

1 **Molecular mechanisms and tissue targets of brominated**
2 **flame retardants, BDE-47 and TBBPA, in embryo-larval life**
3 **stages of zebrafish (*Danio rerio*).**

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28 **Abstract**

29 Brominated flame retardants are known to disrupt thyroid hormone (TH) homeostasis
30 in several vertebrate species, but the molecular mechanisms underlying this process
31 and their effects on TH-sensitive tissues during the stages of early development are
32 not well characterised. In this study, we exposed zebrafish (*Danio rerio*) embryo-
33 larvae to 2,2',4,4'-tetrabromodiphenyl ether (BDE-47) and tetrabromobisphenol A
34 (TBBPA) via the water for 96 h from fertilisation and assessed for lethality, effects on
35 development and on the expression of a suite of genes in the hypothalamic-pituitary-
36 thyroid (HPT) axis via both real time quantitative PCR (qRT-PCR) on whole body
37 extracts and whole mount *in situ* hybridisation (WISH) to identify tissue targets. The
38 96-h lethal median concentration (96h-LC₅₀) for TBBPA was 0.9 µM and mortality
39 was preceded by retardation of development (smaller animals) and morphological
40 deformities including, oedemas in the pericardial region and tail, small heads,
41 swollen yolk sac extension. Exposure to BDE-47 did not affect zebrafish embryo-
42 larvae survival at any of the concentrations tested (1 – 100 µM) but caused yolk sac
43 and craniofacial deformities, a curved spine and shorter tail at the highest exposure
44 concentration. TBBPA exposure resulted in higher levels of mRNAs for genes
45 encoding deiodinases (*dio1*), transport proteins (*ttr*), the thyroid follicle synthesis
46 protein paired box 8 (*pax8*) and glucuronidation enzymes (*ugt1ab*) and lower levels
47 of *dio3b* mRNAs in whole body extracts, with responses varying with developmental
48 stage. BDE-47 exposure resulted in higher levels of *thrb*, *dio1*, *dio2*, *pax8* and
49 *ugt1ab* mRNAs and lower levels of *ttr* mRNAs in whole body extracts. TBBPA and
50 BDE-47 therefore appear to disrupt the TH system at multiple levels, increasing TH
51 conjugation and clearance, disrupting thyroid follicle development and altering TH
52 transport. Compensatory responses in TH production/ metabolism by deiodinases

53 were also evident. WISH analyses further revealed that both TBBPA and BDE-47
54 caused tissue-specific changes in thyroid receptor and deiodinase enzyme
55 expression, with the brain, liver, pronephric ducts and craniofacial tissues appearing
56 particularly responsive to altered TH signalling. Given the important role of TRs in
57 mediating the actions of THs during key developmental processes and deiodinases
58 in the control of peripheral TH levels, these transcriptional alterations may have
59 implications for TH sensitive target genes involved in brain and skeletal
60 development. These findings further highlight the potential vulnerability of the thyroid
61 system to disruption by BFRs during early developmental windows.

62

63 Keywords: brominated flame retardant, thyroid, endocrine, fish, toxicology,
64 development

65 **1. Introduction**

66 Over the last two decades, there has been growing concern over the levels of
67 brominated flame retardants (BFRs) in the environment due to their ability to disrupt
68 the thyroid hormone (TH) system [1]. THs play a key role in a wide range of
69 vertebrate physiological functions both during early development and in adult life
70 stages, including influencing the maturation of bones [2], the gonads [3] and the
71 central nervous system [4]. Given the detrimental effects that can result from subtle
72 changes in TH status, particularly during crucial developmental windows [5], even
73 relatively low environmental levels of BFRs pose a potential risk to the health of
74 humans and wildlife.

75

76 BFRs have been commercially important high production compounds since their
77 introduction to the global market in the 1970s and are used routinely in industrial and
78 consumer products in an effort to reduce fire-related injury and property damage [6].
79 Polybrominated diphenyl ethers (PBDEs) and tetrabromobisphenol A (TBBPA) are
80 amongst the most extensively used BFRs worldwide [6]. PBDEs are a family of 209
81 possible congeners that can be divided into 10 congener groups (mono- to
82 decabromodiphenyl ethers). Commercial PBDE mixtures are made up of congeners
83 with varying numbers of bromine atoms on their two phenyl rings and are classified
84 according to their average bromine content; penta-, octa-, and deca-BDEs. The
85 congener BDE-47 (2,2',4,4'-tetrabromodiphenyl ether) was one of the main
86 components of the now banned commercial penta-BDE mixtures, accounting for
87 approximately 40% of the product by weight [6, 7].

88

89 Since PBDEs are blended physically rather than bonded chemically to the polymer,
90 they subsequently migrate into the environment and over the past four decades they
91 have become ubiquitous environmental contaminants. BDE-47 is often the most
92 commonly detected PBDE congener in the aqueous phase of environmental water
93 samples [8] and often dominates the PBDE profiles of human tissues, marine
94 mammals, birds and birds eggs, invertebrates and fish [9-13]. The production and
95 usage of penta-BDE commercial mixtures were prohibited globally in 2004 and
96 officially labelled as Persistent Organic Pollutants (POPs) in 2009 [14]. Controls on
97 the use of the penta-BDE products are only just beginning to yield declines in the
98 lower brominated congener concentrations in environmental samples [15].
99 Nonetheless, global contamination occurs today as a result of the continued use and
100 disposal of older products, manufactured prior to the restrictions, which still contain
101 PBDEs. In 2016, over a decade since the introduction of restrictions, there is a
102 continued presence of penta-BDE constituents in indoor air and dust [16] and they
103 remain the dominant PBDEs detected in both human samples [up to 141 µg/kg lipid
104 weight (lw) in placental tissue] and wildlife samples (up to 1 mg/kg lw in freshwater
105 fish) [17-20].

106

107 TBBPA is used as a reactive and additive flame retardant in printed circuit boards
108 and electronic enclosures, respectively. It is the most widely used BFR (representing
109 approximately 60% of the total BFR market) [6], with global production estimated to
110 be over 200,000 metric tonnes a year [21]. TBBPA has been identified in dust,
111 sewage sludge, river sediments and the water phase of lakes and rivers [22-26].
112 Despite TBBPA's relatively short half-life [27] which suggests little potential for
113 bioaccumulation, it has been detected globally, in some cases at high levels, in many

114 biotic samples including human breast milk and plasma (up to 37.3 µg/kg lw) [28],
115 suggesting both recent and continuous use.

116

117 Several BFRs are structurally similar to thyroxine (T4), the precursor of the
118 biologically active TH 3,3',5-triiodo-L-thyronine (T3), and concerns have been raised
119 about their effect on the thyroid system of both mammalian and non-mammalian
120 vertebrates [1, 29]. Indeed, numerous *in vivo* studies have shown that both TBBPA
121 and PBDEs can alter the circulating levels of both T4 and T3 in a range of vertebrate
122 species, including fish [30, 31]. The thyroidal system is centrally driven by the
123 hypothalamic-pituitary-thyroid (HPT) axis and under peripheral control in all
124 vertebrates. The HPT axis is regulated as a negative feedback mechanism in which
125 the level of thyroid-stimulating hormone (TSH) secreted by the pituitary controls the
126 production and release of T4 by the thyroid follicles. In the peripheral system, TH
127 activity is tightly regulated by the metabolising enzymes iodothyronine deiodinases,
128 type I, II and III (D1, D2 and D3), which can modulate TH signalling in individual
129 tissues as well as controlling serum TH concentrations. D2 catalyses the outer ring
130 deiodination (ORD) of T4 to produce the bioactive T3. In contrast, D3 catalyses the
131 inner ring deiodination (IRD) of T4 and T3 producing the inactive metabolites reverse
132 T3 (rT3) and 3, 3'-diiodo-L-thyronine (T2), respectively. D1 is a kinetically inefficient
133 enzyme that is capable of catalysing both IRD and ORD [32, 33].

134

135 Until recently, evaluating thyroid disruption by environmental chemicals mainly relied
136 on measures of circulating TH levels, thyroid size or histopathology. It is important,
137 however, to note that the thyroid system can maintain normal physiological functions
138 in response to TH level perturbations by changing the production and/or metabolism

139 of THs by the thyroid gland itself, in target tissues or in the liver [34, 35]. Thyroid-
140 disrupting chemicals (TDCs) may not cause obvious changes to TH levels, but may
141 nonetheless alter TH homeostasis. Consequently, identifying tissue-specific
142 responses to environmental contaminants is fundamental to building our
143 understanding on the effect mechanisms of BFRs. Several studies have
144 demonstrated that BFRs can elicit localised effects on the expression of thyroid
145 related genes in adult fathead minnows (*Pimephales promelas*) and Chinese rare
146 minnows (*Gobiocypris rarus*) [30, 36], however isolating organs and/or tissues of
147 interest from embryos and larvae for studies on gene expression is more
148 challenging. Whole mount *in situ* hybridisation (WISH), however, allows for sites of
149 expression of target genes to be detected in whole zebrafish embryo-larvae and this
150 approach has been used to illustrate a significant up-regulation of deiodinase
151 encoding *dio1* mRNA levels in the periventricular region of the brain and *dio3b*
152 mRNA in the pronephric ducts of zebrafish embryos exposed to the hydroxylated
153 metabolite 6-OH-BDE-47 [37].

154

155 The overall aim of this study was to assess the toxicological effects of two important
156 BFRs, TBBPA and BDE-47, on zebrafish during early development. The first
157 objective was to examine the acute toxicity of both compounds on zebrafish embryo-
158 larvae, in terms of induced mortalities and deformities, in order to calculate lethal and
159 effect concentration (LC and EC values). The second objective was to examine the
160 effect of these compounds on the expression of a suite of genes in the HPT axis,
161 with the goal of highlighting potential effect mechanisms and target tissues of thyroid
162 disruption. Given the plasticity of the TH system, we further undertook to examine
163 the effects of TBBPA across different developmental life stages. We used a

164 combination of real time quantitative PCR (qRT-PCR) assays to quantify changes in
165 gene transcript levels in whole body extracts and WISH to assess changes in tissue
166 gene expression patterns.

167

168

169 **2. Materials and Methods**

170 **2.1 Materials and reagents**

171 Tetrabromobisphenol A (TBBPA; CAS 79-94-7; purity 97%) and 3,3',5-Triiodo-L-
172 thyronine (T3; CAS 6893-02-3; purity \geq 95%) were purchased from Sigma-Aldrich
173 (Gillingham, UK). 2,2',4,4'-tetra-bromodiphenyl ether (BDE-47; purity 99.8%) was
174 provided by Ulrika Winnberg, Jorke Kamstra and Kees Swart on behalf of Dr. Juliette
175 Legler from VU University Amsterdam, The Netherlands. Stock solutions of
176 chemicals were prepared by dissolving them in dimethylsulfoxide (DMSO).

177

178 **2.2 Zebrafish maintenance**

179 Adult zebrafish [*casper* (*mitfa*; *roy*) mutant strain] were obtained from breeding
180 stocks at the University of Exeter. The *casper* mutants lack melanocytes and
181 iridophores, thus facilitating the visualisation of gene expression via WISH. Fish
182 were maintained at 28 ± 1 °C in a 12:12 h light: dark cycle in a closed flow-through
183 system. Embryos were collected approximately 1 hour post fertilisation (hpf) from
184 breeding colonies, washed twice with embryo culture water with the addition of
185 methylene blue (10^{-5} %) to prevent fungal growth [38]. We found no effects of
186 methylene blue on thyroid signalling in fish reported in the literature. Eggs were
187 incubated in culture water without methylene blue. Embryo culture water was
188 aerated artificial freshwater made according to the ISO-7346/3 guidelines (ISO water

189 diluted 1:5, pH 6.5-7.5, air saturation 95-100%) [39]. Embryos were examined under
190 a stereomicroscope and only those fertilised were selected for subsequent
191 experimental work. All fish were maintained under approved protocols, according to
192 the UK Home Office regulations for the use of animals in scientific procedures.

