixed models (LMM) were used to analyse time to exit in the emergence test, 299 movement duration and number of escape attempts in the open arena test, recovery time 300 in the tonic immobility test, plasma TH concentrations and qPCR results. Generalised linear 301 mixed model (GLMM) with a binomial error distribution was used to analyse the proportion 302 of individuals leaving the dark chamber in the emergence test. Fixed effects were treatment, 303 sex, and test number/age (in repeated tests). Interactions between sex and treatment, and 304 between treatment and age (in repeated tests), were also analysed. Random effects were 305 individual ID (in repeated tests) and parental ID (to account for relatedness between siblings 306 in all experiments). F- and p-values for LMMs were obtained using the Kenward-Roger 307 approximation on the denominator degree of freedom, implemented in the package 308 ImerTest (Kuznetsova et al., 2017). For binomial regression models, main effects and 309 interactions were tested by comparing the model with and without the factor. Non-

310 significant (p > 0.05) interactions were excluded from the final models. Post-hoc tests were

311 performed using Tukey tests where appropriate.

312

## 313 Results

# 314 Effects of prenatal thyroid hormone supplementation on behaviour

In general, no statistically significant differences in chick or adult behavioural traits 315 316 were observed between the egg TH manipulation groups (T3, T4, T3 + T4, control). Elevated 317 THs had no significant effect on social motivation or boldness (Table 2, Fig. 1-2), nor on 318 fearfulness to predators (Table 2, Fig. 3). A statistically significant interaction between 319 treatment and age was found in the emergence test (F = 3.58, ddf = 67.2, p = 0.018, Fig. 2): 320 In all treatments except the control group (Tukey t=0.59, p=0.56), chick exit times were significantly longer than those of adults (all t > 3.21, p < 0.002) while no differences between 321 322 treatments were found within each age-category (all t < 2.48, all p> 0.07). There was also a 323 statistically significant interaction between treatment and sex in the tonic immobility test (F 324 = 2.92, ddf = 37.5, p = 0.047, Fig. 3): In males, the recovery time was longer in the T4-group 325 (marginal mean  $\pm$  SE: 177.1  $\pm$  47.8 s) than in the T3-group (marginal mean  $\pm$  SE: 56.1  $\pm$  20.8 326 s, Tukey t = 2.71, p = 0.047), whereas no differences between treatments were observed in 327 females (all t < 2.29, all p >0.12). In the T4-group, the recovery time of males (marginal 328 mean  $\pm$  SE: 177.1  $\pm$  47.8 s) was longer than that of females (marginal mean  $\pm$  SE: 85.2  $\pm$  19.9 329 s) (t=2.16, p= 0.037), while no sex differences were observed within the other treatments 330 (all t<1.80, all p>0.079).

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**Table 2.** Marginal means and standard errors of behavioural tests. Results for chicks (age: 3d, 6-9d, 12d) and adults (age: 4.5 months) are shown separately for each variable. F- and p-values for treatment from linear models (ANOVA, Kenward-Roger approximation). Test statistics ( $\chi^2$ ) and p-values for binomial regressions (model with treatment compared to model without treatment) were calculated with ANOVA.

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Test: measurement Age Control		Control	T3	T3T4	T4	$F_{ddf}/\chi^2$ (df)	P
Tonic immobility:	12 d	92.0±25.5	84.6±21.1	68.4±13.1	123±24.7	2.52 <sub>36.2</sub>	0.073
Recovery time (s)	4.5 m	90.6±25.3	83.3±21.4	67.3±13.6	121±25.5		
Open arena:	3 d	77.2±31.1	100±27.1	73.3±19.1	63.3±20.8	0.44 <sub>46.6</sub>	0.72
Movement duration	4.5 m	181±31.6	204±28.1	177±19.3	167±21.3		
(s)							
Open arena:	3 d	5.85±3.87	4.14±3.53	8.25±2.14	7.21±2.43	0.35 <sub>41.7</sub>	0.79
Escape attempts							
(number)							
Emergence:	6–9 d	34.8±14.7	42.7±16.4	74.7±20.9	121±36.1	1.92 <sub>39.6</sub>	0.14
Exit time (s)	4.5 m	25.6±12.5	6.59±2.75	23.0±7.64	12.6±4.30		
Emergence:	6–9 d	0.80±0.11	0.70±0.12	0.62±0.10	0.58±0.10	2.71 (3)	0.44
Likelihood of	4.5 m	0.91±0.06	0.86±0.08	0.81±0.08	0.78±0.09		
emergence (0-1)							



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Figure 1. Movement duration for chicks (3 d) and adults (4.5 months) in the open arena test
for different prenatal thyroid hormone treatment groups. The lines represent the medians of
the treatments, and the bars represent the interquartile range of the original observations.
Sample size (chicks/adults): CO 7/7, T3 11/11, T3T4 20/21 and T4 16/17.



