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## Decreased axonal density and altered expression profiles of axonal guidance genes underlying lead (Pb) neurodevelopmental toxicity at early embryonic stages in the zebrafish

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### Abstract

Previous studies have reported that environmental lead (Pb) exposure can result in neurological alterations in children leading to reduced IQ, attention deficit hyperactivity disorder, and diminished reading and learning abilities. However, the specific alterations in neurodevelopmental morphology and the underlying genetic mechanisms of these alterations have not yet been thoroughly defined. To investigate alterations in neurologic morphology and test the hypothesis that developmental Pb neurotoxicity is partially mediated through alterations in neuronal growth and transport function of axons, the changes of specific axon tracts in the embryonic zebrafish brain were observed with anti-acetylated  $\alpha$ -tubulin staining at several developmental time points through 36 hours post fertilization (hpf). In addition, the role of a subset of axonogenesis-related genes including *shha*, *epha4b*, *netrin1b*, *netrin2*, and *noi* were investigated with real-time quantitative PCR (qPCR). Pb treatment resulted in decreased axonal density at 18, 20, and 24 hpf for specific axon tracts in the midbrain and forebrain. These observations corresponded to an observed down-regulation of *shha* and *epha4b* at 14 and 16 hpf, respectively. The axonal density in Pb exposed individuals at later stages (30 and 36 hpf) was not significantly different from controls. An overexpression of *netrin2* at these two developmental stages suggests a novel role for this gene in regulating axonal density specific to Pb neurotoxicity. Although no significant differences in axonal density was observed in the two later developmental stages, further studies are needed to determine if the morphologic alterations observed at the earlier stages will have lasting functional impacts.

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#### Conflict of interest statement

The authors declare that there are no conflicts of interest.

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## Keywords

lead; Pb; neurodevelopment; axon tract; embryo; zebrafish; neurotoxicity

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## 1. Introduction

Lead (Pb) exposure can result in a myriad of adverse health effects that are dependent on dose. The central nervous system is the most sensitive target of Pb toxicity and epidemiological studies have reported several neurological alterations in children including reduced IQ, attention deficit hyperactivity disorder, diminished reading and learning abilities, hearing loss and other health and behavioral disruptions [2,9,24,34,42]. Researchers have found that even low-dose Pb exposure (below 10µg/dl, the current action level of CDC) can induce neurodevelopmental alterations in children [5,6,16,33]. However, the specific alterations in neurodevelopmental morphology in the brain and the genetic mechanisms underlying these alterations are not yet completely understood.

Multiple mechanisms have been implicated in Pb neurotoxicity including cell type-specific responses and various molecular targets involving cell signaling and other functions [32]. One of the suggested mechanisms specific to developmental Pb neurotoxicity is the impairment of neuronal morphogenesis. In a study investigating the effects of Pb in the developing retinotectal system of tadpoles, all tested Pb treatments (PbCl<sub>2</sub>, ranging from 10<sup>-10</sup> to 10<sup>-6</sup>M, surgically implanted over optic tectum) were observed to significantly reduce the area and branchtip number of retinal ganglion cell axon arborizations within the optic tectum after 6 weeks treatment [10]. In addition, Pb exposure (0.1 mM or 2 mM Pb acetate administered in drinking water) has been reported to result in neuritic beading in auditory axons of postnatal day 21 Balb/c mice and *in vitro* in differentiated SH-SY5Y cells [19]. These two findings suggest that developmental Pb exposure may impair axonal growth and transport function, but how these alterations progress over the developmental time course and the underlying genetic mechanisms are not well defined.

In this study to further investigate the morphologic and genetic alterations associated with developmental Pb neurotoxicity, the zebrafish vertebrate model system was used. The zebrafish model system provides an informative tool to investigate neurogenesis from the one cell stage to fully-developed organs and systems. A strength of the zebrafish vertebrate model system utilized in this study is the rapid *ex utero* embryonic development to elucidate brain morphological alterations associated with developmental Pb exposure. In a previous toxicogenomics study in our laboratory [37], zebrafish embryos exposed to 100 parts per billion (ppb) Pb from ~2 hours post fertilization (hpf) through 72 hpf resulted in gene alterations associated with neurodevelopment. This gene list included numerous genes involved in neuronal ontogenesis and synapse formation. Thus in this current study, the morphological alterations in axon tracts associated with the Pb exposure were investigated to link genetic and morphologic data at earlier developmental time points.