193

194 **2.3 Acute toxicity of TBBPA and BDE-47**

195 TBBPA exposures were carried out at 0, 0.18, 0.46, 0.92, 1.38 and 2.7 μM (in
196 DMSO, not exceeding 0.01% of the culture medium). BDE-47 exposures were
197 carried out at 0, 1, 5, 10, 50 and 100 μM (in DMSO, not exceeding 0.1% of the
198 culture medium). Controls were incubated in DMSO at 0.01% (TBBPA) and 0.1%
199 (BDE-47). BDE-47 has a low water solubility hence the requirement for a higher
200 DMSO concentration. Twenty fertilised embryos were randomly allocated into
201 glass tanks containing 50 ml of each treatment concentration and half of the
202 exposure solutions were replaced every 24 h with freshly prepared solutions.
203 Exposures were conducted for 96 h starting from approximately 1-2 hpf. The number
204 of dead embryo-larvae and phenotypic deformities (compared to normal
205 development [40]) were recorded every 24 h before removing any dead embryo-
206 larvae. Experiments were carried out in triplicate and repeated three times.

207

208 **2.4 Effect of TBBPA and BDE-47 on gene transcripts in the HPT axis**

209 TBBPA exposures were carried out at concentrations of 0, 0.04, 0.18 and 0.46 μM
210 (in DMSO, not exceeding 0.01% of the culture medium) and BDE-47 exposures were
211 carried out at concentrations of 0, 0.1, 1 and 10 μM (in DMSO, not exceeding 0.1%
212 of the culture medium). Again, a higher DMSO concentration was used in the BDE-
213 47 exposure due to its low water solubility. These concentrations were chosen

214 based on range-finding tests to determine concentrations that were sub-lethal and
215 they are lower than, or similar to, the concentrations used in previous studies
216 reported in the literature [41, 42]. Control groups were incubated in DMSO at 0.01%
217 (TBBPA) and 0.1% (BDE-47). Fifty fertilised embryos were randomly allocated into
218 glass tanks containing 50 ml of each treatment concentration and half of the
219 exposure solutions were replaced every 24 h with freshly prepared solutions. TBBPA
220 exposures were conducted for 48, 96 and 120 h from fertilisation. BDE-47 exposures
221 were conducted for 96 h from fertilisation only, as this compound was obtainable in
222 amounts sufficient for analyses at one life developmental stage only.

223

224 At the desired developmental stage for the TBBPA and BDE-47 exposures, 40
225 individuals from each treatment were fixed in 4% paraformaldehyde (PFA) overnight
226 at 4 °C, washed and dechorionated in PBS and stored at -20 °C in 100% methanol
227 for WISH experiments. The remaining 10 embryo-larvae from each treatment group
228 were pooled, frozen in liquid nitrogen and stored at -80 °C for qRT-PCR analyses.
229 Experiments were carried out in triplicate and repeated three times. An overview of
230 the experimental design is provided in Figure 1.

231

232 **2.5 Transcript profiling by quantitative real-time PCR (qRT-PCR)**

233 qRT-PCR was used to quantify the transcript profiles of several target genes in the
234 HPT axis of zebrafish (whole body samples) including: thyroid receptors (*thraa* and
235 *thrb*), thyroid-stimulating hormone (*tshb*), deiodinases type I, II and III (*dio1*, *dio2*,
236 *dio3b*), transthyretin (*ttr*), corticotropin-releasing hormone (*crhb*), paired box 8 (*pax8*)
237 and uridine diphosphate-glucuronosyltransferase (*ugt1ab*). Ribosomal protein I8
238 (*rpl8*) was used as a control gene for normalisation purposes and its stable

239 expression has been validated (Fig. S1). qRT-PCR assays for each target gene were
240 optimised as previously described [43] and detailed information for each assay is
241 provided in the supplementary information (Table S1). Results are expressed as
242 mean fold changes \pm standard error of the mean.

243

244 **2.6 Whole mount *in situ* hybridisation (WISH)**

245 WISH was used to examine tissue-specific changes in gene expression for several
246 genes of interest in the HPT axis including: thyroid receptors (*thraa* and *thrb*) and
247 deiodinases (*dio1*, *dio2* and *dio3b*), with modifications to the protocol of Thisse and
248 Thisse (2008) [44] For details on the methodologies for gene probe synthesis
249 (including vector information) and WISH are provided in the supplementary material
250 (Supplemental Material and Methods, Tables S2 & S3).

251

252 **2.7 Statistical analyses**

253 All statistical analyses were conducted in R (R Studio, 1.1.423) [45]. LC_x and EC_x
254 values and their 95% confidence intervals (CI), for TBBPA and BDE-47 were
255 calculated using generalised linear models (GLM) with binomial error structures and
256 probit links according to Finney, 1971 [46]. Concentrations were log transformed
257 (\log_{10}) to linearise the data.

258

259 To examine the effect of TBBPA and BDE-47 exposures on gene transcripts in the
260 HPT axis, qRT-PCR data was analysed using general linear mixed models (GLMM)
261 with Gaussian error structures. GLMMs were performed with the *lme4* package
262 within R [47]. P values were obtained using maximum likelihood tests of the full
263 model (with BFR treatment incorporated as a fixed effect) against a reduced model

264 (without BFR treatment incorporated). BFR concentrations were incorporated as a
265 fixed effect into the model. As a random effect, random intercepts for each
266 experiment (since each experiment was repeated 3 times) were incorporated into the
267 model. When an overall significant effect of the BFR concentrations was identified,
268 pairwise comparisons to determine which groups differed were conducted using a
269 multiple comparison analysis of means (Tukey contrasts) with the *multcomp* package
270 within R [48]. Prior to analysis, gene expression data was scrutinised by Chauvenet's
271 criterion to detect outliers for each gene and these were subsequently removed [49].
272 In addition, data were tested for equal variance and for normality using the Shapiro–
273 Wilk test. Non-normal data were subjected to variance-stabilising log
274 transformations. All statistical models were checked for homoscedasticity and
275 normality of residuals. For all statistical analyses, differences were considered
276 significant at $p < 0.05$. All graphed data were plotted using the *ggplot2* R package
277 [50].

278

279

280 **3. Results**

281 **3.1. Acute toxicity of TBBPA and BDE-47**

282 Exposure to TBBPA for 96 h led to a significant increase in mortalities and
283 deformities of zebrafish embryo-larvae ($p < 0.001$; Fig 2; Table S4). The 96h-LC₅₀ with
284 95% confidence intervals for TBBPA was 0.9 μM (0.8 – 1 μM : Fig. 2A). The 96h-
285 EC₅₀ with 95% confidence intervals for TBBPA, based on deformities, was 0.7 μM
286 (0.6 – 0.9 μM : Fig. 2B). At the higher TBBPA concentrations ($>0.92 \mu\text{M}$), mortality
287 was preceded by retardation of development (smaller animals) and morphological
288 deformities (oedemas in the pericardial region and tail, small heads, swollen yolk sac

289 extension; Fig. 2C). Deformities were observed in $86 \pm 11\%$ of surviving individuals
290 exposed to $0.92 \mu\text{M}$ TBBPA, including oedema, short body, curved spine, swollen
291 yolk sac, small eyes and craniofacial deformities (Fig. 2D).

292

293 Exposure to BDE-47 ($1 - 100 \mu\text{M}$) had no significant effect on the mortality of
294 zebrafish embryo-larvae compared with controls (Fig. 3A). Exposure to BDE-47 for
295 a 96-h period led to a significant increase in the proportion of zebrafish deformities
296 ($p < 0.05$). The 96h-EC₅₀ with 95% confidence intervals for BDE-47, based on
297 deformities, was $38.2 \mu\text{M}$ ($10.4 - 475.9 \mu\text{M}$; Fig. 3B). The mean number of
298 deformities were $42 \pm 11\%$, $51 \pm 10\%$ and $68 \pm 12.0\%$ in groups exposed to BDE-47
299 at 10, 50 and $100 \mu\text{M}$, respectively. At these higher BDE-47 concentrations,
300 deformities included oedema, short tail and head deformities in 24 hpf embryos (Fig.
301 3C). At 96 hpf, curved spines, yolk sac deformities and craniofacial deformities were
302 also observed (Fig. 3D). See Table 1 for the full list of LC_x and EC_x values for both
303 TBBPA and BDE-47.

304

305 **3.2. Effect of TBBPA on gene transcripts in HPT axis**

306 3.2.1. TBBPA: Whole body transcript levels (qRT-PCR)

307 The effect of TBBPA (pools of $n=10$ larvae) on the transcription of genes in the HPT
308 axis were quantified at 48, 96 and 120 hpf via qRT-PCR and results are shown in
309 Figure 4 (A-J). TBBPA had no effect on *thraa*, *thrb*, *dio2*, *crhb* and *tshb* mRNA levels
310 in whole body extracts at any of the developmental stages examined relative to
311 controls. There was no effect of TBBPA on the expression levels of *dio1* at 48 and
312 120 hpf in embryo-larvae samples. After 96 h, however, exposure to $0.18 \mu\text{M}$ TBBPA
313 resulted in a significantly higher (1.5-fold) level of *dio1* mRNA in compared with

314 controls ($p < 0.01$). After 48- and 120-h exposures, TBBPA had no significant effect
315 on the levels of *dio3b* at any of the concentrations tested. After 96 h, however,
316 TBBPA exposure resulted in significantly reduced mRNA levels of *dio3b* in a
317 concentration-dependent manner (1.7, 2.8- and 3.2-fold lower in 0.04, 0.18 and 0.46
318 μM treatment groups) compared to the control, respectively ($p < 0.05$). Exposure to
319 0.46 μM TBBPA resulted in significantly higher levels of *ttr* after a 48-h exposure,
320 with mRNA levels 2-fold higher compared with the control ($p < 0.05$), but with no
321 effects thereafter (96- and 120-h exposures). *Pax8* transcript levels were higher (1.5-
322 fold) after a 48-h exposure to all TBBPA concentrations tested compared with the
323 control ($p < 0.01$), but this was not the case after 96- and 120-h exposures. *Ugt1ab*
324 mRNA levels increased, in a concentration-dependent manner, following 48- and 96-
325 h exposures to TBBPA ($p < 0.05$). After 48 h, *ugt1ab* mRNA levels were 1.7- and 5.1-
326 fold higher in the 0.18 and 0.46 μM TBBPA treatment groups, respectively,
327 compared to controls and 2-fold higher in the 0.18 and 0.46 μM treatment groups
328 compared with controls after 96 h. No significant effect of TBBPA exposure on
329 *ugt1ab* transcription was detected after 120 h.

