

Early exposure to atrazine in zebrafish: intergenerational effects and animal personality

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Abstract

Animal personality (consistent inter-individual differences in behaviour over time and context) is important for an individual's fitness. Acute exposure studies in adults show that endocrine disrupting chemicals (EDCs) can deteriorate behavioural consistency and directionally alter personality phenotypes, which may decrease fitness. Early life development is sensitive to environmental stressors and exposure to EDCs may permanently alter phenotypes through to adulthood. In addition, the effects of environmental stressors may not be isolated to a single generation – offspring may indirectly be impacted, via non-genetic processes. Epigenetic mechanisms (e.g. DNA methylation) that help to regulate the genome, can become permanently altered via EDC exposure during early life and alterations have the potential to be inherited in the germ cells. The aims of this thesis were to investigate direct effects of the EDC, atrazine, on animal personality in the zebrafish (*Danio rerio*) and the indirect effects on personality in their offspring. Recent studies suggest that zebrafish inherit the methylome (total DNA-methylation marks across the genome) in an unchanged state from their fathers, suggesting a possible route for environmental specific information to be inherited. Hence, I focus on paternal effects in this thesis. In addition, I investigate the underlying basis of mRNA transcript number differences in F₁ zebrafish brain tissue, with the aim to determine whether observed changes in behaviour are underpinned by changes in gene expression. This is the first study to examine the effects of atrazine on personality, for both direct and indirect exposure.

Juvenile zebrafish were exposed to atrazine at typical environmental concentrations: 0.3, 3 or 30 part per billion (ppb) through sexual differentiation and their personality phenotypes were assayed (activity, aggression, boldness, anxiety and exploration) at adulthood. F₁ offspring were produced from atrazine exposed and control males, mated with unexposed females, creating full-sibling families. Progeny personality was tested at adulthood. Brain samples of a subset of F₁ progeny (based on anxiety and aggression phenotypes) were taken to determine mRNA transcript number of candidate genes involved in the regulation of the serotonergic system (*slc6a4a*, *slc6a4b*, *htr1Aa*, *htr1B*, *htr2B*), an underlying component involved in anxiety and aggressive-related behaviours, and a potential target of atrazine.

I found evidence that direct and paternal atrazine exposure significantly influences aspects of boldness, aggression, anxiety and exploration, though effects were not consistent across all doses. Changes in boldness were sex-dependent, specifically, F₀ female boldness was reduced in a dose-dependent manner and F₀ males exhibited a threshold increase in boldness. F₁ females exhibited a similar dose-dependent reduction in boldness but in the opposite direction to F₀ females and there was little change amongst F₁ males. There was some evidence of reduced aggression after direct exposure, but paternal atrazine exposure significantly reduced aggression in the 0.3ppb and 3ppb groups. No F₀ treatment differences in activity were observed, however, significant F₁ decreases in activity were observed during aggression testing in the 0.3ppb and 3ppb groups. Anxiety increased and exploration decreased significantly in the F₀ 0.3ppb group only, while in the F₁'s there was some evidence of alterations in anxiety and exploration, but not significantly so.

Paternal atrazine exposure significantly down-regulated *htr1Aa* mRNA expression in females and slightly up-regulated it in males; moreover, *htr1Aa* mRNA (and *slc6a4a* - but not significantly so) was found to correlate positively with anxiety levels in controls; however, this relationship was disrupted in the atrazine treatment group. The candidate genes tested here did not explain the significant reduction in aggression observed in the offspring of atrazine treated fathers, though both *slc6a4b* and *htr1B* mRNA correlated negatively with an aggressive personality phenotype.

In conclusion, early developmental exposure to environmental doses of atrazine resulted in persistent changes in personality phenotypes through to adulthood and effects were present in the offspring of atrazine exposed fathers. Moreover, some aspects of the serotonergic system were disrupted in the progeny. Overall, these results add to the ecological consequences of environmental contaminants, most importantly, that effects may be further propagated down the germ line.

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Chapter 1

General introduction

1.1 Animal personality

1.1.1 An overview

Animal personality is defined as consistent behavioural variation at the level of the individual that occurs across time and in different behavioural and environmental contexts (also previously known as temperament, emotionality or coping styles; Koolhaas et al. 1999, Sih et al. 2004a, Réale et al. 2007). Behavioural context describes an aspect of a behavioural function e.g. anti-predator, courtship, or dispersal. Environmental context or situation, describes the conditions experienced by an organism during a particular time point it is being measured under, such as levels of predator density (Sih et al. 2004a; Oers et al. 2005). Historically, behavioural ecology theory posited that behavioural variation of individuals within a given population of animals was due to random error (or non-adaptive variation) that lead towards a central tendency around an adaptive population mean (Dall 2004; Nettle et al. 2010). Today, however, consistent inter-individual differences in behaviour are readily observed across the animal kingdom from all levels of vertebrate systems (e.g. mammals, reptiles, fish birds and amphibians; Smith et al. 2008; Bell et al. 2009) and even to invertebrates (e.g. insects; Jandt et al. 2014).

Animal personality is typically characterised by five broad axes that encapsulate the response of an individual relative to other individuals. These axes being: boldness (also known as the boldness-shyness axis), exploration (also known as the exploration-avoidance axis), activity, aggressiveness and sociability (Réale et al. 2007). Boldness-shyness describes an individual's propensity to take risks (e.g. an animal's latency to emerge from a shelter, latency to approach a novel object or the duration spent in areas of high versus low risk; Wilson et al. 1994; Frost et al. 2007; Réale et al. 2007; Harris et al. 2010; Raoult et al. 2012).

Exploration describes the individual's propensity to seek out information of an unfamiliar space in a new environment (e.g. the duration, frequency or latency visiting distinct areas of a novel environment; Dingemanse et al. 2002; Réale et al. 2007). Activity is defined by the consistency of movement (or the consistency of the rate of movement) constrained by the costs of metabolism (Montiglio et al. 2010; Schuster et al. 2017). Aggressiveness is defined by the degree of antagonistic interactions with conspecifics (Réale et al. 2007) and can be measured by dyadic fights or by measuring interactions with a mirror image (Oliveira et al. 2005; Ariyomo et al. 2013a; Way et al. 2015). Finally, sociability is considered the propensity for an individual to seek out conspecifics (e.g. the cohesion of a group of animals; Miller et al. 2007; Réale et al. 2007). Recently, anxiety is considered to be an additional personality trait and is defined as anticipatory fear caused via uncertainty between a potential threat (that may not be specifically present, but is anticipated to occur), and lack of information needed to ascertain the actual risk of an action or situation (Blanchard et al. 1989; Belzung et al. 2007; Maximino et al. 2012). Anxiety may be measured by an animal's behavioural reaction in uncertain novel environment, such as erratic behaviour, freezing or residing in areas perceived to be of low risk (Egan et al. 2009; Maximino et al. 2012). These axes of variation may also co-vary together resulting in what is termed a behavioural syndrome. For example, on average, individuals that are more aggressive tend to also be bolder (Sih et al. 2004a).

Several aspects of ecology and evolution can be influenced by personality traits (see Wolf et al. 2012). Predator-prey interactions, for example, can shape personality traits and impact how individuals cope with predators. For instance, an aggressive-boldness syndrome in three spined stickleback (*Gasterosteus aculeatus*) is dependent on exposure to predation attempts (Bell et al. 2007) and pike (*Esox lucius*; an ambush predator) selectively target shy roach (*Rutilus rutilus*), whereas Eurasian perch (*Perca fluviatilis*; an active forager) tend not to discriminate between behavioural phenotypes (Blake et al. 2018). Shyer three-spined sticklebacks also preferentially associate with smaller sized individuals and tend to have higher social interactions, whereas bolder fish interacted less and lacked discrete associations across their social network (Pike et al. 2008). Moreover, less sociable individuals are more likely to disperse from their population (Cote et al. 2007; Cote et al. 2010).

Animal personality can contribute to an individual's survival and life-time reproductive success (Smith et al. 2008; Réale et al. 2010; Sih et al. 2012; Wolf et al. 2012). Given the limited behavioural variation of individuals, fitness may shift according to changing contexts, such as predator density or food availability (Réale et al. 2003; Dingemanse et al. 2004). For instance, asocial common lizards (*Lacerta vivipara*) have higher fitness in low-density populations, but when population density increases, the fitness for asociality decreases (Cote et al. 2008). When environmental contexts remain relatively constant over time, variation

at the behavioural extremes may be more beneficial, for example, bolder individuals may trade-off lower survival for more immediate reproductive opportunities when predator abundance is high, whereas shyer individuals may forgo short-term reproduction in order to increase survival. In this situation, individuals at either ends of the boldness-shyness axis theoretically maintain higher life-time reproductive success than individuals that are intermediate in their behavioural phenotype (Smith et al. 2008; MacPherson et al. 2017).

1.1.2 Anthropogenic drivers of animal personality

Several factors are observed to contribute to an individual's personality, these being a combination of social-context (and social interaction), life-time experience, feedback from the environmental context, the early-life environment, and in some instances a heritable, genetic (and possibly non-genetic) component has been attributable (Drent et al. 2003; Frost et al. 2007; Chapman et al. 2010; Reddon 2011; Ariyomo et al. 2013a). These factors together influence the intertwined neural, genetic (including epigenetic) and endocrine processes which govern the expression of behaviour and therefore, the extent to which variation can exist within an individual. What is less understood is how environmental contaminants (that are able to interfere with these processes) may influence and shape an individual's consistency in behaviour along their axes of personality. Of particular interest and concern are endocrine (and neuro-endocrine) disrupting chemicals (EDCs), which interfere with key molecular mechanisms that underlie behaviour (i.e. hormones, gene expression and neural physiology: Vandenberg et al. 2012; Volkova et al. 2015a). These chemicals are predominantly implicated in the disruption of typical reproductive behaviours (i.e. courtship and parental behaviour, see review Söffker et al. 2012), but some studies have shown effects on animal personalities (Dzieweczynski 2011; Dzieweczynski et al. 2013; Dzieweczynski et al. 2014; Hebert et al. 2014; Dzieweczynski et al. 2016; Porseryd et al. 2017b). For example, adult male and female Siamese fighting fish (*Betta splendens*), acutely exposed to the oestrogen mimic 17 α -ethinylestradiol, exhibited reductions in boldness, aggression (males only), activity and the individual consistency of these traits. Moreover the behavioural syndrome between boldness and activity (in females) and boldness and aggression (in males) was disrupted (Dzieweczynski et al. 2014; Hebert et al. 2014).

To date, much work on how EDCs may affect personality has focused on acute adult exposures (e.g. Dzieweczynski 2011; Dzieweczynski et al. 2013; Dzieweczynski et al. 2014; Hebert et al. 2014; Dzieweczynski et al. 2016; with some exceptions, Volkova et al. 2015a; Volkova et al. 2015b; Porseryd et al. 2017a). What is less understood is how the early life environment in combination with EDCs may influence animal personalities through to adulthood. The early life environment refers to critical developmental periods of

an organisms life, i.e. the embryonic and juvenile period, when major development occurs (Burton et al. 2014). During this early developmental period, organisms have greater plasticity but are more susceptible to environmental pressures altering their course of development (reviewed by Jonsson et al. 2014). For example, guppies (*Poecilia reticulata*) raised with an unpredictable food supply became bolder, more exploratory, and tended to be less social at adulthood (Chapman et al. 2010). Early life exposures to EDCs routinely find that effects become either fixed, or manifest permanently later in life (Shenoy 2014; Volkova et al. 2015a; Wirbisky et al. 2015; Wirbisky et al. 2016c). However, little research on animal personalities has been done under an early life exposure framework (e.g. Volkova et al. 2015a).

1.1.3 Non-genetic inheritance and animal personality

Offspring phenotype is not solely derived from the genetic code it inherited from the gametes of its parents. Non-genetic inheritance describes the transmission of parental phenotypic or environmental variation to offspring that do not stem from the inheritance of the DNA sequence (Bonduriansky et al. 2012; O’Dea et al. 2016). Parental effects, a subset of non-genetic inheritance, shows that parents may mediate their offspring’s phenotype through factors such as behaviour (e.g. nurturing behaviour) or physiology (e.g. the transfer of hormones and proteins into eggs; Badyaev et al. 2009; Curley et al. 2011; Sopinka et al. 2017). Moreover, there is some evidence that non-genetic inheritance may additionally attribute to animal personality (Reddon 2011). For instance, personality differences in magpie (*Pica pica*) offspring are predicted by laying order, which is also associated with altered hormone composition in eggs (Rokka et al. 2014). And hatching asynchrony by zebra finch (*Taeniopygia guttata*) mothers interacts with offspring sex to influence exploratory behaviour of offspring at adulthood (Mainwaring et al. 2013). In addition, Taylor et al. (2012) found evidence for maternal effects (as well as genetic effects) correlating with activity and aggression in North American red squirrels (*Tamiasciurus hudsonicus*). Recently, studies have found that personality differences associated with non-genetic inheritance may change with ontogeny. For instance, incubation temperature in bearded dragon (*Pogona vitticeps*) can influence boldness over the short term (but not long term; Siviter et al. 2017), and maternal effects in guppies are associated with risk-taking individual differences in juveniles, but the association is absent at adulthood (White et al. 2018). In general, non-genetic inheritance is an under-appreciated source for influencing animal personality, but also requires further research to validate its potential role in shaping personality traits.

Recent work is recognising that sperm can be a source contributing to non-genetic inheritance (see Immler 2018). Aside from inheritance of the paternal genome, sperm also carries epigenetic modifiers (e.g. DNA-methylation, non-coding RNAs and histone modifications; reviewed in Casas et al. 2014; Rando 2016).

Epigenetic modifiers are molecules that act as an interactive mechanism between environment and the genome by regulating gene expression (see review; Jablonka et al. 2009). These molecules regulate the genome as the environment shifts, but exposure to strong environmental stresses (especially during early life) may leave epigenetic marks that, in turn, permanently alter the phenotype of the adult (see reviews; Barker 2004; Jablonka et al. 2009). Studies are increasingly documenting non-genetic inheritance due to epigenetic effects (see review Jablonka et al. 2009; Curley et al. 2011; O’Dea et al. 2016). For example, mice raised with chronic and unpredictable maternal separation (between 1-14 days after birth, a high stress environment for new born pups) caused depressive-like behaviours and altered behavioural responses to aversive environments through to adulthood. The effects of this high stress and inconsistent nurturing environment during early life were also transferred to descendants, despite cross fostering and continuous maternal care during the offspring upbringing (Franklin et al. 2010). Dias et al. (2014) found that male adult mice subjected to odour fear conditioning produced descendants with increased behavioural sensitivity to the same odour and increased neuroanatomical structuring with the odorant receptor (that was specific to that particular odour). In addition, EDCs may be a potential source for producing maladaptive phenotypes through non-genetic inheritance mechanisms. Guppies exposed to *17alpha*-ethinyl estradiol have a heightened anxiety like phenotype (observed through to two generations; Volkova et al. 2015a) and female F₃ descendants of mice embryonically exposed to the EDC, vinclozolin, prefer males without a history of exposure (Crews et al. 2007). Thus, EDCs may be a potential source for producing maladaptive phenotypes through non-genetic inheritance mechanisms.

Given the increased prevalence of EDCs in the environment (Colborn et al. 1993; Vos et al. 2000; Jenssen 2006; Diamanti-Kandarakis et al. 2009) it is important to understand the consequences of their contamination. Behaviour of the individual, through many interactions, may cascade up to the population level and dynamically affect population stability, persistence, structure or dispersal (Wolf et al. 2012; Araújo et al. 2018). Behaviour may therefore help to serve as preliminary guide to predict how populations might react to EDC exposure (Clotfelter et al. 2004; Araújo et al. 2018). Because animal personality is partly a function of the early life environment (among other factors), the sensitivity observed by organisms to EDCs during this period suggests that animal personality is likely to be affected (Jonsson et al. 2014). Furthermore, since non-genetic inheritance in general appears to be a mediator for altering the trajectories of offspring to suit the predicted local environment (based on the environment of the parent; Marshall et al. 2007; Burton et al. 2014), EDC induced pathological personality phenotypes may additionally be propagated indirectly to offspring. As animal personality contributes to fitness, non-genetic propagation of EDCs may also affect offspring fitness (Smith et al. 2008; Réale et al. 2010; Sih et al. 2012; Wolf et al. 2012). Overall, there is

a lack of research on animal personalities, EDCs and non-genetic inheritance; therefore, this thesis aims to better understand the relationship between these factors.

1.2 Thesis objectives and rationale

The aim of this thesis was to investigate non-genetic inheritance of animal personality under an EDC stressor in the zebrafish (*Danio rerio*; Figure 1.1). Zebrafish are an interesting model to investigate non-genetic inheritance because studies suggest that embryos inherit the methylome (total DNA-methylation marks across the genome) from their fathers, thereby suggesting a pathway for environmental signals to be inherited (Jiang et al. 2013; Potok et al. 2013). In addition, zebrafish have a short 3-4 month generation time (Kimmel et al. 1995) and their genetics are well understood (Norton et al. 2010; Howe et al. 2013). This species can act as a complementary model to mammalian studies (Norton et al. 2010), as well as providing a model for the potential effects of toxicants on other aquatic species. Researchers have established catalogues of complex zebrafish behaviours (Kalueff et al. 2013) and detailed insight into animal personality of zebrafish is becoming increasingly understood (e.g. Thomson 2017).



Figure 1.1: Example of an AB wildtype zebrafish female. Size of the animal is approximately ~35mm.

I utilised the EDC, atrazine (2-chloro-4-ethylamino-6-isopropylamino-1,3,5-triazine; Figure 1.2) as a non-genetic source of environmental variation. Atrazine is a common herbicide used to control broadleaf and weeds in a variety of crops, such as maize and sugarcane, via the disruption of photosynthetic pathways (Barr et al. 2007, Solomon et al. 2008). Previous research has demonstrated that atrazine can reduce mating behaviour (male-male aggression in a courtship context in guppies; Shenoy 2012; Shenoy 2014) and

alter some aspects of sociability (Schmidel et al. 2014). In mice, atrazine has been observed to increase anxiety-like behaviours after short term acute exposure (Lin et al. 2013) and after early life exposure as well (Belloni et al. 2011; Lin et al. 2014a). But no study has yet explored how atrazine may affect personality traits.

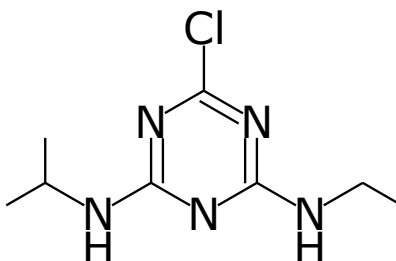


Figure 1.2: Atrazine chemical structure, modified from Graymore et al. 2001.

Atrazine is able to cross the blood brain barrier (Ross et al. 2009), but the mechanisms behind its action within the brain are not fully resolved. Developmental transcriptomic studies in zebrafish embryos at environmental doses highlight interference at the level of transcription of nervous system development and function, tissue development and organismal development through to adulthood (Wirbisky et al. 2015; Wirbisky et al. 2016a). Many studies in rodents implicate the dopaminergic system after adult or developmental exposure (Coban et al. 2007; Lin et al. 2013; Rodriguez et al. 2017). Recent studies in zebrafish and rodents also implicate disturbance of the serotonergic system (Lin et al. 2014a; Wirbisky et al. 2015) and by proxy, the hypothalamic-pituitary-adrenal/interrenal (Heisler et al. 2007; Fraites et al. 2009). A study of the male mice metabolome (composition of all metabolites in plasma) after acute exposure found disruption of several metabolic pathways such as tyrosine (precursor to dopamine), tryptophan (precursor to serotonin; 5-HT), linoleic acid and α -linolenic acid (poly-unsaturated fatty acids, which play a role in normal brain development and can modulate expression of 5-HT and dopamine; Das 2013; Lin et al. 2014b). Atrazine has also been found to act transgenerationally on sperm biomarkers of disease in mice (McBirney et al. 2017) and can modify epigenetic molecules transgenerationally (Hao et al. 2016; Wirbisky et al. 2016b; Wirbisky-Hershberger et al. 2017). These studies indicate that sperm are able to carry epigenetic markers associated with atrazine exposure through the germ line.

The general frame work of this thesis is summarised in Figure 1.3 below. The experiments in Chapter 2 were specifically aimed at investigating to what extent the EDC, atrazine, may directly affect personality in adults and indirectly in offspring. Specifically, this study investigated the effect of atrazine on five personality traits: activity, anxiety, aggression, exploration and boldness. In Chapter 3, I investigate the underlying basis of

mRNA transcript number differences in F₁ zebrafish brain tissue, with the aim to combine observed changes in offspring behaviour with an epigenetic (assessed through gene expression) explanation of inheritance. I specifically targeted candidate genes involved in the regulation of the serotonergic system, the two serotonin (5-HT) transporters (*slc6a4a*, *slc6a4b*) and three 5-HT receptors (*htr1Aa*, *htr1B*, *htr2B*). These candidate genes are implicated as underlying components involved in anxiety and aggressive-related behaviours, and are also a potential target of atrazine. Lastly, Chapter 4 summarises and discusses the key findings of this work in the wider context of the literature.

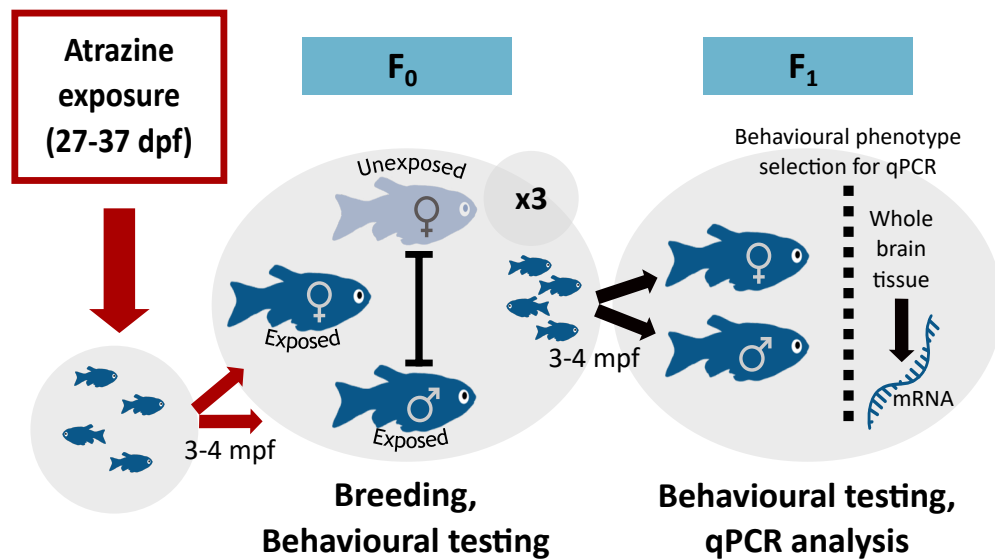


Figure 1.3: Experimental overview of thesis; first, juvenile zebrafish are exposed to atrazine from 27 to 37 days post fertilisation (dpf), then between 3-4 months post fertilisation (mpf), males are bred with unexposed females producing three full-sibling families. F₀ males and females then undergo behavioural testing. Between 3-4 mpf, F₁ fish have their behaviour tested the same way as the F₀'s. Whole brains of F₁'s are then tested for gene expression differences that correspond with differences in behaviour.

Chapter 2

Intergenerational effects of early life atrazine exposure on zebrafish behaviour

2.1 Introduction

Environmental contamination by anthropogenic pollutants is higher than any time in history, and continues to be a growing concern for wildlife, human health and the ecosystem (Vos et al. 2000; Noyes et al. 2015; Wilcox et al. 2016). One group of chemicals that have increased in interest and notoriety within the last decade are endocrine disrupting chemicals (EDCs; Colborn et al. 1993; Vos et al. 2000; Jenssen 2006; Diamanti-Kandarakis et al. 2009). These chemicals interfere with the endocrine system by blocking or mimicking hormone receptors (among a plethora of other complex interactions, see reviews Hotchkiss et al. 2008; Diamanti-Kandarakis et al. 2009; Vandenberg et al. 2012), leading to a range of negative effects ranging from disruption of morphology, behaviour, physiology and reproduction. Endocrine disrupting chemicals resemble aspects of how hormones function at minute concentrations and have the ability to exert different effects at varying levels (Vandenberg et al. 2012). As a result, EDCs frequently exhibit unusual dose-response relationships, (e.g. non-monotonic response Lin et al. 2014b; Vandenberg 2014), and can yield different effects depending on sex, age and exposure duration (Gioiosa et al. 2013; Palanza et al. 2016; see review by Söffker et al. 2012).

The direct effects of EDCs on behaviour has largely focused on the more immediate, reproductive behaviours

(and has been limited to a few EDCs; Söffker et al. 2012), such as reproductive success, sexual displays, nesting behaviours and competition between individuals for mates (e.g. Brian et al. 2006; Saaristo et al. 2010; see review Söffker et al. 2012). However, several recent studies have demonstrated that anthropogenic stressors, including EDCs, can also negatively affect animal personalities under acute exposures (Dziewieczynski et al. 2014; Dziewieczynski et al. 2016; Grunst et al. 2018). Animal personality is characterised as behaviours produced at the level of the individual, which are consistent over time and across different environmental contexts (Dall et al. 2004; Sih et al. 2004b; Réale et al. 2007). These traits include broad patterns of behaviour: anxiety, boldness-shyness, exploration, activity, sociability and aggressiveness (Sih et al. 2004b; Réale et al. 2007). Personality traits have been observed across a variety of taxa (e.g. insects, reptiles, birds, fish, amphibians and mammals; Bell et al. 2009). Personality traits are ecologically significant and can indirectly have major influence over life-time reproductive success, survival of individuals and social interactions (see review Smith et al. 2008). For example, male sleepy lizards (*Tiliqua rugosa*) that are less aggressive tend to associate with more females than more aggressive males (Godfrey et al. 2012). Likewise, bolder homing pigeons (*Columba livia*) are more likely to occupy higher ranks in their leadership hierarchy of the flock, influencing the direction of the flock's collective movement (Sasaki et al. 2018).