## 2. Methods

### 2.1. Zebrafish husbandry

Zebrafish used in this study were of the wild-type AB strain and were housed on a 14:10 hour light:dark cycle in a Z-Mod System (Aquatic Habitats, Apopka, FL) at 28°C. Fish were maintained and bred according to protocols approved by the university's Institutional Animal Care and Use Committee with all fish treated humanely and with regard to alleviation of suffering [44]. Adults were bred in spawning cages to obtain staged embryos following established protocols [44]. Based on our previous study [37], a 100 ppb Pb

treatment (Pb acetate, Sigma, St. Louis, MO) or a control treatment (filtered fish water) were initiated at ~2 hpf and embryos were collected at numerous developmental stages from 12 to 36 hpf for experimental procedures.

## 2.2 Pb concentration of tissue and treatment water determined by inductively coupled plasma - mass spectrometry (ICP-MS)

Zebrafish embryos were collected, washed, weighed, and digested for Pb analysis using an Elan DRC-e ICP-MS (PerkinElmer Sciex, Shelton, CT) at multiple developmental time points from 12 to 36 hpf. For the digestion, whole zebrafish embryos were submerged in 50 to 200  $\mu$ l 50% nitric acid (volume dependent on weight/developmental stage) and digested overnight at 65°C in a water bath. Samples were diluted with deionized water to reach a final 1% HNO<sub>3</sub> concentration and analyzed by ICP-MS following standard protocols [7]. Thallium (ICP-081, ULTRA Scientific, North Kingstown, RI) was used as an internal standard and Pb standard curves were performed simultaneously. The concentration of Pb in zebrafish tissue was calculated as ng/g tissue. In addition, Pb stock solutions and treatment water were collected and analyzed by ICP-MS following a similar procedure. To observe Pb distribution in the chorion and fish body, embryos at the 24 hpf stage were dissected using a tungsten needle and chorion and fish body processed separately.

## 2.3 Immunofluorescence staining

Embryos were collected at multiple developmental time points (18, 20, 24, 30 and 36 hpf) following Pb treatment, washed, and dechorionated. Embryos were fixed overnight with 4% paraformaldehyde (PFA) at 4°C. The eyes and skin in the brain region were removed and embryos were blocked overnight in blocking buffer (PBS, 1% Triton, 10% FBS, 0.5% BSA, 1% DMSO). Embryos were then incubated in 1:1000 mouse monoclonal anti-acetylated  $\alpha$ -tubulin antibody (Catalog No.T6793; Sigma-Aldrich, St. Louis, MO) overnight at 4°C, washed four times with PBS, 1% Triton, and incubated in 1:1000 Alexa Fluor® 546 rabbit anti-mouse IgG (H+L) (Catalog No.A11060; Invitrogen, Carlsbad, CA) overnight at 4°C. Whole mount embryos were laterally oriented and scanned in serial 2  $\mu$ m optical sections to observe the axon tracts using a laser scanning confocal microscope (Nikon, Japan). Z-stack images were merged using ImageJ 1.44p (NIH, Bethesda, MD). After background subtraction, the fluorescence density (in arbitrary unit, AU) of the region of interest was measured using Image-pro plus 6.0 (Media Cybernetics, Silver Springs, MD).

## 2.4 Real-time quantitative PCR

Embryos were collected at multiple developmental time points from 14 to 36 hpf following Pb treatment. RNA was isolated and cDNA was synthesized as described in Peterson and Freeman [36]. qPCR analysis was performed on a Stratagene MX3000P (Agilent, La Jolla, CA) using the iQ SYBR green kit (Biorad, Hercules, CA) following similar methods as described in Peterson et al. [37]. The cycling parameters included a 3 minute incubation phase at 95°C, 40 cycles of 95°C for 10 seconds, 60°C for 30 seconds and 72°C for 30 seconds with a dissociation curve to check for non-specific amplification. Efficiency and specificity were checked with melting and dilution curve analysis and no-template controls. Quantitative copies were calculated by standard curve and individual gene expression was normalized to *gapdh* (ratio of gene/*gapdh*). Details of the primers are listed in Table 1.