330

331 3.2.2. TBBPA: Tissue specific transcript expression (WISH)

332 Following a 48-h exposure to TBBPA, *thraa* expression appeared to be higher in the
333 brain for all three concentrations and in the branchial arches for the 0.18 and 0.46
334 μM exposures, compared with controls (Fig. 5A-D). In 96 hpf larvae, exposure to
335 0.18 and 0.46 μM TBBPA resulted in higher (and concentration related) expression
336 of *thrb* mRNA in the liver, brain, jaw cartilage and swim bladder. There was also an
337 enhanced expression in the otic vesicles in animals treated with 0.46 μM TBBPA
338 compared with the controls (Fig. 5E-H). *Thrb* expression in the pronephric ducts

339 appeared to be reduced in TBBPA treated individuals (Fig. 5E-H) with 56% of control
340 larvae exhibiting *thrb* expression in the pronephric ducts compared with only 15% of
341 larvae exposed to 0.46 μ M TBBPA. TBBPA had no apparent effect of the expression
342 pattern of *thraa* and *thrb* in 120 and 48 hpf embryos-larvae, respectively, compared
343 with controls (Fig. S2A-H).

344

345 For all TBBPA exposure concentrations, WISH analysis illustrated that *dio1*
346 expression was enhanced in the brain of TBBPA exposed animals compared with
347 controls at 48 hpf (Fig. 6A-D). At 96 hpf, *dio1* expression was enhanced in the liver
348 and intestinal bulb in TBBPA exposed larvae (Fig. 6E-H). After a 48-h exposure to
349 TBBPA, a greater proportion of embryos displayed a detectable *dio2* expression in
350 the pituitary gland; 52% in control versus 74% of embryos treated with 0.46 μ M
351 TBBPA (Fig. 6I-L). After 96 h, TBBPA exposure (0.04, 0.18 and 0.46 μ M) resulted in
352 enhanced *dio2* signalling in the liver, brain and intestinal bulb compared with control
353 larvae (Fig. 6M-P). At 96 hpf, enhanced *dio2* signalling was also observed in the
354 pituitary gland of TBBPA treated larvae; 17% in control versus 34% of larvae treated
355 with 0.46 μ M TBBPA (Fig. S3A-D).

356

357 Via WISH, no effect of TBBPA was seen on the expression pattern of *dio3b* in 48 hpf
358 embryos (Fig. S3E-H) but in 96 hpf larvae expression of *dio3b* was reduced in the
359 liver (0.46 μ M; Fig. 7A-D).

360

361 **3.3 Effect of BDE-47 on gene transcripts in HPT axis**

362 3.3.1 BDE-47: Whole body transcript levels (qRT-PCR)

363 The effect of BDE-47 exposure on the transcription of genes in the HPT axis of
364 whole body 96 hpf larval samples is shown in Figure 8 (A-J). BDE-47 had no
365 significant effect on the expression of *thraa*, *crhb* and *tshb*. Compared with controls,
366 exposure to BDE-47 significantly induced whole body transcript levels of *thrb* (1.9-
367 fold at 10 μ M; $p < 0.001$), *dio1* (1.6-fold at 1 μ M; $p < 0.001$), *dio2* (1.4-fold at 1 μ M;
368 $p < 0.01$), *dio3b* (1.5-fold at 10 μ M; $p < 0.01$), *pax8* (2.2- and 2.3-fold at 1 and 10 μ M,
369 respectively; $p < 0.01$) and *ugt1ab* (2.1-fold at 1 μ M; $p < 0.01$). Transcript levels of *ttr*
370 were significantly reduced in BDE-47 exposed larvae compared with controls (1.4-
371 fold at 10 μ M; $p < 0.05$).

372

373 3.3.2. BDE-47: Tissue specific transcript expression (WISH)

374 Via WISH, enhanced *thraa* expression was observed in the brain, liver, pronephric
375 ducts and branchial arches of larvae exposed to BDE-47 (1 and 10 μ M) (Fig. 9A-D).
376 *Thrb* expression on the other hand was seen to be suppressed in the brain, otic
377 vesicle, intestinal bulb and liver, while expression in the pronephric ducts was
378 enhanced in larvae exposed to 1 and 10 μ M BDE-47 (Fig. 9E-H).

379

380 *Dio1* expression was enhanced in the pronephric ducts and liver of BDE-47 exposed
381 larvae compared to controls (1 and 10 μ M; Fig. 10A-D). A greater proportion of BDE-
382 47-exposed larvae showed *dio3b* expression in the pronephric ducts compared to
383 controls as determined by WISH; 28% and 7% of larvae exposed to 1 and 10 μ M
384 respectively versus none in controls (Fig. 10E-H).

385

386

387

388 **4. Discussion**

389 This study adopted zebrafish embryo-larvae as a biological model to evaluate
390 TBBPA and BDE-47 induced developmental toxicity and gene expression changes in
391 the HPT axis.

392

393 **4.1. Acute toxicity of TBBPA and BDE-47**

394 We show that exposure to high concentrations of TBBPA for 96-h was lethal to
395 zebrafish embryo-larvae, with an LC₅₀ value of 0.9 µM that is in general accordance
396 with similar studies conducted previously [51, 52]. Neither of those studies however
397 calculated an LC₅₀ value to allow for direct toxicity comparisons with our findings.
398 Lee *et al.*, (1993) reported an 96h-LC₅₀ value of 3 mg TBBPA L⁻¹ (5.5 µM) for adult
399 zebrafish suggesting a greater sensitivity to TBBPA in early life stages [53].
400 Consistent with previous studies, here we also observed various morphological
401 deformities for exposure to TBBPA, including pericardial and tail oedemas, small
402 heads and swollen yolk sac extensions, which may be a result of apoptosis and/or
403 oxidative damage [51, 52]. Based on deformities in surviving zebrafish, we estimated
404 an 96h-EC₅₀ of 0.7 µM for TBBPA, slightly lower than the 96h-EC₅₀ (based on
405 hatching) of 1.1 mg L⁻¹ (2.0 µM) reported previously [41].

406

407 BDE-47 up to 100 µM was not lethal to zebrafish larvae in the 96-h exposures,
408 consistent with the low acute toxicity of PBDEs observed in rodents [54]. There
409 appear to be considerable differences in sensitivities between fish species with
410 regards to lethal concentrations of BDE-47. In previous studies, zebrafish embryo-
411 larvae showed significant mortalities after exposure to BDE-47 up to 5 mg L⁻¹ (10.3
412 µM) albeit that the exposure period was longer (168 h) and the mortalities occurred

413 after 100 h [55]. In contrast, a relatively low 96h-LC₅₀ of 14.13 µg L⁻¹ (0.03 µM) has
414 been reported for turbot (*Psetta maxima*) larvae [56], which may indicate
415 considerable differences in sensitivities between fish species. We found, however,
416 deformities such as short tails, oedemas, curved spines and craniofacial deformities
417 for exposure to BDE-47, as also reported previously [55, 57] with a comparable 96h-
418 EC₅₀ for deformities (38.2 µM) with that reported for hatching (20.3 mg L⁻¹ [41.8 µM])
419 [41].

420

421 **4.2. Effects of TBBPA and BDE-47 on TR expression**

422 Here we found that *thraa* mRNA levels in whole body extracts were unaffected by
423 exposures to TBBPA and BDE-47 during early zebrafish development, comparable
424 to previous studies with zebrafish larvae exposed to BDE-47 and DE-71 for 96 hpf
425 and 14 dpf, respectively [41, 58]. In contrast, TRα transcript levels were enhanced in
426 zebrafish larvae exposed to TBBPA for 96 hpf [41] as well as larvae exposed to
427 TBBPA and BDE-47 later in development (from the point of hatching) [41]. Exposure
428 to TBBPA did not alter *thrb* RNA levels at any of the developmental stages examined
429 in the present study, consistent with previous finding with similar stage zebrafish
430 embryo-larvae [41]. We did, however, observe that exposure to BDE-47 resulted in
431 elevated *thrb* mRNA levels in 96 hpf larvae. While previous studies found no effect of
432 BDE-47 or DE-71 on TRβ transcript levels in whole body zebrafish larvae samples
433 exposed from fertilisation, others observed reduced mRNA levels in larvae exposed
434 to BDE-47 from the point of hatching [41, 58]. Taken together, these results suggest
435 that TR regulation by BFRs may vary depending on the specific developmental
436 window in which exposure occurs [59] and may differ depending on the receptor and
437 exposure compound in question.

438

439 Through WISH analyses, we observed tissue-specific alterations to both *thraa* and
440 *thrb* expression in zebrafish embryo-larvae following exposure to TBBPA and BDE-
441 47. Following a TBBPA exposure, *thraa* expression was higher in the brain and
442 branchial arches of exposed 48 hpf embryos compared to controls, while *thrb*
443 expression was enhanced in the liver, brain, jaw cartilage, otic vesicle and swim
444 bladder of 96 hpf larvae. As far as we are aware, no study to date has examined
445 tissue specific changes to TR gene expression in fish (either in adults via PCR or in
446 larvae via WISH) exposed to TBBPA. Similar to that seen for the TBBPA exposures,
447 *thraa* expression was found to be enhanced in the brain and branchial arches, as
448 well as the liver and pronephric ducts, of 96 hpf larvae exposed to BDE-47. These
449 results are consistent with the findings of increased *thra* transcript levels in the brain
450 of adult fathead minnows following dietary exposure to BDE-47 and BDE-209 for 21
451 and 28 days, respectively [30, 60]. In contrast, *thrb* expression was suppressed in
452 the liver, brain, intestinal bulb and otic vesicle of 96 hpf larvae exposed to BDE-47 in
453 our study, though expression was induced in the pronephric ducts. Interestingly, a
454 dietary exposure to BDE-47 reduced *thrb* mRNA levels in the brains of adult fathead
455 minnows, but no changes were observed in the liver [61], whilst in a similar study
456 with BDE-209 the opposite occurred [60]. TRs mediate the thyroid system's genomic
457 control over several key developmental processes, including growth [62], auditory
458 [63], eye [64], brain [65] and skeletal [66] development [67]. Whether the BFR-
459 induced changes in TR transcripts observed here translate to altered TH-dependant
460 developmental processes remains unclear. As far as we are aware, only one study
461 has examined the effect of overexpression of TR α 1 in zebrafish and report disrupted
462 hindbrain patterning in embryos [68]. Based on the observed transcriptional changes

463 of TRs, our results suggest that TH signalling in the brain, as well as the liver,
464 intestine and jaw cartilage are sensitive to TBBPA and BDE-47 during early
465 zebrafish development.

466

467 The mechanisms by which TBBPA and BDE-47 altered both TR transcripts in this
468 study are not fully understood. Changes may be a result of BFR-induced alterations
469 in circulating and/or local TH levels or a result of a direct interaction with TRs or their
470 co-repressors/activators (since the genes encoding TRs themselves contain TREs).
471 Since several *in vitro* studies have demonstrated that BFRs have limited agonistic
472 effects on TR-mediated gene transcription [69-71], it seems plausible that the
473 increases in *thraa* and *thrb* transcription following TBBPA and BDE-47 exposures
474 were a result of increased local T3 levels.