Interestingly, the effects of EDC exposure are not consistent across an organisms lifespan. During critical windows of early development (i.e. during early life, embryonic or juvenile development), the environment imparts a stronger pressure on organisms and increases the likelihood of altering developmental trajectories in a range of endpoints such as morphology, physiology and behaviour to mediate environmental pressure (Madsen et al. 2000; reviewed by Jonsson et al. 2014). In addition, developmental plasticity attenuates with age, such that environmentally influenced trajectories tend to become permanent (see Jonsson et al. 2014). For example, guppies raised with an unpredictable food supply became bolder, more exploratory, and tended to be less social at adulthood (Chapman et al. 2010). The plastic ability to respond is particularly adaptive if the early life environment predicts the later life environment, but may be detrimental if a miss-match between phenotype and environment occurs (e.g. Fleming et al. 1997; Visser 2008; Reed et al. 2010). In addition, the lasting impact of strong environmental stressors may be more severe or become pathological later in life. For instance, continuous maternal separation in rodents rapidly augments the stress axis to pathological levels, resulting in depressive, anxiety-like behaviours and changes to neural functioning through to adulthood (Nishi et al. 2013; reviewed by Nishi et al. 2014). In a similar way, exposure to EDCs during early life risks the development of pathological phenotypes, that have the potential to be permanent (Shenoy 2014; Porseryd et al. 2017a).

As well as considering direct effects of EDCs over a single life-time, offspring may indirectly be impacted by

exposure through their parents (Anway et al. 2005; Burton et al. 2014). Non-genetic inheritance is defined as the transmission of traits to offspring. These traits are components of either the parental phenotype or environmental variation, but are not attributable to the inheritance of DNA sequence (Bonduriansky et al. 2012). Presumably this is a proximate mechanism to adjust offspring phenotypes to expected local environments (Burton et al. 2014; O’Dea et al. 2016). A variety of natural environmental stressors including, but not limited to, parental behaviour and social stress have been documented to alter phenotypes in their descendants (see review O’Dea et al. 2016). Studies increasingly find the effects of EDCs can be propagated through generations (see reviews Skinner et al. 2011; O’Dea et al. 2016). For instance, zebrafish developmentally exposed to 17 α -ethinyl estradiol and their offspring exhibit heightened anxiety-like phenotypes and increased shoaling intensity (Volkova et al. 2015b). In another study, F₃ descendants of mice embryonically exposed to vinclozolin during sexual differentiation had an altered (and sex dependent) anxiety phenotype (increased in females and decreased in males; Skinner et al. 2008). Intergenerational phenotypes in general may be produced via a range of non-genetic mechanisms. Maternal effects are particularly well documented (e.g. maternal glucocorticoids influencing offspring fitness; Sopinka et al. 2017), and to some extent so too are paternal effects (e.g. paternal care; Curley et al. 2011), but increasingly, studies are documenting non-genetic inheritance due to epigenetic effects (see reviews Jablonka et al. 2009, Curley et al. 2011, and O’Dea et al. 2016 for examples). Epigenetic inheritance, where the patterning of epigenetic mechanisms (such as DNA-methylation or histone modifications) across the genome (i.e. the epigenome) is inherited, may be a causal mechanism underlying changes associated with parental or ancestral EDC exposure (Skinner et al. 2008; Skinner et al. 2010; Skinner et al. 2011; Wolstenholme et al. 2012). For instance, atrazine exposure induced multiple, transgenerational diseases in rat testes that were associated with sperm DNA-methylation mutations (McBirney et al. 2017). Variation in gene expression is typically associated with behavioural variation (Alter et al. 2008; Jöngren et al. 2010; Theodoridi et al. 2017). As epigenetic mechanisms mediate between the genetic background and the internal/external environment, epigenetic mechanisms may be important for personality differences (Riyahi et al. 2015; Verhulst et al. 2016; Bierbach et al. 2017). For example, DNA-methylation variation at the dopamine receptor D4 gene in great tit (*Parus major*) was associated with personality difference in exploration (Verhulst et al. 2016). Though, more studies are needed to understand the role of how epigenetics (and other factors such as neuroendocrine factors) contributes to animal personality (Trillmich et al. 2018).

In the present study, I investigated if early life exposure to the neuroendocrine disruptor atrazine (also a common herbicide; see Chapter 1) at environmentally relevant doses affects the personality traits (activity, aggression, exploration, boldness and anxiety) of adult zebrafish and their F₁ progeny. Zebrafish make a

good model system for behavioural testing as methods are well developed from the toxicology and behaviour literature (Egan et al. 2009; Maximino et al. 2010) and have since been extensively appropriated into personality research (e.g. Martins et al. 2014; Way et al. 2015; Thomson 2017). Furthermore, zebrafish are a model for research into testing non-genetic inheritance (Wang et al. 2017) and zebrafish males exhibit interesting epigenetic mechanisms dissimilar to other known organisms. After conception by the egg and sperm, studies suggest that the pattern of methylation across the genome (i.e. the methylome) is stably inherited via the father's sperm whereas the mother's methylome is quickly degraded and 're-programmed' (Jiang et al. 2013; Potok et al. 2013), suggesting a possible route for environmental signals to be inherited.

Only a handful of studies have investigated behavioural changes at maturity as result of developmental exposure (e.g. Belloni et al. 2011; Bardullas et al. 2013; Lin et al. 2014a; Shenoy 2014) and no studies, to my knowledge, have investigated specific personality changes due to exposure of atrazine (either acutely or during development). I hypothesised that exposure to atrazine would affect the personality of directly exposed zebrafish and their F_1 progeny, through indirect paternal effects. I predicted that anxiety levels would be increased and that the effect would be stronger in females than males due to the sex specific, neurological effects of atrazine on the serotonergic system from transcriptomic studies (Wirbisky et al. 2015). I predicted that an increase in anxiety would result in an inverse decrease in boldness because these two traits tend to be negatively correlated with one another (Thomson 2017). Furthermore, I predicted that aggression levels would be negatively affected by atrazine exposure based on the studies by Shenoy (2012; 2014), whom found decreased aggression levels in the guppy during a mating context. Exploration is closely related to boldness (Thomson 2017), therefore I predicted that I would observe a decrease in exploration. Overall, activity was expected to be higher based on larval and gestational studies exposed to atrazine at environmental doses in fish and rodents respectively (Carmen Alvarez et al. 2005; Lin et al. 2014a) – though how activity levels might translate to adulthood was unknown.

2.2 Methods

2.2.1 Experimental design

To investigate if early life exposure to atrazine affects the personality of adult zebrafish and their F_1 progeny, I exposed F_0 juvenile zebrafish to three environmentally relevant concentrations, 0.3ppb, 3ppb, and 30ppb, of atrazine during sexual differentiation (27-37 days post fertilisation (dpf); Takahashi 1977; Uchida et al. 2002; Lee 2015). For an ecological perspective, the concentrations used in this study are typical environmental ranges and were selected based on previous use (e.g. Hayes et al. 2002; Weber et al.

2013; Wirbisky et al. 2015). A random sample of three F_0 males per treatment were bred to produce F_1 full-sibling families (Figure 2.1). Both F_0 and F_1 fish were phenotyped at ~ 3 months of age using three behavioural tests: a novel arena test, a novel object test and a mirror assay (see details below). Brains of F_1 fish were further analysed for changes in gene expression of genes involved in the serotonergic system (Chapter 3). All procedures were approved by the University of Otago Animal Ethics Committee, protocol 44/16.

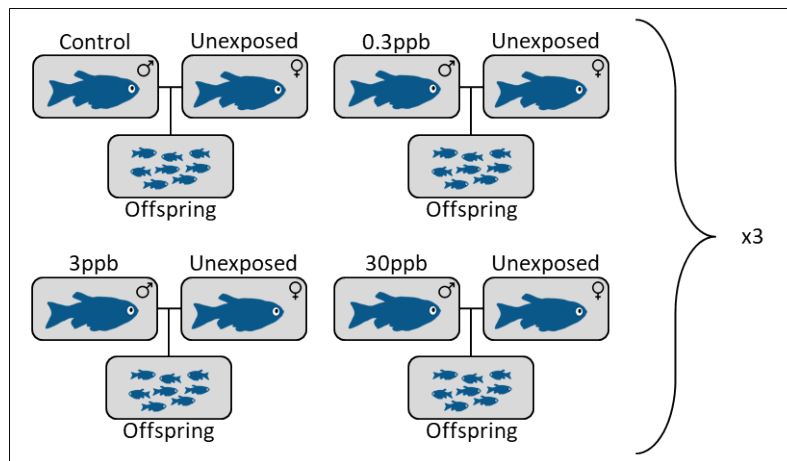


Figure 2.1: Design for assessing the effects of atrazine on F_1 progeny. A male from each treatment is bred with an unexposed female creating a full-sibling family. There were three families per treatment produced for behavioural phenotyping, 12 full-sibling families in total.

2.2.2 Zebrafish husbandry

Breeding, husbandry and atrazine exposure took place within the Otago Zebrafish Facility (OZF). The OZF is a temperature controlled facility with a 14 hr (0800-2200 hr) dawn-dusk light cycle. The conductivity, temperature and pH were maintained at range of 390-458 μ S, 25.2-26.1 $^{\circ}$ C and 7-7.8 pH, respectively. Phenotyping was performed in the Zoology Department. Room temperature was controlled at 25 $^{\circ}$ C and had a 13.5 hr (0700-2030 hr) light cycle with 30 min of simulated dawn and dusk at the start and end of each day.

F_0 embryos were produced by group spawning wildtype zebrafish (strain AB; 24 females; date of birth (DOB); 22/3/16 and 34 males; DOB; 26/2/16) using a Techniplast iSpawn Breeding System. Embryos were collected 30 min to 1 hour post fertilisation (hpf) and incubated in petri dishes (90mm diameter) with E3 media (Cold-Spring-Harbor-Protocols 2008) at 28.4 $^{\circ}$ C until 4 dpf. Dead or unfertilised embryos were removed from petri dishes once every day and E3 media was replaced every day until hatching. After four

days, hatched fry were moved into tanks with AquaOne, 5 parts per trillion (ppt) Aquaria Salt solution (synthetic sea salt), at a density of about 12 fry per litre. From 4-10 dpf zebrafish larvae were fed ad libitum, twice daily with dry food ZM000 (ZM systems; morning and afternoon) and once daily with rotifer (*Brachionus* spp) at midday. At 10 dpf water flow was turned on. After 10 dpf, fry were fed ad libitum, twice daily with dry food mixture of ZM000 and ZM100 (morning and afternoon), and once daily with and live Artemia (*Artemia salina*; midday).

2.2.3 Atrazine exposure

Atrazine was purchased from Sigma-Aldrich (Auckland, New Zealand; CAS 1912-24-9), and a stock solution was prepared three days before exposures started (23/02/17) by dissolving 5mg of atrazine powder in 200mL of purified water to achieve solubility. Ten day exposures to atrazine took place in 44mm x 600mL glass beakers. Eight beakers of 500mL system water were prepared; two served as controls and replicated atrazine concentrations of 0.3ppb, 3ppb or 30ppb were administered to the other six beakers. Forty zebrafish fry (27dpf) were randomly assigned to each of the eight beakers. The atrazine exposure regime lasted 10 days (26/02/17 through to the 8/03/17). This 10 day exposure regime was chosen because it covered the early-mid period of sexual differentiation in zebrafish (Takahashi 1977; Uchida et al. 2002; Lee 2015), where the probability of epigenetic information transfer is hypothesised to be high (based on studies of sexual differentiation and epigenetic inheritance in mammals; it is unknown whether this period is equivalent in zebrafish; Hackett et al. 2013). The juvenile period was also chosen to examine if the sensitivity window (where effects become permanent) of atrazine exposure extends beyond traditional embryonic (or prenatal) exposure regimes, resulting in behavioural changes at adulthood (e.g. Belloni et al. 2011; Shenoy 2014). Fry were fed once a day with ZM100 dry food during the exposure regime and water was changed every day (from day 2 until day 10); thus, concentrations of atrazine were also renewed every day. At the end of 10 days, fry were moved to new tanks in a Techniplast toxicology filtration system, which filtered water every 30min. After three days, fry were re-transferred onto the main OZF system and left to grow.

2.2.4 *In vitro* fertilisation, to produce F₁ progeny

At sexual maturity, ~3 months post fertilisation (mpf; 106–141 dpf), three males per treatment were sampled to produce three F₁ full-sibling families per treatment (Figure 2.1) using *in vitro fertilisation* (IVF) with untreated AB females (DOB=13/05/16). Breeding was done by setting up two males per treatment with an untreated female in breeding boxes (length=18.5cm, width=11cm, depth=8.5cm, water depth=6cm), with the sexes separated by a divider. This setup enables the pheromones of males to contact

the female, synchronising their spawning behaviour and stimulating female zebrafish to release oocytes in preparation for mating (Hisaoka et al. 1962; Gerlach 2006). Breeding boxes were setup in the afternoon and left overnight. The following morning IVF was performed using a modified protocol (see Appendix I; Johnson et al. 2018). Data on survival at 24 hpf and hatching at 48-96 hpf was also collected (see Appendix II). Embryos were transferred into petri dishes (90mm diameter) with E3 media and left to grow, following the same protocol described above.

2.2.5 Personality assays employed: descriptions and rationale

There are several ways in which to measure personality, but it must be suited to the system in question. Below I describe the three methods employed in this thesis for delineating personality phenotypes between zebrafish individuals.

The novel arena test also known as the novel tank diving test is a routinely used assay to test changes in levels of anxiety in response to pharmacological agents, as well as other behavioural measures, such as the activity profile, exploration and boldness (Egan et al. 2009; Maximino et al. 2010). Zebrafish, when exposed to an unfamiliar environment, tend to dive to the bottom of the test arena and tend to remain in the lowest portion (bottom dwelling or freezing), avoiding the higher portions of the water column, and after a few minutes the fish usually begin to explore the other areas of the tank (Egan et al. 2009; Maximino et al. 2010). Bottom dwelling is a common measure of anxiety and correlates negatively with the neurotransmitter serotonin (5-HT; Maximino et al. 2012). In fish, exploration may be tested by the amount of time an animal spends in different areas of the tank (Thomson 2017). Boldness in the novel arena test is based on the fish's risk-taking behaviour in relation to areas of higher risk such as entering or spending more time in areas of higher risk. For instance, spending more time in portions of the water column closer to the surface would constitute a greater boldness phenotype (Maximino et al. 2010).

The novel object test is adapted from rodent studies (Ennaceur et al. 1988) and is considered a complementary assay to the novel arena test to examine differences in boldness and anxiety in fish (Wright et al. 2006a; Ogwang 2017; Maximino et al. 2010). When exposed to a novel object (after an acclimation period within a novel environment), zebrafish show a tendency to avoid the foreign object, exhibit bottom dwelling behaviour or freezing, reduce their activity profile, and after a certain amount of time the zebrafish will begin to occasionally inspect it (Wright et al. 2003; Wright et al. 2006a). Approaching within 1-1.5 body lengths of the novel object is considered a measure of boldness (Wright et al. 2003; Wright et al. 2006a), and is interpreted as a predator inspection behaviour (Wright et al. 2003; Wright et al. 2006a; Ogwang 2017; Maximino et al. 2010).

The mirror test is typically used in assessing an individual's level of aggression (e.g. Ariyomo et al. 2012; Ariyomo et al. 2013a; Way et al. 2015). The mirror test is an alternative to dyadic fights (fights between two individuals) as both assays elicit similar antagonistic responses to one another. The fish are unable to recognise their reflection and are thought to perceive the mirror image as an intruder (Oliveira et al. 2005; Way et al. 2015). Zebrafish naturally form hierarchies and both males and females will be aggressive to one another to maintain dominance and monopolise resources (Spence et al. 2008). The mirror produces a size matched individual removing the variation of the size of an opponent. Mirror images also differ from dyadic fights on a molecular and hormonal level. A dyadic fight always results in either a winner or loser, influencing the level of whole body androgens (11-ketotestosterone and testosterone) and cortisol which in turn leads to transcription changes (depending on the outcome), whereas a fight between a mirror opponent does not resolve, allowing insight into individual level of antagonistic behaviour without the likely confounding effects of transcription and hormonal changes (Oliveira et al. 2005; Oliveira et al. 2016; Teles et al. 2016). This method was adapted from Ariyomo et al. (2013b) and Way et al. (2015), and though only capturing a smaller portion of the repertoire of zebrafish aggressive behaviour (see Oliveira et al. 2011; Kalueff et al. 2013), the mirror test was considered over a dyadic fights for logistical reasons, to limit possible confounding changes in gene expression and hormones (Oliveira et al. 2005; Oliveira et al. 2016; Teles et al. 2016) and because the correlation between an individual's behaviour in a flat mirror assay and a dyadic fight is moderately high (Ariyomo et al. 2013b; Way et al. 2015).

2.2.6 Behavioural phenotyping: experimental protocols

Prior to behavioural assays, all fish were transferred from the OZF to the animal rearing facility in the Otago Zoology Department. Zebrafish were left to acclimate to the new facility for 7 days before behaviour tests began. Fish were fed as described above, except on the day of behavioural phenotyping, where fish were fed after filming finished. The F_0 fish were transferred at 113dpf and the F_1 's at 110-131dpf.

The novel arena (Figure 2.2), the novel object (Figure 2.3) and the mirror tests (Figure 2.4). The novel arena (Figure 2.2), the novel object (Figure 2.3) and the mirror tests (Figure 2.4) were run consecutively in the same tank with each assay lasting 10min, with three fish assayed simultaneously in three separate tanks (length=30cm, width=15.5cm, height=27cm filled with 8L of system water to a depth of 19.5 cm). White coloured white-board film was fixed to the back, bottom and side of each tank wall to limit light and reflection during filming. A flat mirror (vertical length = 19cm, width=15cm) was fixed to the outside wall of each tank (mirror placed on the left side of one tank, and the right side of the other two tanks). During the novel arena and novel object tests, the mirrors were covered with a removable, hard plastic opaque

barrier. Tanks were lighted from 30cm above the tanks with one 240V 48LED aluminium light strip and from behind using three Godox LED170 lights (31-35.5cm away) to provide diffusing light to increase fish contrast during filming. Fish were filmed with a Basler acA1300-60/gc GigE camera with a 4.4-11mm lens placed about 112cm away from the row of tanks and live-tracked using EthoVision XT behavioural tracking software version 11.5 (Noldus et al. 2001).

Previous work in our lab has shown that the repeatability (the intra-class correlation values, i.e. animal personality) of zebrafish behaviours is dependent on an initial exposure to the assay regime (including exposure to the novel arena and novel object) and thereafter, in any subsequent behavioural assays, the exhibited behaviour during the trial is typical of the individual's personality phenotype (Thomson 2017). In light of these findings, a 'tank experience' was provided after 7 days of acclimation. The repeatability of behaviours exhibited during the mirror tests is not dependent on an initial exposure (see Appendix IV), but was included to maintain consistency with the other behavioural tests.

Novel arena test

The first test, 'novel arena' (Figure 2.2), consisted of carefully netting the fish from their home tank into the novel, empty tank. Live-tracking with EthoVision started 10-30sec after the fish were placed in the tanks. Within EthoVision, the novel arena was divided into 12 square zones (four across, three down, approximately 5x5cm) to measure exploration. The total distance moved (cm), velocity (cm/s), the frequency of transitions into the top third, time spent moving/ not moving and the cumulative duration (sec) in each zone was recorded. The time spent within each third of water column was calculated by the sum of time spent in each zone of the same level. The latency to enter the top zone was taken as the first moment that a fish entered into any one of the four zones in the top third. Exploration was calculated as the standard deviation of time spent in each of the 12 zones (i.e. an explorative fish would have a standard deviation closer to zero, meaning a fish spent an equal amount of time in all 12 zones, a high standard deviation indicates the fish was less explorative, preferring to spend the majority of the assay in a few zones; Thomson 2017).

Novel object test

After filming the novel arena, the novel object test was conducted (Figure 2.3). An orange rubber bung (measuring 3.2cm long, 4.3cm wide at the bottom and 3.7cm wide at the top) attached to plastic fishing wire was placed slowly into the middle of the tank, hanging in the middle of the arena (at an approximate depth of 6.9cm below the surface of the water). Live-tracking started 10-30sec after the rubber bung was positioned, once the water settled. Within EthoVision, the novel object arena was divided into an upper

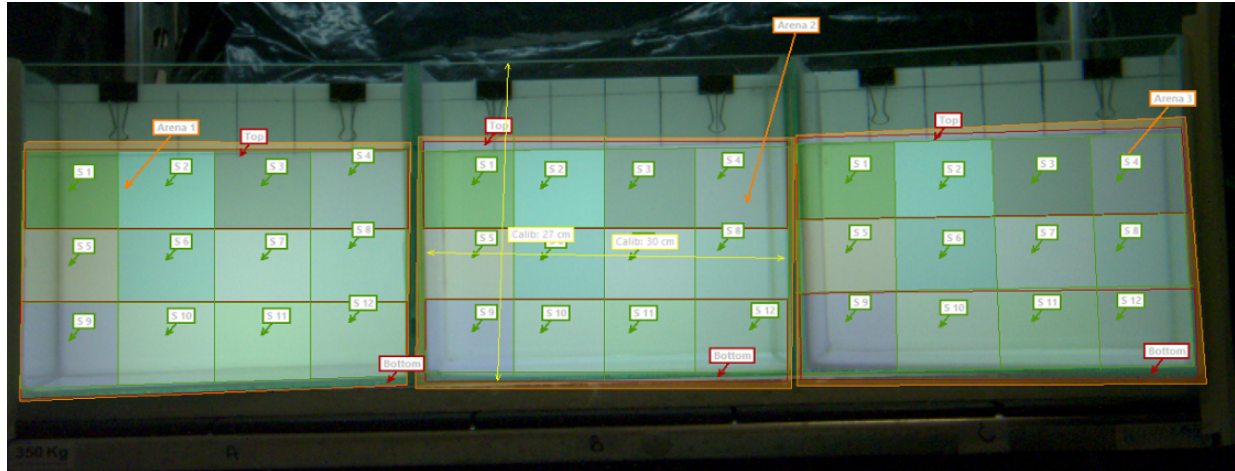


Figure 2.2: The settings for quantifying behaviours during the novel arena test. Within EthoVision XT, a four by three square grid is superimposed over the three test tanks. Each differently coloured square indicates a different zone (of $\sim 5\text{cm}$ wide and long).

and lower zone and a zone around the novel object. The total distance moved (cm), the mean velocity (cm/s), whether or not the fish entered the novel object zone (0,1), and the cumulative duration (sec) in the bottom half of the tank was recorded.

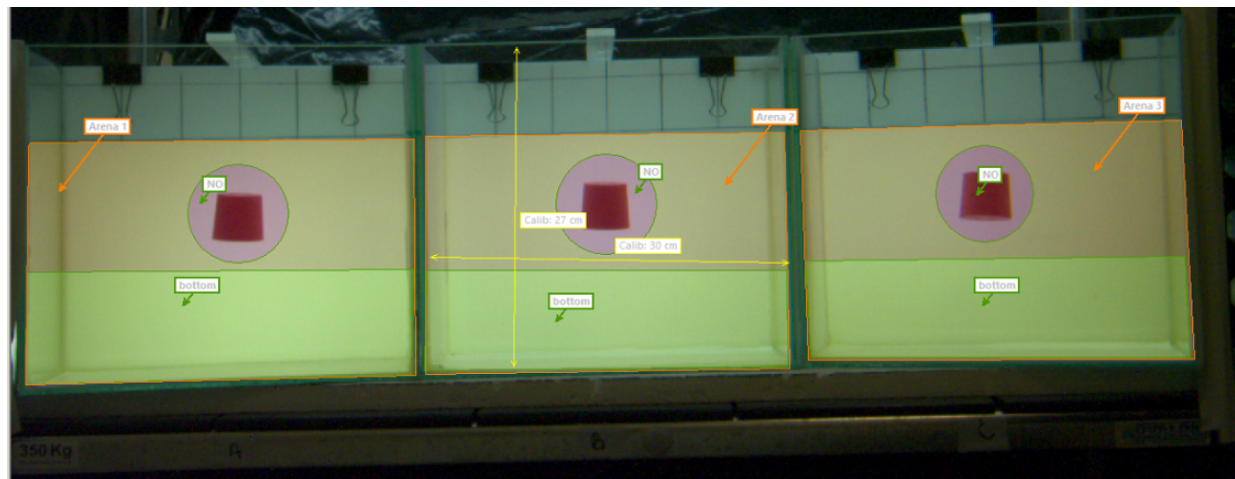


Figure 2.3: The settings for quantifying behaviours during the novel object test. Within EthoVision XT, a circular purple zone, measuring ~ 1.5 body lengths is superimposed over the novel object that hangs from the top of the tank. A green zone superimposed onto the tank indicates the bottom zone.

Mirror test

After filming the novel object assay, the mirror test was conducted (Figure 2.4). The novel object and opaque barrier were removed, and live-tracking started 10-30sec once the water had settled. The total

distance moved (cm), velocity (cm/s), the frequency of times the fish entered into mirror zone and the time spent interacting with the mirror (sec) was recorded.