## 2.5 Statistical analysis

The quantitative data are expressed as mean  $\pm$  SD. For the comparison of Pb concentration in tissue throughout development, data were analyzed with a two-way ANOVA and a post hoc least significant difference (LSD) test. A Student's *t*-test was used to compare Pb concentration in chorion and embryo within treatment. Anti-acetylated  $\alpha$ -tubulin staining

data were analyzed by time point with a Student's *t*-test. The gene expression data was analyzed by gene at each time point with a Student's *t*-test. The level accepted for statistical significance in all cases was  $p < 0.05$ .

### 3. Results

#### 3.1 Time course absorption of Pb in zebrafish tissue

To confirm Pb exposure and estimate the actual dose of Pb in zebrafish tissue, ICP-MS was employed to analyze the Pb concentration of zebrafish tissue in a series of developmental time points (12, 14, 16, 18, 20, 24, 30 and 36 hpf). Compared to the control group at each time point, Pb treated embryos showed an obvious increase of Pb accumulation in tissue ( $p < 0.0001$ , for developmental time point, Pb treatment, and the interaction between time point and Pb treatment, Fig. 1A). The values at all time points ( $\geq 404.39 \pm 57.49$  ng/g) was  $>6$ -fold higher than the measured Pb concentration in the original treatment solution ( $65.64 \pm 1.45$  ppb). From 12 to 30 hpf, Pb concentrations increased from  $404.39 \pm 57.49$  to  $778.11 \pm 23.27$  ng/g, and then decreased slightly at 36 hpf ( $635.25 \pm 42.14$  ng/g). The comparative data among the Pb concentration in the original treatment water and the increasing Pb concentration in the embryos indicate Pb accumulates in zebrafish tissue. At 24 hpf, bodies of Pb treated embryos contained a much higher amount of Pb than the chorions (body:  $184.94 \pm 26.30$  ng/g; chorion:  $32.93 \pm 8.09$  ng/g,  $p = 0.001$ , Fig. 1B). The average concentration of Pb in the body and chorion of the treatment group at 24 hpf is less than the average amount of Pb measured in the complete embryo at this same time point (Fig. 1A). This unaccounted Pb is suspected to be in the chorionic fluid and thus lost when fish body and chorion were separated for this measurement. No obvious differences for Pb distribution were found between fish bodies and chorions in the control group ( $p = 0.549$ ).

#### 3.2 Changes of specific axon tracts in forebrain and midbrain at 18, 20, 24, 30, and 36 hpf

Anti-acetylated  $\alpha$ -tubulin antibody is commonly used to stain early axon tracts. Specific axon tracts including the anterior commissure (AC), post-optic commissure (POC), supra-optic tract (SOT), tract of post-optic commissure (TPOC), tract of the posterior commissure (TPC) and dorsal-ventral diencephalic tract (DVDT) in the forebrain and midbrain in lateral view were observed in this study (Fig. 2). In the control group, the AC and POC were slightly stained at 18 hpf (Fig. 2A). These two commissures extended longer and showed brighter staining at 20 hpf, while the TPOC also appeared at this time point (Fig. 2D). At 24 hpf, considerably more axon tracts, including the SOT, TPC, DVDT, and epiphyseal cluster (epi) had formed (Fig. 2G). Further development of axon tracts was observed at 30 hpf (Fig. 2J) and 36 hpf (Fig. 2M). Density data showed that 100 ppb Pb treatment significantly decreased the density of axon tracts at 18, 20 and 24 hpf ( $p = 0.030$ ,  $0.039$  and  $0.029$ , respectively, Fig. 2B, 2C, 2E, 2F, 2H, and 2I), but not at 30 and 36 hpf ( $p = 0.729$  and  $0.996$ , respectively, Fig. 2K, 2L, 2N, and 2O).