475

476 **4.3. Effects of TBBPA and BDE-47 on deiodinase expression**

477 Exposure to both TBBPA and BDE-47 resulted in higher levels of *dio1* transcripts in
478 whole body samples of 96 hpf zebrafish larvae, in agreement with results observed
479 in zebrafish larvae and adults exposed to DE-71, BDE-209 and BDE-47 [58, 72, 73].
480 In contrast, other studies found that *dio1* transcript levels were not altered in
481 zebrafish juveniles exposed to BDE-47 or zebrafish larvae exposed to TBBPA,
482 respectively, though the exposure period in the BDE-47 study began at a later stage
483 in development [42, 74]. Interestingly, here TBBPA had no effect on *dio2* levels in
484 whole zebrafish embryo-larvae at any stage examined, while exposure to BDE-47
485 led to higher *dio2* levels in 96 hpf larvae. These results are in line with previous
486 studies, where no change in *dio2* levels were detected in zebrafish larvae exposed to
487 TBBPA for 5 d from fertilisation [42], while increased *dio2* levels in larvae were

488 detected following exposure to BDE-47 and DE-71 for 14 d from fertilisation [58, 73].
489 Whilst D1 is capable of both activating and deactivating THs, in fish it has been
490 suggested that the former is its primary function, in contrast to its regulation and
491 function in mammals [75, 76]. D2 is considered, however, to be the major TH-
492 activating enzyme. Therefore, the increased levels of *dio1* and *dio2* transcripts
493 observed here provide strong evidence of an induced systemic hypothyroid condition
494 in the TBBPA and BDE-47 exposed zebrafish larvae [76-78]. This likelihood is
495 supported by the reduced *dio3b* mRNA levels observed in whole larval (96 hpf)
496 samples exposed to TBBPA (0.18 and 0.46 μ M groups), since D3 is the major
497 inactivating pathway of THs and thyroidal status often parallels hepatic D3 activity,
498 increasing during hyperthyroidism and decreasing during hypothyroidism [33, 79].

499

500 WISH analysis provided us with a more detailed insight into the tissue specific
501 changes and the mechanisms underlying thyroid disruption to *dio1*, *dio2* and *dio3b*
502 expression in zebrafish embryo-larvae following BFR exposure. We observed higher
503 *dio1* expression in the brain of embryos (48 hpf) and in the liver and intestinal bulb of
504 larvae (96 hpf) exposed to TBBPA, while higher levels of *dio2* expression were
505 observed in the pituitary of embryos (48 hpf) and in the brain, pituitary and intestinal
506 bulb of larvae (96 hpf). As far as we aware, no previous studies have examined
507 tissue specific changes in the expression of genes encoding the deiodinase
508 enzymes following exposure to TBBPA. The increased expression of *dio2* in the
509 pituitary of TBBPA exposed embryo-larvae is not fully understood. In fish under
510 normal physiological regulations, reduced circulating TH levels lead to increased
511 stimulation and secretion of TSH from the pituitary through the negative feedback
512 system of the HPT axis [35]. An increased conversion of T4 into T3 in the pituitary

513 (catalysed by increased D2 levels) may deplete local T4 levels, inducing the negative
514 feedback loop and increasing the production of THs by the thyroid follicles. In
515 support of the qPCR results, the higher expression of *dio1* and *dio2*, along with the
516 reduced expression of *dio3b*, in the liver of TBBPA exposed larvae provides strong
517 evidence of systemic hypothyroidism [76]. In addition, the elevated *dio1* and *dio2*
518 mRNA levels in the brain and intestinal bulb are likely associated with reduced local
519 TH levels and may be compensatory responses in order to maintain local TH
520 homeostasis during the development of neurological and gastrointestinal tissues. In
521 BDE-47 exposed larvae, enhanced *dio1* expression was observed in the pronephric
522 ducts and liver. This is in contrast to a previous study with zebrafish, which found no
523 significant changes in *dio1* transcript levels in the liver of adults exposed to BDE-47
524 for 180 days from fertilisation, though *dio2* levels were significantly increased [73].
525 Interestingly, we found that BDE-47 had no effect on *dio3b* expression in the liver but
526 higher levels were observed in the pronephric ducts. Increased D3 activity as a result
527 of increased *dio3b* expression in the pronephric ducts may reflect increased local T3
528 production (*dio1* expression also increased in pronephric ducts). Our results suggest
529 that altering the expression of deiodinase genes could be a potential mechanism by
530 which TBBPA and BDE-47 disrupt/correct TH homeostasis, though observed effects
531 vary depending on the life stage examined, exposure period and compound tested.

532

533 **4.4. Effects of TBBPA and BDE-47 on the expression of further genes in the** 534 **HPT Axis**

535 Elevated *ugt1ab* mRNA levels were detected in whole zebrafish samples exposed to
536 TBBPA (at 48 hpf and 96 hpf) and BDE-47 (at 96 hpf), similar to the effect observed
537 with zebrafish exposed to DE-71 [58]. UDPGTs play an important role in TH

538 metabolism in the liver and increased *ugt1ab* expression could lead to the increased
539 elimination of THs, a plausible explanation for the altered TH signalling observed
540 throughout the HPT axis here [80]. Previous findings in mammals and fish have
541 reported reduced circulating levels of T4, combined with increased UDPGT activity
542 and/or transcription, following exposure to BFRs [58, 81].

543

544 Elevated levels of *pax8* transcripts were observed here in zebrafish embryo-larvae
545 following exposures to TBBPA (at 48 hpf) and BDE-47 (at 96 hpf). Similar findings
546 have been reported in zebrafish embryos/larvae exposed to various other pollutants
547 [58, 82], although to our knowledge no studies have yet examined the effect of
548 TBBPA on *pax8* transcription. Pax8 proteins are essential for the late differentiation
549 of the thyroid follicular cells during development [83], therefore our results suggest
550 the promotion of thyroid primordium growth and possibly a compensation
551 mechanism for reduced TH levels, induced by the BRFs tested.

552

553 TTR is proposed to be the major TH carrier protein in fish [84] and plays a key role in
554 maintaining extra-thyroidal stores of TH and regulating its supply to various target
555 tissues [85]. Here we found that TBBPA significantly increased *ttr* transcripts levels
556 in 48 hpf embryos while BDE-47 significantly decreased *ttr* transcript levels in 96 hpf
557 larvae. Our results are consistent with previous studies which have found increased
558 and decreased *ttr* mRNA levels in zebrafish larvae following exposure to TBBPA [41]
559 and BDE-47, respectively [41, 73]. Altered levels of TTR proteins in exposed
560 individuals, may decrease or increase the amount of TTR available to bind free TH.
561 In the case of BDE-47, the reduced *ttr* levels may lead to excess unbound TH, which
562 would be more susceptible to hepatic catabolism, resulting in a greater clearance

563 rate and a decrease in circulating TH concentrations. The impacts of the increased
564 *ttr* levels in TBBPA exposed larvae may be more complex, since TBBPA has been
565 shown to potently compete with T3 for TTR binding sites. Increased TTR protein
566 levels could potentially lead to increased substrate available for TBBPA binding,
567 making the displaced and unbound THs in the plasma more susceptible to hepatic
568 metabolism and suppress circulating levels of T4/T3 [86]. The altered *ttr* mRNA
569 expression by TBBPA and BDE-47, suggest that while induction of UDPGTs may be
570 partly responsible for disrupting systemic TH levels, other mechanisms may also be
571 involved.

572

573 Corticotropin-releasing hormone (CRH) and thyroid-stimulating hormone (TSH)
574 regulate THs synthesis and secretion in fish through a negative feedback
575 mechanism within the HPT axis, triggered via alteration in circulating T4 levels [87].
576 Interestingly, in the present study, both TBBPA and BDE-47 had no effect on *tshb*
577 and *crhb* transcript levels in zebrafish embryo-larvae. This is in agreement with
578 previous studies that found no effect on *tshb* mRNA levels in 5 dpf zebrafish larvae
579 exposed to TBBPA and juvenile zebrafish exposed to BDE-47 for 40 days [42, 74].
580 However, the expression of *crhb* and *tshb* has been shown to be elevated in adult
581 fathead minnows exposed to BDE-47 [61] and both adult and larval zebrafish
582 exposed to various BFRs, including TBBPA and BDE-47 [41, 73]. Increased *crhb*
583 and *tshb* expression in these studies were accompanied by reduced T4 levels.
584 Interestingly, recent studies have reported lower *crhb* and/or *tshb* transcript levels in
585 zebrafish adults, embryos and larvae exposed to TBBPA or BDE-47 [41, 88],
586 associated with increased T4 and T3 levels [88]. From these studies, it is evident
587 that BFRs have varying effects on the expression of *crh* and *tsh* genes in fish,

588 highlighting the importance of examining the effect of TDC on multiple levels of the
589 HPT axis.

590

591 **4.5. Combining qRT-PCR and WISH to enhance our understanding of tissue**
592 **specific targets of BFRs.**

593 Given the wide range of tissues influenced by THs during early development and the
594 complex nature involved in regulating the TH system, it is important to assess the
595 effect of TDCs on the spatial expression of targeted thyroid-related genes. To date,
596 assessing transcriptional changes in fish embryo-larvae following exposure to BFRs
597 have largely relied on qRT-PCR, while tissue-specific effects have been limited to
598 adult studies where isolating organs/tissues of interest was relatively straightforward.
599 In the present study, we simultaneously examined the effects of TBBPA and BDE-47
600 on the transcription of several thyroid-related genes using WISH and qRT-PCR.
601 While qRT-PCR provides quantitative data on gene expression and is sensitive at
602 detecting even small changes in mRNA levels, when carried out on pooled
603 homogenates of whole embryo-larvae, transcriptional changes in small and/or
604 localised tissues may not be detected if diluted by non-responsive organ/tissues of
605 the whole animal. Only recently has WISH been used in ecotoxicology studies as a
606 means of examining gene expression changes on a spatial scale and identifying
607 tissue-specific markers of exposure during early stages of development [37, 89].
608 Interestingly here, the transcription of several genes in the HPT axis appeared
609 unaffected by BFR exposures when examined using qRT-PCR but tissue-specific
610 effects were observed using WISH. These included *thraa* (at 48 hpf; TBBPA and at
611 96 hpf; BDE-47) *dio1* (at 48 hpf; TBBPA), *dio2* (at 48 and 96 hpf; TBBPA) and *dio3b*
612 (at 96 hpf; BDE-47). Here we demonstrate the value of WISH as an approach, in

613 conjunction with qRT-PCR, to examine the mechanistic basis of the effects of TDCs
614 on early-life stage zebrafish.

615

616 **4.6. Developmental stage-specific effects**

617 We also examined the effect of TBBPA exposure on thyroid gene transcripts at three
618 early developmental stages and found that effects were developmental stage
619 specific. TBBPA-induced alterations to *dio1* and *dio3b* transcription were observed in
620 96 hpf larvae while *ttr* and *pax8* mRNAs were altered in 48 hpf embryos only. Finally,
621 *ugt1ab* expression was altered at 48 and 96 hpf. TBBPA had no effect on any of the
622 genes examined following 120-h exposures. This study is one of only a few that
623 examined the effect of BFRs on thyroid-related gene transcription at different life
624 stages. Similar to our findings, several genes in the HPT axis were found to be
625 transiently altered in the liver and/or brain of adult male fathead minnows exposed to
626 BDE-209 [60]. The TH system is regulated by sophisticated compensatory
627 mechanisms both centrally (negative feedback loop in HPT axis) and peripherally (by
628 deiodinases), which, if adequate, can normalise both serum and peripheral TH
629 concentrations [90]. These compensatory mechanisms may explain why no
630 transcriptional effects were detected here in 120 hpf larvae. It is also important to
631 note that increased endogenous T3 levels are associated with the embryo-larval
632 transition of teleost fish, and this natural increase may intensify the effects of BFRs
633 on the thyroid system at this life stage. We observed that zebrafish at 96 hpf, a
634 period coinciding with increased TH levels, were particularly sensitive to TBBPA
635 exposure.

