Transgenerational endocrine disruption? Experimental copper exposure, but not heat stress, leads to elevated egg thyroid hormone levels

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Abstract

Climate change and pollution are some of the greatest anthropogenic threats to wild animals. Transgenerational plasticity – when parental exposure to environmental stress leads to changes in offspring phenotype – has been recently highlighted as a potential mechanism to respond to various environmental and anthropogenic changes across taxa. Transgenerational effects may be mediated via multiple mechanisms, such as transfer of maternal hormones to eggs/fetus. However, sources of variation in such hormone transfer are poorly understood in fish. We studied genetic and environmental variation (in response to temperature and endocrine disrupting copper) in maternal thyroid hormone (TH) transfer to eggs in a common fish model species, the three-spined stickleback (*Gasterosteus aculeatus*) using multiple approaches: (i) We compared egg TH levels among six populations across a wide geographical range in the Baltic Sea, including two populations at high water temperature areas (discharge water areas of nuclear power plants) and we experimentally exposed fish to (ii) environmentally relevant heat stress and (iii) copper for 7 days. We found that populations did not differ in egg TH levels, and short-term heat stress did not influence egg TH levels. However, copper exposure increased both T4 and T3 levels in eggs. These novel results suggest that copper exposure can potentially induce transgenerational endocrine disruption, but the functional consequences of altered THs on offspring phenotype should be further studied.

Capsule: We show that experimental copper exposure, but not heat stress (experimental or among-population variation), leads to elevated egg thyroid hormone levels in sticklebacks, suggesting potential for novel transgenerational endocrine disruption via copper.

Keywords: thyroid hormones, T3, T3, plasticity, maternal effect, endocrine disruption, transgenerational, metal pollution, heat stress

Introduction

Climate change and pollution are some of the greatest anthropogenic threats to wild populations. Organisms may respond to changes in their environment either via changing distributions (range shifts), showing plastic responses, or potentially genetic adaptation (i.e. microevolution)(Habary et al., 2017; Parmesan, 2006; Stillman and Armstrong, 2015). One recently highlighted form of plasticity is transgenerational plasticity, i.e. when variation in parental environment leads to changes in offspring phenotype (e.g. Donelson et al., 2012; Donelson et al., 2018; Metzger and Schulte, 2017; Meylan et al., 2012; Salinas and Munch, 2012; Shama et al., 2014; Shama and Wegner, 2014). For example, Shama et al. (2014) showed in their seminal paper rapid transgenerational acclimation to elevated temperature, although the molecular mechanisms are not yet understood. Transgenerational effects may be mediated via multiple mechanisms, such as epigenetic markers or transfer of maternal hormones or RNAs to eggs/fetus (e.g. Adrian-Kalchhauser et al., 2018; Best et al., 2018; Kim et al., 2019; Metzger and Schulte, 2017; Meylan et al., 2012; Ruuskanen and Hsu, 2018).

Hormones transferred from the mother to embryos and eggs are known to profoundly influence offspring development, physiology, morphology, behavior and even survival across taxa (mammals, Dantzer et al., 2013; fish, McCormick, 1999; birds, Ruuskanen, 2015; Ruuskanen and Hsu, 2018; reptiles, Uller et al., 2007). One class of these hormones could be thyroid hormones (THs, prohormone thyroxine T4, and biologically active tri-iodothyronine, T3) and recent studies suggest that maternal THs transferred to eggs and embryos are important for offspring development across vertebrates (fish, Brown et al., 2014; birds, Hsu et al., 2017; Hsu et al., 2019; mammals, Patel et al., 2011; Ruuskanen et al., 2016a). THs are a key class of hormones that control and regulate vital biological processes such as thermogenesis, reproduction but also growth and metamorphosis (Norris and Carr, 2013). THs are critical regulators of thermal acclimation in fish, increasing in higher temperatures,

(e.g. Little et al., 2013) and plasma THs have been found to fluctuate with varying water

temperature (e.g. Arjona et al., 2010; Cyr et al., 1998; Eales, 1985).