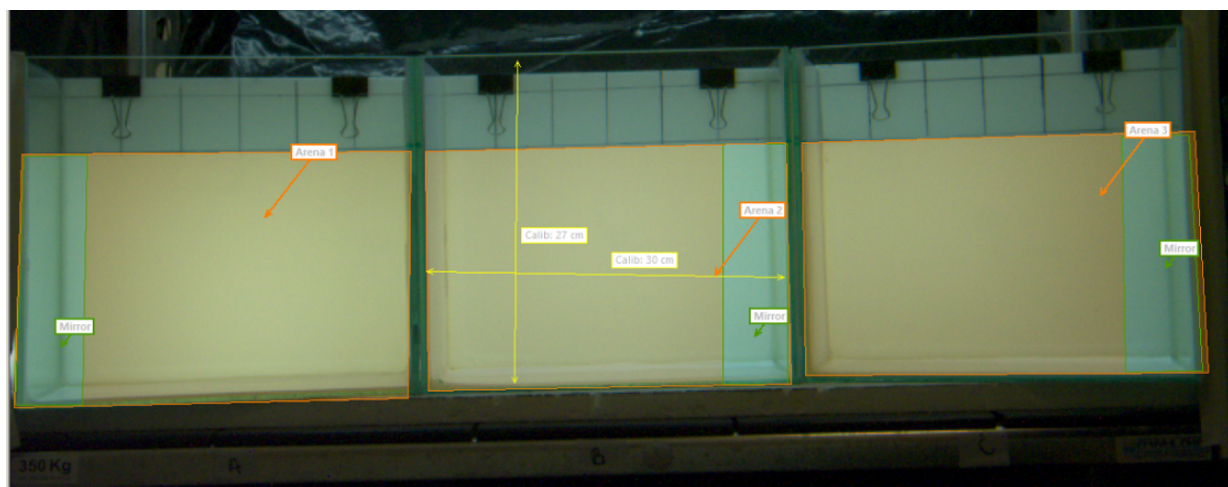


Figure 2.4: The settings for quantifying behaviours during the mirror test. Within EthoVision XT, a blue zone measuring ~ 5 cm wide, along the side of the tank indicates the mirror zone, where the flat mirror was located.

Faulty tracking

Tracking was occasionally lost when fish velocity was lower than 1.5cm/s or stopped moving and settled on the bottom of the tank, this was noted and was corrected for post assay. Tracking continued if the fish increased speed or discontinued resting. Other tracking errors that occurred during filming were manually identified and fixed using EthoVision's 'integrated vision function', and were either corrected or the video was re-analysed using different detection settings. During the behavioural testing of the F_1 generation, there were eight instances during the novel object assay where tracking was unable to be accurately established due to experimental error. All cases were therefore excluded from subsequent analyses; three controls (two male, one female), one 0.3ppb male, three 3ppb fish (one male and two females) and one 30ppb female.

2.2.7 Statistical analysis

All statistical analyses were conducted in R, version 3.5.0 (R Core Team 2018). All analyses described below contained the two main predictors, treatment (i.e. controls, 0.3ppb, 3ppb, 30ppb) and sex (male and female) and an interaction term. The interaction term was removed if non-significant and the data was re-analysed without it. All F_0 behavioural measures were analysed with linear models (LMs) or generalised linear models (GLMs). All F_1 behavioural measures were analysed with linear mixed effects models (LMMs) or

with generalised linear mixed effects models (GLMMs) using the package ‘*lme4*’ version 1.1-1.3 (Bates et al. 2015), with family ID included as a random effect to account for genetic influences (with some exceptions, see Appendix V). Specific details on each behavioural measure analysed can be found in Appendix V.

For all models, results are presented with estimates and 95% confidence intervals (CI). Confidence intervals were calculated using the ‘*confint*’ function. For reference, 95% CI that do not include zero are statistically significant. The full model output, including parameter coefficients (back transformed where appropriate), parameter standard error (SE), test statistics and p-values are reported in Appendix VIII. P-values for LMMs and GLMMs were calculated using the ‘*lmerTest*’ package (Kuznetsova et al. 2017). A subset of the behavioural data is presented in text, other measures can be found in Appendix VI. See Appendix III for specific sample sizes used for each assay, per treatment and sex (sample size ranges are given in figure legends). Descriptive statistics are reported as means \pm SE in Appendix VI.

Lastly, in some instances of F_1 analyses, family level variation was suspected of influencing the model outputs, indicating a genetic component of behaviour. Therefore, likelihood ratio tests (LRT) were used to compare models with and without the random effect. The variance and standard deviation (SD) from the LMM or GLMM model output are reported in text when they occurred (models summarised in Appendix IX, Table A9.1). Family variation was also visually compared across treatments to investigate if atrazine altered the amount of variation present at the family level (Appendix IX).

2.3 Results

2.3.1 Anxiety

Changes in anxiety occurred in the lowest F_0 treatment (0.3ppb) in the novel arena assay. F_0 fish from the 0.3ppb treatment spent significantly longer in the bottom zone of the novel arena compared to controls (Est. 83.03 [1.18, 164.87 CI]; Figure 2.5 A), with no other treatment or sex differences occurring. In the F_1 's there was significant sex difference; males spent less time in the bottom zone (Est. -86.78 [-128.30, -45.25 CI]) than females (Figure 2.5 B). F_1 controls (both males and females), on average spent less time in the bottom zone than all other treatment groups (Figure 2.5 B), however no significant treatment effects were observed owing to the large variance produced at the family level (Appendix IX). The family level variance was 11425 sec with a SD of 106.9 sec and explained ($\chi^2 = 33.6$, $df = 1$, $p < 0.001$) a significant portion of variation present in the LMM model (Appendix IX). Nonetheless, there appears to be a suggestive trend of more time spent in the bottom zone with fish from treated males.

2.3.2 Exploration

In the novel arena assay, F_0 fish from the 0.3ppb treatment were significantly less exploratory than controls (Est. 15.12 [2.94, 27.30 CI]; Figure 2.5 C), but no other treatment effects were observed. Sex differences were marginally non-significant, with males tending to be more exploratory than females (Est. -7.705 [$-16.17, 0.76$]). In the F_1 's there was no difference amongst treatments in exploratory behaviour, but there was a significant sex difference observed, with males being more exploratory than females (Est. -12.43 [$-20.77, -4.09$ CI]; Figure 2.5 D).

2.3.3 Activity

There was a significant sex difference in activity observed in both generation during the novel arena assay, with males travelling a greater total distance than females (F_0 males Est. 532.19 [198.37, 866.02 CI]; F_1 males Est. 1255.64 [892.14, 1619.14 CI]); however, no treatment differences were observed in either generation in the total distance moved (Figure 2.6 A and B).

During the novel object assay, males from both generations travelled a significantly greater total distance than females (F_0 males Est. 631.47 [284.32, 978.62 CI]; F_1 males Est. 1257.18 [917.76, 1596.60 CI]). But no treatment differences were observed in either generation (Figure 2.6 C and D).

During the mirror test assay, males travelled a significantly greater total distance than females (F_0 males Est. 700.55 [400.53, 1000.57 CI]; F_1 males Est. 689.30 [393.78, 984.89 CI]). In the F_0 's there was no treatment effect observed (Figure 2.6 E). However, activity levels were significantly different amongst F_1 fish from the atrazine treatment compared to controls. F_1 fish from both 0.3ppb (Est. -110.63 [$-175.35, -45.91$]) and 30ppb (Est. -95.98 [$-160.24, -31.72$]) travelled a significantly less total distance than controls (Figure 2.7 F). F_1 3ppb fish also travelled less than controls, but this observation was marginally non-significant (Est. -56.32 [$-120.25, 7.61$] Figure 2.7 F).

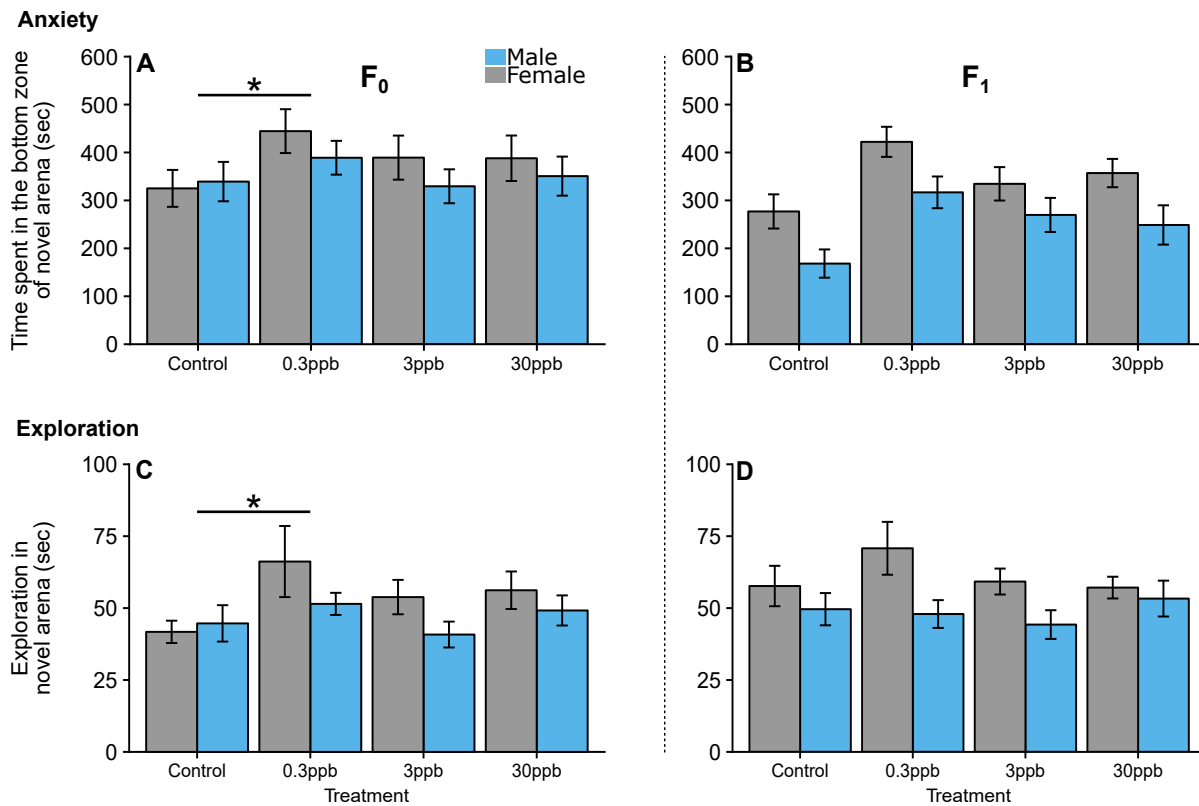


Figure 2.5: Measures of anxiety and exploration of atrazine exposed F₀ fish (left) and F₁ offspring from exposed males (right). (A) and (B) the time spent in the bottom zone of the novel arena (sec) during the novel arena test; (C) and (D) the standard deviation of exploration (sec) during the novel arena test. Bars represent means, with error bars representing standard errors of the mean. An asterisk indicates a significant difference between controls and atrazine treatment ($p < 0.05$). For each sex and treatment, sample sizes for F₀'s ranged from $n=9-12$; F₁'s ranged from $n=18-30$. Total sample size for F₀'s were $n=93$; F₁'s were $n=190$, with $n=3$ families per treatment.

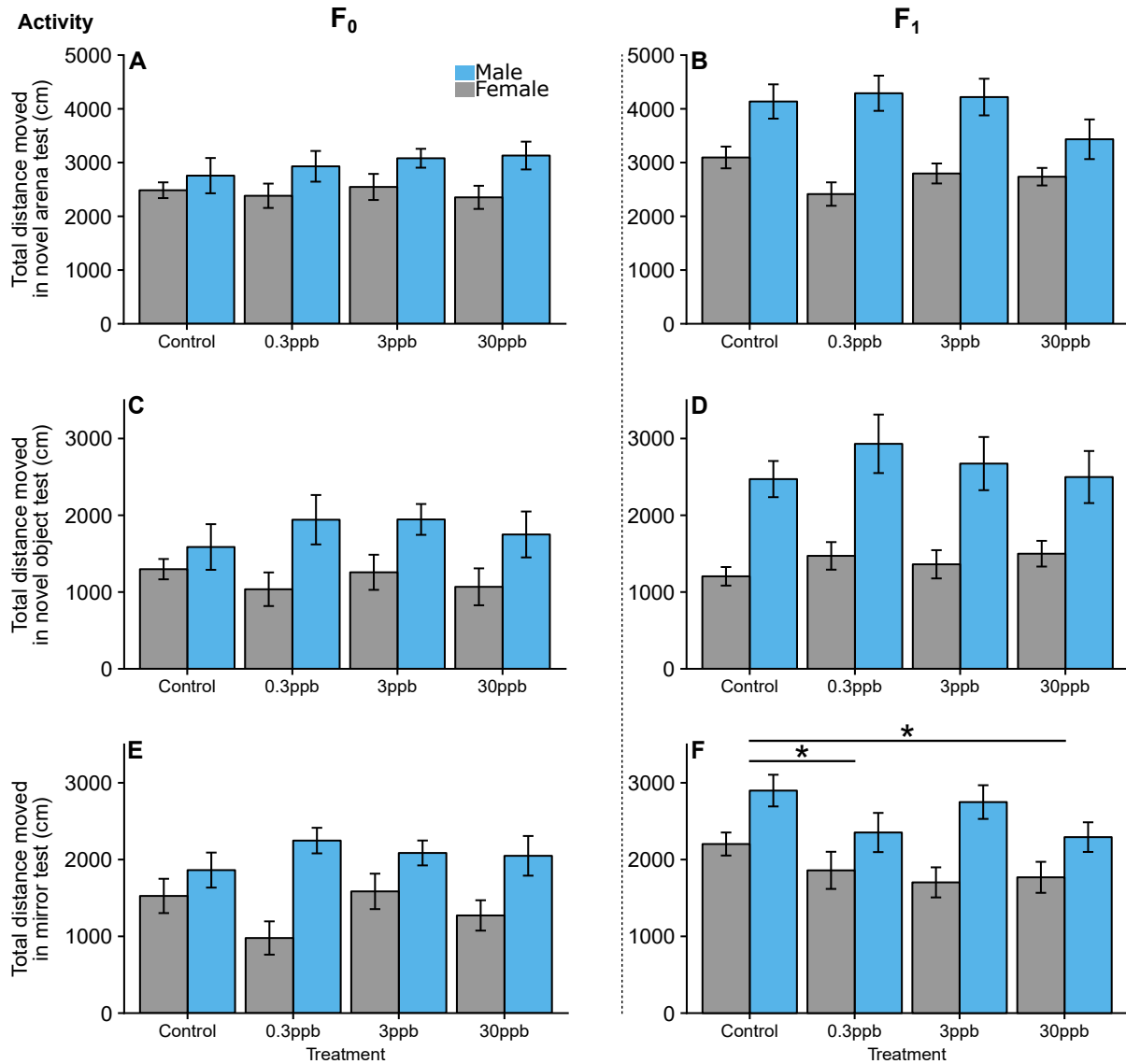


Figure 2.6: Measures of activity of atrazine exposed F₀ fish (left) and F₁ offspring from exposed males (right). The total distance moved (cm); (A) and (B) during the novel arena test; (C) and (D) during the novel object test; (E) and (F) during the mirror test. Bars represent means, with error bars representing standard errors of the mean. An asterisk indicates a significant difference between controls and atrazine treatment ($p < 0.05$). For each sex and treatment, sample sizes for F₀'s ranged from $n=9-12$; F₁'s ranged from $n=18-30$. Total sample size for F₀'s were $n=93$; F₁'s were $n=190$, with $n=3$ families per treatment.

2.3.4 Boldness

Developmental atrazine exposure altered measures of boldness in both the F_0 and F_1 generation in the novel arena assay (Figure 2.7 A and B). Direct F_0 atrazine exposure significantly increased latency in the 30ppb concentration (Est. 2.36 [1.17, 6.96 CI]) and non-significantly increased latency in the 3ppb treatment (Est. 2.36 [-1.03, 5.75 CI]; Figure 2.7 A), but no significant treatment differences were observed in the 0.3ppb treatment (Figure 2.7 A). Sex itself did not influence overall latency in the F_0 's, however several treatment:sex interactions were observed. F_0 males from both the 3ppb (Est. -3.88 [-13.46, -1.12 CI]) and the 30ppb (Est. -5.39 [-18.73, -1.55 CI]) treatment showed a significantly lower latency in comparison with females from the same treatment when compared to controls. Differences between males and females from the 0.3ppb treatment were marginally non-significant (Est. -3.54 [-13.20, 1.02 CI]). On average, F_0 female latency increased with increasing atrazine concentration, whereas treated F_0 males exhibited lower latency than controls, but then plateaued regardless of the atrazine concentration (Figure 2.7 A).

In the F_1 generation, all treatments exhibited significantly higher latency compared to controls (0.3ppb Est. 6.16 [2.01, 18.73 CI]; 3ppb Est. 4.07 [1.34, 12.30 CI]; 30ppb Est. 3.71 [1.22, 11.25 CI]; Figure 2.7 B), but no interactive effects were observed. Males, regardless of treatment took significantly less time to enter the top zone compared to females (Est. -2.3 [-2.32, -2.27 CI]). On average, F_1 female latency was highest in the lowest treatment (0.3ppb atrazine) and then declined with increased atrazine concentration. Control F_1 females displayed the lowest latency on average compared to all other females (Figure 2.7 B). The variance produced at the family level was 1.85 sec with a SD of 1.36 and significantly influenced the model ($\chi^2 = 157.4$, $df = 1$, $p < 0.001$; Appendix IX).

In both generations there was no treatment effect observed in whether or not an individual approached the novel object (Figure 2.7 C and D). In the F_0 's there was no sex effect observed, whereas in the F_1 generation, males approached the novel object significantly more so than females did (Est. 0.87 [0.76, 0.93 CI]; Figure 2.7 D).

2.3.5 Aggression

There was a non-significant trend of reduced time spent interacting with the mirror with increasing atrazine concentration amongst the F_0 males, however no specific treatment effects nor sex differences were observed (Figure 2.7 E). Significant treatment effects were observed amongst the F_1 's, where F_1 fish from the 0.3ppb (Est. -110.63 [-45.91, -3.35 CI]) and 30ppb (Est. -95.98 [-31.72, -2.93 CI]; Figure 2.7 F) treatments spent significantly less time in the mirror zone compared to controls. Fish from the 3ppb treatment also

spent less time in the mirror zone but this was marginally non-significant (Est. -56.32 [$-120.25, 7.61$ CI]; Figure 2.7 F). There was also a clear sex difference in the F_1 's, with males spending significantly more time in the mirror zone than females (Est. 74.42 [$27.96, 120.89$ CI]).

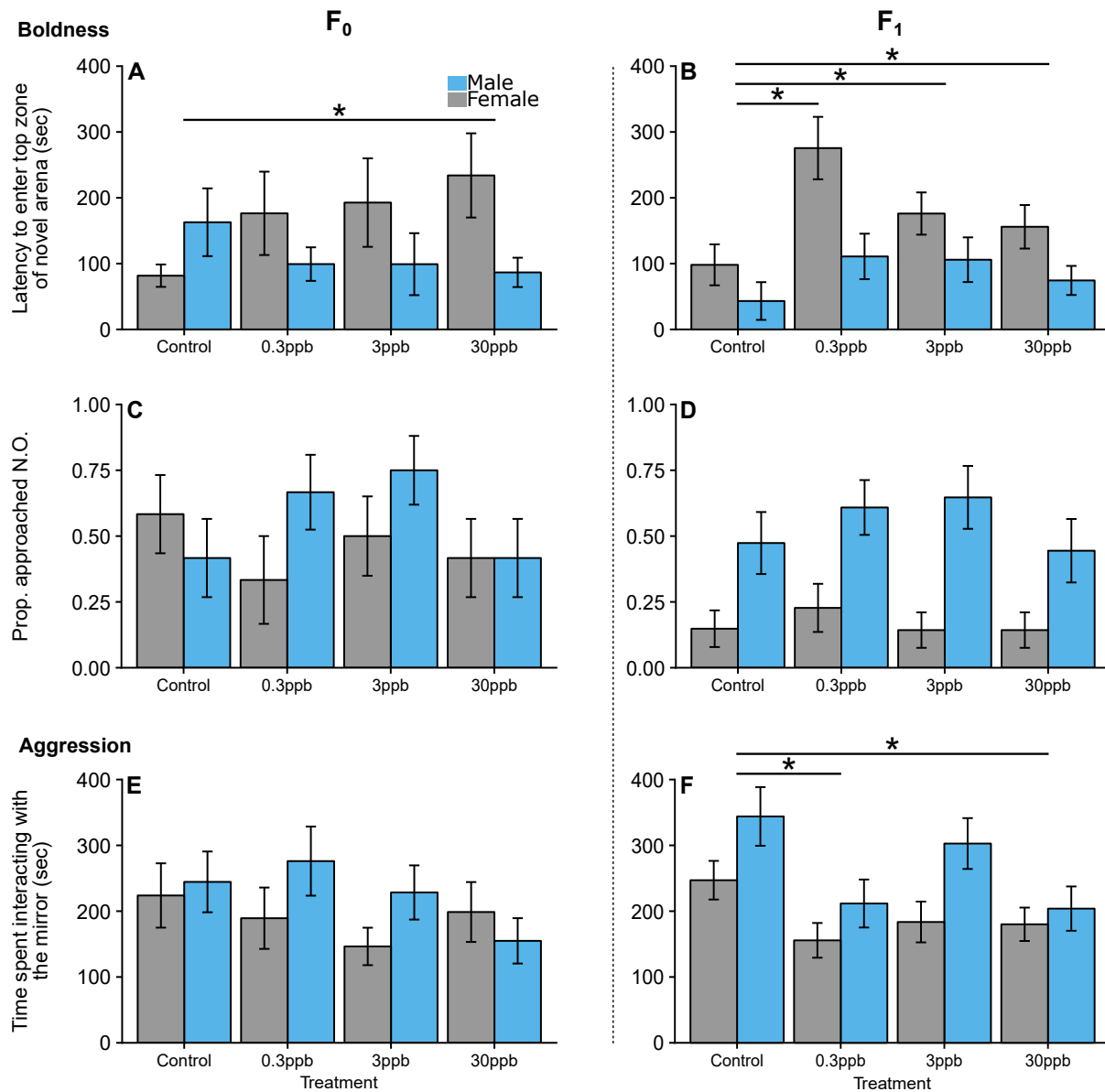


Figure 2.7: Measures of boldness and aggression on atrazine exposed F₀ fish (left) and F₁ offspring from exposed males (right). (A) and (B) the latency to enter the top zone during the novel arena test (sec); (C) and (D) the proportion of fish that approached the novel object; (E) and (F) the time spent interacting with the mirror (sec). Bars represent means, with error bars representing standard errors of the mean. An asterisk indicates a significant difference between controls and atrazine treatment ($p < 0.05$). For each sex and treatment, sample sizes for F₀'s ranged from $n=9-12$; F₁'s ranged from $n=18-30$. Total sample size for F₀'s were $n=93$; F₁'s $n=182-190$, with $n=3$ families per treatment.

2.4 Discussion

2.4.1 Revisiting aims

The aims of this study were two-fold: first, to identify if environmentally relevant concentrations of atrazine exposure during zebrafish juvenile development altered the personality phenotype in adulthood, and secondly, if the offspring of exposed males also showed an altered phenotype. I predicted that direct and indirect exposure to atrazine would increase anxiety levels and decrease boldness, aggression and exploration, and potentially alter activity levels in both generations. Furthermore, I predicted that increases in anxiety would be female biased.

Overall, several behavioural measures were altered under juvenile atrazine treatment in adulthood and manifested in the adult offspring, though not all effects were consistent across all doses. Several significant changes to behaviour occurred in either generation. Changes to the latency to enter the top zone of the novel arena (boldness) were particularly altered. F_0 females exhibited increased latency with increasing concentration and F_0 males exhibited decreased latency in a threshold-like pattern compared to controls. In the following generation, F_1 females exhibited a significant increase in latency that decreased with increasing concentration, but there was little change amongst F_1 males compared to controls. The time spent interacting with the mirror (aggression), and the total distance moved during the mirror test (activity) were significantly reduced amongst the F_1 progeny in the 0.3ppb and 30ppb treatment groups, with some evidence for reduced aggression but no evidence for altered activity in the mirror test amongst the F_0 's. Lastly, the time spent in the bottom zone (anxiety), and the time spent exploring the novel arena (exploration) were significantly reduced in the F_0 0.3ppb group only, while in the F_1 's there was some evidence of alterations in anxiety and exploration. These results suggest that atrazine disrupts aspects of anxiety, boldness, aggression and exploratory personality phenotypes at environmentally relevant concentrations, and these effects can be transferred via sperm to the following generation.

2.4.2 Effects of atrazine exposure on boldness, anxiety and exploration

One of the strongest effects of the atrazine treatment was observed in the latency to enter the top zone (latency) of the novel arena (a measure of boldness; Figure 2.7 A and B). Changes in latency occurred in different directions depending on the sex, with phenotypes more pronounced in females (in both generations) than males. F_0 females exhibited increased latency with increasing atrazine concentration compared to controls, whereas F_0 males displayed reduced latency across all treatments compared to controls. This phenotype was inconsistent between generations, female F_1 offspring from atrazine treated males exhibited

decreased latency with increased atrazine concentration (though all higher than control females).

