

1 The Utility of Zebrafish to Study the Mechanisms by which Ethanol Affects
2 Social Behavior and Anxiety During Early Brain Development

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27 **Abstract**

28 Exposure to moderate levels of ethanol during brain development has a number of
29 effects on social behavior but the molecular mechanisms that mediate this are not well
30 understood. Gaining a better understanding of these factors may help to develop
31 therapeutic interventions in the future. Zebrafish offer a potentially useful model in this
32 regard. Here, we introduce a zebrafish model of moderate prenatal ethanol exposure.
33 Embryos were exposed to 20mM ethanol for seven days (48hpf – 9dpf) and tested as
34 adults for individual social behavior and shoaling. We also tested their basal anxiety
35 with the novel tank diving test. We found that the ethanol-exposed fish displayed
36 reductions in social approach and shoaling, and an increase in anxiety in the novel tank
37 test. These behavioral differences corresponded to differences in *hrt1aa*, *slc6a4* and *oxtr*
38 expression. Namely, acute ethanol caused a spike in *oxtr* and *ht1aa* mRNA expression,
39 which was followed by down-regulation at 7dpf, and an up-regulation in *slc6a4* at
40 72hpf. This study confirms the utility of zebrafish as a model system for studying the
41 molecular basis of developmental ethanol exposure. Furthermore, it proposes a putative
42 developmental mechanism characterized by ethanol-induced OXT inhibition leading to
43 suppression of 5-HT and up-regulation of 5-HT_{1A}, which leads, in turn, to possible
44 homeostatic up-regulation of 5-HTT at 72hpf and subsequent imbalance of the 5-HT
45 system.

46 **Keywords:** social; anxiety; zebrafish; moderate prenatal ethanol; serotonin; oxytocin;
47 vasopressin

48 **1. Introduction**

49 Maternal alcohol consumption during pregnancy results in a range of effects on the
50 developing fetus, collectively referred to as Fetal Alcohol Spectrum Disorders (FASD;
51 Paintner et al., 2012). FASD are characterized by a range of teratogenic and
52 psychological defects, and represent the leading non-hereditary cause of mental
53 retardation, with the prevalence estimated at between 2 and 5% of the population of
54 the USA and western Europe (May et al., 2009). At the extreme end of the spectrum,
55 when mothers drink heavily during pregnancy, fetal alcohol syndrome (FAS) typically
56 results in gross skeletal and craniofacial abnormalities, and severe CNS dysfunction
57 (Hanson et al., 1976). Moderate alcohol consumption (e.g., equivalent to 1-2 drinks/day,
58 average BAC ~0.01-0.04g/dL Valenzuela et al., 2012) is associated with a range of more
59 subtle cognitive and behavioral defects, including aggression and depression (Sood et
60 al., 2001), vulnerability to stress (Hellemans et al., 2008), impulsivity and inattention
61 (Streissguth et al., 1989;Suess et al., 1997), and poor scholastic performance (Olson et
62 al., 1998).

63 Social behavior in offspring exposed to alcohol during gestation (PNE) has been
64 extensively studied, with deficits ranging from problems forming social relationships to
65 severe antisocial behavior (Thomas et al., 1998;Roebuck et al., 1999;Kelly et al.,
66 2000;McGee et al., 2008;Keil et al., 2010;Rasmussen et al., 2011). The heterogeneous
67 nature of social relationships in humans is such that social deficits observed in PNE
68 children are likely to be the result of numerous additive or interactive factors, ranging
69 from insecure attachment styles in both the offspring (O'Connor et al., 2002) and
70 caregivers (Swanson et al., 2000), to deficits in perception of social cues or the ability to
71 sustain functional relationships (Kelly et al., 2000).

72 Preclinical models have typically used animals to gain insight into neurobiological
73 processes underlying the social deficits associated with PNE. For example, recent work
74 from Hamilton et al. (2010) demonstrated that the adult offspring of rats exposed to
75 moderate levels of ethanol (PNE rats) during pregnancy exhibited changes in some
76 aspects of social behavior (social investigation and aggression), especially in males.
77 These changes appeared to be related to alterations in experience-dependent structural
78 plasticity in frontal cortical regions (agranular insular cortex [AID], the rat homologue

79 of the primate orbital-prefrontal cortex). These data strongly suggest that structural
80 and synaptic plasticity, particularly in brain regions associated with social behavior
81 (e.g., areas of the neocortex), is affected by moderate PNE. The cellular and molecular
82 factors that underpin and modulate this, however, are less well understood.