Surprisingly, studies characterizing the environmental or genetic sources of among-

female (or within-female) variation in egg TH levels are rare (Ruuskanen and Hsu 2018), yet,

such variation could contribute to variation in offspring phenotype and fitness and help

understand the scope for transgenerational plasticity. In birds, egg THs are associated with

food availability (Hsu et al., 2016) and temperature (Ruuskanen et al., 2016c), while T3 (but

not T4) also shows heritable variation (Ruuskanen et al., 2016b, Hsu et al. 2019). In fishes,

there is indication for variation in egg THs among stocks (rainbow trout, *Oncorhynchus*

mykiss Leatherland et al., 1989), which could reflect either genetic or environmental

variation. Interestingly, McComb et al. (2005) reported that the egg T3 and T4 concentration

in bonnethead sharks (*Sphyrna tiburo)* from Tampa Bay was consistently higher than from

eggs from Florida Bay. The authors suggested that this may be due to higher temperatures in

Tampa Bay, and speculated that egg THs might, thus, explain the faster growth rates and

metabolic rates at this site. However, no systematic investigation into the sources of variation

in egg THs have been conducted in fish (or other vertebrates).

Moreover, transfer of maternal hormones to offspring may also be affected by

disruption in female hormonal status via endocrine disrupting chemicals (EDC), such as

metals. This phenomenon is called transgenerational endocrine disruption (Ruuskanen et al.,

2019). In one of the rare examples, an experimental exposure to pollutants (lead and

polybrominated diphenyl ethers) resulted in decreases plasma and in egg THs in zebrafish

(*Danio rerio*)(Chen et al., 2017). Alteration of the early-life hormonal environment via

maternal exposure to EDCs could have detrimental consequences on offspring development

and phenotype.

We studied environmental (including endocrine disruption) and potential genetic

variation in maternal TH transfer using both correlational and experimental approach in a

common fish model, the three-spined stickleback (*Gasterosteus aculeatus*), to understand the

scope for transgenerational plasticity via hormone transfer. This species was selected as it is

an abundant and wide-spread species across the Northern Hemisphere, and an important

species in biomonitoring and ecotoxicology as well as in behavioural and ecological studies

(e.g. Scholz and Mayer, 2008). First, we sampled fish in six populations in the Baltic Sea (Fig

1), of which two were from discharge water areas of nuclear power plants, and four reference

populations. Areas used for nuclear power discharge waters function as a natural experiment

for long-term temperature increases, mimicking the effects of global warming, as discharged

water is about 10-12 °C warmer than the intake water, and the discharge has continued for

decades (Keskitalo and Ilus, 1987). This higher water temperature at the nuclear power plants

could be a potential selecting agent on thermal physiology, including TH levels. We

measured intraovarian THs of the fish from these six populations after an acclimation period

in the laboratory. If there has been selection for altered TH metabolism and TH transfer in

wild populations in these warmer water areas, we expect to see altered egg TH levels in the

populations at the vicinity of the nuclear power plants compared to reference areas. Given

that stickleback populations across the Baltic Sea are genetically different from each other,

and the populations differ in response to thermal habitats and salinity they encounter (Guo et

al., 2015), we may also expect overall differences in THs among the six populations.

Secondly, we experimentally exposed the fish from each population to (i) mild

temperature treatment (10°C increase), mimicking a heat wave or (ii) to environmentally

relevant dose of copper (Cu), to study transgenerational TH disruption, and again measured

intraovarian THs. The contamination load in the Baltic is quite high with metals such as

copper, zinc, lead, cadmium and mercury reported in abiotic materials like sediments and

seawater, but also in biota like liver tissues of herring and cod (Perttila et al., 1982). Copper

exposure has been found function as an EDC and both increase and decrease plasma THs in

fish, depending on species and timing of exposure (Eyckmans et al., 2010; Hoseini et al.,

2016; Oliveira et al., 2008), while transgenerational endocrine disruption by copper has not

been studied to date. To this end, we predict that intraovarian T3 and T4 should be (i) higher in

the heat stress treatment compared to controls, due to increasing metabolic rates; and (ii)

altered in the copper treatment compared to controls due to EDC effects of copper.