In an ecological setting, entering the top portion of the water column where little vegetation occurs likely carries a higher risk of avian predation and may increase conspicuousness of the individual to fish predators (Werner et al. 1983; Rypel et al. 2007; Pink et al. 2018). Avian predation is not well documented in wild zebrafish, but populations tend to live in tandem with the common kingfisher (*Alcedo atthis*) and the Indian pond heron (*Ardeola grayii*), which are hypothesised to be the main avian predators (Spence et al. 2006). When confronted by an avian predator simulation above the tank (e.g. a bird silhouette or a black dot increasing in size, simulating an approaching bird), defensive behaviours such as bottom dwelling are triggered (Luca et al. 2012a; Luca et al. 2012b). The costs of entering the top portion of the water column may be offset by access to resources such as surface dwelling invertebrates, which make up the largest portion of wild zebrafish diet (Pitcher et al. 1988; McClure et al. 2006; Arunachalam et al. 2013; but also see; Spence et al. 2007). If the highest portion is perceived to carry a higher risk, then changes in latency observed from atrazine treated fish might reflect a distortion in an individual's ability to either assess risk, or a change in propensity to take risks regardless of the risk entailed. Based on this data, it is difficult to tease apart these two components as they are theoretically both tied to one another, though the consequences of either remain the same. The lack of difference (amongst the F_0 's) in the time spent in the top zone or in the frequency to enter the top zone might indicate a stronger hypothesis for the latter. However, the non-significant reduction amongst the treated F_1 's might suggest more weight for the former. Changes in perceived risk is not the only factor impacting survival that needs to be considered (see General Discussion).

In addition to changes in boldness, developmental atrazine exposure altered zebrafish anxiety and exploratory phenotypes in the lowest dose (Figure 2.5 A and C). Previous findings have suggested that atrazine exposure produces an anxiety-like behaviour in fish (Steinberg et al. 1995) and in rodents (Lin et al. 2013; Walters et al. 2015). Changes in exploration profiles have also been observed in rodents (Belloni et al. 2011) as well as in insects (Figueira et al. 2017), but not so in fish (Schmidel et al. 2014).

Risk assessment of the environment is crucial during foraging and exploration (Pitcher et al. 1988). Individuals that under assess risk, or are generally bolder, increase their susceptibility to predation (Smith et al. 2008; MacPherson et al. 2017). Whereas over assessment or more anxious individuals may increase long-term survival, but may place costs on other functions, such as the ability to forage (Wolf et al. 2007; Biro et al. 2008b). Furthermore, in a given context, individuals that are less willing to explore their environments may be at risk of starvation if resources are generally scarce, limit the ability to find suitable mates, or to seek out pertinent information relating to survival if new threats arise (e.g. increased predator

prevalence or territorial intruders; MacPherson et al. 2017). Populations naturally contain an assortment of individuals that display risk-taking or risk averse like personality phenotypes (Bierbach et al. 2017), with each strategy hypothesised to be maintained by life-history trade-offs (i.e. between early vs later life reproduction; Schuett et al. 2015). Populations may also differ in the extent of phenotypes produced, depending on factors such as differences in predation rates or food and mate availability, and may change across space and through time if these factors also change (Bierbach et al. 2015; Barbosa et al. 2018b). It stands to reason that the consequences of EDCs influencing risk assessment, anxiety or exploratory behaviour are wide reaching at the individual level by influencing their probability of survival (MacPherson et al. 2017), and may also extend to the population level by pushing groups of exposed individuals in one direction or possibly pushing each sex in differing or converging directions (or by disrupting signals; Senior et al. 2014; White et al. 2017). For instance, diminished behavioural variation commonly found in exploited fishing populations, due to the tendency of bolder individuals to take bait, have led to a skewed distribution of behavioural phenotypes comprising a greater proportion of anxious individuals (Biro et al. 2008a).

2.4.3 Possible mechanisms behind altered boldness, exploration and anxiety

Mechanistic explanations behind this distorted boldness, exploratory and anxious phenotype are difficult to pinpoint, but several neurotransmitter systems have been implicated in atrazine exposure across taxa (fish, mammals and insects), including the cholinergic (Schmidel et al. 2014), dopaminergic (Walters et al. 2015; Figueira et al. 2017) and serotonergic (Wirbisky et al. 2015) systems. A cell culture study found that atrazine causes neuronal damage and augments neurotransmitter release (Peña-Contreras et al. 2016). In particular, these aforementioned studies have found greater distortion towards females than males. For example, Wirbisky et al. (2015) found that there was a significant reduction in neurotransmitter turnover of serotonin (5-HT; i.e. the rate of 5-HT synthesis, indicative of reduced serotonergic activity), in adult zebrafish females exposed as embryos, but not in males. Brain transcriptome studies also implicate several genes involved in the serotonergic pathway, steroidogenesis and neurotransmission within the hypothalamus-pituitary-interrenal and hypothalamus-pituitary-thyroid axes (Weber et al. 2013; Wirbisky et al. 2015; 2016a; 2016c).

Another possible reason for a distortion in risk assessment might be a result of atrazine's immunotoxic action. Atrazine, as well as a neuroendocrine disruptor, is also immunotoxic acting via the dysregulation of specific cytokine genes, leading to an immunosuppression profile (Devos et al. 2003; Wang et al. 2011; Kirsten et al. 2017). Studies in mammals and now recently in fish have shown a relationship between immune system function and neural activity that reflects changes in behaviour (Moon et al. 2015; Filiano

et al. 2016; Kirsten et al. 2018). Cytokines are a broad group of protein molecules functioning within the immune system (for review see; Dinarello 2007).

Studies have shown that cytokine expression is related to individual behavioural patterns (e.g. anxiety-like behaviour in mice; Moon et al. 2015; Kirsten et al. 2018). Moreover, cytokines appear to be involved or associated with sociality, social status (through contests of aggression), defensive behaviours and exploratory behaviour (among others; Moon et al. 2015; Filby et al. 2010a; Kirsten et al. 2018). Fish with an immune suppression cytokine profile exhibit alterations in these aspects of behaviour (Moon et al. 2015; Kirsten et al. 2018). Atrazine at high concentrations (1000ppb) has been shown to disrupt defensive behaviours which include shoaling (Schmidel et al. 2014) as well as reduce male-male competition, and interactions with other conspecifics (Shenoy 2014). It is also hypothesised that immune-suppressed fish might lead to an erroneous ability to assess risk (Kirsten et al. 2018). Given that neural disruption is frequently observed at these same concentrations used in this study (Weber et al. 2013; Wirbisky et al. 2015; Wirbisky et al. 2016a; Wirbisky et al. 2016c), it stands to reason that immunodisruption could also occur, and that these systems together may influence behaviour.

2.4.4 Diminished aggression response

Juvenile exposure to atrazine did not result in a statistically clear reduction in time spent interacting with the mirror; however, there appeared to be a suggestive negative trend as concentration increased amongst the F_0 males (Figure 2.7 E). Interestingly, the suggestive trend observed in the F_0 males was significantly pronounced in the F_1 males, though not in the similar dose-dependent manner suggesting a non-monotonic effect (Figure 2.7 F). Furthermore, the reduction in mirror interaction in the F_1 males was paralleled with reductions in their activity levels (Figure 2.6 F). Previous studies have found impairment in reproductive behaviours after direct atrazine exposure, in particular with reductions in male-male aggression and sexual displays (Hayes et al. 2002; Hayes et al. 2010; Shenoy 2012; Shenoy 2014); however, this is the first study to document altered aggression as a function of an individual's personality, due to paternal atrazine exposure.

Mechanisms behind this change in aggression via atrazine exposure are conflicting throughout the literature. Primarily, it is hypothesised that atrazine influences levels of androgens, testosterone and 11-ketotestosterone (11-KT; in fish), which are important for mediating aggression (Bell 2001). Atrazine exposure is hypothesised to increase the expression of aromatase, an enzyme that converts testosterone to oestrogen (Hecker et al. 2005; Fan et al. 2007; Hayes et al. 2010; Fa et al. 2013). Atrazine has been shown to lower testosterone levels in mammals (Trentacoste et al. 2001; Gely-Pernot et al. 2015), amphibians (Hecker et al. 2005) and

lower both testosterone and 11-KT in goldfish (Spanò et al. 2004), though this finding is not always consistent (Wirbisky et al. 2016a). In addition, the effects through aromatase expression are less clear, especially in fish when compared to other vertebrates (Papoulias et al. 2014).

In the wild, aggression in zebrafish is important for establishing dominance hierarchies, which in the wild enables the monopolisation of foraging resources and the maintenance of territories for spawning sites (Spence et al. 2008). In addition, more aggressive zebrafish males exhibit higher reproductive success (Ariyomo et al. 2012). The consequences of reduced aggression may therefore limit the ability to compete for access to foraging sites and limit the ability to maintain territories. Aggression is also important in females, as either sex can establish a dominance hierarchy (Grant et al. 1992, Spence et al. 2008). Males and females also tend to exhibit similar levels of aggression (Ariyomo et al. 2013a; Ariyomo et al. 2013b), though this was not observed in this study. In this study, F_0 female aggression levels tended to decrease with increasing concentration, though F_0 30ppb females appeared to be more aggressive than their male counterparts. In the F_1 's, control females exhibited the highest levels of aggression compared to other F_1 females. Previous works have identified similar reduced aggression responses from other EDCs: chronic BPA (Wang et al. 2015), ethinyloestradiol (Bell 2001) and nonylphenol (Xia et al. 2010).

2.4.5 Potential of non-genetic environmental inheritance

The data shown here demonstrates that paternal exposure to atrazine (at environmental doses) has the potential to produce intergenerational effects on behaviour. Indeed, as similar (or dose dependent) changes in boldness and aggression (and to some extent, anxiety and exploration) occurred across differing concentrations, there is a stronger implication that effects were transferred via exposed sires. Moreover, it appears that intergenerational behavioural changes associated with atrazine are sex dependent.

In general, the process of producing a phenotype in zebrafish offspring is a complex combination of the inherited genetics from both parents, maternal factors deposited in the egg (Ariyomo et al. 2013a), experience during life and possibly environmentally induced epigenetic factors derived from paternal ancestors (e.g. the global methylome pattern; Jiang et al. 2013; Potok et al. 2013). Recent work has shown that direct atrazine exposure can interfere with the epigenetic machinery itself (e.g. regulation and activity of *dmnt* expression and maintenance DNMT respectively), and by altering global DNA-methylation levels (Wirbisky-Hershberger et al. 2017). However an epigenetic inheritance explanation requires transgenerational observations i.e. the effect must be observed in the first generation where the germ line was not developmentally exposed (Szyf 2015; Jacobs et al. 2017). As this study only investigated intergenerational effects (i.e from F_0 to F_1), it is not possible to discern whether the effect of the atrazine on the F_1 's was

an effect of the ‘experience’ of exposure to atrazine of the F_0 's (i.e. the distortion to neural pathways, leading to an alteration in behaviour, that then lead to a change in the germ line) or whether it was the direct exposure of the F_0 's developing germ cells (the future cells of F_1 's), that gave rise to the changes in behavioural phenotypes observed in the F_1 's. Nonetheless, these results suggest that the transgenerational effects on behaviour may be possible. Previous research has shown that atrazine (at high doses) can induce transgenerational effects of disease in mice (Hao et al. 2016; McBirney et al. 2017), and indeed it appears that disease phenotypes can be exacerbated with each generation (McBirney et al. 2017). Timing and logistical reasons did not allow transgenerational effects to be tested for, but should an interesting prospect for future studies.

Differences in behaviours observed between each generation (e.g. changes in boldness Figure 2.7 A and B), are also likely to be a symptom of atrazine causing a germ line epigenetic effect (that would be passed to their progeny) in combination with a direct effect on other (somatic) tissues in the F_0 's. Whereas in the F_1 's (and beyond), epigenetic changes in somatic cells derive their origin from the germ cells (Jacobs et al. 2017). This effect is more readily witnessed with increasing generation distances, i.e. F_1 to F_3 , rather than intergenerationally. For instance, bisphenol-A exposure of mice *in utero* lead to dampened social interactions in F_1 juveniles, but juvenile social interactions significantly increased in the F_2 - F_4 generations (Wolstenholme et al. 2012). Furthermore, as males were randomly selected for breeding, it is not possible to discern whether or not the phenotype of the offspring were similar to the father.

2.4.6 Genetic influence

Several behavioural measures showed signs of a strong genetic component, i.e. there were large amounts of family level variation present (Appendix IX). Environmental inheritance studies utilise families (the inclusion of individuals from the same parents) to limit the genetic variability and to emphasise an epigenetic effect if it was passed on. If an inherited effect was present in the F_1 's, the effect should supersede the variation between families of the same treatment. In this respect, the lack of the number of families was a limitation in this study. In some cases an inherited effect was observed despite the presence of family level variation, such as in the latency to enter the top zone (Appendix IX, Table A9.1). However, there were several endpoints where the genetic component superseded an environmental inheritance effect, for instance in the time spent in the bottom zone of the novel arena and novel object assays (Appendix IX). Overall, F_1 offspring from atrazine treated fathers showed more variation at the family level between each treatment level (0.3ppb, 3ppb and 30ppb), than the family level variation observed between F_1 control fish and F_1 treated fish. However, there was no indication that atrazine treatment influenced the level of variation, due to a lack of

a patterning in rank orders. Therefore the variation present at the family level is likely more attributable to genetic differences, rather than atrazine (Appendix IX, Table A9.1).

2.4.7 Conclusion

In conclusion, the aims of this study were to determine if direct and paternal exposure to atrazine altered personality phenotype in adulthood of zebrafish (*Danio rerio*). I found evidence for changes in boldness, aggression, anxiety and exploratory personality phenotypes due to direct and paternal atrazine exposure. Furthermore some behavioural changes exhibited a sex-specific interaction (e.g. stronger effects in females for changes in boldness), while the majority of behavioural effects did not (e.g. aggression, anxiety and exploration). Personality traits can be associated with fitness costs and benefits (Smith et al. 2008; MacPherson et al. 2017), therefore understanding how common environmental contaminants may shape behavioural phenotypes might be potentially informative to understand consequences up to the population level. The findings from this study contribute to the growing body of literature that finds that EDCs have the potential to affect non-reproductive behaviours and effects can be propagated beyond the generation that was directly exposed, suggesting potential transgenerational effects. From the basis of behavioural alterations observed in this study, the following chapter (Chapter 3) investigates mRNA expression differences in the brains of F₁ progeny. Specifically, genes involved in regulating the serotonergic, which are key genes thought to be underpinning aspects of anxiety and aggression.

Chapter 3

The influence of atrazine on the serotonergic system in F₁ progeny

3.1 Introduction

Endocrine disrupting chemicals are reported to induce a variety of aberrant behaviours (see reviews Clotfelter et al. 2004; Söffker et al. 2012). How EDCs mechanistically exert their effects (e.g. in the brain) that in turn, alter behaviour of an organism is one of the fundamental challenges of EDC research (see review Nesan et al. 2018). Any complex behaviour, including animal personality, is understood to be underpinned by a combination of molecular mechanisms such as changes in gene expression, hormones, neurotransmitter levels and molecular machinery that mediates between these inputs (e.g. Whitfield et al. 2003; Filby et al. 2010b; Thörnqvist et al. 2015; Wiese et al. 2018). Alterations in gene expression (that then leads to changes in behaviour) from EDC exposure may be a result of how an EDC interacts with hormone receptors, interferes with hormone metabolism, alters structuring of the brain during developmental or possibly by altering the epigenetic machinery that controls gene expression (see review Nesan et al. 2018). For example, one specific effect of the EDC, bisphenol-A (BPA; see review Nesan et al. 2018), causes hyperactivity in larval zebrafish, via an induced up-regulation of aromatase transcripts (the aromatase enzyme converts androgens to oestrogens) mediated by agonism of androgen receptors in the developing hypothalamus, resulting in an unexpected early neurogenesis (Kinch et al. 2015; Le Fol et al. 2017).

As discussed in the previous chapter, non-genetic factors may play a role in facilitating behavioural phenotypes through to the following generation (see Chapter 2), and as a consequence, may influence the underlying genetic components that reflect changes in expression of the behavioural phenotype (Burton

et al. 2014; Dias et al. 2014). In the previous chapter I showed that the offspring of males exposed to the neuro-endocrine disruptor, atrazine (at concentrations 0.3ppb, 3ppb and 30ppb) during juvenile development exhibited reduced aggression (0.3ppb and 30ppb significantly reduced), and also showed some evidence that paternal atrazine may also affect anxiety (see Chapter 2). Aggression plays a crucial role across the animal kingdom (Bell et al. 2009) and is characterised as antagonistic interactions between conspecifics (Réale et al. 2007). Animals use aggression as a means to outcompete other individuals for resources such as food, mates, territory or social rank (Spence et al. 2008). Anxiety on the other hand, is anticipatory fear when uncertainty of a potential threat is high (Maximino et al. 2012). In the presence of uncertainty of an unknown threat, animals behave in a manner that maximises their self preservation, but the strength or duration of these behaviours will tend to differ between individuals and across contexts, such as a novel environment (Thomson 2017).

A first step to understand the functional outcomes of atrazine exposure on behaviour is to investigate the associations exposure has on molecular endpoints. Recent transcriptomic studies from adult zebrafish embryonically exposed to atrazine have highlighted the serotonergic system as a potential target of developmental atrazine exposures (Weber et al. 2013; Wirbisky et al. 2015; 2016). Moreover, Wirbisky et al. (2015) demonstrated that atrazine exposure altered serotonin (5-HT) turnover in adult females brains, indicating increased serotonergic activity. 5-HT is one of the primary neurotransmitters present in the brain and has been implicated in the facilitation of anxiety and aggression behaviours (among a myriad of other endpoints; see review Backström et al. 2017). In addition, developmental atrazine exposure has been previously shown to alter anxiety-like behaviours in rodents (Lin et al. 2013; Walters et al. 2015) and aggression in fish (Shenoy 2014). The aims of this study were two-fold: firstly, to investigate if atrazine influenced the level of transcript number in the offspring of exposed males; and secondly, if changes in transcript number occurred, are these related to changes observed in personality, particularly aggression and anxiety behavioural phenotypes?

Understanding the way in which 5-HT regulates anxiety or aggression appears complex (Clotfelter et al. 2007; Maximino et al. 2012; Maximino et al. 2013; Herculano et al. 2014), even in different strains of zebrafish (Barbosa et al. 2018a). The role 5-HT plays is not straight forward, but generally, 5-HT activity tends to repress aggression (Popova 2008). In mice, the knock-out of the 5-HT transporter results in reduced aggression (Holmes et al. 2002; Heiming et al. 2013). For measures of anxiety, the understanding is less clear. For example, two distinct behavioural tests that are both considered paradigms for testing anxiety in zebrafish (novel tank compared to the light/dark test) result in altered 5-HT levels, but in opposite directions to one another (Maximino et al. 2012). The behavioural and functional endpoints of

5-HT neurotransmitter system appears to be strictly conserved across vertebrate systems (and apparently in invertebrate systems; Fossat et al. 2014) with differences occurring in the organisation in the brain (e.g. in teleosts compared to mammals; Lillesaar 2011). In comparison to other vertebrates, a duplication event in the teleost evolutionary history lead to duplications in 5-HT_{1A} receptor and the 5-HT transporters (Norton et al. 2008), leading to possible innovations and increased complexity in how the serotonergic signalling system may operate within the teleost brain.

As candidate genes, this study selected three 5-HT receptor genes *htr1Aa*, *htr1B*, and *htr2B*, as well as the two 5-HT transporter genes *slc6a4a* and *slc6a4b* for their involvement in the regulation of the serotonergic system and in anxiety and aggression behaviours. The three 5-HT receptors are G-protein-coupled receptors that function by propagating neurotransmission in the post-synaptic membrane by relaying 5-HT activity and are involved in controlling 5-HT release in the pre-synaptic membrane (Piñeyro et al. 1999; Aghajanian et al. 1972; Norton et al. 2008). The two 5-HT transporters (Wang et al. 2006) function by mediating the removal of 5-HT from the synaptic cleft and transporting it back into the presynaptic neurons, thereby regulating the strength and duration of neurotransmission (Norton et al. 2008). Drugs that target the functional proteins of these genes and expression levels of these genes are implicated in anxiety-like and dominance (via aggression) behaviours (Filby et al. 2010b; Maximino et al. 2013). Moreover single nucleotide polymorphism (SNPs) of 5-HT transporters have been associated with personality differences in other taxa (Craig et al. 2009; Miller-Butterworth et al. 2007; Müller et al. 2013; Holtmann et al. 2016). But it is not well understood to what extent these genes together influence consistent differences between individuals.

I predicted that aggressive personality would correlate negatively with mRNA transcript number of all candidate genes based on previous experiments in dominant and subordinate relationships in zebrafish and their mRNA transcripts (e.g. Filby et al. 2010b; Theodoridi et al. 2017). I predicted that an anxious personality would correlate positively with mRNA levels (in particular with *htr1Aa*; Maximino et al. 2013). I also predicted that offspring from atrazine treated males would have higher transcript number of the candidate genes compared to controls (Wirbisky et al. 2015), indicative of their higher anxiety and lower aggression behavioural profiles.

3.2 Methods

3.2.1 Phenotype selection, tissue extraction, RNA extraction and purification

After behavioural phenotyping (see Chapter 2), the two zebrafish that spent the most and the least amount of time at the bottom in the novel arena (a high and low anxious phenotype) from each treatment per family

and per sex, were selected (Appendix X). Once selected, fish were then euthanised in an ice water bath for 10 min. Fish were then dissected in PBS and whole brain tissue was extracted and immediately placed in eppendorf tubes with 50-100 μ L of RNA Later (*in vitro*) and subsequently stored at -30°C . Based on the measures of anxiety (Figure A10.1), F₁ fish from controls and from the 0.3ppb treatment were selected for qPCR analysis due to the greater difference between these treatment groups. Additionally, as there was a lack of a strong relationship found between the anxiety phenotype and the aggression phenotype (time spent interacting with the mirror; Figure A11.1), this method of selection enabled coverage of both animal personalities.

RNA was extracted using a modified protocol from the Norgen Biotek RNA kit and using Trizol reagent (Ambion). Briefly, tissues were homogenised with 1mL of Trizol reagent to separate the RNA phase from that of the protein and DNA. The RNA phase was then transferred to the Norgen column and processed according to the Norgen protocol. RNA concentration was measured using a ND-1000 nanodrop spectrophotometer. Samples were eliminated of contaminants using Turbo DNase (Invitrogen) and then re-nanodropped to confirm 260/280 ratio was between 1.8-2. The RNA was reverse transcribed to obtain cDNA using 400ng of RNA using a high capacity cDNA reverse transcription kit (Applied Biosystems). The parameters for reverse transcription consisted of one cycle of 25°C for 10 min, 37°C for 120 min and 85°C for 5 min. The cDNA was then diluted into 10ng/ μ L for qPCR.

3.2.2 Quantitative real-time PCR (qPCR)

The mRNA expression of genes encoding for *slc6a4a*, *slc6a4b*, *htr1Aa*, *htr1B*, and *htr2B* (see Table 3.1 for primers used) was determined in brain samples of adult offspring from atrazine exposed and non-exposed fathers with quantitative polymerase chain reaction (qPCR) assays. Primers were ordered from Integrated DNA Technologies (DIT; Singapore). qPCR was done with Quant studio 5 (Thermofisher). The total reaction volume was 10 μ L, of which contained 5 μ L SYBR green (Takara), 0.5 μ L forward primer, 0.5 μ L reverse primer, 2.8 μ L MilliQ water, 0.2 μ L ROX and 1 μ L of cDNA. For each primer, a gradient temperature assessment was carried out from 60-64 $^{\circ}\text{C}$ in order to find the optimal annealing temperature (Table 3.1). Each cycle of qPCR was 95°C for 2 min, 95°C for 5 sec, optimal annealing temperature of gene primer for 10 sec, 72°C for 5 sec, 95°C for 1 min, 55°C for 30 sec, 95°C for 30sec, for 30-40 cycles per plate. qPCR standards for each gene were made using pooled cDNA (of all 46 samples) at a ratio of 1:3 dilution, and serially diluted 6 times. Efficiency for all qPCR runs was between 95-105% (see Appendix XII). *b-actin* was validated and proved to be a suitable reference gene (Appendix XIII).

Table 3.1: Primers used for qPCR analysis, validated by Theodoridi et al. (2017).

Gene	Forward primer	Reverse primer	Annealing temperature
<i>b-actin</i>	5' TGTCCCTGTATGCCTCTGGT 3'	5' AAGTCCAGACGGAGGATGG 3'	62°C
<i>slc6a4a</i>	5' GTCTCCAATGGTTATCGCAGTA 3'	5' GATGACCGACAACAGGAAGT 3'	60°C
<i>slc6a4b</i>	5' GAATCCTCTGGGCTTGGTAATG 3'	5' GCTGAAGTAGACAATGGTGAAGAT 3'	60°C
<i>htr1Aa</i>	5' CAGAGCAGAGCAGCACAAG 3'	5' TGGTCTGAGAGTTCTGGTCTAATC 3'	60°C
<i>htr1B</i>	5' GTGTCGGTGCTCGTGATG 3'	5' CAGCCAGATGTCGCAGATG 3'	60°C
<i>htr2B</i>	5' GCTGCTCATTCTTCTGGTCAT 3'	5' GTTAGTGGCGTTCTGGAGTT 3'	60°C

3.2.3 Statistical analyses

All statistical analyses were conducted in R, version 3.5.0 (Team 2018). To test if paternal atrazine exposure altered mRNA transcript number in the brains of F₁ offspring, I ran several linear mixed effect models (LMMs) using the package ‘*lme4*’ version 1.1-1.3 (Bates et al. 2015). In all models described below, family ID was used as a random effect to account for genetic differences and all candidate genes were normalised by *b-actin* expression. Treatment group (control and 0.3ppb), sex (male and female) and an interaction term were predictors and gene mRNA were response variables. To investigate associations between mRNA transcripts of candidate genes and behaviour, LMMs were conducted between behavioural measures (time spent in the bottom zone and time spent interacting with the mirror) as response variables and candidate genes, treatment and an interaction term as predictors (see Appendix XVI for additional analyses where fish were categorised as the least and most anxious/aggressive). Sex as a main effect was non-significant and thus removed from further analyses between behavioural phenotypes. For these tests, after normalising, the candidate genes were further centred using the ‘*scale*’ function. If an interactive effect occurred, control and 0.3ppb offspring were additionally analysed separately.