83 Zebrafish are a widely used model system in developmental neuroscience. This is due
84 predominantly to their small size, prolific breeding and unparalleled genetic tractability
85 (Guo, 2004). Zebrafish offer a potentially excellent model for studying the molecular
86 processes resulting from PNE because, a) embryos develop *ex utero* meaning that very
87 precise volumes of ethanol can be added to the embryo medium, and b) the embryos are
88 completely transparent, facilitating real-time visual inspection of developing cells.
89 Zebrafish are also a social (shoaling) species, and provide a potentially excellent model
90 for studying the social aspects of PNE (Fernandes and Gerlai, 2009; Buske and Gerlai,
91 2011; Pham et al., 2012; Oliveira, 2013). Previous work by Buske and Gerlai (2011) and
92 Fernandes and Gerlai (2009) showed that brief (1-2hr) exposure to high concentrations
93 of ethanol (50mM [0.25% v/v], 100mM [0.5% v/v], 200mM [1% v/v]) alters adult social
94 behavior (operationalized by nearest neighbor and by proximity to a virtual fish) and
95 reduces levels of 2-(5-Hydroxy-1H-indol-3-yl)acetic acid (5HIAA; a 5-HT metabolite) in the
96 adult brain. An assessment of more moderate levels of ethanol on these aspects of
97 zebrafish behaviour, however, has not been previously carried out. Therefore, here, we
98 examined the effect of moderate developmental ethanol exposure on mRNA expression
99 of genes that code for components of neurotransmitter systems implicated in the
100 control of social behavior, namely serotonin receptor 1a (5-HT_{1A}; *htr1a*, Bell and
101 Hobson, 1993; Strobel et al., 2003), serotonin transporter (5-HTT; *slc6a4* Wendland et
102 al., 2006; Canli and Lesch, 2007) and receptors for the neuropeptides oxytocin and
103 vasopressin (OT; *oxtr* and *AVP*; *avpr* Winslow et al., 1993; Heinrichs et al., 2009).

104

105 **2. Materials and Methods**

106 *2.1. Subjects*

107 Adult Tubingen (mixed male/female) zebrafish were kept in a re-circulating system, on
108 a 14/10-hour light/dark cycle, at 28.5°C within our zebrafish aquarium. Fish were fed

109 with a mixture of flake food, fresh brine shrimp and bloodworm. Adults were bred in
110 house and fry reared according to the above protocols. Larvae from each condition
111 (20mM ethanol and control) were sacrificed at 24hpf (i.e., before ethanol exposure),
112 50hpf (acute ethanol exposure: 2hrs after ethanol added), 72hpf and finally at 7dpf. All
113 animal work was carried following approval from the Queen Mary Research Ethics
114 Committee, and under licence from the Animals (Scientific Procedures) Act 1986. Care
115 was taken to minimize the numbers of animals used in this experiment in accordance
116 with the ARRIVE guidelines (<http://www.nc3rs.org.uk/page.asp?id=1357>). Specifically,
117 we examined data from previous pilot studies and studies with other species to carry
118 out a power calculation and assess the minimum number of animals necessary for the
119 expected effect size with power of 0.8.

120

121 *2.2. Developmental ethanol exposure*

122 Tubingen zebrafish embryos were treated by transferring them into 20mM (0.12% v/v,
123 equating to ~0.04g/dL BAC [see below]) ethanol in aquarium water at 48hpf, the long-
124 pec stage (Kimmel et al., 1995). All embryos were carefully staged before treatment.
125 Prior to adding the ethanol, it was our policy to dechorionate the larvae if they had not
126 hatched (in practice, this was rarely necessary as most embryos had hatched by this
127 point). This developmental stage was chosen as it represents a key stage in brain
128 ontogeny, including the development of monoaminergic neurons (Guo et al., 1999).
129 Previous research in adult zebrafish found brain concentrations of ethanol following
130 chronic exposure typically reach ~80-90% bath concentration in adults (Dlugos and
131 Rabin, 2003; Mathur et al., 2011) and ~30-40% in embryos and larvae (Reimers et al.,
132 2004). The relationship between brain and blood alcohol content is not straightforward,
133 and estimates for the ratio of brain: blood ethanol range from 0.6-1.5 (Moore et al.,
134 1997). Therefore, we estimate that the larvae would have had a blood alcohol
135 concentration (BAC) of ~0.02-0.07g/dL putting the zebrafish model in the moderate
136 prenatal ethanol classification (Valenzuela et al., 2012).