636

637

638 **Summary**

639 In summary, the present study demonstrates that the BFRs, TBBPA and BDE-47,
640 can disrupt the thyroid axis of zebrafish embryo-larvae at multiple levels by altering
641 mRNA transcript levels of genes encoding deiodinases, thyroid synthesis proteins,
642 transport proteins and glucuronidation enzymes. We have also shown that both
643 compounds induced tissue-specific transcriptional changes for several genes in the
644 HPT axis, with TH signalling in the brain, liver, pronephric duct and craniofacial
645 tissues appearing particularly sensitive to TBBPA and BDE-47 exposures (Fig. 11).
646 Furthermore, TH disruption by TBBPA appeared more pronounced in larvae at 96
647 hpf, compared to 48 and 120 hpf. These results demonstrate the effects of BFRs on
648 the thyroid system should not be generalised across tissues or developmental
649 stages in fish species. Future work should focus on understanding the
650 consequences of chronic low level exposure to BFRs and examine whether the BFR-
651 induced changes in TR transcript levels in the brain and skeleton as observed here
652 translate into health consequences for wild fish populations and humans.

653

654

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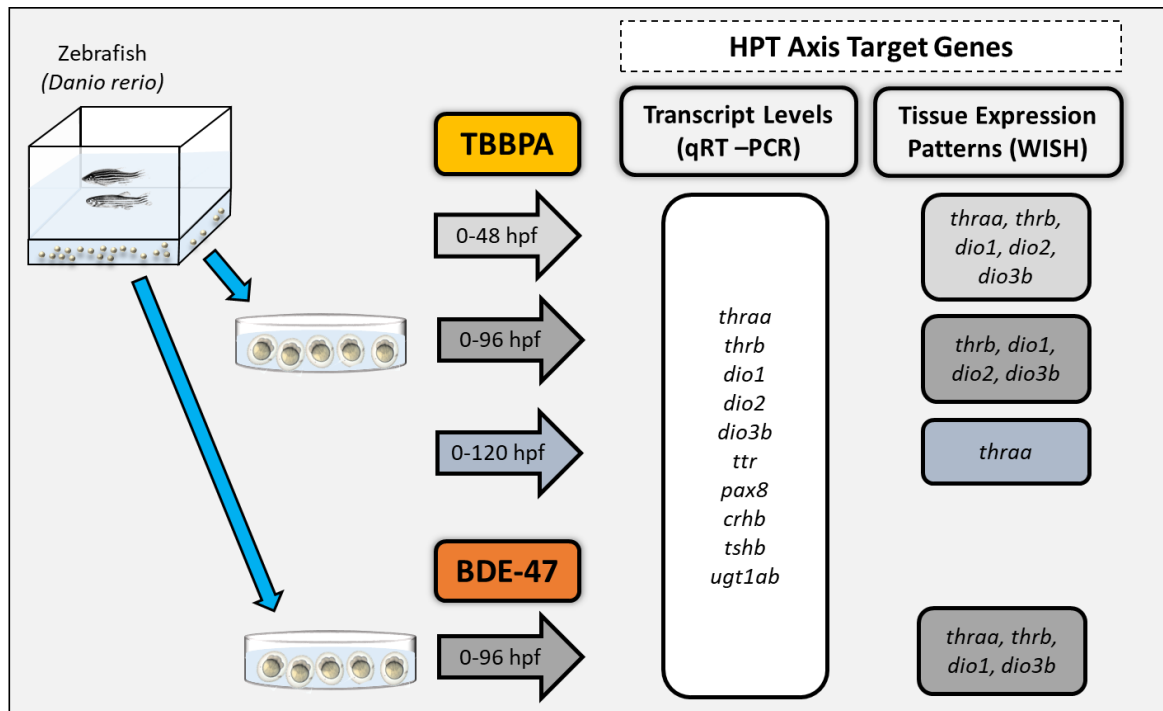
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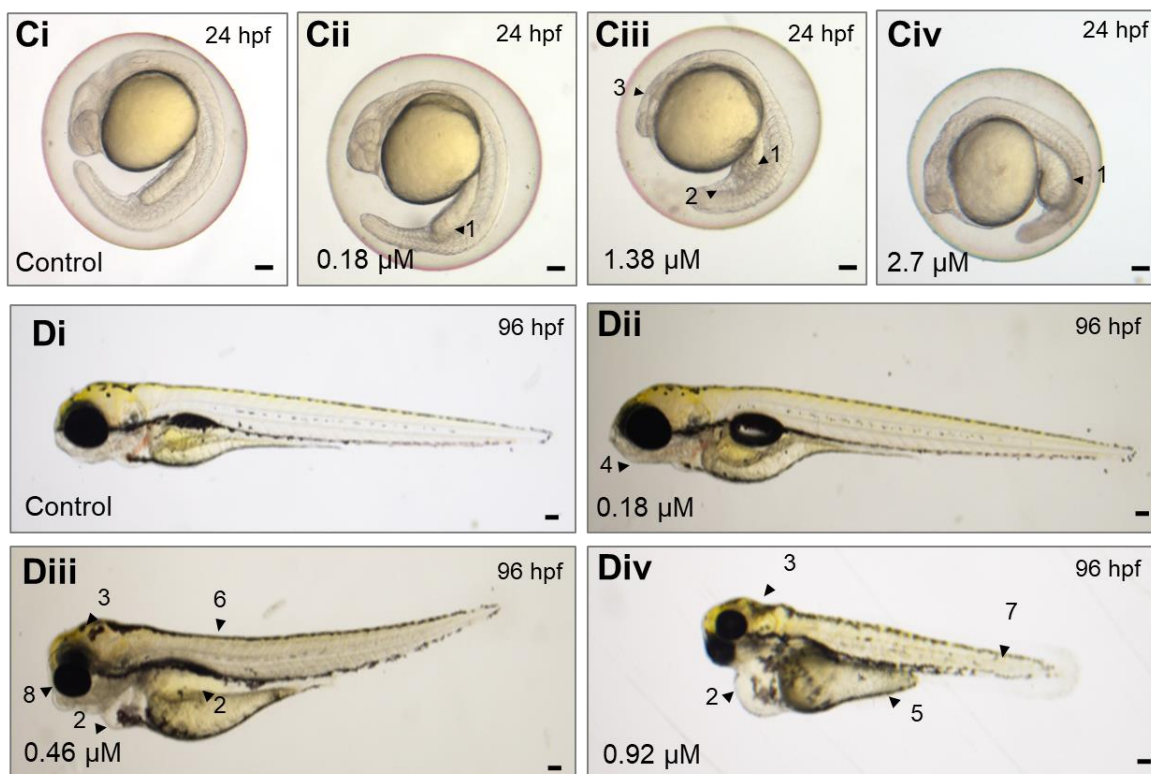
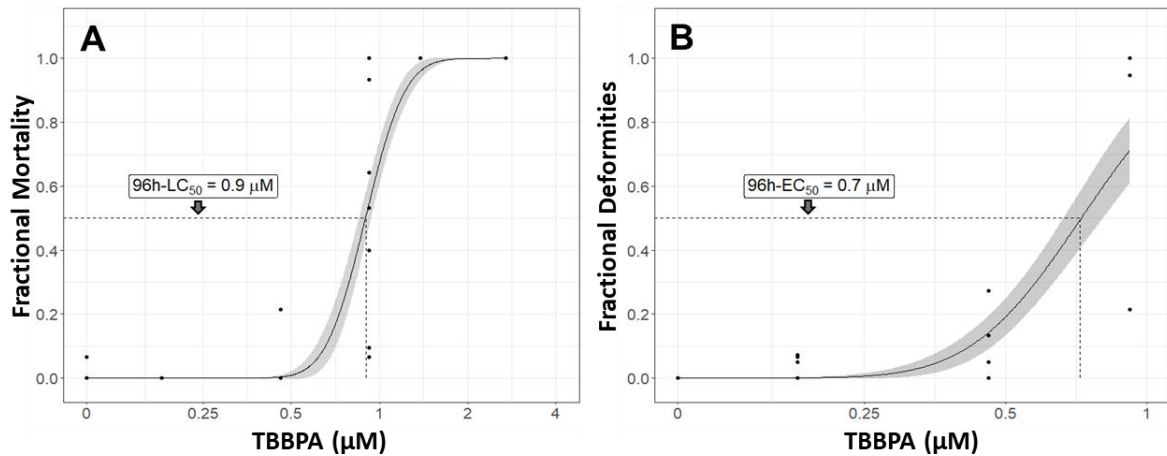
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889 **Figures**



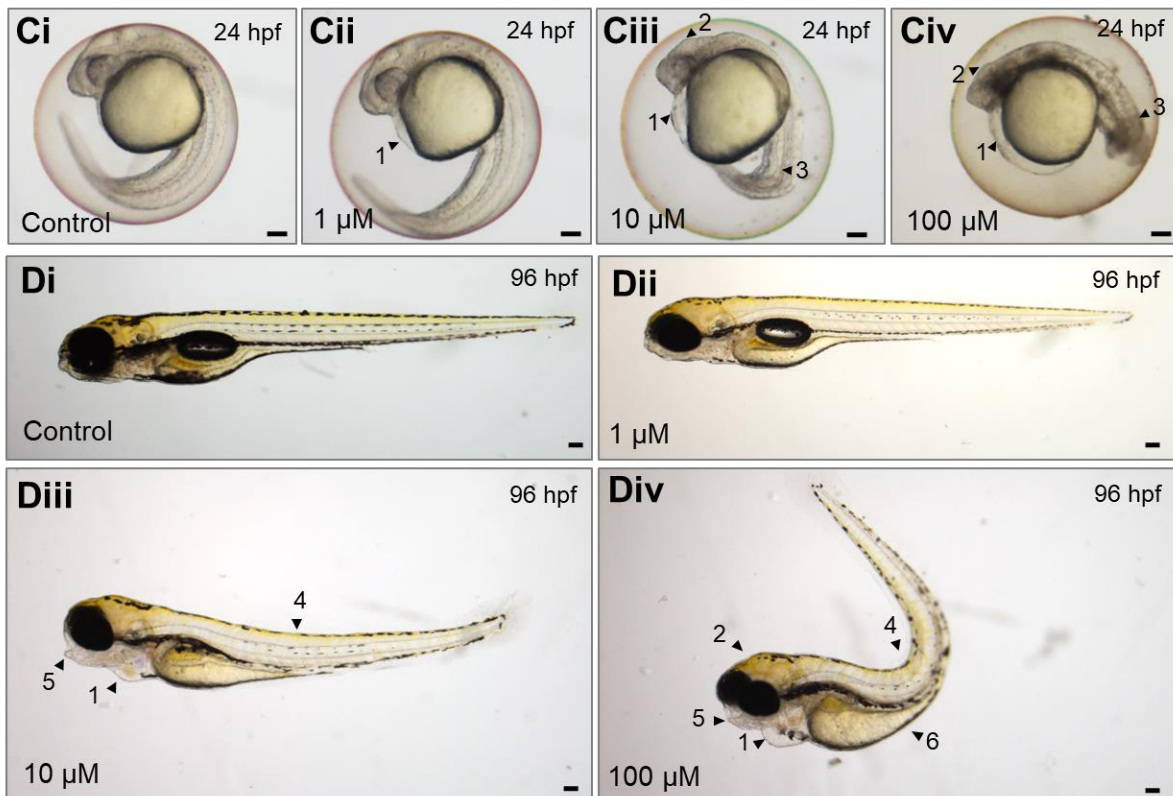
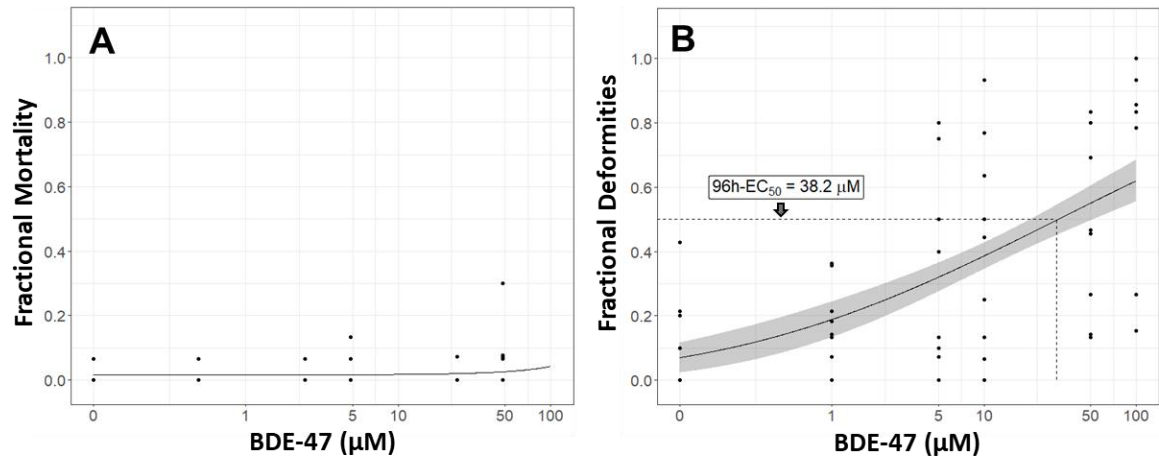
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891 Figure 1. Overview of the experimental design of this study, showing the HPT axis
 892 target genes analysed with WISH and qRT-PCR assays following exposures of
 893 zebrafish embryo-larvae to TBBPA or BDE-47 exposures.