Model residuals were checked for normality and models involving *slc6a4a* and *htr1Aa* were log transformed. The interaction term was removed if non-significant and re-analysed. Occasionally, when the interaction term influenced model outcomes (i.e. main predictors statistical significance was $p < 0.05$), but no significant interactive effect was present ($p > 0.05$), small-sample-size-corrected Akaike Information Criterion (AIC_C; Burnham et al. 2003) was run to determine the inclusion or exclusion of the interaction term using the ‘*dredge*’ function implemented in the ‘*MuMIn*’ package (Bartoń 2017). Parameters were model-averaged within 6 AIC_C units (when $\Delta\text{AIC}_C \leq 6$; Symonds et al. 2011) using the natural average method, ‘*model.avg*’ function (Burnham et al. 2003) also available in the ‘*MuMIn*’ package (Bartoń 2017). AIC_C tables are provided in Appendix XIV Table A14.1. For all models, results are presented with estimates and 95% confidence intervals (CI). Confidence intervals were calculated using the ‘*confint*’ function within base R

(Team 2018). For reference, 95% CI that do not overlap with zero are statistically significant. The full model output, including parameter coefficients, parameter standard error (SE), adjusted SE (for models that underwent AIC_C analyses), test statistics and p-values are reported in Appendix XV. P-values for LMMs were calculated using the ‘*lmerTest*’ package (Kuznetsova et al. 2017).

3.3 Results

3.3.1 mRNA expression levels between offspring from treated and non-treated fathers

Overall, *htr1Aa* mRNA was significantly down-regulated in control males compared to females (Est. -0.71 [$-1.39, -0.02$ CI]); however, the overall pattern was significantly reversed in atrazine 0.3ppb treatment offspring with *htr1Aa* mRNA significantly up-regulated in males instead (Est. 1.05 [$0.11, 2.00$ CI]; Figure 3.1 C). Similarly, control females had, on average, higher *slc6a4a* mRNA than control males. The trend was reversed between the sexes of offspring from treated fathers, suggesting an interactive treatment:sex effect, but no significant differences in either treatment, sex or an interaction between treatment and sex were observed (Figure 3.1 A). On average, there was a down-regulation of *slc6a4b* mRNA in control males compared to control females, whereas the pattern between sexes was absent in the atrazine treated offspring. After model averaging, however, no significant differences were observed (Figure 3.1 B). There was no statistically significant difference in the mRNA transcripts of *htr1B* and *htr2B* between fish from either atrazine treated or non-treated fathers (Figure 3.1 D and E respectively).

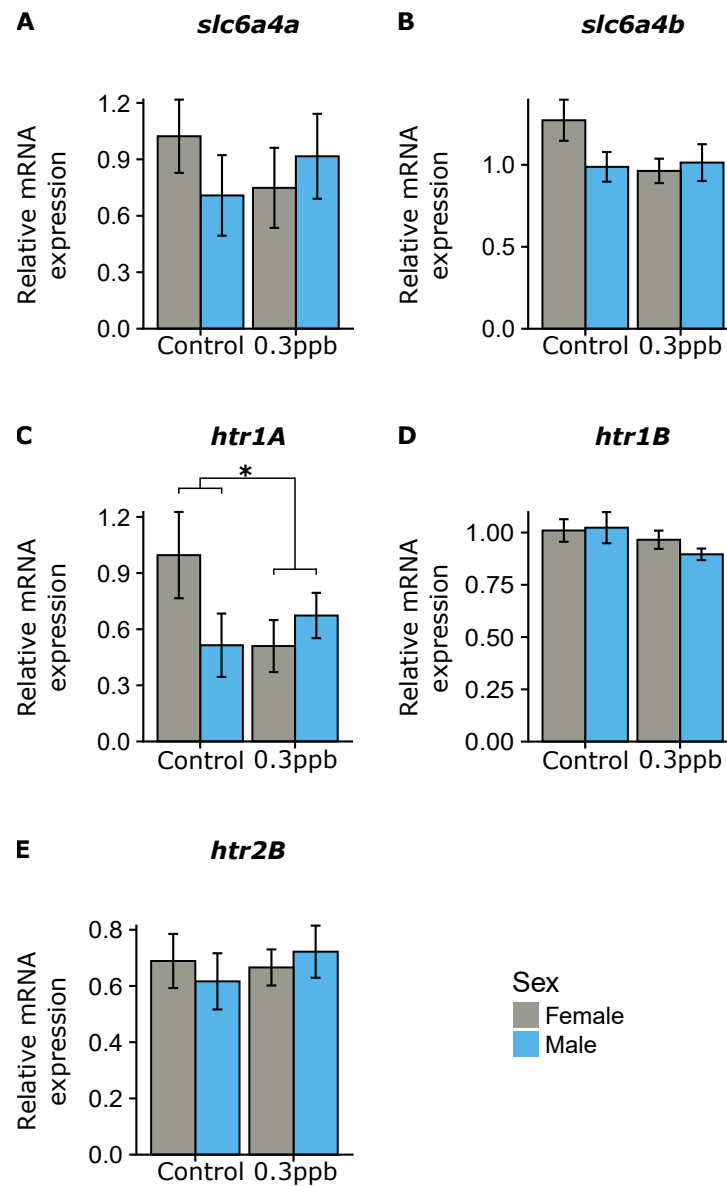


Figure 3.1: Atrazine effects on the relative mRNA levels of the candidate genes (A) *slc6a4a*; (B) *slc6a4b*; (C) *htr1Aa*; (D) *htr1B* and (E) *htr2B* of F₁ offspring from control and 0.3ppb atrazine treated males normalised to *b-actin* (from whole brain samples). Bars represent means, with error bars representing standard errors of the mean. Sample sizes for control offspring were n=22 (males=10; females=12; progeny per family=6-8; families n=3), and for 0.3ppb offspring were n=24 (males=12; females=12; progeny per family=8; families n=3).

3.3.2 Behavioural phenotype and corresponding mRNA expression levels

Anxiety

Time spent in the bottom zone did not significantly correlated with *slc6a4a* or *slc6a4b* mRNA transcript number; however, a marginally non-significant interactive relationship was observed between *slc6a4a* mRNA and treatment group (Est. -22.27 [$-46.64, 2.09$ CI]; Figure 3.2 A; B). On further inspection, no significant relationships were observed in time spent in the bottom zone and *slc6a4a* mRNA when controls and atrazine treated fish were analysed separately, owing to high variation present in both treatment groups. Paternal atrazine exposure altered the relationship between *htr1Aa* mRNA and time spent in the bottom zone (Figure 3.2 C). Overall, a significant positive relationship between time spent in the bottom zone and *htr1Aa* mRNA was observed (Est. 17.28 [$2.42, 32.13$ CI]) and a marginally non-significant interaction was observed between *htr1Aa* mRNA and the treatment group (Est. -21.08 [$-43.98, 1.82$ CI]). On further analysis, control fish drove the significant positive relationship (Control; Est. 17.28 [$2.31, 32.24$ cI]), whereas the relationship was absent in the 0.3ppb atrazine treatment fish (0.3ppb; Est. -3.80 [$-21.12, 13.51$ CI]; Figure 3.2 C). Lastly, no relationship between time spent in the bottom zone and *htr1B* and *htr2B* mRNA transcript was observed (Figure 3.2 D; E).

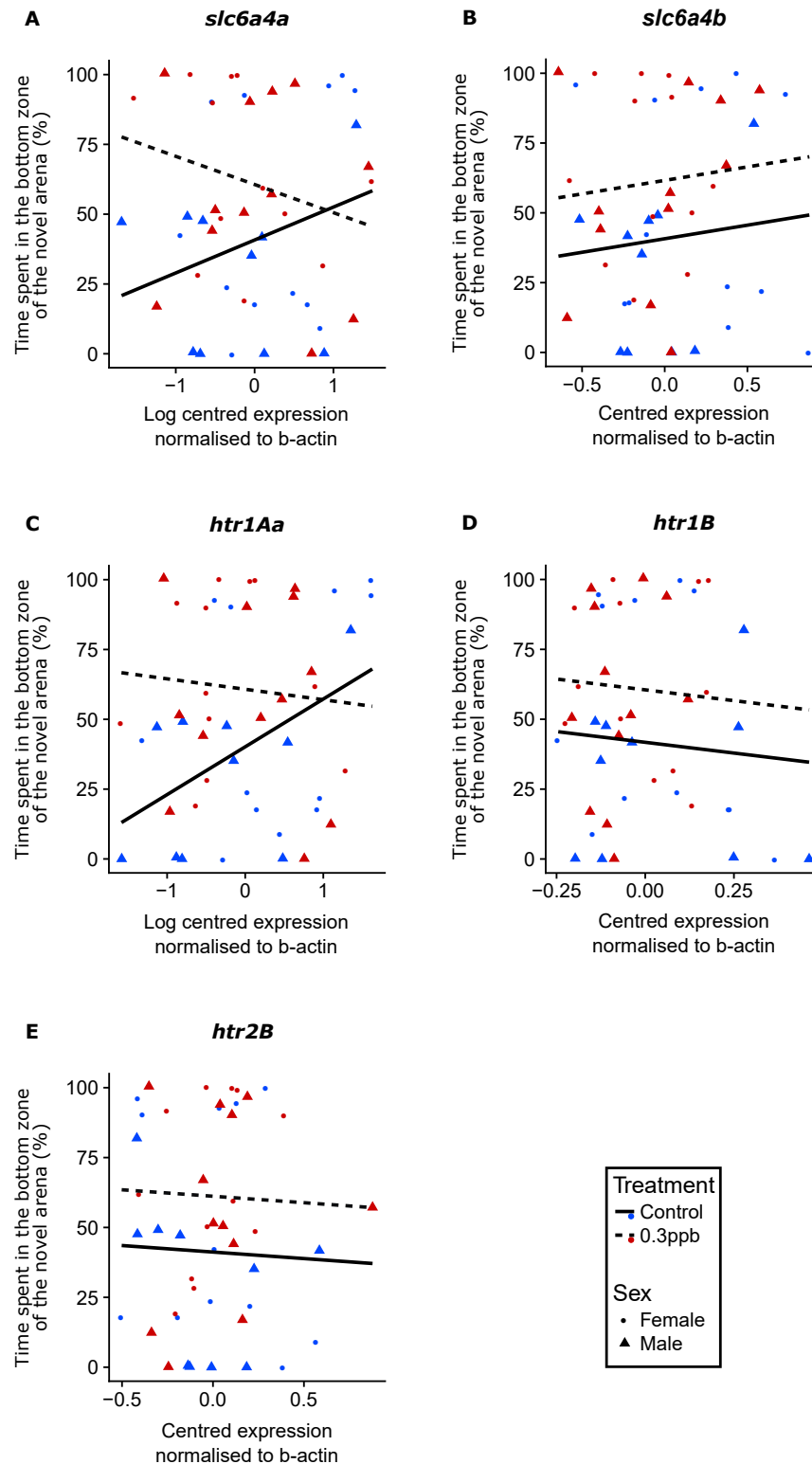


Figure 3.2: Relationship between anxiety (time spent in the bottom zone of the novel arena, sec) and relative mRNA levels of the candidate genes (A) *slc6a4a*; (B) *slc6a4b*; (C) *htr1Aa*; (D) *htr1B* and (E) *htr2B* of F₁ offspring from control (solid line) and 0.3ppb atrazine treated (dashed line) males normalised to *b-actin* and then further centred (from whole brain samples). Both lines are based on model predictions. Control (blue) and 0.3ppb treated (red) offspring for both males (triangles) and females (circles) raw data is overlaid. Sample sizes for control offspring were n=22 (males=10; females=12; progeny per family=6-8; families n=3), and for 0.3ppb offspring were n=24 (males=12; females=12; progeny per family=8; families n=3).

Aggression

No relationship with the mRNA transcripts of *slc6a4a* or *htr1Aa* and the time spent interacting with the mirror was observed in either candidate (Figure 3.3 A and C respectively). But both *slc6a4b* and *htr1B* mRNA transcript number significantly predicted the amount of time a fish spent interacting with the mirror (*slc6a4b* Est. -69.07 [-116.11, -22.03 CI]; *htr1B* Est. -408.952 [-683.82, -134.08 CI]; Figure 3.3 B and D respectively), with higher amounts of mRNA transcripts (in both *slc6a4b* and *htr1B*) correlating with a less aggressive personality. A high amount of variation precluded a significant correlation between *htr2B* mRNA transcript number and the time a fish spent interacting with the mirror, though a suggestive negative relationship was observed (Figure 3.3 E).

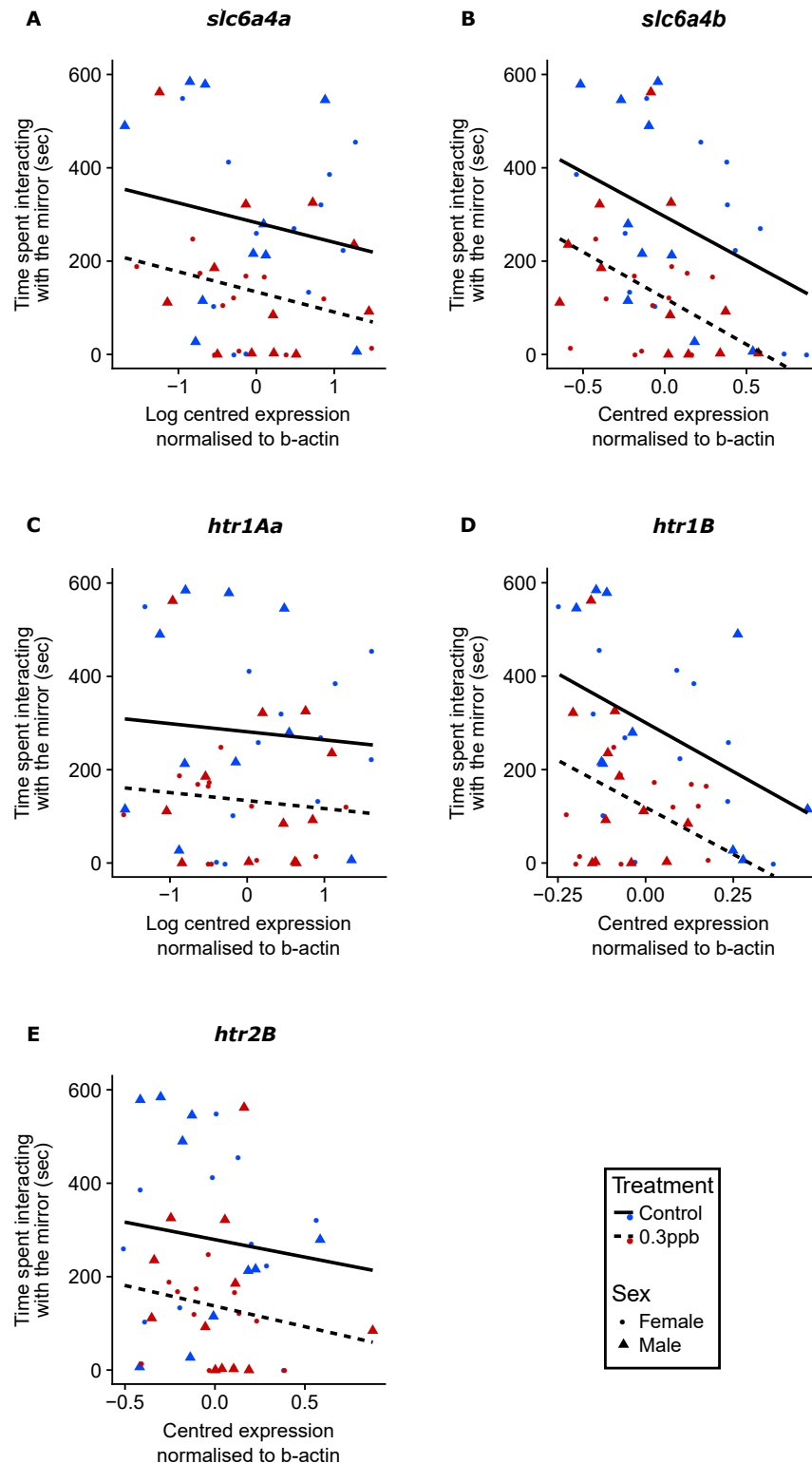


Figure 3.3: Relationship between aggression (time spent interacting with the mirror, sec) and relative mRNA levels of the candidate genes (A) *slc6a4a*; (B) *slc6a4b*; (C) *htr1Aa*; (D) *htr1B* and (E) *htr2B* of F₁ offspring from control (solid line) and 0.3ppb atrazine treated (dashed line) males normalised to *b-actin* and then further centred (from whole brain samples). Both lines are based on model predictions. Control (blue) and 0.3ppb treated (red) offspring for both males (triangles) and females (circles) raw data is overlaid. Sample sizes for control offspring were n=22 (males=10; females=12; progeny per family=6-8; families n=3), and for 0.3ppb offspring were n=24 (males=12; females=12; progeny per family=8; families n=3).

3.4 Discussion

The aim of this study was to investigate if F₁ behavioural differences in anxiety and aggression (observed in Chapter 2) from atrazine treated and non-treated fathers, could be explained by differences in mRNA transcript number of genes involved in the regulation of the serotonergic system. Specifically, I investigated the two serotonin transporters genes, *slc6a4a*, *slc6a4b* and three serotonergic receptor genes, *htr1Aa*, *htr1B*, *htr2B*. Irrespective of behaviour, I found that *htr1Aa* mRNA was down-regulated in atrazine females and slightly up-regulated in atrazine males, compared to controls. There was similar relationships altered in other genes (e.g. *slc6a4a* and *htr2B*), but not significantly so. A positive relationship was found for *htr1Aa* (and non-significantly with *slc6a4a*) mRNA with increasing time spent in the bottom zone. However, paternal atrazine exposures appears to have deteriorated the relationship between time spent in the bottom zone and *htr1Aa* mRNA transcript number (and non-significantly with *slc6a4a* mRNA levels as well). Time spent interacting with the mirror was negatively associated with increased *slc6a4b* and *htr1B* mRNA, regardless of paternal atrazine exposure.

The results of this study support previous research associating atrazine exposure with disruption of some aspects of the serotonergic system (Lin et al. 2013; Lin et al. 2014a; Wirbisky et al. 2015). Indeed, the significant decrease in *htr1Aa* mRNA levels (females in particular) is also observed in adult zebrafish females directly exposed during embryonic development (at the same concentrations; Wirbisky et al. 2015). Wirbisky et al. (2015) also found altered 5-HT turnover in the brain (in females only), so it would be interesting to test for a similar response here. For the most part, it appears that the effects of atrazine on the serotonergic system are sex, development and dose specific with some overlap between these parameters. For example, adult mice exposed to atrazine (at 3mg/kg) during gestation and lactation, via atrazine in the mothers drinking water, had sex specific decreases in 5-HT levels in specific brain regions (both sexes exhibited 5-HT decreases in the striatum, whereas only females exhibited decreases in the perirhinal cortex; Lin et al. 2014a). Another study found disparities in the metabolite profile of exposed male mice, including tryptophan (precursor to 5-HT) and other metabolites important in normal 5-HT function (e.g. linoleic acid and α -linolenic acid), at concentrations >5mg/kg (Lin et al. 2014b). At higher doses, (>125 mg/kg; but not at concentrations tested between 0-25 mg/kg) atrazine increased the 5-HT metabolite, 5-hydroxyindoleacetic acid (5-HIAA) levels and altered 5-HT turnover in the brain (Lin et al. 2013).

It is interesting to note that while sperm are the source of non-genetic atrazine exposure (on the assumption that both male and female offspring inherited the same environmental epigenetic marks), the EDC still appears to differentially affect males and females (at the mRNA transcript level; Figure 3.1 C, and

sometimes at the behavioural level e.g. Figure 2.7 B). This finding indicates that indirect effects of atrazine are also likely to be dependent on developmental and sex specific interactions, just as they are under direct exposure (e.g. Wirbisky et al. 2015; 2016). Other EDCs have seen similar sex-dependent intergenerational/transgenerational effects (Crews et al. 2007). For example, paternal bisphenol-A exposure decreases spatial memory retention in F₁ female rats (Fan et al. 2013), and F₃ female rats ancestrally exposed to vinclozolin discriminate against potential mates that have a history of vinclozolin exposure, an effect that was absent in males (Crews et al. 2007).

In the previous chapter (Chapter 2) I identified that F₁ fish from the 0.3ppb treatment spent more time, on average, in the bottom zone (indicative of a heightened anxiety phenotype). Though this pattern was not observed consistently across treatments (Figure 2.5 B), the selection criteria (for further qPCR analysis) at the extremes of this phenotype (i.e. the least and most anxious fish) suggests that paternal atrazine exposure accentuates the ranges and shifts the medians of this anxiety phenotype (Figure A10.1). When fish were ranked as either the least or most anxious behavioural phenotype from their family cohorts, high variation precluded observable differences of mRNA levels between treatment (Figure A16.1). When continuous data was considered, it revealed that *htr1Aa* mRNA was associated with anxiety (time spent in the bottom zone; Figure 3.2 C). In addition, paternal atrazine treatment appears to have significantly deteriorated the pattern between time spent in the bottom zone and *htr1Aa* mRNA transcript number (and non-significantly in *slc6a4a* mRNA levels as well).

This study provides supporting evidence that up-regulation of *htr1Aa* mRNA is associated with increased anxiety levels, as are the functional proteins of this gene (Maximino et al. 2013). Previous work has shown that bottom dwelling behaviour is positively correlated with 5-HT levels, but that drugs that interfere with *htr1Aa* protein products can alter the anxiety phenotype (Maximino et al. 2013). For example, blocking 5-HT from binding to HT_{1A}R receptor on post synaptic membranes by the drugs, busporine (an agonist of the pre-synapse and antagonist on the post synapse) and WAY 100 635 (antagonist at both pre and post synapse), is associated with producing an anxiolytic effect in zebrafish (Maximino et al. 2013). HT_{1A}R knockout mice exhibit increased anxiety at adulthood (Zhuang et al. 1999) and it appears that retaining proper function of HT_{1A}R during development is critical for normal anxiety behavior as an adult (Gross et al. 2002). Disruption of HT_{1A}R during adolescence in mice is enough to sustain increased anxiety levels through to adulthood (Garcia-Garcia et al. 2016). While this study did not test for effects over development, it is likely that *htr1Aa* mRNA disruption was present over ontogeny and possibly lead to perturbations in anxiety. Future studies should examine ontogenic features of atrazine, anxiety and their relationship to *htr1Aa* mRNA, either directly or indirectly.

One reason why paternal atrazine exposure appears to have been associated with *htr1Aa* mRNA only could be due to the methodology used here. Using a whole brain approach, rather than investigating specific regions of the brain, may not truly reflect the nuances of gene expression activity, especially in regard to behavioural changes. For instance, 5-HT activity is non-uniform across the brain, and so too are receptor and transporter gene expression (Wang et al. 2006; Norton et al. 2008; Filby et al. 2010b). The non-significant interaction between *slc6a4a* and paternal atrazine exposure is likely due to lack of sample size and increased variability, precluding observable differences, as no significant correlation was observed when controls and atrazine treated fish were tested separately (Figure 3.2; Table A15.4). Fluoxetine (a 5-HT transporter antagonist) is shown to produce an anxiolytic phenotype (Maximino et al. 2013), indicating some involvement of *slc6a4a* and *slc6a4b* in anxiety. Other drugs such as SB 224289 (an inverse agonist to the HT_{1B}R receptor) also produces an anxiolytic effect in a novel arena assay (Maximino et al. 2013), further suggesting that this study failed to capture involvement of the genes tested. Another reason for lack of a trend may be that an anxiety phenotype is produced via epistatic interactions between genes (many genes of small effect culminating together) and that mRNA levels of these candidate genes (except for *htr1Aa* and *slc6a4a*) in isolation do not overall reflect the many processes that lead to an anxiety phenotype. Indeed, quantitative trait loci mapping in wild and laboratory strain zebrafish show that some measures of boldness (among other behaviours) are epistatic in nature (Wright et al. 2006b).

The relationship between aggression and *slc6a4b* expression was partially expected. Previous work by Theodoridi et al. (2017) demonstrated that whole brain *slc6a4b* expression was related to dominant and subordinate positions in male zebrafish hierarchies (subordinates express *slc6a4b* mRNA relatively more than dominant individuals). The authors further suggested that differences in expression may be due to associated coping strategies, i.e personality differences. Moreover, the lack of a significant relationship between *slc6a4a* and aggression is further supported by Theodoridi et al. (2017) who found no difference between dominant and subordinate individuals (though in that study fluoxetine treatment still altered expression levels and caused behavioural changes). Other work has found similar patterns in rainbow trout (*Oncorhynchus mykiss*) between low reactive (consistently more aggressive) and high reactive (consistently less aggressive) individuals; low reactive individuals have down-regulated *slc6a4b* expression and no difference in *slc6a4a* expression (Rosengren et al. 2017). A negative relationship between 5-HT transporter mRNA and aggression levels in zebrafish was proposed by Filby et al. (2010b). However, Filby et al. (2010b) did not investigate *slc6a4b* levels, and presided this hypothesis on *slc6a4a* mRNA levels only. Filby et al. (2010b) found evidence of altered expression of *slc6a4a* between dominant and subordinate females only, but when the authors compared day 1 expression to day 5 expression of dominant and subordinate fish, the relationship in

females with *slc6a4a* expression levels disappeared and appeared in males (higher *slc6a4a* levels in dominant individuals). This research by Filby et al. (2010b) suggests a more temporal component of *slc6a4a* mRNA, that might in function, be involved in reactive aggression, whereas *slc6a4b* might be involved in a more stable phenotype as it correlates with aggression personality (in this study). Overall, this study suggests that *slc6a4b* transcripts might be more important for personality differences than *slc6a4a* mRNA levels.