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Figure 2. Time to exit the dark chamber for chicks (6-9 d) and adults (4.5 months) in the emergence test for different prenatal thyroid hormone treatment groups. The test was performed twice for chicks and once for adults. The lines represent the medians of the treatments, and the bars represent the interquartile range of the original observations. Sample size (chicks/adults): CO 11/6, T3 15/9, T3T4 24/13 and T4 22/13.



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Figure 3. Time to resume standing position for females and males, chicks (12 d) and adults (4.5 months) combined, in the tonic immobility test for different prenatal thyroid hormone treatment groups. The lines represent the medians of the treatments, and the bars represent the interquartile range of the original observations. Sample size (female/male: CO 6/8, T3 13/7, T3T4 14/23 and T4 18/14.

358

359 Effects of prenatal thyroid hormone supplementation on plasma thyroid

Increasing TH levels in eggs did not significantly affect plasma TH levels in 4.5month-old birds (T3: F = 0.13, ddf = 37.7, p = 0.94, T4: F = 2.1, ddf = 41.5, p = 0.12, Fig.
4). Treatment also had no sex-dependent effect on plasma T3 or T4 (treatment-sex
interactions, p > 0.36).

<sup>360</sup> hormone levels



<sup>365</sup> 

Figure 4. Plasma T3 and T4 concentrations in 4.5-month-old birds between different prenatal thyroid hormone treatment groups. The lines represent the medians of the treatments, and the bars represent the interquartile range of the original observations. Sample size (T3/T4 concentrations): CO 6/7, T3 10/10, T3T4 14/15 and T4 14/16.

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371 Effects of prenatal thyroid hormone supplementation on pallial gene 372 expression

Increasing TH levels in eggs did not affect pallial gene expression of TH-related genes in 10-month-old birds (*DIO2*: F = 2.49, ddf = 17.3, p = 0.09, *NCOA1*: F = 0.092, ddf = 21.1, p = 0.96, *THRA*: F = 0.47, ddf = 18.9, p = 0.71, Fig. 5). Treatment also did not have sex-dependent effect on gene expression (treatment-sex interactions, p > 0.26). Females had higher expression of *NCOA1* (F=7.68, p=0.012) and *THRA* (F=5.17, p=0.034) than males.



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Figure 5. Relative expressions of *DIO2*, *THRA* and *NCOA1* in the pallium of 10-month-old birds
between different prenatal thyroid hormone treatments. The lines represent the medians of
the treatments, and the bars represent the interquartile range of the original observations.
Sample sizes (CO/T3/T3T4/T4): DIO2 5/7/8/7, THRA 6/4/8/8 and NCOA1 6/7/8/7.

384

# 385 Discussion

In this study, we tested for long-term changes in behaviour, plasma TH levels and pallial gene expression following increased exposure to maternal THs. In terms of behaviour, we observed no significant changes in boldness, social motivation or fearfulness to predators in chicks (3-12 d) or adults (4.5 months) when the egg TH level was increased. To our knowledge, there are no similar previous studies, and it is

391 therefore difficult to assess whether there may have been changes in other 392 behavioural traits. The detection of changes was also partly limited by the small 393 sample size, which meant that we may not have had enough statistical power to detect 394 subtle changes in behaviour. Nevertheless, we found some age-dependent and sex-395 dependent effects of TH treatment on behaviour. The age-dependent effect in the 396 emergence test suggests that elevated maternal TH leads to a decrease in boldness 397 and social motivation with age (from chick to adult), whereas such a decrease is not 398 observed in controls. However, the lack of differences between control and treatment 399 within a given age suggests that such an effect (if not associated with type I statistical 400 error) is rather mild. The sex-dependent effect in the tonic immobility test also seems 401 relatively unconvincing because no sex differences were observed within treatments, 402 nor did T3T4 females or T4 males differ from the same-sex controls. Given the 403 important role of THs in brain development, we expected that elevated maternal THs 404 would have led to at least some behavioural changes. Yet, our data suggest that an 405 experimental increase in maternal THs within the natural range has no clear effect on 406 the behavioural traits measured. However, in contrast to maternal THs, post-hatch TH 407 treatment has been shown to affect imprinting and learning (Yamaguchi et al., 2012). 408 The discrepancy between our results and Yamaguchi et al. (2012) could suggest that 409 the sensitive period during which THs can influence behaviour might occur at a later 410 timing of development, but this requires further investigation.

We found that elevated maternal THs did not significantly affect plasma TH levels in adulthood (4.5 months of age). This was the first study to test for long-term changes, whereas previously changes in response to elevated prenatal THs have been observed at younger ages: Experimental elevation of egg THs was found to increase male and decrease female plasma TH levels in rock pigeons at 14 days of age (Hsu et

al., 2017), but such effects were not observed in two other wild passerine species (Hsu
et al., 2023b, 2020). In contrast to these studies in altricial species, not only was this
study conducted on a precocial species, but our measurements were also made at a
different life stage (adulthood). Overall, this suggests that an increase in maternal THs
may only lead to transient changes in offspring plasma TH levels and that the effects
may not be universal across species.