894
 895 Figure 2. Effects of TBBPA on developing zebrafish embryo-larvae. Dose response
 896 curves showing (A) mortality and (B) deformities in zebrafish embryo-larvae after 96-
 897 h exposures to TBBPA (at nominal concentrations between 0.18 and 2.7 μM). Each
 898 point on the graphs represents the relative mortality and relative deformities in an
 899 individual replicate glass dish containing 15-20 individuals and the lines represent
 900 the best-fit model for the data, calculated using generalised linear models in R
 901 (model output summarised in Table S5A). Representative images of zebrafish
 902 embryos at 24 hpf exposed to (Ci) Control (Cii) 0.18 μM , (Ciii) 1.38 μM and (Civ) 2.7

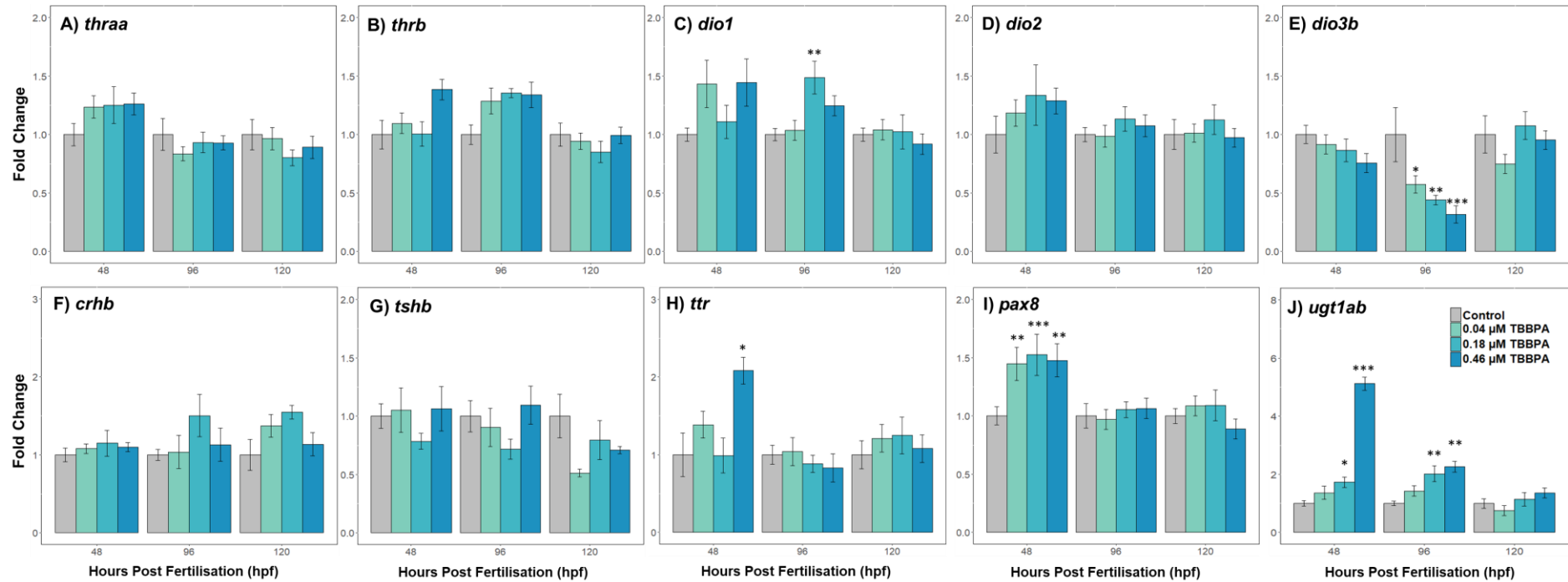
903 μM and at 96 hpf exposed to (Di) Control (Dii) 0.18 μM , (Diii) 0.46 μM and (Div) 0.92
904 μM showing- 1. swollen yolk sac extension, 2. oedema, 3. head deformity, 4. lower
905 jaw deformity, 5. yolk sac deformity, 6. bent spine, 7. short tail and 8. smaller than
906 normal eyes. Scale bar=100 μm .



907

908 Figure 3. Effects of BDE-47 on developing zebrafish embryo-larvae. Dose response
 909 curves showing (A) mortality and (B) deformities in zebrafish embryo-larvae after 96-
 910 h exposures to BDE-47 (at nominal concentrations between 1 and 100 μM). Each
 911 point on the graphs represents the relative mortality and relative deformities in an
 912 individual replicate glass dish containing 15-20 individuals and the lines represent
 913 the best-fit model for the data, calculated using generalised linear models in R
 914 (model output summarised in Table S5B). Representative images of zebrafish

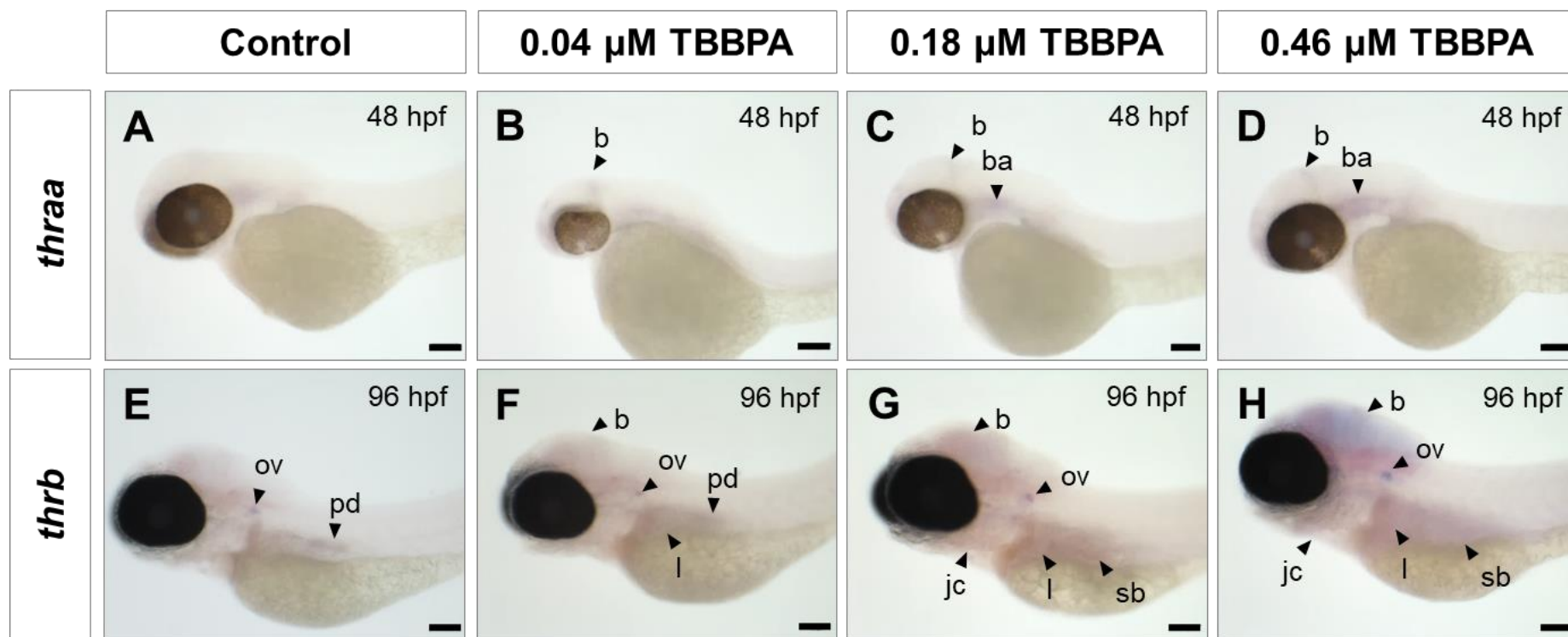
915 embryo-larvae at (C) 24 hpf and (D) 96 hpf following exposure to (i) Control (ii) 1 μ M,
916 (iii) 10 μ M and (iv) 100 μ M showing— 1. oedema, 2. head deformity, 3. short tail, 4.
917 bent spine, 5. lower jaw deformity and 6. yolk sac deformity. Scale bar=100 μ m.



918

919 Figure 4. Transcript profile of genes in the HPT axis of zebrafish embryo-larvae exposed to TBBPA. Changes in (A) thyroid receptor
 920 alpha (*thraa*), (B) thyroid receptor beta (*thrb*), (C) deiodinase type I (*dio1*), (D) deiodinase type II (*dio2*), (E) deiodinase type III
 921 (*dio3b*), (F) corticotropin-releasing hormone (*crhb*), (G) thyroid-stimulating hormone (*tshb*), (H) transthyretin (*ttr*), (I) paired box 8
 922 (*pax8*) and (J) uridine diphosphate-glucuronosyltransferase (*ugt1ab*) mRNA levels in whole zebrafish following exposure to TBBPA
 923 (0, 0.04, 0.18 and 0.46 μM) for up to 120 hpf. Transcript profiles were determined using qRT-PCR and the relationship between
 924 transcript expression and TBBPA concentration was assessed using linear mixed models (model output summarised in Table S6).

925 Plotted data are presented as mean fold changes (normalised against the expression of the control gene *rp18*) \pm SEM compared to
926 the corresponding control group. Outliers, identified as described in the text, were excluded from the analysis, resulting in a
927 replication of n= 6-9 samples per treatment group. Significance codes: *p<0.05, **p<0.01, ***p<0.001.