The 5-HT receptors HT_{1B}R and HT_{2B}R are shown to be involved in offensive and defensive aggression behaviours (Fish et al. 1999; Miczek et al. 2001; Almeida et al. 2002; Takahashi et al. 2011; Theodoridi et al. 2017). The lack of relationship between aggression levels and *htr2B* mRNA expression in this study is in contrast with what has been found previously. For instance, dominant zebrafish exhibit significantly lower transcripts of *slc6a4b* and *htr2B* in comparison to subordinate conspecifics (with suggestible evidence for lower *htr1B* expression levels in dominants; Theodoridi et al. 2017). Aggressive encounters in zebrafish involve two transient phases; first an appetitive element (overt offensive engagement, e.g. displaying, circling and biting, by both parties), followed by a resolution element (dominant or subordinate specific behaviours, e.g. chases or flights/freezing respectively; Kalueff et al. 2013; Theodoridi et al. 2017). The lack of relationship between mirror induced aggression and *htr2B* mRNA is possibly a reflection of the methodology used to capture zebrafish aggression in this study. The mirror test offers a proxy look at aggression (theoretically restricting aggression to its appetitive element) without the dynamic effect of winner or loser outcomes (Kalueff et al. 2013; Oliveira et al. 2016). The relationship between mirror induced aggression and *htr1B* mRNA, and lack thereof with *htr2B* mRNA, possibly reflects *htr1B* mRNA is important for personality differences pertaining to an individual's engagement in aggression. Whereas *htr2B* may be more involved in defensive aggression behaviours when there is an ability to process feedback information about an opponent. This remains speculation and more work is needed to clarify how offensive and defensive aggression behaviours may be related to gene expression.

Although paternal atrazine exposure altered aggression personality in this study, the lack of an effect on these genes suggests other factors are responsible for the phenotypic differences. The system that leads to aggression is complex (encompassing more than 40 genes of at least eight pathways; Filby et al. 2010b). In general, this study is limited by the whole brain approach of which there are clearly massive differences in expression across distinct regions of the brain (Filby et al. 2010b). Also, compared to Filby et al. (2010b), who used the WIK zebrafish strain, this strain is wildtype AB. Different strains of zebrafish appear to differ in their behaviour, in baseline levels of 5-HT and also react differently to chemicals that interfere with the serotonin system differently (Barbosa et al. 2018a). This study was not able to account for the dominance of individuals in their home tank, as social experience can influence future aggressive encounters

(Kloke et al. 2011). As stated before, the mirror test used here offers a limited proxy for aggression, though repeatability of this measure was observed (see Appendix IV), indicating this method likely captures personality differences between individuals. A recent study provides improved methodology to quantify the full range of aggression in zebrafish and other fish species (e.g. Li et al. 2018).

Overall, paternal atrazine exposure appears to alter offspring anxiety and aggression. These data suggest that disruption in *htr1Aa* mRNA levels (and possibly *slc6a4a* mRNA levels) contribute to an increase in time spent in the bottom zone (a shift in anxiety phenotype). In contrast, paternal atrazine exposure decreased offspring aggression, but the atrazine treatment did not appear to influence expression of any of the candidate genes and as such, shifts in behavioural phenotype appear to be under the control of different systems. Additional candidates are likely to be numerous due to the complexity of the aggressive phenotype (Filby et al. 2010b). Moreover, transcriptome and DNA methylation studies suggest a wide array of pathways are affected by early developmental exposure to atrazine (Wirbisky et al. 2015; Hao et al. 2016; Wirbisky et al. 2016a; McBirney et al. 2017). Changes in gene expression underlying anxiety and aggression may also be potentially propagated transgenerationally. Hence, future studies combining changes in behaviour and gene expression due to atrazine exposure should frame their research with transgenerational effects in mind, as exposure to atrazine has been demonstrated to induce transgenerational effects (Hao et al. 2016; McBirney et al. 2017). Indeed, McBirney et al. (2017) consistently found that the systems associated with signalling, transcription, metabolism and receptors (among others) to be differentially methylated after transgenerational atrazine exposure (despite DNA-methylation profiles being discordant between generations, F₁-F₃). These studies suggest that mechanisms behind atrazine induced behaviours are likely to be affected transgenerationally as seen in other EDCs (e.g. Crews et al. 2007; Volkova et al. 2015a; see O’Dea et al. 2016). This study is one of a few to investigate paternal atrazine exposure on offspring phenotype, but more studies are needed to identify the extent of paternal contribution after atrazine exposure.

Chapter 4

General discussion

4.1 Thesis objectives

The main objective of this thesis was to investigate the influence of an endocrine disruptor, atrazine on non-genetic inheritance of personality traits in zebrafish. In Chapter 2, I used a series of behavioural assays (novel arena test, novel object test and a mirror test) to investigate if exposure to atrazine during juvenile development affected the five behavioural traits (aggression, anxiety, activity, boldness and exploration) of the exposed individuals and more importantly, their unexposed progeny. Chapter 3 used whole brain tissue to investigate if behavioural changes observed in the F₁ progeny were able to be explained by changes in mRNA transcript number in genes involved in regulating the serotonergic system (two serotonin (5-HT) transporters, *slc6a4a* and *slc6a4b*, and three 5-HT receptors: *htr1Aa*, *htr1B*, *htr2B*). I focused on two personality traits, anxiety and aggression, because I found evidence for an intergenerational effect on these two traits. Moreover, anxiety and aggression related behaviours are strongly associated with the serotonergic system, and developmental atrazine exposure is associated with the serotonergic system dysregulation (Chiavegatto et al. 2001; Gross et al. 2002; Filby et al. 2010b; Wirbisky et al. 2015). Therefore, this study also sought to extend previously observed direct serotonergic system effects to a further generation (Wirbisky et al. 2015).

For the experiments in Chapter 2, I predicted increases in anxiety and activity levels and decreases in boldness, aggression and exploration levels in both generations. Furthermore I predicted that increases in anxiety would be more prominent in females than males. For the experiments in Chapter 3, I predicted that F₁ progeny aggression levels would correlate negatively with mRNA transcript number and that anxiety levels would correlate positively with all candidate genes. In particular, I predicted a stronger positive

association of *htr1Aa* mRNA with anxiety levels. I also predicted that F₁ progeny from atrazine treated males would have higher transcript number at the candidate genes, compared to controls, indicative of their higher anxiety and lower aggression behavioural profiles.

4.2 Summary of main findings

Several behavioural measures across a range of personality phenotypes were altered by either developmental exposure or paternal exposure to the herbicide, atrazine, at typical environmental concentrations (summarised in Table 4.1). These findings support the hypothesis that atrazine exposure has the potential to alter personality traits. The personality trait most affected by developmental atrazine exposure was boldness (measured by latency to enter the top zone of the novel arena) - F₀ female latency increased with increasing atrazine concentration, whereas males exhibited a threshold drop in latency compared to controls. F₀ fish exhibited increased anxiety (measured by the time spent in the bottom zone of the novel arena) and decreased exploration (measured by the standard deviation of time spent in each zone of the novel arena) in the 0.3ppb group only. There was some evidence for decreased aggression (measured by time spent interacting with the mirror) amongst F₀ fish, but not significantly so. No treatment differences in activity levels (measured by the total distance moved) were observed across any of assays in the F₀'s.

Despite never being directly exposed, several behavioural changes were observed amongst the F₁ progeny, supporting the hypothesis that effects of atrazine may be further transmitted down the germ line. Paternal atrazine exposure significantly increased the latency of the F₁ progeny, in all treatment groups compared to controls. In addition, there was a suggestive pattern of decreasing latency to enter the top zone with increasing atrazine concentration amongst the F₁ progeny from atrazine treated males. Significant decreases in aggression were observed amongst the treated F₁'s (in the 0.3ppb and 30ppb groups) that was paralleled with significant decreases in activity during the mirror test in the treatment groups, 0.3ppb and 30ppb, compared to controls. No other treatment differences in F₁ activity were observed during the novel arena test or in the novel object test. There was no treatment differences in F₁ exploration, but there was some evidence of increased anxiety across the treatment groups compared to controls, but not significantly so. These findings suggest that the offspring of exposed males have decreased aggression, decreased boldness and potentially increased anxiety, which can have consequences for how these fish interact with predators, conspecifics and their environment, such as competition for food, mates, territories or social rank (Spence et al. 2008; Paull et al. 2010).

Paternal atrazine exposure significantly down-regulated *htr1Aa* mRNA expression in females and slightly

up-regulated it in males, compared to controls. Moreover, *htr1Aa* mRNA (and *slc6a4a* - but not significantly so) was found to correlate positively with anxiety levels in controls, but this relationship was disrupted in the atrazine treatment group. While the candidate genes tested here did not explain the significant reduction in aggression observed in the offspring of atrazine treated males, both *slc6a4b* and *htr1B* mRNA correlated negatively with an aggressive personality phenotype.

Though several studies before me have previously identified a range of behavioural changes associated with atrazine exposure (e.g. male-male aggression, shoal-cohesion, anxiety-related behaviours; Belloni et al. 2011; Shenoy 2012; Lin et al. 2013; Lin et al. 2014a; Schmidel et al. 2014; Shenoy 2014), this is, to my knowledge, the first study to investigate if behavioural changes from atrazine exposure are associated with personality differences. These results also support the hypothesis that non-genetic inheritance can contribute to animal personality (Reddon 2011) and that mRNA transcript of genes (i.e., a measure of gene expression) involved in regulating the serotonergic system are associated with personality differences.

Table 4.1: Summary of significant personality differences from direct and indirect atrazine exposure for F₀ and F₁ zebrafish. Each concentration where a significant treatment effect was observed, relative to controls, is specified. Arrows indicate either a positive (up) or negative (down) direction of effect. Sex differences are presented as females (F) compared to males (M). For treatment:sex interactive differences, the first arrow indicates the direction of effect for males and second arrow indicates female direction. To remind the reader, an increased SD (standard deviation) of exploration indicates a less exploratory phenotype.

Personality trait	F ₀			F ₁		
	Treatment	Sex	Interaction	Treatment	Sex	Interaction
Boldness						
Latency to enter the top zone	↑(30ppb)	NS	↓(M)↑(F)(3, 30ppb)	↑(0.3, 3, 30ppb)	↑(F)	NS
Anxiety						
Time spent in bottom zone of novel arena	↑(0.3ppb)	NS	NS	NS	↑(F)	NS
Exploration						
Exploration SD of the novel arena	↑(0.3ppb)	NS	NS	NS	↑(F)	NS
Aggression						
Time spent interacting with the mirror	NS	NS	NS	↓(0.3, 30ppb)	↓(F)	NS
Total distance moved in mirror test	NS	NS	NS	↓(0.3, 30ppb)	↓(F)	NS

4.3 Secondary findings

While not a specific aim of this thesis, I additionally assessed the early development of F₁ progeny, 24hpf survival and hatching at 48hpf, to examine how atrazine might influence these important early life parameters. I found that paternal atrazine exposure reduced survival at 24hpf in the 3ppb treatment group and significantly reduced the proportion of offspring that hatched by 48hpf in the 3ppb and 30ppb treatment groups, compared to controls (Appendix II). Interestingly, these results are in contrast to those of Wirbisky et al. (2016c), who found no difference in survival at 24hpf or hatching at 48hpf of progeny from male and female zebrafish that were embryonically exposed to atrazine (at the same concentrations; 0.3ppb, 3ppb and 30ppb). This study out-crossed atrazine exposed males with unexposed females (using IVF) whereas Wirbisky et al. (2016c) employed natural matings and crossed both exposed males and females. The results found in this study may therefore be a consequence of either the difference in the timing of exposure (during juvenile development i.e. during sexual differentiation in this study versus during embryonic development in Wirbisky et al. (2016c)). Alternatively, atrazine has been shown to induce a variety of testicular and sperm quality issues (Abarikwu et al. 2010; Feyzi Dehkhargani et al. 2011; Farombi et al. 2013; Abarikwu et al. 2015); if the probability of defective sperm contacting released oocytes is lower during natural mating conditions compared to *in vitro* fertilisation conditions, then this could explain the contrasting findings between this study and Wirbisky et al. (2016c).

4.4 Limitations of the study

No study is without its limitations, as there were several aspects of each experiment that could have been improved or where the assumptions needed further testing. One of the key assumptions was that zebrafish exhibit long-term repeatable behaviour when given a single ‘tank experience’ followed by testing one week later (Thomson 2017). In our lab, this regime has been demonstrated to be able to discriminate between individual behavioural phenotypes (Thomson 2017). However, it is also known that several EDCs have the ability to disrupt individual consistency and disrupt behavioural syndromes between related traits (e.g. Dzieweczynski et al. 2013; Dzieweczynski et al. 2014; Dzieweczynski et al. 2016). So it may be probable that the behavioural data shown here is not the full extent to which atrazine is capable of altering behavioural phenotypes. Moreover, it remains to be seen whether or not behavioural consistency (i.e. the repeatability of traits) is also affected indirectly. Understanding early life exposure and intergenerational (or transgenerational) effects on behavioural consistency is a promising gap in the literature that should be explored with additional studies. If so, it is likely to have ramifications for how researchers are able

to predict long term behavioural outcomes. Nevertheless, exposure to atrazine either directly or indirectly appears to alter a variety of personality traits in a directional manner, based on treatment group averages and reasonably minimal error bars (for behavioural data).

As males were randomly selected for breeding (via IVF) and there was no markers for individual identity during behavioural testing, the ability to investigate if offspring phenotype was similar to that of the father was not possible. Zebrafish paternal personality and social status appears to non-genetically influence the activity of their offspring (at least during the larval stage; 7 and 10 dpf; Zajitschek et al. 2017). So it may be possible that this phenomena extends to other personality traits and lasts through to adulthood.

Timing and logistical reasons did not allow the possibility to test for transgenerational effects. Indeed, it is this limitation that prevents this study from discriminating between a true epigenetic explanation of inheritance from the possibility that effects were caused via exposure of the F_0 developing germ cells i.e., the future cells of the F_1 progeny (Szyf 2015; Jacobs et al. 2017). Nonetheless, these results suggest that transgenerational effects on behaviour may be possible.

There will always be important lessons to be learned with hindsight from the testing regimes researchers use to understand personality and treatment differences in a particular species. The novel object and mirror test were two such cases that could have been improved. The aim of the novel object test was to incorporate an additional measure to investigate boldness/anxiety in zebrafish (see Chapter 2; Methods); however these data suggest that the novel object itself was inappropriate for assessing effects on zebrafish behaviour, and instead possibly induced a fear response (see Appendix VI). Furthermore, the novel object test may have influenced the mirror test that followed. These data further reinforce the need for methodological validation on zebrafish (and other fish species; as mentioned by Maximino et al. 2010). Nevertheless, the reduction (near significance) in cumulative duration spent in the bottom zone by F_1 fish from the 0.3ppb and 3ppb treatments suggest that treatment effects that influence fear responses might be revealed with methodological improvements to this assay, or with greater sample size as other attempts (using different novel objects) have been successful (Fangmeier et al. 2018; Kirsten et al. 2018).

The mirror test was able to capture repeatable differences between individuals (Appendix IV), and though it is known to correlate with other behavioural measures of aggression (Ariyomo et al. 2013b; Way et al. 2015), it must be noted that the mirror test provides a partial evaluation of aggressive behaviour (Oliveira et al. 2011; Kalueff et al. 2013). The mirror test is also sometimes used as a measure for sociability rather than aggression (e.g. Cattelan et al. 2017, and references therein), though in the assay utilised here, aggressive behaviours were observed and were directed towards the mirror image e.g. bites and lateral displays. A

recent study provides improved methodology to quantify the full range of aggression in zebrafish and other fish species in improved predictability of winner-loser fights (e.g. Li et al. 2018).

Another limitation was the number of full-sibling families used in this study. Full-sibling families are used to separate the environmental variation attributable to the stressor of interest and the genetic signal inherited from the genetics of the parents. This study used three families per treatment; 12 different families in total, with around 13-18 fish per family. Despite this, in many cases, the genetic contribution superseded an environmental inheritance effect even when an inherited effect was suspected (Appendix IX, Table A9.1). Therefore it is recommended that researchers should configure their experimental design of their future studies to maximise the number of families utilised. Based on the limitations of this study alone, researchers should employ at least four or more families at minimum (per treatment) as a recommendation. However, a comprehensive study comparing minimal family sample size requirements would be valuable to the epigenetics field.

While the objectives of the study were aimed at getting a more general idea for atrazine's effects on the serotonergic system, as mentioned previously in Chapter 3, using a whole-brain approach to investigate mRNA differences likely confounded the nuances of how intergenerational atrazine might have affected different brain regions. The serotonergic system is peppered throughout the zebrafish brain (see Wang et al. 2006; Norton et al. 2008) and behavioural differences resulting from knock out in some areas of the brain (but not others) are not equal (Zhuang et al. 1999; Gross et al. 2002). Therefore, future studies should aim for testing similar patterns across different brain regions. In addition, the results presented in Chapter 3 clearly show a wide amount of variation present, possibly resulting from a lack of power, which is likely due to sample size.

One particular draw back of a lab based approach to understand behavioural changes is that extrapolation to individuals in the wild must be taken with caution. The environment comprises numerous and complex interactions (e.g. prey, predators, shelter, seasonal components etc.) however, employing controlled lab experiments allow the construction of base-line effects to which effects could be expected in the wild. Testing wild population for causal effects is a greater challenge, but an intermediate step would be use mesocosm experiments. Additionally, exposure to a single contaminant in the wild is not often the norm; instead, organisms are much more likely to face exposure from a multitude of xenobiotics (Vos et al. 2000; Swanson et al. 2018).

The outcome of exposure to multiple EDCs may not be entirely predictable as EDCs in combination may result in a variety of effects, such as augmentation of one effect, suppression of another, EDCs may synergise together or possibly lead to new effects than when observed singularly (Vos et al. 2000; Hayes et al.

2006; Pérez et al. 2013). Indeed, atrazine has been known to synergise with chlorpyrifos (an organophosphate; acetylcholinesterase inhibitor), increasing the prevalence of disturbed locomotor behaviour in larval zebrafish (Pérez et al. 2013). Future studies should make use of testing later life effects (and intergenerational/transgenerational) of atrazine in combination with other likely to encounter pollutants (Swanson et al. 2018).

4.5 Future directions

In the present study, changes in personality traits and mRNA differences observed in Chapter 2 and Chapter 3 should be verified under differing environmental contexts to give additional weight to the hypothesis that atrazine alters personality traits. For instance, future studies should aim to provide more ecological relevance to these findings by investigating the same or related behavioural measures under differing ecological pressures, such as increased predation risk, social density, food resources or environmental complexity (e.g. Luca et al. 2012b; Luca et al. 2012a). Additionally, future studies should aim to extend personality testing under atrazine exposure to other taxa, in order to identify how broadly atrazine may affect personality traits. In Chapter 2, I proposed two different scenarios that atrazine may be affecting boldness (i.e. risk-taking) behaviour in zebrafish; 1) atrazine is altering the propensity to take risks or 2) atrazine is altering the ability to process risk. Researchers could utilise zebrafish's inclination for shoaling to disentangle these two hypotheses by measuring group dynamics of a focal (developmentally exposed to atrazine) individual within a group of unexposed individuals (Spence et al. 2008). Zebrafish use shoaling as a means of predator defence (Olst et al. 1970; Landeau et al. 1986; Wright et al. 2006c) and will alter their shoal cohesion depending on ecological pressures such as predation or food availability (Pitcher et al. 1988; Bass et al. 2008; Miller et al. 2007). If atrazine is altering the propensity to take risks then I predict that focal individuals will try to maintain shorter nearest neighbour distances and actively try to maintain a centred position in the safest part of a shoal. I also predict that the focal individual will exhibit riskier behaviour (e.g. leaving the shoal) when risk decreases. On the other hand, if atrazine is altering the ability to process risk then I predict the focal individual will show less structuring in the shoal as risk is manipulated. When exposed to high doses of atrazine (1000ppb) as adults, group cohesion decreases, so it might be more probable for an altered ability to process risk (Schmidel et al. 2014).

Only a handful of studies have investigated atrazine's role in altering male (but not female) aggression (Shenoy 2012; Shenoy 2014). Further studies should investigate how atrazine exposure alters aggression in a social or dominance context. It remains to be seen what the specific consequences of reduced aggression

in females would be; however, I posit that atrazine exposed (and indirectly exposed) females (and males) will be less likely to achieve a higher social ranking, and be less likely to maintain a higher social rank for longer if able to outcompete other individuals. Since body size is a predictor of social rank (Paull et al. 2010), I further posit that smaller unexposed females would be able to outrank larger directly or indirectly exposed females.

Several intergenerational effects on behaviour and mRNA transcript number were observed throughout this study, suggesting future research into transgenerational effects would be valuable. Though transgenerational effects have been observed previously in mice after atrazine exposure (e.g. Hao et al. 2016; McBirney et al. 2017), the studies were not conducted at environmentally relevant concentrations as in this study design. Additionally, atrazine differentially affects male and female behaviour and the mRNA transcript level after indirect exposure, suggesting that if epigenetic markers are being inherited equally (between the sexes), that atrazine is acting upon developmental and sex specific factors, such as hormones. It would be valuable for future studies to address this interesting gap and attempt to isolate what molecular components are interacting together.

The findings here exemplify the importance and sensitivity of development during early life exposure to environmental stressors, and the potential for effects to be carried through to, or manifest, at maturity (Jonsson et al. 2014). The majority of work in investigating the relationship between atrazine and the early life environment in fish have usually limited their exposure regimes to the embryonic period (e.g. Wirbisky et al. 2015; Wirbisky et al. 2016a). Therefore, this work is important because it extends the sensitivity window for later life consequences of atrazine exposure. It is not unsurprising, however, as juvenile development is a period of increased brain growth and is when numerous factors, including a variety of hormones, start the process to differentiate zebrafish males from females (Jonsson et al. 2014; Lee 2015). Interference during this critical developmental period is therefore predicted to result in adverse outcomes (Burton et al. 2014; Jonsson et al. 2014). In future, it would be valuable to investigate the difference between embryonic and juvenile development (and time points in between and after) on adult behaviours, as it might give further insight into predicting potential ecological consequences and possibly into atrazine's mechanisms of action.

Lastly, while researching, a prominent gap identified in the literature was the lack of understanding to which atrazine mechanistically exerts its effects in the nervous system and other tissues. Previous research indicates that atrazine has the potential to affect a wide variety of systems, directly or indirectly (e.g. Wirbisky et al. 2015; Wirbisky et al. 2016a; McBirney et al. 2017); therefore, this is a crucial area that needs to be addressed. Indeed, this is a common problem throughout the endocrine disrupting field (Gore

et al. 2015). This gap is likely owing to several difficulties arising from the properties of EDCs in which effects are usually sex, timing, and dose specific and are typically non-monotonic in nature (Vandenberg et al. 2012), as is the case with atrazine (e.g. this study; Shenoy 2012; Brodeur et al. 2013; McCallum et al. 2013; Riffle et al. 2014; Marcus et al. 2016). As a first step, a review of potential mechanisms is sorely needed in the atrazine field in order to collate potential hypotheses and identify where specific research gaps in knowledge are. There is also increasing realisation that EDC (and other contaminant) exposure at the individual level could translate to consequences at the population level (Crews et al. 2007; Shenoy et al. 2011; Senior et al. 2014; Araújo et al. 2018).

4.6 Conclusions

The major findings from this study were that exposure to environmentally relevant concentrations of atrazine during early life development influences a variety of personality traits in the direct exposed individuals and in the progeny of exposed males, despite the progeny never being directly exposed to atrazine. The affected personality traits include boldness, aggression, anxiety and exploration, with sex-specific effects on boldness observed in both F_0 and F_1 . In addition, I found evidence that paternal atrazine exposure appears to have disrupted aspects of the serotonergic system; specifically, *htr1Aa* mRNA levels between the sexes was reversed in the 0.3ppb treatment, relative to controls, and that the positive relationship between *htr1Aa* mRNA and a more anxious personality was perturbed. It appears that other factors not studied here are responsible for alterations in aggression, but *sl6a4b* and *htr1B* mRNA are associated with an aggressive personality. A secondary finding was that paternal atrazine exposure can reduce early life survival and hatching of progeny. This study contributes to the growing understanding that EDCs have the ability to affect behavioural traits and shows the sensitivity of the early life environment. Moreover this research shows that effects of an EDC in one generation may be further propagated down the germ line.

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Appendix I: IVF protocol

The following protocol describes the method of *in-vitro-fertilisation* (IVF) written by Dr Sheri Johnson. This method is also available from Johnson et al. (2018). IVF was chosen over other natural breeding methods, because the time of conception can be easily manipulated. Doing IVF over natural spawning also removes the behavioural uncertainty of successful mating between zebrafish males and females. The process describes several steps needed to ensure successful IVF.

Squeezing

Day before experiments:

1. Put one vial of Tricaine in fridge to thaw overnight. Check that have buffers for ZSI (see recipe below). Note that 1 X PBS can be used in place of ZSI, if necessary.

Day of experiment:

1. Prepare zfish sperm immobilizer (ZSI; see recipe below)
2. Prepare tubes with 10 uL of ZSI in each tube (close tubes so no ice or water get in tubes)
3. Set up two 250 mL beakers with anaesthetic: add 200 mL of fish water to the 8.2 mL aliquot, and then split between two beakers.
4. Anesthetise males until breathing slows, remove beaker, blot off excess anesthetic on paper towel, rinse fish in fish water and then place on paper towel and blot gonadal region with Q-tip dipped in ZSI.
5. Collect sperm in capillary tube and mouth pipette into pre-prepared tubes. Make a note on datasheet whether you collected 0.5-1 uL or more than 1 uL.
6. Once all males are squeezed, add 10 uL more of ZSI to males that gave more than 1.5-2 uL, 20 uL to males that give 2.5-3 uL and so on.