Materials and methods

Study areas, catching and maintenance

The experiments were conducted with wild, adult female three-spined stickleback caught

from a wide geographical range, from six different locations from the Baltic Sea (see Fig 1).

Three of the locations were in Gulf of Finland and three in Gulf of Bothnia. In both Gulfs one

of the locations was in the cooling water discharge area of a nuclear power plant, where

discharged water is about 10-12 °C warmer than the surface water and the discharge has

continued for decades(Keskitalo and Ilus, 1987). In Gulf of Finland the areas were Loviisa

(LOV, Nuclear Power Plant area, N6021.928; E2622.228), Kotka (KOT, reference site,

N6026.103; E2652.181) and Porvoo (POO, reference site, N6015.090; E2546.134). In Gulf

of Bothnia the areas were Olkiluoto (OLK, Nuclear Power Plant area, N61.2360278;

E21.4347222), Pyhäranta (PYH, reference site, N6057.149; E2125.986) and Pori (POR,

reference site, N6130.140; E2135.675). Temperatures, salinity and pH of the locations during

catching in May 2018 are presented in Supplementary Table 1. Temperature loggers (three

per area, HOBO Water Temp Pro v2 Logger, U22, Onset Computer Corporation, Bourne,

MA, USA) were situated directly at the catching locations at the sea bottom (depth 1.5-2 m)

and water temperature was recorded 16/5/2018 – 31/8/2018 from all the locations four times

per day. The average daily temperatures ± SD are presented in Supplementary Table 1.

The fish were caught with beach seine net and transferred to University of Turku for

rearing. No mortalities were observed during the transfer and the acclimation period in

laboratory facilities. The fish (N=100 per population, mixed sexes, some of the fish were

used also in other experiments) were let to acclimate into 180 L tanks at +16°C for two weeks

(each population in its own tank). Water salinity was 4 ppt (filtered water with 76% NaCl;

20% $MgSO_4$; 3.5% $CaCl_2$; 0.5% $KHCO_3$), pH=8 and oxygen saturation over 80%.

Photoperiod was set to 17L:7D. All fish procedures were performed according with Finnish

Animal Care permission (ESAVI/2867/2018). Fish were fed with frozen bloodworms

(Delang & Ekman AB/ Akvarieteknik, Sweden) five times per week. One third of the water

was changed once a week. Upon arrival the fish were treated against nematodes using

Nematol (Sera GmbH, Heinberg, Germany) according to the instructions of the manufacturer.

In order to reduce any tank effects the fish were tagged intraperitoneally with 1.35×7 mm

RFID subcutaneous microchips (Loligo® Systems, Viborg, Denmark) under anesthesia (100

ppm MS-222 in 4 ppt brackish water buffered with 6 ppm HCO_3) after two weeks'

acclimation period. The populations were mixed into nine tanks (with density of 2L/fish) and

were let to recover for three weeks before further testing.

Experimental treatments

After the recovery period, the fish were exposed to three different environmental conditions

in their rearing tanks: control (CTRL7D), sublethal level of copper (Cu 7D) or heat stress (HS

7D) for one week. Equal numbers of fish from each sampling location were distributed to all

three treatments. Fish were sampled before and after the exposures (see below). The sub-

lethal copper exposure (CU 7D) was done for total number of 127 fish (three replicate tanks,

20 female fish from them were used for the current study). Fish in three replicate control

tanks (CTRL 7D) were treated similarly but no copper was added to tanks (N=120 in total, 25

female fish used for the current study). For exposure, Cu^{2+} was added manually as copper (II)

sulphate pentahydrate solution (nominal: 100µg/L of $CuSO_4 \cdot 5H2O$, Merck, Darmstadt,