137 Fish water was changed on alternate days and the tanks were cleaned in order to
138 reduce the buildup of yeast and control variation in the ethanol concentration due to
139 evaporation. Embryos were kept in Petri dishes until they were five days old, after

140 which they were transferred into tanks with dimensions 10 x 10 x 20 cm (depth x width
141 x length cm) and a volume of 500ml with airlines. A maximum of 40 fish were kept in
142 each tank and numbers of ethanol and control fish were balanced; dividing equally the
143 quantity of fish receiving ethanol or aquarium water for the controls between the tanks.
144 Feeding of the embryos ZM000 and paramecium commenced at five days. After seven
145 days swimming in fish water containing ethanol (i.e., aged nine days), the fry were
146 transferred back to pure aquarium water. At this stage, we photographed a selection of
147 larvae from each treatment (ethanol and control) and measured their size (analysis of
148 pixel density) to ensure there were no gross morphological differences. At age three
149 weeks the volume of water in the tanks was increased to 1L to provide more space for
150 growth and the ZM000 was replaced with ZM100 and artemia. At age five weeks they
151 were transferred into 7L tanks and cleaning was reduced to once a week. For the qPCR
152 analysis, embryos were removed at 24hpf, 50hpf, 72hpf, and 7dpf (see below for
153 details).

154 *2.3. Stress Reactivity (Tank Diving)*

155 [FIGURE 1 HERE]

156

157 The tank diving task was carried out in 1.5 L trapezoid tanks (15.2 height x 27.9 top x
158 22.5 bottom x 7.1 width cm) filled with aquarium treated water from the main
159 aquarium supply. Prior to tank diving, all fish were pair housed for 2-weeks as
160 previously described (Parker et al., 2012). For housing and transport, fish were placed
161 into individual holding tanks, measuring height x width x length: 10 cm x 11 cm x 20 cm.
162 All fish were transported from the aquarium to the test room the day before testing in
163 order to acclimate them to the test-room conditions. Within the holding tanks were
164 located tank inserts with perforated bases. This allowed the fish to be removed easily
165 for testing, thus controlling for the potentially confounding factor of difficulties netting
166 the fish for testing. The order in which the fish were tested was fully counterbalanced
167 according to both pre- and post-natal exposure to ethanol. Testing was carried out
168 during the light phase (i.e., between 9am and 5pm) over a four-day period. During the
169 procedure, each fish was individually placed in the novel tank. They were filmed over a
170 five minute period, during which time we recorded the duration of time spent in the
171 bottom third of the tank, as well as the distance travelled (see Fig 1). The filming and

172 analysis was carried out using Noldus Ethovision XT software (TrackSys, Nottingham,
173 UK). Following the tank dive, the fish was removed from the novel tank and placed back
174 in its holding tank.

175

176 *2.4. Individual social behavior*

177 [FIGURE 2 HERE]

178 Figure 2 displays the apparatus used for the individual fish social behavioral assay. Five
179 adult zebrafish were placed into one side of the tank, in the other side was the test
180 zebrafish. Perforated sheets divided the two segments such that the test zebrafish could
181 both see and smell the group of conspecifics. The tank was filled with aquarium treated
182 water. The five fish in the group were either PNE fish or controls, matching the test
183 fish's developmental treatment. Using Etho-Vision XT 5.0, the duration out of a 20-
184 minute trial that the test zebrafish spent swimming in the segment closest to the divide
185 was recorded for adult control and ethanol treated zebrafish. This assay was performed
186 on four-month-old fish (~50% male/female) with $n = 33$ fish in the control and $n = 30$
187 fish in the PNE group on both occasions.