#### 3.3 Observation of axon tracts and neurons in hindbrain at 20 and 24 hpf

Hindbrain axon tracts were also observed in this study as anti-acetylated  $\alpha$ -tubulin stains Mauthner neurons and other axon tracts including the dorsal longitudinal fasciculus (DLF) and ventral longitudinal fasciculus (VLF) in the zebrafish hindbrain. No significant difference between control and Pb treatment were observed at either time point ( $p = 0.516$  and  $0.374$ , respectively).

#### 3.4 Changes of axonogenesis genes in Pb neurotoxicity at 14, 16, 18, 20, 24, 30 and 36 hpf

The expression of a subset of genes involved in axonogenesis [*sonic hedgehog a (shha)*, *ephrin type-A receptor 4b (epha4b)*, *netrin (netrin1b, netrin2)*, and *no-isthmus (noi)*] were

evaluated by qPCR analysis at multiple developmental time points from 14 to 36 hpf. In Pb treated embryos the *shha* transcript level was significantly decreased at 14 hpf (decreased 30%,  $p=0.018$ , compared to control, Fig. 3A). At 16 hpf, a significant decrease in transcript level was observed for *epha4b* (decreased 24%,  $p=0.043$ , compared to control, Fig. 3B). The expression of *netrin2* was significantly upregulated at 30 and 36 hpf ( $p=0.022$  and  $0.026$ , respectively, compared to control at each time point, Fig. 3C and 3D). Significant differences in expression of these genes were not observed at the intermediate developmental time points analyzed (i.e., 18, 20, or 24 hpf). Moreover, no significant difference in transcript level was observed for *netrin1b* or *noi* at any of the developmental time points assessed in this study.

## 4. Discussion

The early development of embryonic zebrafish primarily occurs in a chorionic membrane. After fertilization, embryos grow rapidly through several stages with differences in morphology denoting progression through each developmental stage. These developmental stages include the zygote period (0–0.75 hpf), cleavage period (0.7–2.2 hpf), blastula period (2.25–5.25 hpf), gastrula period (5.25–10 hpf), segmentation (10–24 hpf), pharyngula period (24–48 hpf) and hatching period (48–72 h) [21]. At 16 hpf, the first post-mitotic neurons appear in the forebrain and midbrain in three distinct clusters, including the dorsorostral cluster (drc), ventrorostral cluster (vrc) and ventrocaudal cluster (vcc) [38]. These first primary neurons extend long axons to pioneer pathways and make small local connections [18,20]. Another cluster, epi, can be found from 18 hpf [38]. By 24 hpf, these clusters have extended axon tract commissures including AC, POC, SOT, TPOC, TPC, and DVDT forming a simple scaffold of axon tracts (Fig. 2G). Specific and simple axon tracts can be studied in embryonic zebrafish and are a good model to study the axonogenesis process. By using anti-acetylated  $\alpha$ -tubulin staining, our results indicate that 100 ppb Pb treatment decreases axon density at early embryonic stages (18, 20 and 24 hpf), but this difference is no longer observed at 30 and 36 hpf.

To investigate the genetic pathways associated with the decreased axonal density associated with developmental Pb neurotoxicity observed at 18, 20 and 24 hpf in this study, the expression level of a subset of genes expressed in the regions adjacent to the AC and POC and involved in axonal guidance and migration were evaluated [45]. These genetic targets included *shha* [1,4], *epha4b* [29,46], *netrin1b* [30,41], *netrin2* [30,41], and *noi* [3,29]. *Shh* is a morphogen that plays an important role in the patterning of multiple organs, including the nervous system [17,26]. *Shh* can act as an axonal guidance cue on spinal cord commissural axons and retinal ganglion cell (RGC) axons [8,22]. In zebrafish, a previous study reported that *shh* is expressed at 6 hpf (embryonic shield), 11 hpf (notochord, floor plate, and ventral forebrain), 48 hpf and 3 weeks (whole nervous system) [13]. In an additional study, a decreased *shh* transcript level was found at 5 and 6 days post fertilization (dpf) compared to that at 1 dpf [14]. In the present study, Pb exposure down-regulated *shha* levels at 14 hpf. *Shh* is reported to collaborate with Netrin1 to guide commissural axon extension [39] and overexpression of *shh* impairs *netrin1b* expression in the ventral neural tube [41]. Although *netrin1b* expression was not altered in Pb treated embryos in this study, decreased *shha* transcript levels may relate to limited growth of axon tracts observed at 18, 20 and 24 hpf.