Germany) to the experimental tanks. This concentration of copper represents environmentally

relevant concentrations encountered in polluted waters (Sanchez et al., 2005). Water samples

were taken both from exposure and control tanks (i) 2 hrs and (ii) one week after the release

of copper in order to measure the copper concentration during one week of exposure (no

water changes were done during the exposure). Fifty millilitres of water was sampled in

polypropylene Falcon tubes from tanks. In order to keep water samples fresh prior to

analyses, 1 ml of concentrated HNO_3/100ml was added to the samples and samples were kept

at +4°C before analyses at SYNLAB Analytics & Services Finland Oy (Karkkila, Finland).

The copper levels were measured by using inductively coupled plasma mass spectrometry

(ICP-MS) (ThermoFisher Scientific, MA, USA). 5 fish died during the one week of exposure.

A third group of fish (N=125 in total, 20 female fish of them used for the current

study) was exposed to heat stress (HS 7D) (three replicate tanks) for a week during the same

time as the other fish were exposed to copper/control treatment. For simulating an

environmental heat wave the water temperature in the experimental tanks (16°C) was

gradually increased by 1°C every 30 minutes until reaching 26°C and kept at this temperature

for a week. The warming experimental set-up was done with chiller-heater (Julabo, Model:

F32; AC: 230V 50/60 Hz 12A, Julabo GmbH, Seelback, Germany) connected to stainless

steel coils. Six fish died during the exposure to heat stress.

Sampling procedure

Before exposing fish to heat stress or copper, 17 female fish were sampled from a control

tank (hereafter CRTL). Fish were sacrificed with cranial percussion. Tag code, weight (g),

length (cm) were recorded from each fish. For the current study the ovaries were collected

and flash frozen in liquid nitrogen. Samples were stored at -80°C for further molecular

analyses. Similar sapling of the fish and ovaries was done also seven days after exposure for

control (CTRL7D), copper exposure (Cu7D) and heat stress treatment (HS7D) for a total

number of 65 female fish.

Thyroid hormone analyses

The whole content of thawed ovaries (i.e. egg follicles and associated ovarian fluids) was

gently squeezed out of the ovaries directly into a microcentrifuge tube and the sample was

weighed (~0.001g). Samples were analysed for T3 and T4 at the University of Turku.

LCMS/MS was conducted at the facilities of Turku Center for Biotechnology. T4 and T3

were extracted from yolk following previously published methods (de Escobar et al. 1985,

Ruuskanen et al. 2018). In short, samples were homogenized in methanol in tissue lyser

(Quiagen, Retsch GmbH, Haan, Germany). As an internal recovery tracer, a known amount

of $^{13}C_{12}$-T4 (Larodan, Sweden) was added to each sample. This allowed us to control for the

variation in recovery (i.e. extraction efficiency) for each sample. Next, 400 µl of chloroform

was added to sample. After centrifugation (15 min, 1900 g, +4°C), the supernatant was

collected and the pellet was re-extracted in a mixture of chloroform and methanol (2:1).

Back-extraction into an aqueous phase (0.05% $CaCl_2$) was followed by a re-extraction with a

mixture of chloroform:methanol: 0.05% $CaCl_2$ (3:49:48) and this phase was further purified

in-house on Bio-Rad AG 1-X2 (USA) resin columns. The iodothyronines were eluted with

70% acetic acid, and evaporated under N_2. Blanks (plain reagents without any sample) were

analysed in each extraction batch to detect any contamination. Samples from different

populations and treatments were equally distributed across four extraction batches. There was

no difference among the batches in hormone concentrations ($F < 1.0$, $p > 0.38$). T3 and T4

were quantified using a nanoflow liquid chromatography-mass spectrometry (nano-LC-

MS/MS) method, developed and validated in Ruuskanen et al. (2018). Briefly, before the

analysis, the dry samples were diluted in ammonium (NH_3). Internal standards $^{13}C_6$-T_3 and