928

929

Figure 5. WISH images of TR mRNA expression patterns in zebrafish embryo larvae following exposure to TBBPA. Representative images of (A-D) *thraa* mRNA expression patterns in zebrafish embryos at 48 hpf and (E-H) *thrb* mRNA expression patterns in zebrafish larvae at 96 hpf treated with TBBPA. Lateral (A-D) and dorsal (E-H) views of whole embryo-larvae are shown with anterior positioned to the left. Focal areas of expression are indicated by black arrowheads. b=brain, ba=branchial arches, ov=otic vesicle, l=liver, sb=swim bladder, pd=pronephric duct, jc=jaw cartilage. Scale bar=100 μm . A summary of the variability in mRNA expression in the different zebrafish tissues is provided in Table S7.

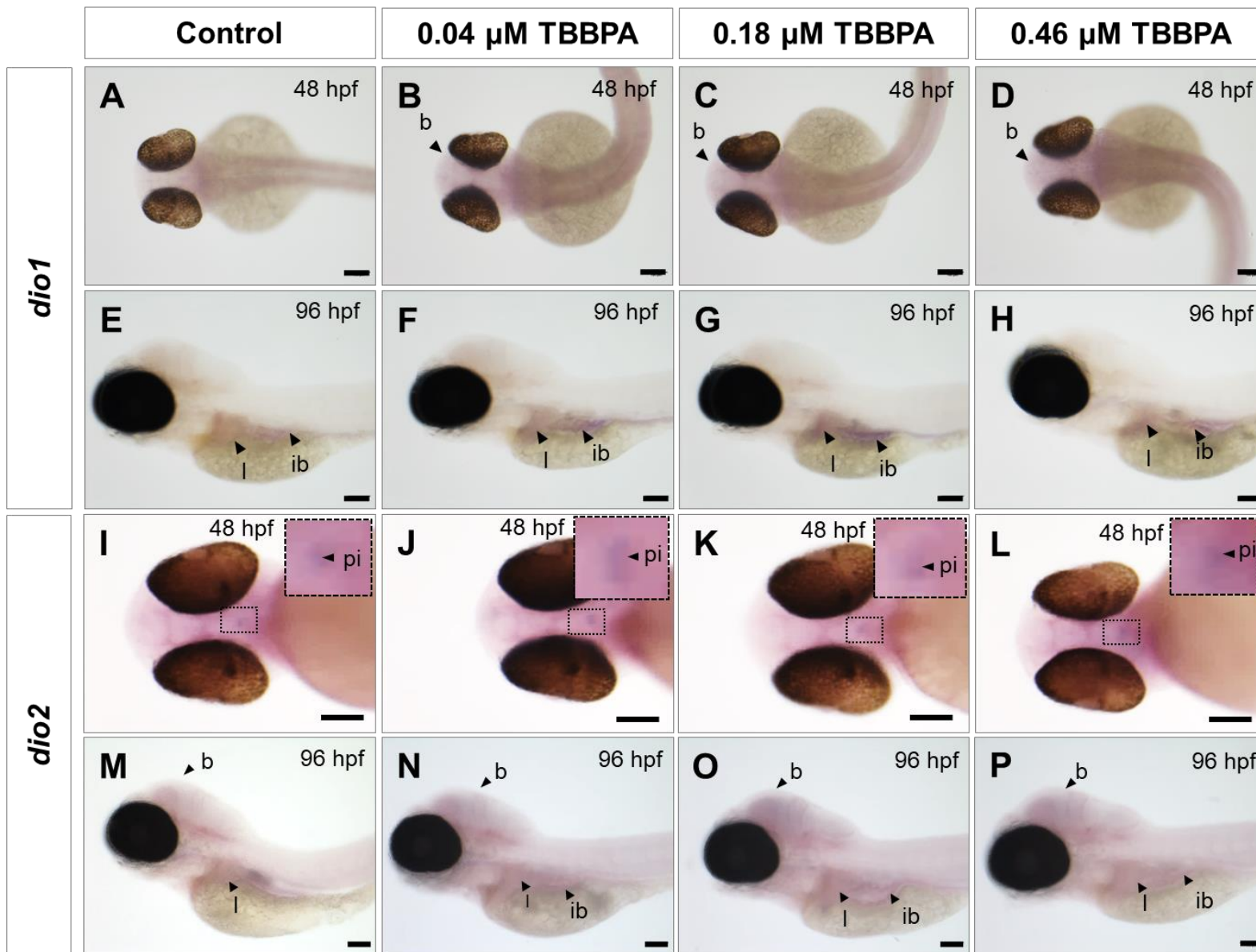
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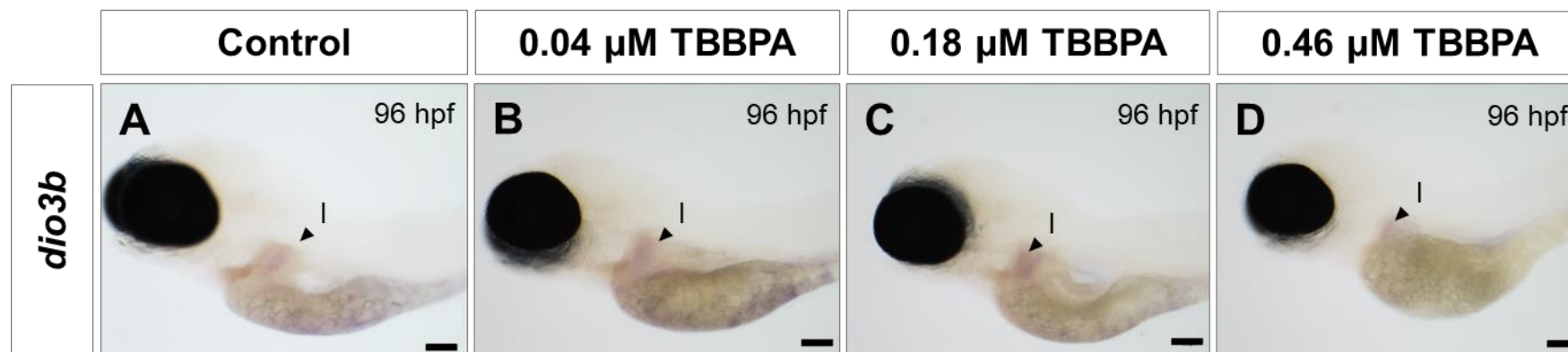
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936 Figure 6. WISH images of deiodinase type I and type II mRNA expression patterns in zebrafish embryo larvae following exposure to
937 TBBPA. Representative images of mRNA expression patterns in zebrafish embryo-larvae treated with TBBPA showing (A-D) *dio1*
938 at 48 hpf, (E-H) *dio1* at 96 hpf, (I-L) *dio2* at 48 hpf and (M-P) *dio2* at 96 hpf. Dorsal (A-D), lateral (E-H, M-P) and ventral (I-L) views
939 of whole embryo-larvae are shown with anterior positioned to the left. Focal areas of expression are indicated by black arrowheads.
940 l=liver, ib=intestinal bulb, b=brain, pi=pituitary. Scale bar=100 μ m. A summary of the variability in mRNA expression in the different
941 zebrafish tissues is provided in Table S8.



942

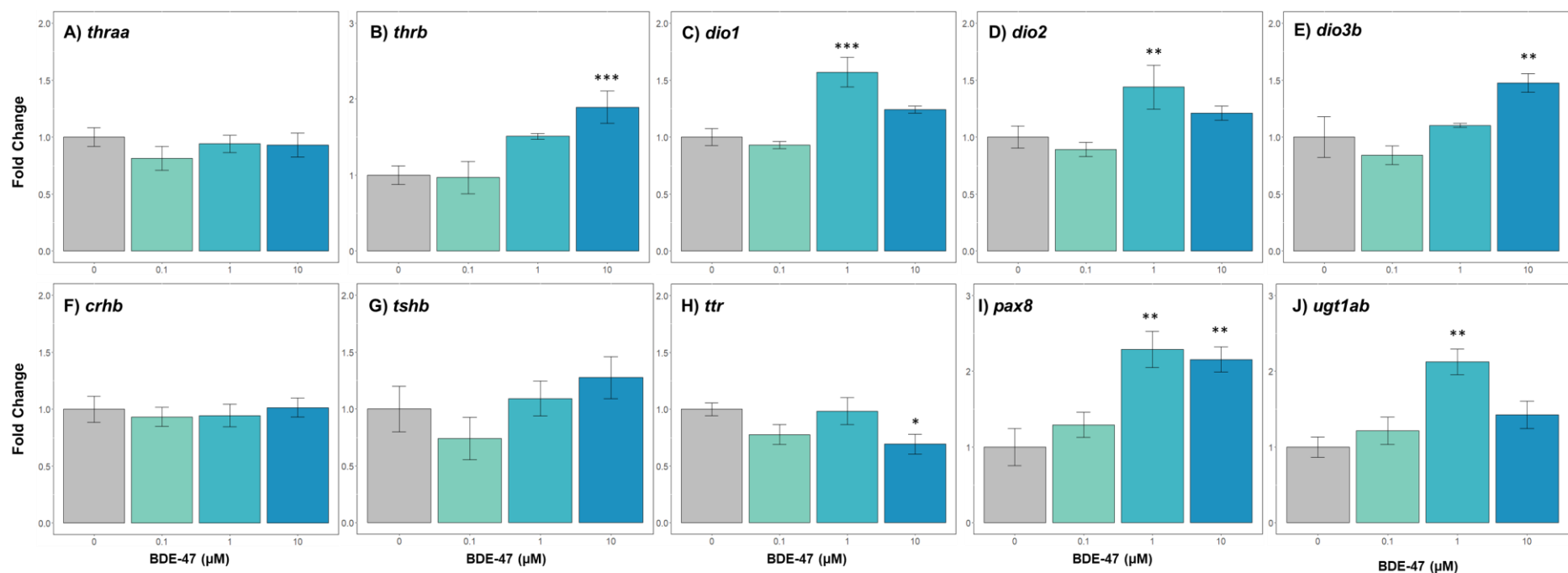
943 Figure 7. WISH images of deiodinase type III mRNA expression patterns in zebrafish larvae following exposure to TBBPA. (A-D)

944 Representative images of *dio3b* mRNA expression patterns in 96 hpf zebrafish larvae treated with TBBPA. Lateral views of whole

945 larvae are shown with anterior positioned to the left. Focal areas of expression are indicated by black arrowheads. l=liver. Scale

946 bar=100 μm . A summary of the variability in mRNA expression in the different zebrafish tissues is provided in Table S8.