*Epha4* is a member of the ephrin receptor subfamily of protein-tyrosine kinases and is involved in mediating developmental events, particularly in the nervous system [15]. In mouse, *Epha4* has been implicated in the restriction of axons from entering the anterior branch of the AC [11]. As evidenced by an *Epha4* mutant mouse, knocking out *Epha4* causes failure of axon guidance of corticospinal neurons and a defect in formation of the AC [12]. Kullander et al. [23] demonstrated that *Epha4* has both kinase-dependent and kinase-

independent functions in the formation of major axon tracts, including the corticospinal tract and the AC. In zebrafish, *epha4* has been related to the segmental restriction of gene expression in the hindbrain and is required in patterning the developing zebrafish forebrain (diencephalic territory and structures) [46]. In our study, the transcript level of *epha4b* following Pb treatment was significantly decreased at 16 hpf indicating *epha4b* may also play a role in the decreased axonal density observed at 18, 20 and 24 hpf.

The netrin family of laminin-related molecules is reported to be involved in axon guidance and cell migration during development [27,41]. In zebrafish, two *netrin* homologues, *netrin1b* and *netrin2*, are mainly expressed in the floor plate and the anterior ventral neural tube (including ventral to the AC) [29,30,35,41]. It has been demonstrated that netrins act to stimulate axonal outgrowth in addition to attracting and repelling growth cones in zebrafish embryos [25]. Although there is no direct evidence of *netrin* knockout defects in zebrafish, Serafini et al. [40] reported that *Netrin1* deficient mice exhibit defects in spinal commissural axon projections and in several forebrain commissures. In the present study, Pb treatment resulted in a significant increase in expression of *netrin2*, but not *netrin1b* at 30 and 36 hpf. Considering the role of netrins on stimulating axonal outgrowth, our results indicate that *netrin2* may play a role in regulating axonal growth in the response mechanism to Pb neurodevelopmental toxicity.

*noi* is a zebrafish gene that belongs to the Pax family of transcriptional regulators which is important in early development for differentiation of multiple tissues, including nerve development [43,47]. In the zebrafish, *noi* has important functions at multiple stages of embryogenesis [28,29,31] and is expressed in restricted domains of the developing zebrafish central nervous system, such as midline cells dorsal/rostral to the POC [3,29]. *noi* normally discourages axonal extension into the *noi* expressed domain [29,45]. Early decussating axon tracts are prevented from entering the territory that expresses *noi* protein in wild-type embryos, and *noi* mutant zebrafish embryos show severe defects in the rostral forebrain. These defects include axon tracts that are no longer tightly fasciculated in POC and many axon tracts that have migrated inappropriately [29,45]. Based on our data, *noi* expression was not significantly changed due to Pb treatment at any of the developmental time points tested in this study.

In summary, our results showed that Pb exposure (100 ppb) decreased the density of axon tracts in midbrain and forebrain at 18, 20 and 24 hpf, corresponding to the down-regulation of *shha* and *epha4b* genes at 14 hpf and 16 hpf, respectively. Alternatively, no significant difference in axon density was observed at 30 and 36 hpf. The increased expression level of *netrin2* at these two developmental stages, indicates *netrin2* may play a role mediating axon density alterations caused by Pb toxicity. Although no significant difference in axon density was detected at the later stages in this study, further studies need to be conducted to determine if lasting functional impacts may result from the lag in axonal development.