$^{13}C_6$-T_4 (Sigma-Adrich, St.Louis, USA) were added to each sample to identify and quantify

the THs. A triple quadrupole mass spectrometer (TSQ Vantage, Thermo Scientific, San Jose,

CA) was used to analyse the samples. For the chromatographic separation of hormones, a

nanoflow HPLC system Easy-nLC (Thermo Scientific) was applied. On-column

quantification limits were 10.6 amol for T4 and 17.9 amol for T3. MS data was acquired

automatically using Thermo Xcalibur software (Thermo Fisher Scientific) and analysed using

Skyline (MacLean et al. 2010). For the analyses, peak area ratios of sample to internal

standard were calculated. TH concentrations are expressed as pg/mg fresh mass. Few samples

failed in the extraction, see samples sizes in Fig and 3.

Statistical analysis

All statistical analyses were conducted with SAS Enterprise Guide version 7.1. T4

concentration (pg/mg) was log-transformed to reach normality. We first analysed differences

in intraovarian T3 and T4 among long-term heat exposed (i.e. populations in the vicinity of

the nuclear power plants that are exposed to warm discharge waters; pooled LOV and OLK,

N = 13 individuals) and reference populations (pooled POO, POR, KOT, PYH, N = 24

individuals) using a linear model, using fish mass as a covariate. Data from individuals from

control treatments (CRTL and CRTL7D) were only included in this analysis. We then

analysed differences in intraovarian T3 and T4 concentration among all populations, again

using data only from the control treatments. From POO, we only had 1 individual for this

analysis, and thus this population was excluded from the analysis. Population and fish mass

were included as fixed effects.

234   Differences in ovarian T3 and T4 concentration between treatments (CRTL 7D, Cu

7D, HS 7D) were analysed using linear mixed models where treatment and fish body mass

were included as fixed effects and population as a random intercept to account for non-

independence of fish from the same population. Finally, we also repeated the above models

for fish size (mm). Post-hoc tests were further used to test pair-wise differences among

treatments. Models were reduced by removing non-significant factors ($\alpha = 0.05$). Normality

and heteroscedasticity of the residuals were visually inspected. Degrees of freedom were

estimated with Satterthwaite estimation method. Means and standard errors (SE) are shown in

the text and in the figures.

Results

When using data from control treatments only (pooled CRTL and CRTL 7D), we found no

differences in intraovarian T3 or T4 concentration in sites exposed to warm discharge waters

(pooled LOV and OLK) compared to reference sites (pooled four reference sites, T3: $F_{1,36} =$

0.88, p = 0.18, T4: $F_{1,36} = 0.25$, p = 0.62; Mean (pg/mg) ±SE: T3 exposed populations:

1.60±0.19, T3 reference populations: 2.21±0.24; T4 exposed populations: 0.57±0.08; T4

reference populations 0.53±0.04). When further comparing all populations, we found no

strong statistical evidence for differences among populations in intraovarian T3 or T4

concentration (T3: $F_{4,30} = 0.50$, p = 0.73, T4: $F_{4, 30} = 2.38$, p = 0.08; Fig 2a, b). Body mass

correlated negatively with intraovarian T3 concentration (estimate±SE: -0.842±0.363, $F_{1,30} =$

5.37, p = 0.027), but not with T4 concentration ($F_{1, 30} = 0.0$, p = 0.96). Fish size did not differ

among exposed and reference populations ($F_{1,35}$ = 1.89, p = 0.18), while there were small

differences among populations ($F_{4,32}$ = 6.92, p = 0.004): fish from KOT were significantly

smaller than those from OLK, PYH and POR (see Suppl Table 2).