IVF Initial Prep:

1. Label small petri dishes (on the bottom and the top).
2. Dilute sperm collected from males. In a new tube, add 5 uL of diluted sperm to 45 uL of ZSI and mix. Return rest of sperm dilution to fridge for CASA analysis, etc. Keep IVF aliquot on ice at all times.

IVF:

1. Squeeze females after 5-10 min in Tricaine
2. Check for quality of eggs under microscope
3. Add 45 uL sperm (gently pipette before first replicate) on top of eggs and IMMEDIATELY add 1 mL of fish water, THEN proceed to the next replicate and switch tips between replicates
4. Wait 5 minutes and then add 5 mL of fish water
5. After 60 min, transfer to incubator @ 28.5 C
6. Assay fertilization 4 hpf and then transfer to labelled large petri dishes. The bottom should just have the male ID and replicate (e.g., AB 7A) that corresponds with that IVF sample. The lid should have this same label, plus the fish line, date and researcher initials). Enter cross information in the cross binder
7. Assess hatching rate at 3 dpf and then move to system 5dpf

Zebrafish Reagent Recipes

Zebrafish Sperm Immobilizer (page 114,941 in Mol. Lab book)

- 140 mM NaCl (1M stock: 14.61 g NaCl in 250 mL MQ)
- 10 mM KCl (100 mM stock: 1.86 g of KCL in 250 mL MQ)
- 2 mM CaCl₂ (100 mM stock: 2.77 g of CaCl₂ in 250 mL MQ)
- 20 mM HEPES (100 mM stock: 4.78 g of HEPES in 200 mL of MQ), pH 8.5 (filtered with 0.22 uM syringe filter, kept at 4C)

To prepare working stock morning of sampling:

- 14 mL 1M NaCl
- 10 mL 100 mM KCl

- 2 mL 100 mM CaCl₂
- 20 mL 100 mM HEPES
- 54 mL of DI H₂O
- Keep on ice

Tricaine:

- 400 mg tricaine powder
- 97.9 ml UP H₂O water
- ~2.1 ml 1 M Tris (pH 9).

Tricaine (3-amino benzoic acid ethyl ester also called ethyl 3-aminobenzoate) comes in a powdered form from Sigma (Cat.# A-5040). It is also available as Finquel (Part No. C- FINQ-UE) from Argent Chemical Laboratories, Inc. Make triacaine solution for anaesthetizing fish by combing the following in a glass bottle with a screw cap:

Adjust pH to ~7 and aliquot:

- 8.4 mL tricaine in 15 mL tubes
- Label tubes with instruction to dissolve in 200 mL Zfish system water
- Store aliquots in the -20 C freezer. (Buy the smallest amount possible because tricaine gets old)

Appendix II: Early life effects of offspring from atrazine exposed fathers

Methods

At 24hpf dead embryos were removed from petri dishes and counted. To ascertain hatching success, at 48hpf, 72hpf, 96hpf each petri dish was photographed using a Cannon EOS 700D with a cannon 50mm macro-lens and number of hatched and unhatched fry were counted in ImageJ (version 1.50i; Schneider et al. 2012). Figure A2.1 shows an example of hatched and unhatched embryos.

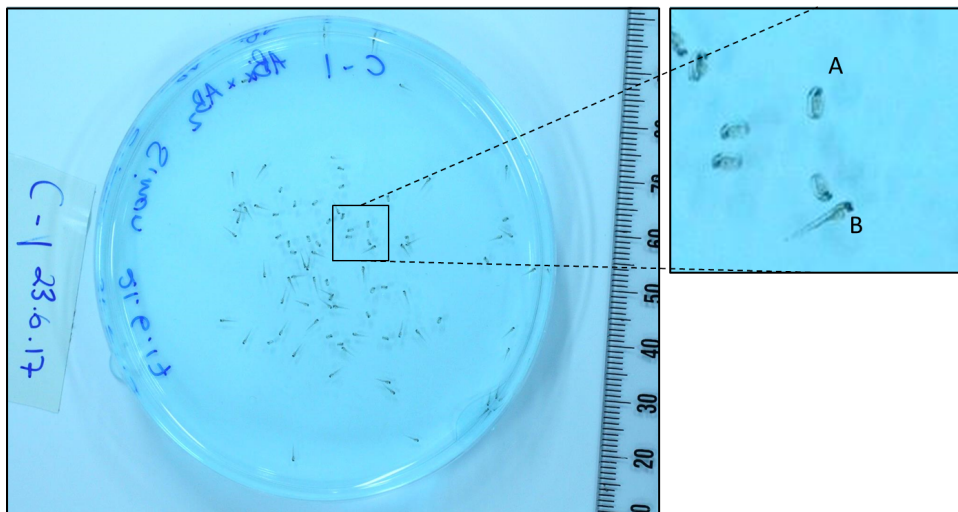


Figure A2.1: Example of zebrafish at 48hpf, photograph taken in order to count hatched and unhatched embryos. Example of A) an unhatched and B) hatched zebrafish.

Results

Embryonic death peaked by 24hpf (see Table A2.1 for numbers) across all treatments. There was a significant reduction in the proportion of offspring that survived to 24hpf in the 3ppb treatment ($z = -6.68$, $p < 0.001$), but no other response was observed in either the 0.3ppb and 3ppb treatments (Figure A2.2 A). Interestingly, control variation was minimal relative to all other treatments, with variation greatest in the 3ppb treatment. Unexpectedly, the majority of survival in the 30ppb was above the mean of controls, whereas the majority of families from both 0.3ppb and 3ppb treatments were below. Deaths that occurred past 24hpf were minimal for each treatment (0 - 2 individuals) or did not occur whilst fish were held within the petri-dish.

The majority of hatching had occurred by 48hpf and by 72hpf $> \sim 97\%$ had emerged (see Table A2.2 for numbers). The proportion of embryos that hatched by 48hpf was significantly less in the 3ppb ($z = -6.50$, $p < 0.001$) and 30ppb ($z = -4.57$, $p < 0.001$) treatments, and was near significance in the 0.3ppb ($p = 0.087$) treatment (Figure A2.2B). Additionally, variation in the proportion of offspring that hatched increased with increasing atrazine concentration. Note that the number of families were seven.

Table A2.1: Numbers of zebrafish offspring survived to 24hpf. Data are summarised across all families per treatment, number of families per treatment, $n=7$.

Treatment	n (Total)	n (Total survived)	n (Total died)
Control	692	608	84
0.3ppb	768	650	118
3ppb	720	529	191
30ppb	614	553	61

Table A2.2: Numbers of zebrafish offspring that hatched at 48hpf. Data are summarised across all families per treatment, number of families per treatment $n=7$.

Treatment	n (Total)	n (Hatched)	n (Did not hatch)
Control	604	416	188
0.3ppb	650	418	232
3ppb	525	261	264
30ppb	552	308	244

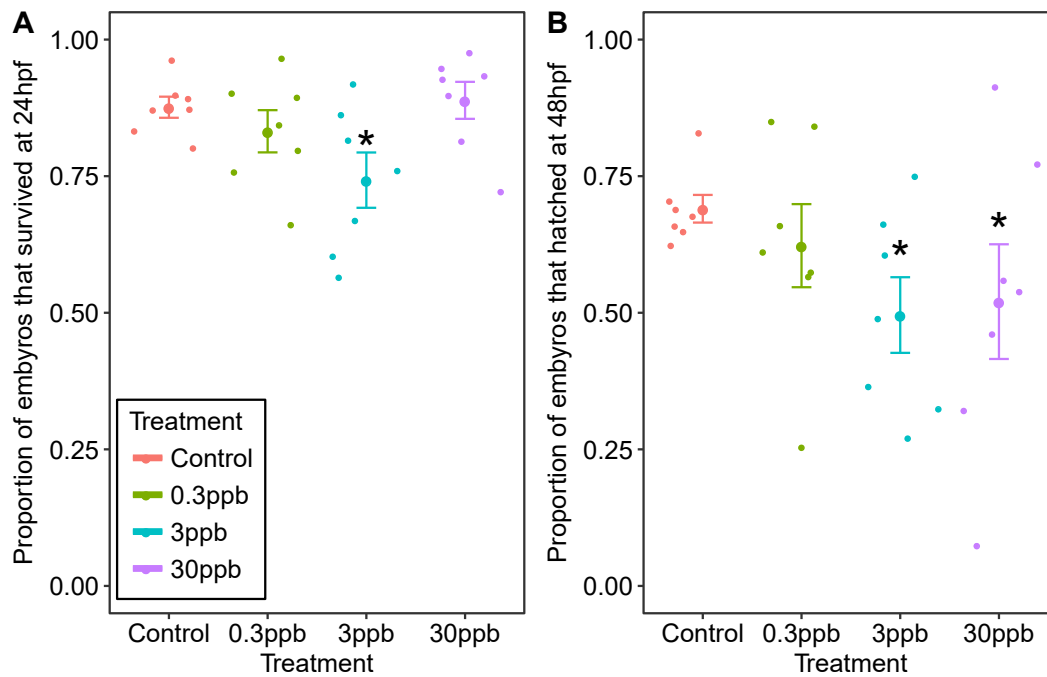


Figure A2.2: Early life parameters of zebrafish embryos from untreated and atrazine treated fathers. A) the proportion of embryos that survived to 24hpf. B) the proportion of embryos that hatched by 48hpf. A * indicates significant differences ($p < 0.05$). Large points represent means, with error bars representing standard errors of the mean. Smaller, randomly jittered (along the x-axis) points are the raw proportions from each family, from each treatment. Total number of families per treatment were $n=7$.

Appendix III: Sample sizes used in behavioural assays

Figure legends from figures depicting behavioural phenotypes (in main text, see Chapter 2) show ranges of the sample sizes used in order to improve general readability. This was due to un-evenness of samples across sex and treatments (in particular amongst the F_1 's), in addition, this was also due to some F_1 individuals (8 fish in total) being excluded from the analyses of behaviours measured in the novel object assay caused by experimental error (see Chapter 2, Methods, Faulty tracking). Therefore, below are the sample sizes used for all behavioural test (Table A3.1).

Table A3.1: Sample sizes of zebrafish used in behavioural assays across each assay.

Novel arena test	F₀		F₁	
	Male	Female	Male	Female
Control	12	12	21	28
0.3ppb	12	9	22	24
3ppb	12	12	18	30
30ppb	12	12	18	28
Novel object test				
Control	12	12	19	27
0.3ppb	12	9	21	24
3ppb	12	12	17	30
30ppb	12	12	18	27
Mirror test				
Control	12	12	21	28
0.3ppb	12	9	22	24
3ppb	12	12	18	30
30ppb	12	12	18	28

Appendix IV: Repeatability of zebrafish aggression using a mirror stimulus assay

Objectives

In order to understand the impact of adult zebrafish aggression under an atrazine exposure during juvenile development, an assessment of a mirror test was undertaken to discern consistent differences between individuals (animal personality). I assessed the repeatability of two behavioural measures thought to be representative of aggression; the time spent interacting with the mirror and the frequency entering the mirror zone in non-exposed adults wildtype (AB) strain over the course of three sampling dates.

Methods

Experimental overview

The zebrafish used in the experiment were part of a long term repeatability project that ran five behavioural sampling periods over the course of 27 weeks, from 13/10/16 to 14/04/17 (Thomson 2017). This study took place in between sampling point four and just after sampling point 5 (week 14 to just after week 27) of that experiment. In total, 41 zebrafish (date of birth; 22/03/2016) were used (comprising 20 females and 21 males). Each fish was uniquely tagged by injecting visible implant elastomers at the start of the long term repeatability experiments, one on either side of the fish's body resulting in a unique and identifiable colour combination (colours used were; yellow, green, red or orange). Zebrafish were fed twice daily with dry food; ZM400 and once a day with live *Artemia salina*. On the day of testing, zebrafish were

fed after the assay. Temperature during housing was maintained at 25°C with a 14:12 hour light:dark, dawn-dusk lighting regime. Three different time points were chosen to assess repeatability; 0 days, 7 days and 28 days, initial behavioural tests were performed on the 18th and 19th of March 2017, after which the seven day repeatability tests were conducted on the 25th and 26st of March, the 28 day repeatability tests were conducted on the 15th and 16th of April 2017.

Experimental setup

The experimental setup follows the design described in methods of Chapter 2 (see Chapter 2, Methods, Behavioural phenotyping; Experimental protocols; adapted from Ariyomo et al. (2012); Way et al. (2015)), with slight differences in filming equipment and volume of water used. In brief, three identical glass tanks (length=30cm, width=15.5cm and height=27cm) were used side by side as test arenas for the assay. A mirror (length=19cm, width=15cm) was fitted externally to the side of the tank; two tanks with mirrors fixed on the right and one mirror placed on the left. White plastic film was fitted to the inside of tanks to block external stimulus between tanks and to minimise unwanted reflections during filming, the side facing the camera was not covered. A removable opaque plastic partition was used to cover up mirrors. Tanks were filled with a combination of 3L system water and 3L aerated adjusted water and left to age overnight. Trials took place over two days, testing half of the fish on one day, and half the next. In preparation for repeatability tests the following day, 3L of tested water was replaced in each tank and left to age overnight. Two-three days before assessment, males and females were separated within their holding tanks (length=18cm, width=11cm and height=8.5cm) to allow easy individual identification after each assay. Before entering the test arena, fish were netted from their holding tanks to an intermediary tank for 2-4min to homogenise entering time. Three fish were filmed simultaneously per assay with one individual per test tank at a time. The aggression behavioural assay started with an initial 10min acclimation period (i.e. a novel arena test), after which, the opaque plastic partition was removed, revealing the mirror stimulus. To minimise the interference of unsettled water in tracking ability, filming commenced 1min after the last mirror was revealed. Filming during the mirror assay lasted for 10min. Fish were netted out of the test arena into their intermediary holding tanks and their unique colour combinations were identified with a UV torch. Trials were filmed using GoPro Hero3+ cameras at a resolution of 1080p at 60 frames per second. Cameras were placed at distance of (~ 15cm away from the tank) along the mirrors edges, capturing roughly 1/2 (~15cm) of the entire tank during filming (see Figure A4.1). Each camera was placed on a make-shift LEGO® platform to achieve required height relative to the distance to maximise vision in the assay. Platforms were covered in white plastic film to block colour variation from the LEGO® blocks. To control for variation in circadian

rhythms, each tank was tested within 30 minutes of the last time it was tested; i.e. tank one was always tested between 09:30 and 12:00, while tank two was always tested between 12:50 and 15:00. After filming, measures of aggression were recorded using EthoVision (Noldus et al. 2001). A mirror zone ~ 5 cm from the mirror was drawn and the time spent in the mirror zone as well as the frequency of times entering the mirror zone (sec) was recorded (Figure A4.1).

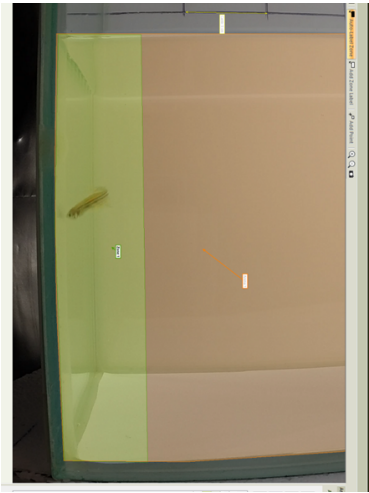


Figure A4.1: EthoVision setup for investigating repeatability during the mirror test. The light green region depicts the mirror zone area, while the orange zone depicts the rest of the testing arena.

Statistical analysis

Repeatability of each behavioural measure was analysed using the ‘*rptR*’ package (Stoffel et al. 2017) in R, version 3.5.0 (Team 2018), with permutations and bootstrap set to 1000. The cumulative duration spent in the mirror zone was normally distributed and analysed under the ‘gaussian datatype’. The frequency of times entering the mirror zone was analysed using the ‘poisson datatype’.

Results

The time spent within the mirror zone was significantly repeatable, for both sexes and across all time points, with the exception between week and a month for males (Table A4.1). Overall, females were more repeatable than males, 0.52 compared to 0.35. Figure A4.2 depicts the raw values for all individuals.

The frequency of entering the mirror zone was significantly repeatable between the initial and one week tests, but was not repeatable overall (Table A4.2). After the week test, the majority of males declined sharply in

their frequency, whereas females showed a combination of increases and decreases in their frequency (Figure A4.3).

Table A4.1: Repeatability estimates (R) of the time spent within the mirror zone, including lower and upper confidence intervals (CI), standard error (SE) and P values. Repeatability estimates are separated by time points and sex. Males, $n_{[\text{individuals}]}=21$, $n_{[\text{total observations}]}=63$; females, $n_{[\text{individuals}]}=20$, $n_{[\text{total observations}]}=60$.

Time Points	Group	R	Lower 95% CI	Upper 95% CI	SE	P value
Initial - Week	Combined	0.67	0.44	0.80	0.09	< 0.001
	Male	0.57	0.22	0.81	0.15	0.001
	Female	0.72	0.42	0.88	0.12	< 0.001
Week - Month	Combined	0.406	0.131	0.631	0.127	0.003
	Male	0.287	0.000	0.647	0.173	0.057
	Female	0.504	0.097	0.762	0.167	0.008
Initial - Month (Overall)	Combined	0.439	0.242	0.616	0.093	< 0.001
	Male	0.351	0.102	0.609	0.132	< 0.001
	Female	0.520	0.245	0.725	0.129	< 0.001

Table A4.2: Repeatability estimates (R) of the frequency entering the mirror zone, including upper and lower confidence intervals (CI) standard error (SE) and P values. Repeatability estimates are separated by time points and sex. Males, $n_{[\text{individuals}]}=21$, $n_{[\text{total observations}]}=63$; females, $n_{[\text{individuals}]}=20$, $n_{[\text{total observations}]}=60$.

Time Points	Group	R	Lower 95% CI	Upper 95% CI	SE	P value
Initial - Week	Combined	0.61	0.38	0.76	0.1	< 0.001
	Male	0.70	0.39	0.85	0.12	< 0.001
	Female	0.53	0.08	0.78	0.17	0.006
Week - Month	Combined	0	0	0.28	0.08	0.5
	Male	0	0	0.33	0.10	1
	Female	0	0	0.41	0.13	0.5
Initial - Month (Overall)	Combined	0.08	0	0.25	0.075	0.176
	Male	0.14	0	0.37	0.11	0.081
	Female	0.06	0	0.33	0.10	0.334

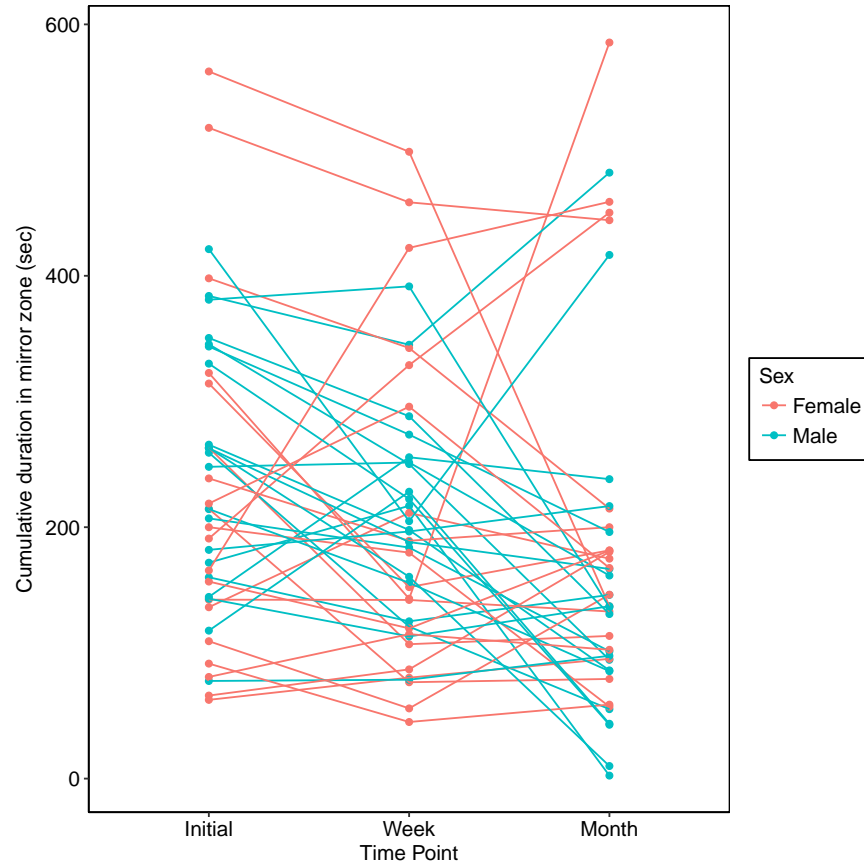


Figure A4.2: Raw values (points) of the cumulative duration spent in the mirror zone of each individual, connected across all three time points (lines); Initial, week and month. Males are coloured in blue, $n_{\text{individuals}}=21$, $n_{\text{observations}}=63$; females in pink, $n_{\text{individuals}}=20$, $n_{\text{observations}}=60$.

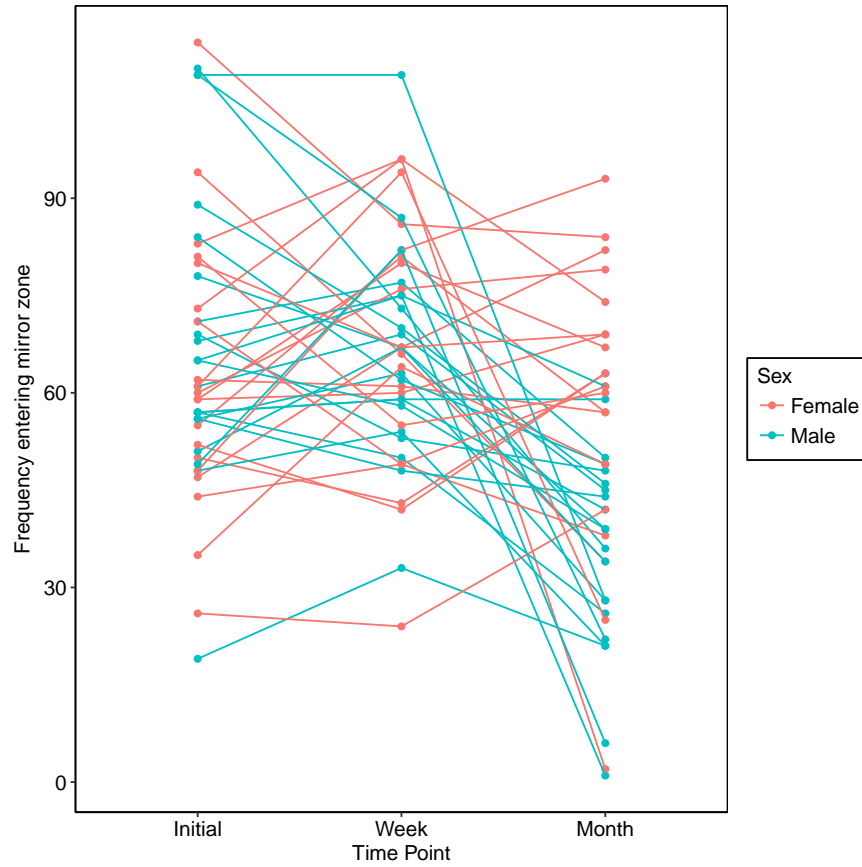


Figure A4.3: Raw values (points) of the frequency entering the mirror zone of each individual, connected across all three time points (lines); Initial, week and month. Males are coloured in blue, $n_{\text{individuals}}=21$, $n_{\text{observations}}=63$; females in pink, $n_{\text{individuals}}=20$, $n_{\text{observations}}=60$.

Appendix V: Statistical analyses for behavioural measures

All statistical analyses were conducted in R, version 3.5.0 (Team 2018). All analyses described below contained the predictors; treatment (i.e. controls, 0.3ppb, 3ppb, 30ppb), and sex (male and female) with an interaction term. The interaction term was removed if there was no influence and was re-analysed without it. All F_0 behavioural measures were analysed with linear models (LMs) or generalised linear models (GLMs). All F_1 behavioural measures were analysed with linear mixed effects models (LMMs) or with generalised linear mixed effects models (GLMMs) using the package ‘*lme4*’ version 1.1-1.3 (Bates et al. 2015) with family id included as a random effect to account for genetic influences (with some exceptions, see Appendix V). Specific details on each behaviour analysed can be seen in Appendix V.

Exploration and measures of anxiety, the time spent in the bottom zone of the novel arena test and the time spent in the bottom zone of the novel object test were analysed using a LM.

Analyses for measures of boldness are described hereafter. The time spent in the top zone of the novel arena test was analysed using a LM. The latency to enter the top zone was right skewed and therefore analysed with a (GLM) using a gamma error structure (log-link function). The assumptions of a GLM with a gamma error structure are that zeros are unable to be present as they skew the results, therefore if a fish was in the top zone at the start of the trial; it was given a latency of 0.1 sec, if the fish never entered the top zone; it was given a latency of 600 sec (the total duration of the assay). For the F_0 's, the frequency entering the top zone of the novel arena was analysed using a GLM with a quasi-poisson error structure (log-link function) to account for over-dispersion. For the F_1 's however, frequency was analysed with a GLMM with a poisson error structure (log-link function), with two random effects; family, to account for genetic influence and fish id, to account for over-dispersion (a quasi-poisson error structure is not possible with random effects). Interacting with the novel object was analysed using a GLM with a binomial error structure (logit-link

function) where the binary response was ‘interacted with the novel object’ (1), or ‘did not interact with the novel object’ (0).