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## Abbreviations

<b>AC</b>	anterior commissure
<b>dpf</b>	days post fertilization

<b>DLF</b>	dorsal longitudinal fasciculus
<b>drc</b>	dorsorostral cluster
<b>DVDT</b>	dorsal-ventral diencephalic tract
<b><i>epha4b</i></b>	<i>ephrin type-A receptor 4b</i>
<b>epi</b>	epiphyseal cluster
<b>hpf</b>	hours post fertilization
<b>ICP-MS</b>	inductively coupled plasma - mass spectrometry
<b><i>noi</i></b>	<i>no-isthmus</i>
<b>Pb</b>	lead
<b>PFA</b>	paraformaldehyde
<b>POC</b>	post-optic commissure
<b>ppb</b>	part per billion
<b>qPCR</b>	real-time quantitative PCR
<b>RGC</b>	retinal ganglion cell
<b><i>shha</i></b>	<i>sonic hedgehog a</i>
<b>SOT</b>	supra-optic tract
<b>TPC</b>	tract of the posterior commissure
<b>TPOC</b>	tract of post-optic commissure
<b>vcc</b>	ventrocaudal cluster
<b>VLF</b>	ventral longitudinal fasciculus
<b>vrc</b>	ventrorostral cluster

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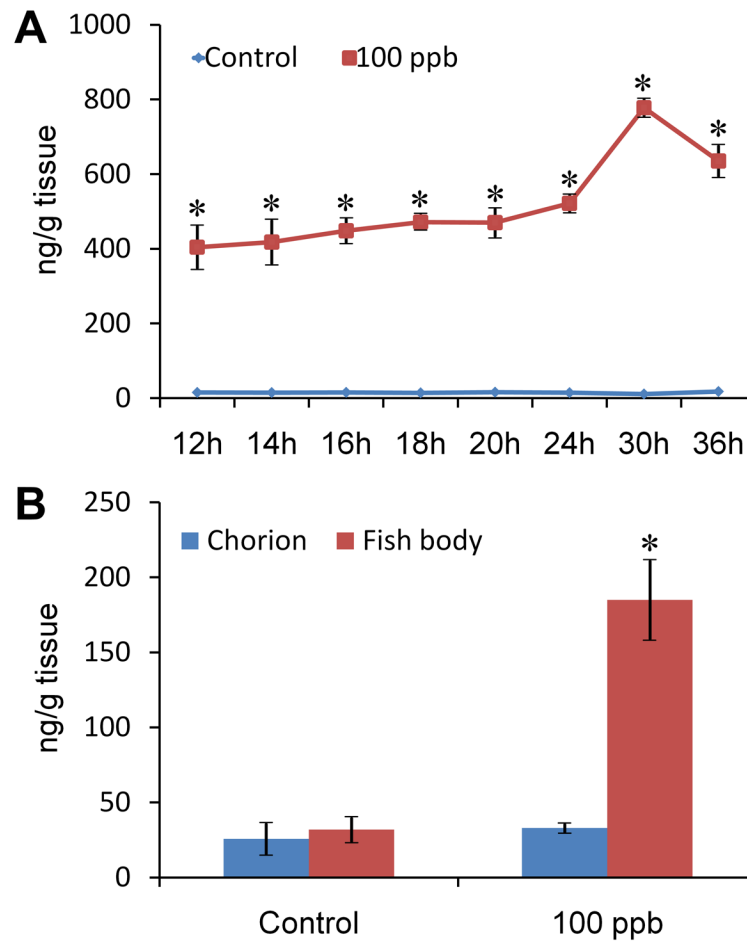