In the copper exposure treatment, the measured concentrations of copper were 110,

120, 110 and 41µg/L after 2 hrs and 54, 67, 56 and 37µg/L after one week in the three

exposure and one control tank, respectively. After experimental exposure to copper for 7

days, fish from copper exposure group had higher intraovarian T3 and T4 concentration

compared to fish from control treatment (T3: overall test $F_{2,53.1}$ = 3.14, p = 0.05, post-hoc

Control vs Cu $t_{52.3}$ = -2.30, p = 0.02; T4: $F_{2, 53}$ = 2.62, p = 0.08, Control vs Cu $t_{53.8}$ = -2.26, p

= 0.027, Fig 3a, b). T3 or T4 concentrations of fish from the heat treatment did not differ

statistically from control treatment (post-hoc p-values >0.45), while ovarian T3 concentration

was lower in heat treatment compared to copper exposed fish (t_{52} = 2.52, p = 0.045, Fig 2a).

Fish size did not differ among the treatments ($F_{2,51}$ = 0.64, p = 0.53).

Discussion

In contrast to our predictions, we found no evidence that egg TH levels of individuals

originating from populations close to a long-term heat source (nuclear power plant discharge

waters), differ from individuals originating from reference populations. Furthermore, the six

populations sampled across a wide geographical range over the Baltic Sea showed similar

egg TH levels after acclimation in captivity. These results suggest that populations do not

show strong genetic (or permanent developmentally induced) variation in egg T4 or T3

levels, although at least some Baltic Sea stickleback populations are known to be genetically

differentiated (Guo et al., 2015). However, we cannot rule out that environmental sources of

variation (temperature, salinity, food availability) could influence egg THs in the wild,

enabling scope for transgenerational plasticity. Among-population variation in egg THs in

field populations was indeed reported by McComb et al. (2005) in bonnethead sharks.

Nevertheless, given that egg THs were not influenced by short-term experimental exposure to

heat, we may conclude that temperature – using the tested experimental duration and scope –

may not influence transfer of THs to eggs and offspring. The sample size was, however,

rather low in the among-population comparison, and the differences in environmental

conditions (see Suppl Table 1) among the populations were rather small at the time of the

sampling, thus the results should be interpreted with caution. Interestingly, stickleback

adaptive divergence from marine to stream environment was recently found to involve

thyroid hormone signalling (Kitano and Lema, 2013; Kitano et al., 2010), thus it would be

interesting to further compare the egg THs from freshwater and marine populations to

understand the role of transgenerational plasticity in the adaptation process.

We found that individuals exposed to environmentally relevant concentrations of

copper for seven days showed higher intraovarian T3 and T4 levels than controls. While

associations between copper and maternal THs have not been studied to date in fish, previous

studies on the associations between copper as an EDC and THs in plasma show complex

patterns depending on the duration of the exposure and species, as reported in Eyckmanns et

al. (2010): In common carp (*Cyprinus carpio*), T3 levels were *elevated* only after long-term

exposure (1 month), while in gibel carp (*Carassius gibelio*) there was a *decrease* in T3 from

24 h to 1 month of exposure. Both species showed *increases* in T4 over short-and long-term

exposure. In rainbow trout, T4 levels were elevated very fast after copper exposure and

remained elevated for 12h, whereas there was no influence on T3. Copper exposure also

increased plasma T4 in the common carp in another experiment (Hoseini et al., 2016).

Finally, copper exposure significantly decreased T3 but not T4 in European eels (*Anguilla*

Anguilla) (Oliveira et al., 2008), suggesting changes in deionization from T4 to T3 in tissues.

These inconsistencies may be explained by experimental conditions and dosages, exposure

time and species-specific responses.

The association between plasma and egg hormone levels in fish has not been fully

elucidated, but if we assume that there is some correlation between TH levels in eggs and

circulating TH levels (Ayson and Lam, 1993; Brown et al., 1988; Brown et al., 2014; Kang

and Chang, 2004; Raine and Leatherland, 2003), our results are in parallel with those of

common carp (see above). The increased T4 and T3 levels in response to copper exposure

reported in this study suggest that T4 biosynthesis or degradation was altered, but also that

conversion of T4 to T3 was potentially increased. Increased TH levels could be explained by

increased metabolic rates and energy expenditure in response to copper exposure, along with

increased oxidative stress (De Boeck et al., 2006; De Boeck et al., 1997; Sanchez et al.,

2005). All in all, this evidence suggests that copper exposure changes plasma and associated

egg TH levels, and can function as a transgenerational endocrine disruptor in fish. The effects

of altered egg TH status on offspring development, and e.g. metabolism and thermotolerance,

need to be further studied.