188

189 *2.5. Shoaling*

190

191 [FIGURE 3 HERE]

192

193 The shoaling procedure (Figure 3) was based on an earlier study (Parker et al., 2013). In
194 order to carry out the shoaling assay, fish from each treatment were split randomly into
195 four groups of five (familiar individuals; mixed male/female). They were placed into an
196 open arena ($W \times L \times H$: $42 \times 49 \times 15$ cm) filled with 6L aquarium treated water. The fish
197 were left for five minutes to acclimate to the arena, and then filmed from above for 10-
198 minutes. Behavioral sampling took place from the video recording. The arena was
199 separated into eight equal sections (see Figure 3). At each 30-second interval during the
200 10-minutes, the maximum number of fish in one section (Max) was divided by the total
201 number of sections occupied by the fish (Total), thus providing a cluster score for each
202 time point (t) (Collins et al., 2011).

203

204 2.6. *Quantitative real-time PCR*

205 Batches of zebrafish larvae (aged 24hpf, 50hpf, 72hpf, and 7dpf) were digested in 200µl
206 Lysis buffer with 2µl Proteinase K for 30-45min (55°C) (n = 3 larvae/sample, n = 4
207 samples/age group). These embryos were checked for any differences in gross
208 morphology and size (quantified by pixel density of photographs) to rule out any
209 potential changes being due to developmental delay caused by the ethanol exposure.
210 mRNA was isolated using 40µl Dynabeads® Oligo(dT)₂₅ according to manufacturer's
211 instructions, and cDNA was synthesized and tested in a quantitative real-time
212 polymerase chain reaction (qPCR). Reference genes (see Table 1 for primer sequences)
213 were chosen according to previous research *β-actin*, *ef1α* and *rpl13α* (Tang et al., 2007).
214 Target genes used were *slc6a4*, *htr1a*, *oxtr*, *avpr* (see Table 1). Absolute quantification
215 was obtained by making standards for each gene, prepared using the relevant primers
216 to amplify fragments from cDNA. Samples were then PCR purified and diluted to 10¹¹
217 fragments using the Avogadro constant. All qPCR reactions were carried out in
218 triplicate. 2µl of cDNA and 2µl each of forward and reverse primers (see Table 1) was
219 added to 10µl SYBR® Green PCR Master mix (Applied Biosystems) on a LightCycler
220 LC480 instrument (Roche Diagnostic). For detailed methods see (Gemenetzidis et al.,
221 2010;Teh et al., 2012).

222

223 [TABLE 1 HERE]

224

225 2.7. *Data preparation and statistical analysis*

226 Tank diving data were fitted to a linear mixed effects model, with group (PNE vs
227 handling control) and time (mins 1-5), and their interaction, as fixed effects. Distance
228 traveled was entered as a covariate to control for individual differences in
229 freezing/darting in the novel tank (Parker et al., 2012). The dependent variable was
230 time spent in the bottom third of the tank. Individual social behavioral data were
231 analyzed using between-subjects t-tests. The independent variable was group (PNE vs
232 handling control) and the dependent variable was total time spent in social zone during
233 test (sec). Cluster analysis data were fitted to a linear mixed effect model, with group

234 (PNE vs handling control) as a fixed effect and time (20-levels) as a repeated effect with
235 a structured identity covariance matrix specified. The dependent variable was cluster
236 score (0.2-5). Relative mRNA expression ratios in the qPCR were calculated with respect
237 to reference gene cycle-threshold (Ct) values, and then subjected to a two-way factorial
238 (between-subjects) analysis of variance (ANOVA). Significant main effects and
239 interactions were followed up with pairwise comparisons. The between-subjects factors
240 were age (4-levels: 24hpf, 50hpf, 72hpf, 7dpf) and ethanol treatment (2-levels: 20mM
241 ethanol vs handling control) and their interaction. Homogeneity and normality were
242 ascertained by visual inspection of quantile-quantile (q-q) plots, and residual vs fitted
243 values. All test statistics were evaluated with respect to a type-1 error rate of 0.05. All
244 descriptive statistics are reported as estimated marginal means \pm SE unless otherwise
245 indicated. Statistical analyses were carried out in IBM SPSS for Macintosh (v. 19).