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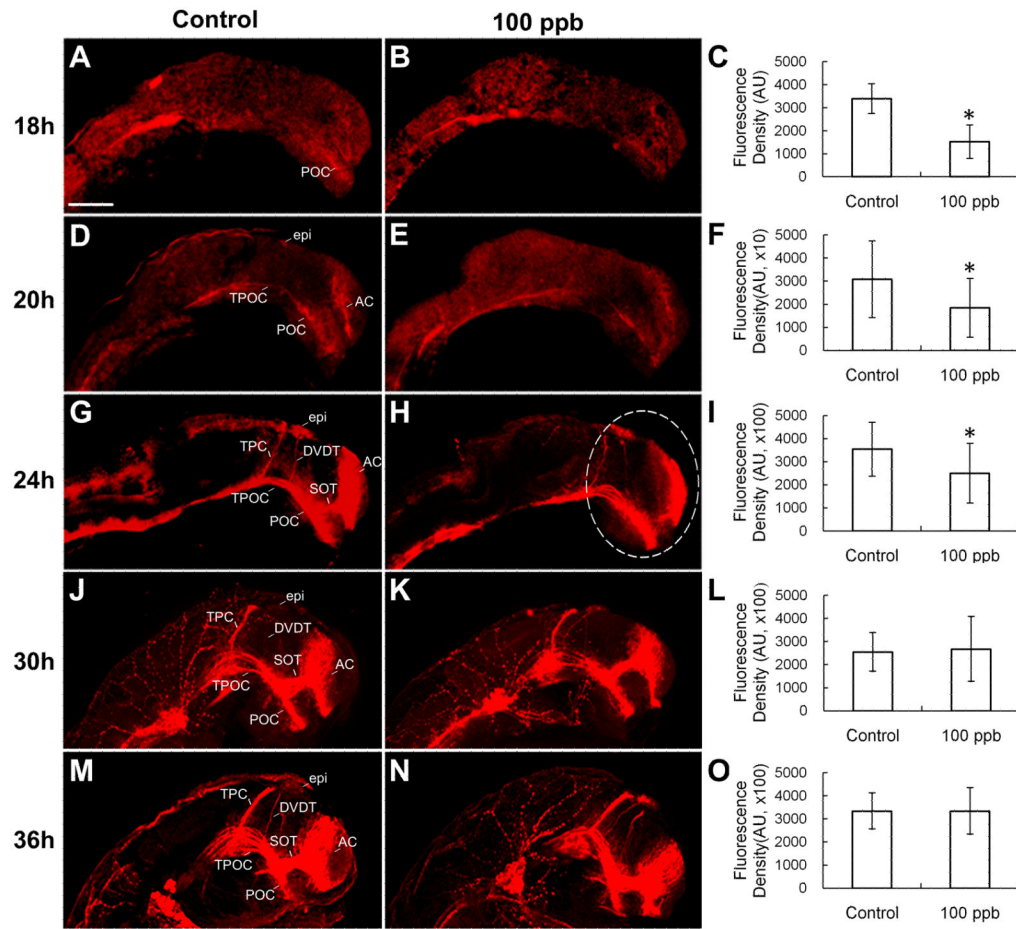
### Highlights

- Decreased axonal density at 18, 20, and 24 hpf in the midbrain and forebrain.
- Decreased axonal density corresponded to down-regulation of *shha* and *epha4b*.
- Axonal density in Pb exposed individuals at later stages was not altered.
- Overexpression of *netrin2* observed at these later developmental stages.



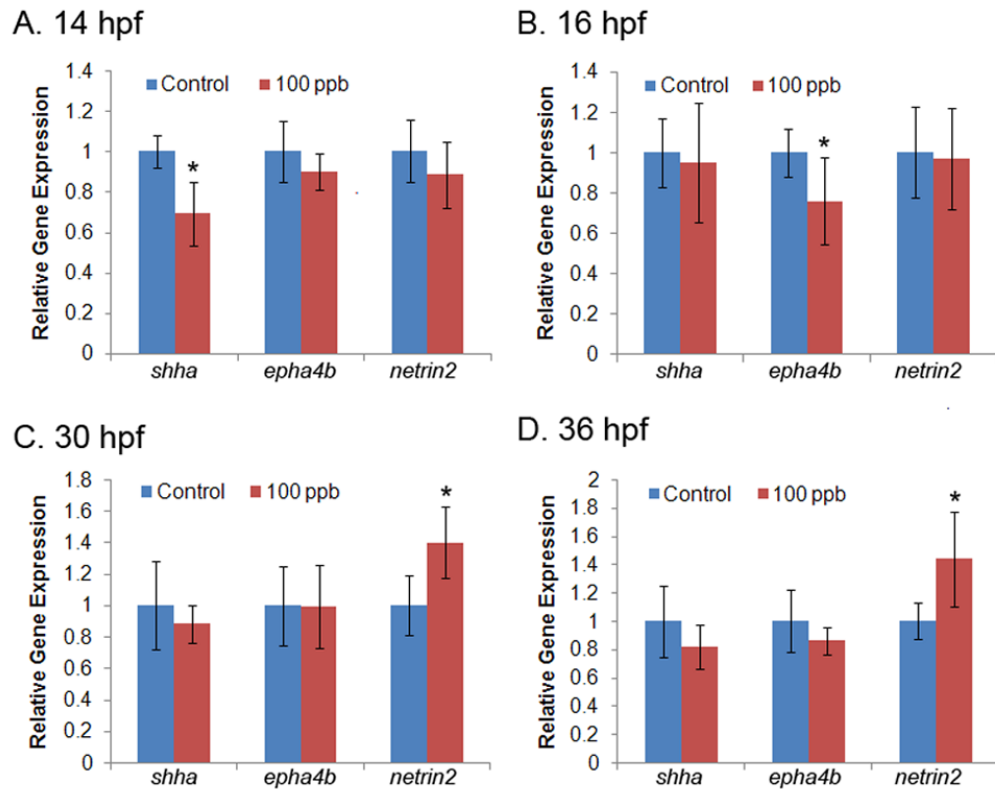
**Figure 1. Pb accumulation in zebrafish tissue during embryonic exposure**