In contrast to our predications, we did not find any differences in TH concentrations

between experimental heat treatment and control. In a previous study it has been found that

three weeks acclimation to high and low temperature changed the muscle TH profile of

zebrafish, warm acclimated fish having higher concentrations of T3 and T2 than cold

acclimated ones (Little et al., 2013). However, in their study the warm acclimated fish were

not sensitive to changes in hormone levels suggesting that with temperature exposure also

e.g. TH transporters and receptors need to be evaluated. This could be one reason why we did

not see any change in egg TH levels with heat stress even though it is well known that heat

stress increases the metabolic rate of fish (e.g. Anttila et al., 2013; Fry, 1947).

We conclude that variation in egg THs was not explained by (genetic) variation

among populations nor short-term heat exposure, thus THs may not mediate transgenerational

plasticity over temperature (under the tested conditions). Given that parental temperature

environment is known to alter offspring phenotype (e.g. Donelson et al., 2012; Donelson et

al., 2018), further studies are needed to elucidate the molecular mechanism of such

transgenerational effects. Copper may lead to transgenerational endocrine disruption in

fishes, as both T4 and T3 levels in eggs were altered in response to moderate copper

exposure. The functional consequences of altered THs on offspring phenotype should be

further studied.

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Figures:

Fig 1. Sampling locations across the Baltic Sea. Yellow-and-black symbols refer to sites with

nuclear power plant discharge water and green symbols to reference populations.

Fig. 2. Intraovarian T3 (a) and T4 (b) concentrations (pg/mg, mean±SEs) in three-

spined sticklebacks from six populations across the Baltic Sea coast. Numbers above

the bars refer to sample sizes. POO was excluded from the statistical analysis due to

low sample size.

a)   b)

Fig.3. Intraovarian T3 (a) and T4 (b) concentrations (pg/mg, mean±SEs) in three-spined

sticklebacks experimentally exposed to copper (100µg/L, CU 7D), warm temperature (10°C

increase, HS 7D) or respective controls (CRTL 7D) for 7 days. Numbers above the bars refer

to sample sizes. Stars represent significant differences between treatments at p < 0.05. For the

statistical analyses the populations were pooled.

a)   b)

Supplementary Table 1. The environmental variables (mean±SD) of catching locations across
the Baltic Sea.

Location	Temperature during catching	Salinity	pH	Average ± SD temperature 16/5/–31/8/2018
LOV	19.9 ± 1.6	3.2 ± 0.2	7.6	19.6 ± 4.4
OLK	20.1 ± 0.1	5.3 ± 0.0	9.0	19.9 ± 3.6
POO	18.8 ± 0.5	4.3 ± 0.0	9.0	19.1 ± 4.2
POR	18.9 ± 0.8	4.9 ± 0.0	9.0	19.7 ± 3.4
PYH	17.6 ± 1.1	4.2 ± 0.4	9.0	19.5 ± 2.5
KOT	18.1 ± 0.7	1.7 ± 0.0	9.0	17.3 ± 4.9

507 Supplementary Table 2. The average lengths (mm, SE) of fish captured from six different

508 locations (populations) across the Baltic Sea. Populations with different letters are

509 statistically significantly different from each other (p <0.05, Tukey post-hoc test). POO was

510 not included in the statistical analyses due to small sample size.

Location	Fish size, mm average (SE)
LOV	6.4 ± 0.4ab
OLK	6.6 ± 0.3b
POO	6.1 ± 0.7
POR	6.9 ± 0.3b
PYH	6.5 ± 0.3b
KOT	5.2 ± 0.2a