246

247

248 **3. Results**

249 *3.1. Tank Diving*

250 [FIGURE 4 HERE]

251 Figure 4 displays the time spent in the bottom third during the five minutes exposure to
252 the novel tank. As is clear, although both groups displayed the typical tank diving
253 response, gradually exploring the upper regions of the tank over the course of the
254 exposure) the PNE fish appeared to spend longer in the bottom third of the tank. This
255 was fitted to a linear mixed effects model, which confirmed a significant main effect for
256 time, $F_{4,179} = 7.24, p < 0.001$ and for group, $F_{1,179} = 25.87, p < 0.001$. There was no time x
257 group interaction ($F < 1$).

258 *3.2. Individual social behavior*

259 [FIGURE 5 HERE]

260 Figure 5 displays the time spent in the social segment according to PNE treatment. The
261 PNE 20mM ethanol fish spent significantly less time in the social zone than the control
262 animals, as confirmed with a between-subjects t-test, $t(61) = 2.35, p = 0.02$.

263 *3.3. Shoaling behavior*

264 [FIGURE 6 HERE]

265 Figure 6 displays the mean cluster scores for each shoal according to developmental
266 ethanol exposure. The PNE shoals showed significantly less group cohesion during the
267 course of the 10-minute observation period, and this difference was confirmed with a
268 linear mixed effects model, $F_{1,158} = 18.34, p < 0.001$.

269

270 3.4. Real-time quantitative PCR

271

272 [FIGURE 7 HERE]

273 Figure 7 displays the mRNA expression ratios for PNE and control embryos for genes
274 relating to social behavior. Visual inspection of the data suggests that for all of the
275 genes we tested, there was a spike in expression at 50hpf (i.e., 2hrs after addition of the
276 ethanol). In addition, for *ht1aa* and *oxtr*, it appears that the difference between ethanol
277 and control embryos reverses by 7dpf. These differences were further characterized
278 with two-way ANOVAs. For *avpr* (Figure 7A), there was a significant effect of age, $F_{3,24} =$
279 $18.49, p < 0.001$, with mRNA expression higher at 50hpf than at any other age ($ps <$
280 0.001). There was no significant effect of treatment, although this approached
281 significance, $F_{1,24} = 3.47, p = 0.07$, nor was there a significant age x treatment
282 interaction, $F_{3,24} = 1.2, p = 0.3$. For *oxtr* (Figure 7B), there was a significant effect of age,
283 $F_{3,24} = 9.37, p < 0.001$, but not significant effect of treatment, $F < 1$. There was an age x
284 treatment interaction, $F_{3,24} = 4.38, p = 0.01$. The interaction was characterized as
285 ethanol treated embryos showing higher *oxtr* mRNA expression at 50hpf than controls,
286 but this effect reversing by 7dpf. For *slc6a4* (see Figure 7C), there was a significant
287 effect for age, $F_{3,24} = 8.18, p = 0.001$, but not for treatment, $F_{1,24} = 1.28, p = 0.27$. There
288 was also a significant age x treatment interaction, $F_{3,24} = 3.57, p = 0.03$, characterized as
289 a significant increase in *slc6a4* mRNA expression at 72hpf in the ethanol treated group,
290 but no differences at other stages. For *ht1aa* (Figure 7D), there was a significant effect of
291 age, $F_{3,24} = 10.94, p < 0.001$, but not for treatment, $F < 1$. There was also a significant age
292 x treatment interaction, $F_{3,24} = 4.82, p < 0.01$, characterized as ethanol increasing *ht1aa*
293 mRNA expression at 50hpf ($p < 0.01$), but this effect reversing at 7dpf ($p < 0.01$).

294

295 **4. Discussion**

296 In both humans and in comparative animal models, exposure to moderate levels of
297 alcohol during early brain development leads to social behavioral deficits. Here, we
298 chronically exposed developing zebrafish embryos to a moderate level of ethanol
299 (20mM, equivalent to a BAC of ~0.04g/dL), and observed both reduced social cohesion
300 (shoaling) and reduced individual social behavior in this species. Analysis of
301 developmental mRNA expression in genes relevant to social behavior revealed transient
302 up-regulation of *slc6a4* at 72hpf, and putative adaptive changes in *oxtr* and *htr1aa*
303 expression following exposure to ethanol during early brain development, suggesting a
304 developmental mechanism by which the observed effects on social behavior may
305 manifest. This supports and extends previous work by Buske and Gerlai (2011) and
306 Fernandes and Gerlai (2009), who showed that brief (1-2hr) exposure to much higher
307 concentrations of ethanol (0.25%, 0.5%, 1% v/v, equivalent to 0.08, 0.16 and 0.32g/dL
308 BAC) alters adult social behavior (operationalized by nearest neighbor and by proximity
309 to a virtual fish) and reduced levels of 5HIAA (5-HT metabolite) in the adult brain. We
310 also observed that PNE fish spent longer in the bottom third in the novel tank diving
311 test, suggesting higher levels of trait anxiety in these fish (Stewart et al., 2011).