A: Pb concentration of zebrafish tissue from 12 to 36 hpf (with chorion).  $*p < 0.0001$ , compared with control by time point; B: Pb distribution in chorion and fish body at 24 hpf.  $*p < 0.05$ , fish body compared with chorion within treatment. For each concentration each time point,  $n = 3$ .



**Figure 2. Effect of Pb on density of axon tracts of embryonic zebrafish brain in lateral view at 18, 20, 24, 30 and 36 hpf**

Anti-acetylated  $\alpha$ -tubulin staining clearly showed axon tracts at 18, 20, 24, 30 and 36 hpf. A, B and C: 18 hpf (n=15–18); D, E and F: 20 hpf (n=14–16); G, H and I: 24 hpf (n=14–16), J, K and L: 30 hpf (n=21–22); M, N and O: 36 hpf (n=13–14). Density data of stained region of interest (white dash line marked) were measured using Image-pro plus (C, F, I, L and O). The units for F are 10 times greater than C while the units for I, L and O are 100 times greater (denoted as  $\times 10$  and  $\times 100$ ). AC, anterior commissure; POC, post-optic commissure; SOT, supra-optic tract; TPOC, tract of post-optic commissure; TPC, tract of the posterior commissure; DVDT, dorsal-ventral diencephalic tract; epi, epiphyseal cluster. Scale bar=100  $\mu$ m. \* $p$ <0.05.



**Figure 3. Altered transcript level of axonogenesis genes following Pb treatment at different developmental time points**

Transcript levels for five genes of interest were measured at seven developmental time points with qPCR. Data were quantified with standard curves using *gapdh* as a reference gene and normalized to control treatment. Transcript levels of *shha* and *epha4b* were significantly altered at 14 hpf (A) and 16 hpf (B), respectively. *Netrin2* was significantly altered at 30 and 36 hpf (C and D). No differences were detected in gene expression at 18, 20 or 24 hpf for *shha*, *epha4b* or *netrin2* (data not shown) or in *netrin1b* or *noi* at any time point analyzed (data not shown). n=6 (50 pooled embryos for each replicate per time point per group). \* $p < 0.05$ , Pb treatment compared to control

**Table 1**

Primer sequences for axonogenesis genes for qPCR analysis

Gene name	Primer sequence	GenBank ID	Amplicon size (bp)
<i>shha</i>	Forward: AGACCGAGACTCCACGACGC Reverse: TGCAGTCACTGGTGCGAACG	NM_131063.1	265
<i>epha4b</i>	Forward: TGCCGAGTGCGAAACCAGT Reverse: TCCCTCCCTCACACCGAGTC	NM_153658.1	219
<i>netrin1b</i>	Forward: TCACCTCGATGGTGTCGG Reverse: GCAGCGTCTAGGGTTCCCGT	NM_130998.1	110
<i>netrin2</i>	Forward: TATAGGGACATGGGCAAAG Reverse: AGCCTTAGCACAGCGGTTA	NM_001037415.1	160
<i>noi</i>	Forward: GGGACTGAGAGACTCGCC Reverse: ACGAAGACACGAGGCTGGTCA	NM_131184.2	291