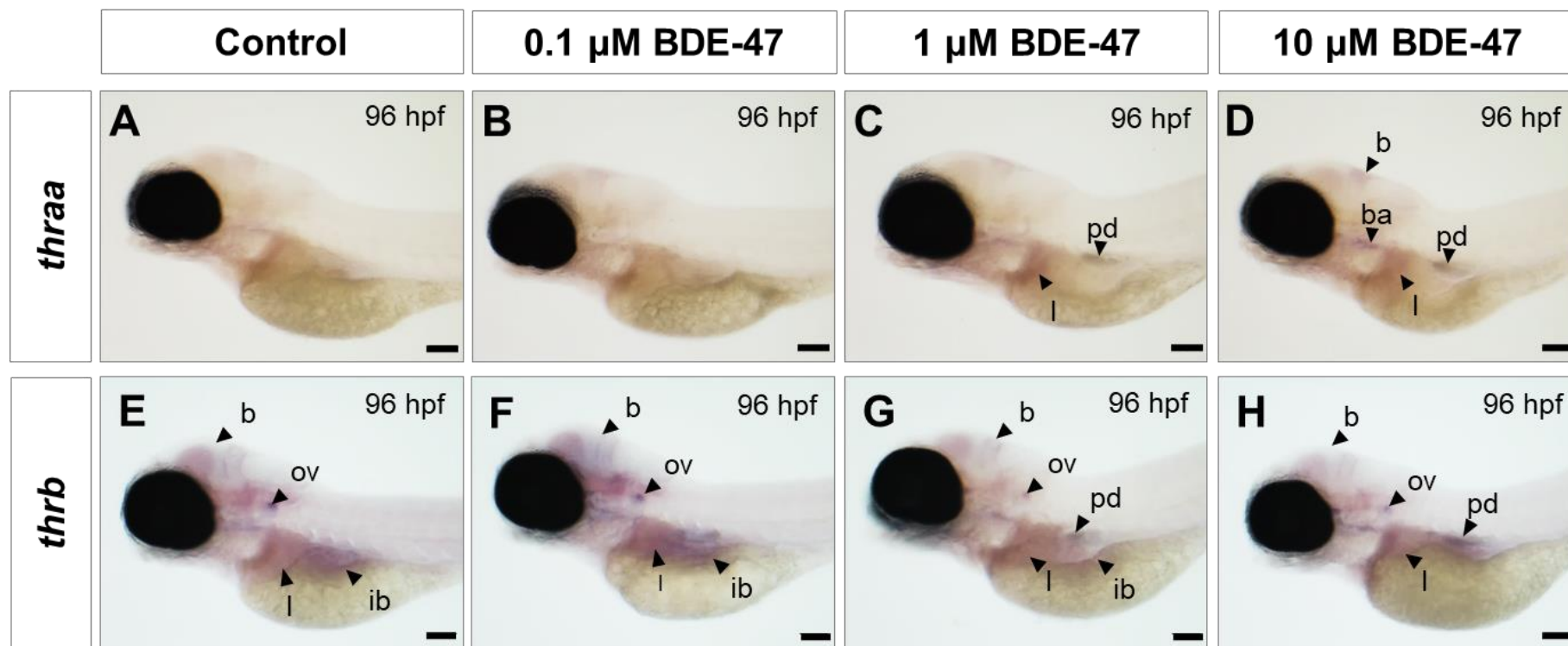
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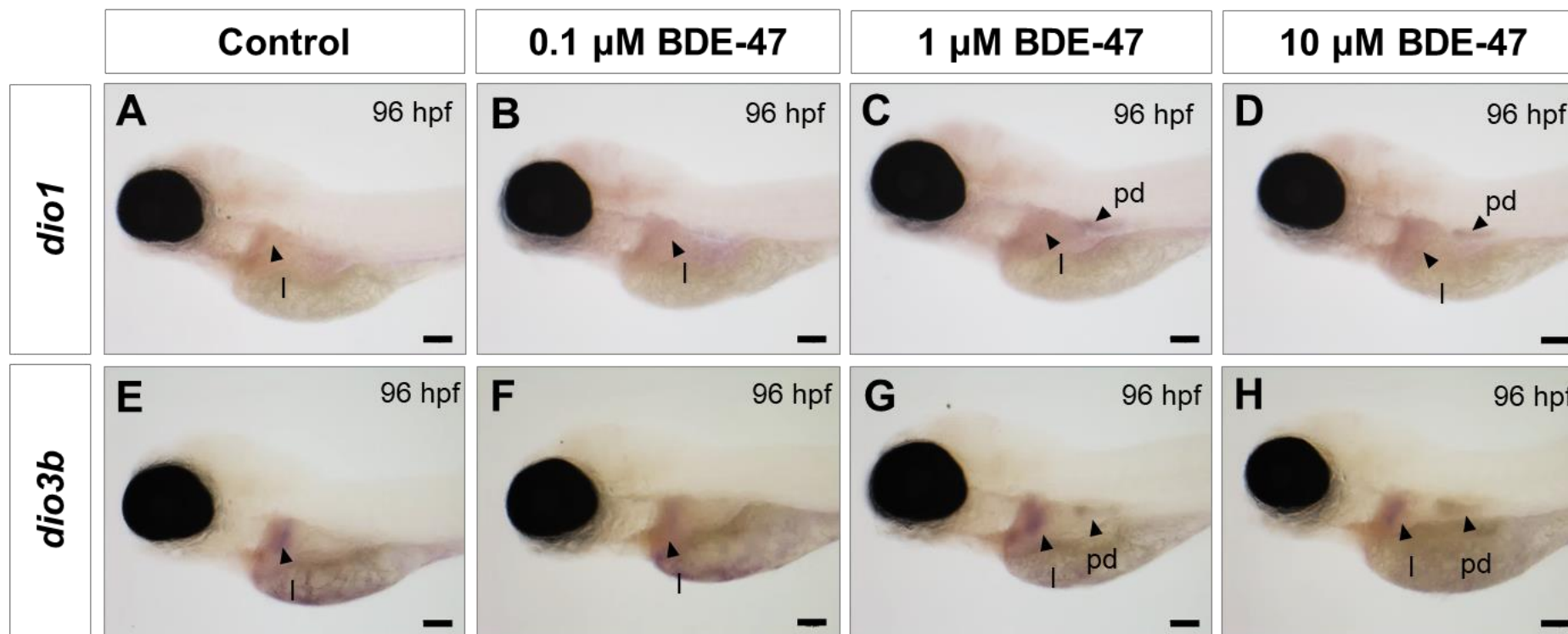
949 Figure 8. Transcript profile of genes in the HPT axis of zebrafish larvae exposed to BDE-47. Changes in (A) thyroid receptor alpha
 950 (*thraa*), (B) thyroid receptor beta (*thrb*), (C) deiodinase type I (*dio1*), (D) deiodinase type II (*dio2*), (E) deiodinase type III (*dio3b*), (F)
 951 corticotropin-releasing hormone (*crhb*), (G) thyroid-stimulating hormone (*tshb*), (H) transthyretin (*ttr*), (I) paired box 8 (*pax8*) and (J)
 952 uridine diphosphate-glucuronosyltransferase (*ugt1ab*) mRNA levels in whole zebrafish following exposure to BDE-47 (0, 0.1, 1 and
 953 10 µM) for 96 hpf. Transcript profiles were determined using qRT-PCR and the relationship between transcript expression and
 954 TBBPA concentration was assessed using linear mixed models (model output summarised in Table S9). Plotted data are presented
 955 as mean fold changes (normalised against the expression of the control gene *rp18*) ± SEM compared to the corresponding control

956 group. Outliers, identified as described in the text, were excluded from the analysis, resulting in a replication of n= 5-6 samples per
957 treatment group. Significance codes: *p<0.05, **p<0.01, ***p<0.001.



958

959 Figure 9. WISH images of TR mRNA expression patterns in zebrafish larvae following exposure to BDE-47. Representative images
 960 of (A-D) *thraa* and (E-H) *thrb* mRNA expression patterns in 96 hpf zebrafish larvae treated with BDE-47 (0.1, 1 and 10 μM). Lateral
 961 views of whole larvae are shown with anterior positioned to the left. Focal areas of expression are indicated by black arrowheads.
 962 b=brain, ba=branchial arches, pd=pronephric ducts, l=liver, ov=otic vesicle, ib=intestinal bulb. Scale bar=100 μm . A summary of the
 963 variability in mRNA expression in the different zebrafish tissues is provided in Table S10.



964
965 Figure 10. WISH images of deiodinase type I and type III mRNA expression patterns in zebrafish larvae following exposure to BDE-
966 47. Representative images of (A-D) *dio1* and (E-H) *dio3b* mRNA expression patterns in zebrafish larvae at 96 hpf treated with BDE-
967 47 (0.1, 1 and 10 μ M). Lateral views of whole larvae are shown with anterior to the left and focal areas of expression are indicated
968 by black arrowheads. b=brain, pd=pronephric ducts and l=liver. Scale bar=100 μ m. A summary of the variability in mRNA
969 expression in the different zebrafish tissues is provided in Table S11.

Key Changes in Transcript Levels (qRT-PCR)

Key Changes in Tissue Expression Patterns (WISH)

TBBPA

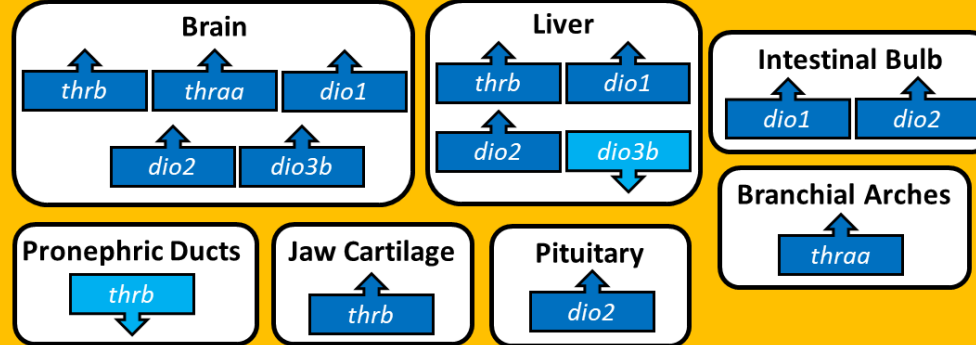
48 hpf embryos



96 hpf larvae

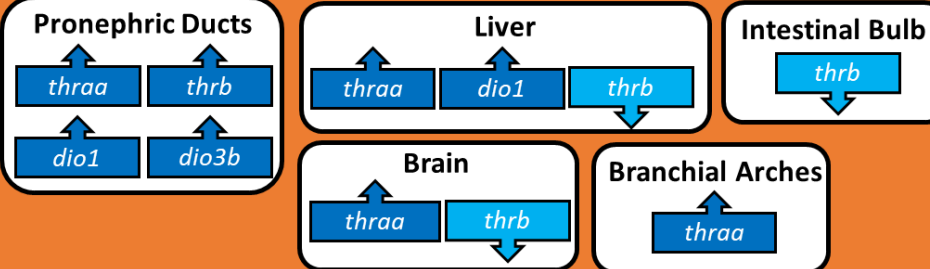
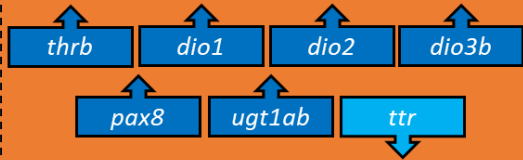


120 hpf larvae



BDE-47

96 hpf larvae



971 Figure 11. Overview of the key changes in the transcript level (measured via qRT-PCR) and tissue expression pattern (measured
972 via WISH) of HPT axis target genes in zebrafish embryo-larvae exposed to TBBPA and BDE-47. Dark blue arrow blocks indicate
973 transcript levels were up-regulated and light blue arrow blocks indicate transcript levels were down-regulated.

974 **Tables**

975 Table 1: LC_x and EC_x values (μM) with corresponding 95% confidence intervals for
 976 zebrafish embryo-larvae following 96-h exposures to TBBPA and BDE-47.

	TBBPA	BDE-47
LC₁₀	0.6 (0.5-0.7)	NC
LC₅₀	0.9 (0.8-1)	NC
LC₉₀	1.3 (1.2-1.5)	NC
EC₁₀	0.4 (0.3-0.5)	0.03 (0-0.23)
EC₅₀	0.7 (0.6-0.9)	38.2 (10.4-475.9)
EC₉₀	1.25 (0.9-2.2)	NC

977