We also found no significant effect of elevated maternal THs on the expression 422 423 of THRA, DIO2 or NCOA1 in the pallium of adult (10 months old) individuals. However, 424 changes in gene expression have been observed in other TH-related genes, but again, 425 at much shorter time spans: In chickens, injection of THs into the yolk decreased the 426 expression of the TH-transporter OATP1C1 and increased the expression of the TH-427 transporter MCT8 in 4-day-old embryos (Van Herck et al., 2012). The expressions of 428 our target genes, THRA, DIO2 and NCOA1, have also been shown to be at least 429 transiently responsive to THs in other species: THRA and DIO2 in zebrafish embryos 430 (Walpita et al., 2007), and NCOA1 in adult rats (Misiti et al., 1998). It is therefore 431 possible that the gene regulation by maternal THs is restricted to early development 432 when the embryo is not yet able to produce THs. As the embryonic TH production 433 begins, the influence of maternal THs might be expected to decrease, but studies on 434 the relative contributions of maternal vs embryonic THs are lacking. Following 435 embryonic TH production, the initially higher manipulated TH concentration may begin 436 to level off and the differences in gene expression may disappear. Differences could 437 still exist in adulthood, if higher maternal THs could lead to epigenetic changes, but 438 this has not been studied with prenatal THs in their natural range. There is also no 439 information on possible long-term changes in other brain regions or genes. Therefore,

440 it remains unknown whether higher maternal THs can lead to long-term changes in

441 gene expression.

442 In general, the lack of changes in the measured traits following the elevation of 443 egg THs could be explained if the higher egg TH levels did not lead to a higher TH 444 concentration in the embryo. This could be due to the precise regulation of the 445 transport of THs from the egg to the embryo. However, we have previously shown in 446 the same experimental birds, that an increase in prenatal THs influences hatching 447 (Sarraude et al., 2020), suggesting that THs do influence prenatal success 448 development. Effects of prenatal THs on offspring growth and physiology have also 449 been reported in other species (Hsu et al., 2023b, 2019a, 2017; Ruuskanen et al., 450 2016b; Stier et al., 2020). In terms of brain gene expression, it could be that tightly 451 regulated TH transport to the developing brain prevents elevated maternal THs from 452 affecting gene expression, but this would contradict the previously observed changes 453 in brain gene expression following TH injection in chickens (Van Herck et al., 2012). It is 454 also possible that embryos could buffer the effect of maternal THs by altering the 455 expression of deiodinases (in the brain or other tissues). It has recently been shown in 456 several species, including Japanese quail, that embryos express DIO2 and DIO3 at very 457 early stages (Ruuskanen et al., 2022; Too et al., 2017). Metabolisation of THs to 458 inactive forms in key tissues could reduce phenotypic variation caused by differences 459 in egg TH levels. However, it is unlikely that buffering alone could explain the complete 460 lack of changes in this study, given the effects on development previously observed in 461 the same experimental birds (Sarraude et al., 2020). As it is, the lack of changes in TH-462 related gene expression in this study is likely to be explained by the stage of ontogeny, 463 prenatal THs not leading to long-lasting effects. However, the lack of prenatal

464 measurements means that this cannot be fully confirmed, and it remains possible that

465 elevated maternal THs have no effect on the measured traits in Japanese quails.

466 In conclusion, we found no evidence that elevated maternal THs cause long-467 term changes in behaviour, plasma TH levels or pallial gene expression. This is 468 consistent with our previous findings in the same experimental birds, where elevation 469 of prenatal THs (T3, T4, or in combination) did not lead to long-term effects on 470 oxidative stress, moult or reproduction (Sarraude et al., 2020). While behavioural 471 changes have not been previously studied, plasma TH concentrations and gene 472 expression have been shown to respond to elevated maternal THs in the short-term 473 (Hsu et al., 2017; Van Herck et al., 2012). However, based on this study, these changes 474 would appear to be limited to early development or shortly after hatching with the 475 offspring's own TH production later replacing the effects of elevated maternal THs. 476 Maternal THs could theoretically influence the offspring's own TH production at a later 477 age by altering the function of the HPT-axis, but no evidence has yet been found to 478 support this hypothesis. The long-term effects of variation in maternal THs remain 479 unclear, but this study has provided new information on some previously untested 480 traits. Understanding long-term effects in addition to short-term effects will help to 481 better understand the evolutionary and ecological importance of maternal THs.

482

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486

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- 493 Data availability statement
- 494 Data will be available on Figshare (DOI: 10.6084/m9.figshare.22266211) at publication.
- 495
- 496 Author contributions
- 497 S.R., T.S., B-Y.H, A.S and K.A. designed the study. S.R. obtained funding. T.S., B-Y.H and
- 498 S.R conducted the experiment. K.A. and T.S. collected the behavioural data from
- 499 videos. K.A., A.S and S.R. collected the physiological data and conducted laboratory

500 work. K.A. analysed all data with support from all co-authors, and wrote the first draft

501 of the manuscript. A.S, B-Y.H, T.S. and S.R commented on the manuscript.

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