312 *ht1aa* mRNA expression initially increased following 2hrs exposure to ethanol, and
313 subsequently decreased after 5 days suggesting adaptation of mRNA expression. *ht1aa*
314 codes for the 5-HT_{1A} receptor, and our findings support previous work demonstrating
315 reduced binding of the 5-HT_{1A} agonist 8-hydroxy-2-dipropylaminotetralin (8-OH-DPAT)
316 to 5-HT_{1A} receptors in frontal cortical regions following PNE (Kim et al., 1997). This
317 suggests that acute ethanol causes a spike in mRNA expression, and subsequent down-
318 regulation at 7dfp. The role of the 5HT_{1A} receptor in mammalian social behavior is well
319 established. For example, the selective 5HT_{1A} receptor agonist 8-OH-DPAT increased
320 social interactions in gerbils (Cheeta et al., 2001), and the 5-HT_{1A} antagonist pindobind
321 decreased defensive and other social behavior in a resident-intruder test in mice (Bell
322 and Hobson, 1993). Our data therefore support the hypothesis that alterations in social
323 behavior observed here, and in many other PNE studies, may be the result of altered 5-
324 HT activity.

325

326 We also found an increase in trait anxiety, as operationalized by time spent in the
327 bottom third of a novel tank (Stewart et al., 2011), in the PNE fish. Previous work in
328 mammals has identified differences in anxiety and stress reactivity following exposure
329 to ethanol during early brain development (Weinberg et al., 1996; Osborn et al., 1998),
330 and our findings confirm that similar mechanisms may be in place in fish. 5-HT_{1A} knock-
331 out mice show increased anxiety (Heisler et al., 1998) and allelic variation at the 5-HT_{1A}
332 locus is related to pathological anxiety and depression in humans (Strobel et al., 2003).
333 It could be hypothesized therefore that the increase in anxiety is the result of decreased
334 *htr1aa* mRNA expression observed here in the 7pdf fish. An alternative interpretation is
335 that the novel tank test is simply an extension of the reduced social interactions
336 observed in the fish. We previously reported that individually housed zebrafish show
337 markedly lower 'anxiety' responses on the novel tank test, while at the same time
338 showed higher basal cortisol (CORT) than group-housed conspecifics (Parker et al.,
339 2012). We suggested on the basis of this that the novel tank test might represent a
340 social 'searching' assay, rather than an assay of anxiety *per se*, which would be
341 consistent with the current data. PNE has a complex effect on stress reactivity and
342 adaptation (Weinberg, 1993; Giberson and Weinberg, 1995; Weinberg et al., 1996). For
343 example, PNE does not cause changes in baseline CORT or adrenocorticotrophic
344 hormone (ACTH) in rats (Weinberg et al., 1995) but does alter hypothalamic pituitary
345 adrenal-cortical axis (HPA) adaptation to repeated stressors (Weinberg et al., 1996). In
346 addition, Osborn et al. (1998) demonstrated sex differences in the physiological and
347 behavioral responses to the elevated plus maze exposure following PNE. Hofmann et al.
348 (2007) also showed that PNE female rats exhibit a decreased ACTH response to the 5-
349 HT_{1A} agonist, 8-OH-DPAT, but an increased ACTH response to the 5-HT_{2A} agonist DOI.
350 Male rats showed no differences. This suggests that, in females at least, there is some
351 disruption by PNE of the interaction between the serotonin system and HPA axis. This
352 would imply a link between the social and stress-reactivity effects of PNE. An
353 examination of the molecular structure of the HPA system (hypothalamic pituitary
354 interrenal axis [HPI] in fish) and how this interacts with the 5-HT system during early
355 brain development following developmental ethanol exposure may help to elucidate
356 this. Although beyond the scope of this paper, our findings here demonstrate that
357 zebrafish offer an excellent model system for examining this mechanism *in vivo*.

358 We also observed transient up-regulation of *slc6a4* at 72hpf, with expression
359 normalizing by 7dpf. *slc6a4* codes for the serotonin transporter molecule (5-HTT),
360 which is located pre-synaptically, and is responsible for 5-HT reuptake (Blakely et al.,
361 1991). Interestingly, we observed that a change in *slc6a4* was preceded by changes in
362 *ht1aa* and *oxtr*, which codes for the oxytocin (OT) receptor. 5-HTT is known to be an
363 important mediator of social behavior and social cognition (Canli and Lesch, 2007). For
364 example, Wendland et al. (2006) demonstrated that allelic variation in the 5-HTT-linked
365 polymorphic region (5-HTTLDR) explained variance in aggression and social cohesion
366 in macaques. Similar variations have subsequently been cited as a potential mediator of
367 impulse control (Retz et al., 2004; Paaver et al., 2007) and a variety of other
368 psychopathologies in humans (Risch et al., 2009). PNE in rats causes permanent
369 alterations in 5-HTT sites in the hypothalamus (Zafar et al., 2000) and further adds to
370 the support for the involvement of the 5-HT system in social behavioral deficits
371 observed in FASD.

372 In addition to the 5-HT system, the hypothalamic neuropeptides OT and AVP modulate
373 and mediate aspects of social behavior (Winslow et al., 1993; Heinrichs et al., 2009).
374 Here we observed a spike in *oxtr* mRNA expression as an acute response to addition of
375 ethanol to the larvae at 50hpf. We then observed what appeared to be adaptation, with
376 expression significantly down-regulated at 7dpf in a similar manner to *ht1aa* expression
377 patterns. We did not observe a significant change in *avpr* expression at 50hpf. In
378 mammals, as both 5-HT and OT regulate social behavior, the mechanisms by which the
379 two interact have been of interest, and the co-localization of 5-HTT and OT expressing
380 neurons in the hypothalamus are thought to constitute a mechanism by which 5-HTT
381 regulates OT release (Emiliano et al., 2007). Acute ethanol directly inhibits OT release
382 (Fuchs and Wagner, 1963; Kalant, 1975; Eisenhofer and Johnson, 1982). Our findings
383 that there is a sharp spike in *oxtr* mRNA at 50hpf (2hrs after ethanol exposure) are
384 consistent with this, and suggest that ethanol initially increases, then ultimately
385 decreases *oxtr* expression. Thus, as OT facilitates 5-HT release in the raphe nucleus in
386 mammals (Yoshida et al., 2009), this suggests that acute ethanol will reduce 5-HT
387 release in 5-HT neurons expressing OT receptors. This could result in initial
388 homeostatic up-regulation of *ht1aa* expression as seen here. As the system re-balances,
389 by 72hpf we saw an increase in *slca6*, possibly as a result of an increase in cells or as

390 adaptation to remove any excess 5-HT. Collectively, this suggests a possible mechanism
 391 by which moderate ethanol exerts its effects on molecular factors associated with social
 392 behavior.

393 In conclusion, we have shown that zebrafish may represent a good model for
 394 translational work on the effects of ethanol exposure during brain development on
 395 behavior and gene expression. We found that moderate levels of ethanol exposure affect
 396 social behavior, and that this appears to be mediated by changes in 5-HT and OT mRNA
 397 expression levels. It also appeared that this reduction specifically in *ht1aa* and *oxtr*
 398 mRNA expression at 7dpf was an adaptation following an initial spike in expression on
 399 acute ethanol exposure. Finally, we found evidence for some role of ethanol during early
 400 brain development on anxiety. More research will be needed to elucidate the
 401 mechanisms by which these changes occurred, but this does suggest that zebrafish may
 402 be a suitable model for examining the molecular aspects of the stress – social
 403 interactions relating to PNE. Recognition and characterization of these processes will
 404 aid in the development of therapeutic interventions to help ameliorate negative
 405 symptoms of FASD and related psychiatric disorders.

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