For measures of aggression, the time spent interacting with the mirror was analysed with a LM. For the frequency entering the mirror zone, the F_0 ’s were analysed using a GLM with a quasi-poisson error structure (log-link function) to account for over-dispersion. The F_1 ’s however, frequency entering the mirror zone was analysed with a GLMM with a poisson error structure (log-link function), with two random effects; family, to account for genetic influence and fish id, to account for over-dispersion.

Analyses for measures of activity are described hereafter. The time spent not moving in the novel arena, total distance moved and the mean velocity in each of the three behavioural tests were analysed with LMs. I also investigated how the influence of the different assays employed might have affected activity levels. I ran a LMM with assay type (i.e. novel arena, novel object and mirror test), treatment and with sex as predictors and the total distance moved as the response (mean velocity reported in Appendix VI). For the F_0 ’s, fish id was included as a random effect to account for pseudo-replication. For the F_1 ’s, fish id nested within family was included as a random effect (Harrison 2014)

I investigated how the influence of the different assays employed might have affected activity levels. Assays were compared against one another using a LMM with assay type (i.e. novel arena, novel object and mirror test), treatment and sex as predictors, and the total distance moved as the response (mean velocity reported in Appendix VIII). For the F_0 ’s, fish ID was included as a random effect to account for repeated measures on the same individuals. For the F_1 ’s, fish ID nested within family was included as a random effect (results can be found in Appendix VI).

In some instances of F_1 analyses, family level variation was suspected of influencing the model outputs. This would indicate a genetic component of behaviour, i.e. siblings regardless of sex would exhibit a similar phenotype to each other than to individuals from another family, even if from the same treatment, and even if the behavioural measure usually exhibited sex differences. Therefore models were additionally tested against itself with and without the random effect included using a likelihood ratio test (LRT). Differences between models are reported below when they occurred with the level of variance and the standard deviation (SD) from the LMM or GLMM model output (all measures are summarised in Appendix VIII, Table A9.1). Family variation was also visually compared across treatments to investigate if atrazine altered the amount of variation present at the family level (Appendix VIII).

Appendix VI: Additional measures of personality

Results

Anxiety

Overall, regardless of treatment F_0 males spent an average of 352.12 ± 18.78 sec (range 76.83–580.92 sec) and F_0 females spent an average 382.87 ± 22.45 sec (range 23.04–600 sec; the entirety of the assay) in the bottom zone during the novel arena test.

During the novel object assay the mean time spent in the bottom zone was 363.3 ± 30.92 sec for F_0 males (range 0–600 sec) and 419.67 ± 27.44 sec (range 0–600 sec) for F_0 females. For F_1 males mean time was 355.45 ± 24.96 sec (range 0–600 sec) and by F_1 females was 437.47 ± 21.80 (range 0–600 sec). There was no difference between treatments or sexes in the amount of time spent within the bottom half for the F_0 's (Figure A6.1). In the F_1 's, fish from the 0.3ppb treatment spent more in the bottom zone than controls (Est. 180.19 [13.18, 347.20 CI]; Figure A6.1) though the difference was only marginal ($p = 0.06$). Fish from the 3ppb treatment also spent more time in the bottom zone, but the difference was marginally non-significant (Est. 163.90 [−2.92, 330.71 CI]). There was no difference observed in the higher 30ppb treatment. F_1 males overall, spent less time in the bottom zone of the novel object test compared to females (Est. −78.44 [−137.31, −19.57 CI]). The family level variance was 8301 sec with a SD of 91.11 sec and explained ($\chi^2 = 7.2$, $df = 1$, $p < 0.007$) a significant portion of variation present in the LMM model (Appendix VIII). Like-wise to the novel arena test, there appears to be a suggestive trend of more time spent in the bottom zone of the novel object assay with fish from treated fathers than from fish from untreated fathers (Figure A6.1).

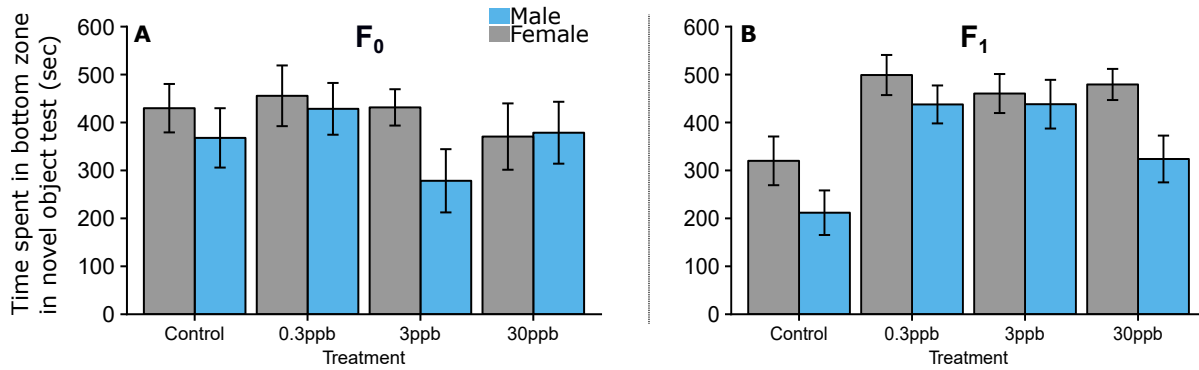


Figure A6.1: Additional measure of anxiety on atrazine exposed F₀ fish (left side) and F₁ offspring from exposed fathers (right side). (A) and (B) the time spent in the bottom zone during the novel object test. Bars represent means, with error bars representing standard errors of the mean. For each sex and treatment, sample sizes for F₀'s ranged from n=9-12; F₁'s ranged from n=17-30. Total sample size for F₀'s were n=93; F₁'s n=190, with n=3 families per treatment.

Exploration

Average exploration regardless of treatment by F₀ males was 46.53 ± 2.52 (range 19.20–89.37) and was 53.71 ± 3.66 sec (range 23.83–158.57) by F₀ females. F₁ males averaged 48.75 ± 2.67 (range 15.38–109.69) and females averaged was 60.60 ± 3.04 (range 19.90–176.93).

Boldness

Overall, mean latency for F₀ males was 111.94 ± 19.26 sec and for F₀ females was 170.82 ± 28.54 sec. In total, 8 fish (8.6%) of fish were in the top zone at beginning of recording and 6 fish 6.45% never entered. Excluding these individuals, latency amongst the F₀'s ranged from 3.50–502.60 sec for males and 2.52–518.92 sec for females. Mean latency for F₁ males was 84.10 ± 15.56 sec and for F₁ females was 170.74 ± 18.32 sec. In total, 47 fish (24.7%) of fish were in the top zone at beginning of recording and 12 fish (6.3%) never entered. Excluding these individuals, latency amongst the F₁'s ranged from 0.26–533.63 sec for males and 0.60–549.20 sec for females.

There was no differences in treatment in either the F₀ or F₁ generation in the frequency to enter the top zone once over-dispersion was accounted for (A6.2). F₀ males overall exhibited a higher frequency of entering the top zone than females (Est. 1.50 [1.08, 2.10 CI]) and so too did F₁ males (Est. 2.04 [1.51, 2.75 CI]). Mean frequency of F₀ males was 23.58 ± 2.64 (range 0–94), and F₀ females was 16.84 ± 1.872 (range 0–49). The mean frequency for F₁ males was 38.0 ± 2.74 (range 0–119) and for F₁ females was 20.90 ± 1.7 (range 0–82).

There was no significant difference between treatments across both generations in the time spent in the

top zone during the novel arena. However, the time spent by F_0 fish from the 0.3ppb treatment was non-significantly less (Est. -62.22 [$-132.79, 8.36$ CI] A6.2). There was no sex difference observed amongst the F_0 's. However, amongst the F_1 's, males spent significantly longer than females in the top zone (Est. 84.21 [$43.09, 125.32$ CI]). Family level variation was 11693 sec with a SD of 108.1 and explained a significant portion of the variance in this model ($\chi^2 = 35.53$, $df = 1$, $p < 0.001$; Appendix III). Overall, F_0 male averaged 164.30 ± 17.025 sec (range $0-451.97$ sec), and female averaged 131.71 ± 18.36 sec (range $0-512.86$ sec) of time spent in the top zone. Male F_1 's averaged 255.87 ± 19.44 (range $0-599.10$) and female F_1 's averaged 167.81 ± 16.06 (range $0-597.0$).

In both generations there was no treatment effect observed in whether or not an individual approached the novel object (A6.2). In the F_0 's there was no sex effect observed, whereas in the F_1 generation, males approached the novel object significantly more so than females did (Est. 0.87 [$1.17, 2.66$ CI]). In total, 48 (52%) F_0 fish and 59 (32%) F_1 fish, regardless of treatment or sex approached the novel object over the course of the novel object assay.

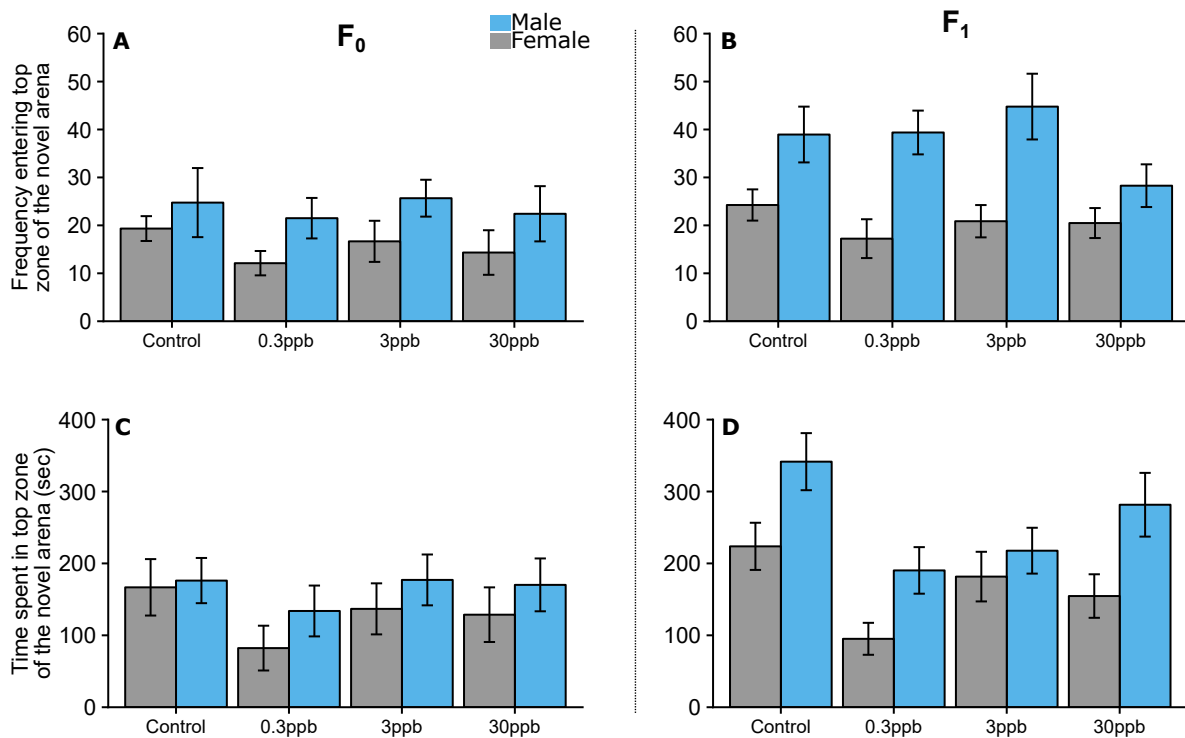


Figure A6.2: Additional measures of boldness on atrazine exposed F_0 fish (left side) and F_1 offspring from exposed fathers (right side). (A) and (B) the frequency entering the top zone of the novel arena; (C) and (D) the time spent in the top zone of the novel arena (sec). Bars represent means, with error bars representing standard errors of the mean. For each sex and treatment, sample sizes for F_0 's ranged from $n=9-12$; F_1 's ranged from $n=17-30$. Total sample size for F_0 's were $n=93$; F_1 's $n=182-190$, with $n=3$ families per treatment.

Aggression

Mean time interacting with the mirror by F_0 males was 225.99 ± 189.6 (range 0–577.96) and was 189.60 ± 21.19 sec (range 0–473.0) by F_0 females.

There was no difference amongst treatments in the frequency a fish entered the mirror zone amongst the F_0 's once over-dispersion was taken into account (Figure A6.3). Overall, F_0 males exhibited a higher frequency of entering the mirror zone than females (Est. 1.33 [1.03, 1.73]). Male F_0 average frequency was 49.46 ± 3.59 (range 0–99) and female F_0 average frequency was 37.04 ± 4.26 (range 0–112). In the following generation, the frequency entering the mirror zone by F_1 females tended to decrease as atrazine concentration increased, whereas males tended to increase (Figure A6.3). However, an interactive difference between sex and treatment was only observed in the 3ppb treatment (Est. 3.75 [1.45, 9.68 CI]) compared to controls, with no other treatment effects observed. F_1 males were slightly more likely to enter the mirror zone at a higher frequency than females but not significantly so (Est. -1.76 , [-3.42 , 1.11]). F_1 mean male frequency was 36.77 ± 2.7 (range 0–87) and mean F_1 female frequency was 36.93 ± 2.44 (range 0–100). Family level variation was 0.03 with a SD of 1.16 and significantly explained a portion of the GLMM ($\chi^2 = 127.84$, $df = 1$, $p < 0.001$; Appendix VIII).

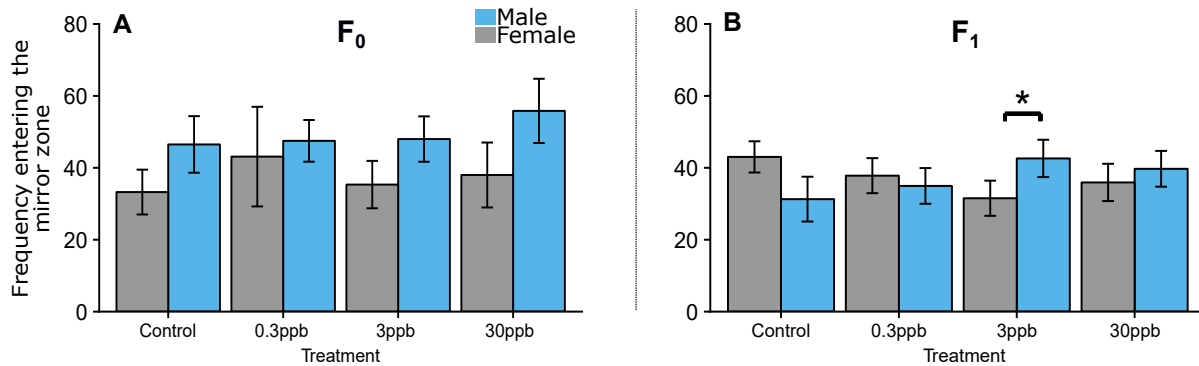


Figure A6.3: Additional measure of aggression on atrazine exposed F_0 fish (left side) and F_1 offspring from exposed fathers (right side). (A) and (B) the frequency entering the mirror zone. Bars represent means, with error bars representing standard errors of the mean. For each sex and treatment, sample sizes for F_0 's ranged from $n=9-12$; F_1 's ranged from $n=17-30$. Total sample size for F_0 's were $n=93$; F_1 's $n=190$, with $n=3$ families per treatment.

Activity

There was a significant sex difference observed in both generation with males spending less time immobile (F_0 males Est. -0.34 [$-0.57, -0.11$ CI]; F_1 males Est. -0.55 [$-0.79, -0.31$ CI]). However no treatment differences were observed in either generation in the total distance moved nor in the time spent immobile (A6.4), though F_1 fish from the 30ppb treatment were marginally less immobile than controls ($p = 0.083$). In addition, despite the measures total distance moved and mean velocity being highly correlated in the novel arena assay (correlation coefficient = 0.87 , $R^2 = 0.76$; Appendix VI), F_1 fish from the 30ppb treatment had a significantly lower mean velocity than controls (Est. -0.83 [$-1.62, -0.03$]). During the novel arena, the mean total distance for F_0 males was 2974.48 ± 131.36 cm (range 1131.24 – 5596.57 cm) and for F_0 females was 2445.70 ± 102.20 cm (range 902.92 – 4643.76 cm). Mean total distance for F_1 males was 4044.20 ± 170.34 cm (range 887.60 – 8536.37 cm) and for F_1 females was 2780.29 ± 96.88 cm (range 11.66 – 5518.67 cm). The mean time spent immobile for F_0 males was 98.11 ± 9.4 sec (range 15.16 – 340.59 sec) and for females was 134.58 ± 13.30 sec (range 50.08 – 452.85 sec). F_1 males was 90.31 ± 11.30 sec (range 10.40 – 480.93 sec) and for F_1 females was 141.58 ± 11.72 sec (4.03 – 600 sec).

Mean total distance moved in the novel object assay for F_0 males was 1806.16 ± 138.92 cm (range 145.78 – 4867.44 cm) and for F_0 females was 1173.47 ± 102.66 (range 85.09 – 2836.30 cm). Mean total distance for F_1 males was 2659.45 ± 166.99 cm (range 0 – 8341.81 cm) and for F_1 females was 1380.96 ± 82.01 cm (range 0 – 3820.19 cm).

Mean total distance for F_1 males was 2569.01 ± 114.80 cm (range 0 – 5580.16 cm) and for F_1 females was 1880.54 ± 98.76 cm (range 0 – 3646.06 cm).

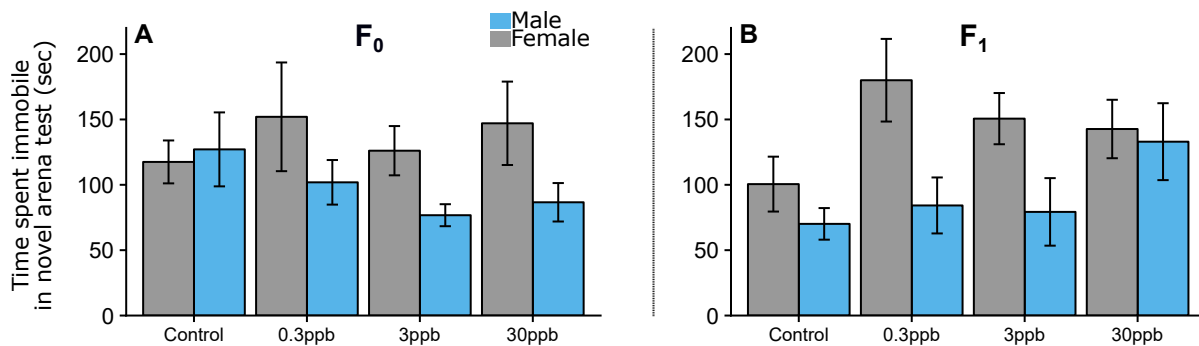


Figure A6.4: Additional measures of activity on atrazine exposed F_0 fish (left side) and F_1 offspring from exposed fathers (right side). (A) and (B) the time spent immobile during the novel arena test (sec); (C) and (D) the total distance moved during the novel object test. Bars represent means, with error bars representing standard errors of the mean. For each sex and treatment, sample sizes for F_0 's ranged from $n=9$ – 12 ; F_1 's ranged from $n=17$ – 30 . Total sample size for F_0 's were $n=93$; F_1 's $n=180$ – 190 , with $n=3$ families per treatment.

Differences between assays in activity

After the conclusion of the novel arena and the start of the novel object assay, the addition of the novel object into the arena caused a strong drop in the total distance moved (F_0 Est. -1218.6 [$-1383.43, -1053.77$ CI]; F_1 Est. -1408.50 [$-1584.37, -1232.64$ CI]). On average F_0 males decreased by 39.3% and females decreased by 52.3%. F_1 males similarly decreased by 34.2% and F_1 females decreased by 48.2%. Activity levels increased across most of the treatments during the mirror test (there some instances of a decrease e.g. in F_0 0.3ppb females, and in F_1 0.3ppb and 30ppb males) but were still significantly lower than during the novel arena assay (F_0 , Est. -994.23 [$-1159.06, -829.40$ CI]; F_1 Est. -1145.07 [$-1315.96, -974.19$ CI]). On average this increase in the mirror assay was relatively, much smaller than the initial decrease between the novel arena and novel object. On average F_0 males increased by 14.4% and females increased by 15.09%. F_1 males however decreased on average by 2.65% whereas females increased by 36.1%. The sexes differed overall in activity (F_0 male Est. 621.40 [$357.44, 885.37$ CI]; F_1 male Est. 1072.51 [$806.94, 1338.07$ CI]) as reported earlier (see section; Activity), but both sexes exhibited a similar reaction to the novel object. Lastly, there were no treatment differences observed in activity when compared across all three assays, though treatment effects were found previously when assays were analysed independently of each other (e.g. in the F_1 mirror test). Amongst the F_0 fish, there was high variation across individuals, with fish identification producing a variance of 310151 cm with a SD of 556.9, that significantly helps explain a portion of the variation ($\chi^2 = 56.5$, $df = 1$, $p < 0.001$; Appendix VIII) in the model. The variance was similarly high in the F_1 's with fish id nested within family id producing a variance of 626963 cm with a SD of 791.8. The inclusion of the nested random factor helped significantly explain a portion of the model's variance ($\chi^2 = 99.2$, $df = 1$, $p < 0.001$; Appendix VIII).

Appendix VII: The correlation between the total distance moved and mean velocity

Across all assays, the total distance moved was highly correlated with mean velocity. I provide the correlation coefficient between the total distance moved and mean velocity for each assay and for each generation. I additionally provide the R^2 from the linear models between these two variables as an another congruent measure, and also to account for pseudoreplication at the individual level and at the family level for the F_1 generation. For the F_1 generation, the R^2 was calculated using the 'MuMin' package (reference).

Table A7.1: Correlation coefficients (cor. coef), R^2 and P -values of the relationship between the total distance moved and the mean velocity during the novel arena, novel object and mirror test assay across both F_0 and F_1 generations.

Novel Arena		F_0			F_1		
	Cor. coef	R^2	P -value	Cor. coef	R^2	P -value	
	0.99	0.98	<0.001	0.87	0.76	<0.001	
Novel object		F_0			F_1		
	Cor. coef	R^2	P -value	Cor. coef	R^2	P -value	
	0.99	0.98	<0.001	0.87	0.76	<0.001	
Mirror test		F_0			F_1		
	Cor. coef	R^2	P -value	Cor. coef	R^2	P -value	
	0.82	0.66	<0.001	0.64	0.41	<0.001	

Appendix VIII: Model parameters of behavioural analyses

Table A8.1: Model coefficients of F_0 behavioural measures.

Parameter	Estimate	Variance	Lwr95%CI	Upr95%	Z/t	P
Anxiety measures						
Time in bottom zone NA, $F_{0,n=93}$						
Intercept	349.13	31.97	286.46	411.79	10.92	<0.001
0.3ppb	83.03	41.76	1.18	164.87	1.99	0.0499
3ppb	27.18	40.29	-51.80	106.15	0.67	0.5018
30ppb	37.09	40.29	-41.88	116.06	0.92	0.3598
Sex(male)	-33.83	29.01	-90.70	23.03	-1.17	0.2467
Time in bottom zone NO, $F_{0,n=93}$						
Intercept	428.73	46.00	338.58	518.88	9.32	<0.001
0.3ppb	45.56	60.08	-72.19	163.30	0.76	0.4500
3ppb	-43.92	57.97	-157.53	69.69	-0.76	0.4510
30ppb	-24.21	57.97	-137.82	89.41	-0.42	0.6770
Sex(male)	-59.72	41.74	-141.53	22.09	-1.43	0.1560
Boldness measures						
Latency to enter top zone $F_{0,n=93}$						
Intercept	81.45	1.38	46.53	162.39	13.96	<0.001
0.3ppb	2.16	1.62	-1.19	5.75	1.60	0.1135

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Table A8.1 – *Continued from previous page*

3ppb	2.36	1.57	−1.03	5.75	1.93	0.0574
30ppb	2.86	1.57	1.17	6.96	2.36	0.0205
Sex(male)	1.99	1.57	−1.22	4.85	1.55	0.1253
0.3ppb:Sex(male)	−3.54	1.93	−13.20	1.02	−1.93	0.0572
3ppb:Sex(male)	−3.88	1.88	−13.46	−1.12	−2.15	0.0345
30ppb:Sex(male)	−5.39	1.88	−18.73	−1.55	−2.67	0.0091
Frequency entering top zone $F_{0,n=93}$						
Intercept	17.62	1.20	12.18	24.78	15.73	<0.001
0.3ppb	−1.30	1.27	−2.10	1.22	−1.09	0.2771
3ppb	−1.04	1.24	−1.60	1.48	−0.19	0.8526
30ppb	−1.20	1.25	−1.88	1.30	−0.81	0.4223
Sex(male)	1.50	1.18	1.08	2.10	2.43	0.0173
Time in top zone $F_{0,n=93}$						
Intercept	153.89	27.57	99.85	207.92	5.58	<0.001
0.3ppb	−62.22	36.01	−132.79	8.36	−1.73	0.0875
3ppb	−14.51	34.74	−82.60	53.59	−0.42	0.6773
30ppb	−22.01	34.74	−90.10	46.09	−0.63	0.5281
Sex(male)	35.09	25.02	−13.94	84.13	1.40	0.1642
Approach NO $F_{0,n=93}$						
Intercept	0.45	0.61	−0.25	0.67	−0.43	0.6700
0.3ppb	0.52	0.65	−0.25	0.78	0.11	0.9100
3ppb	0.63	0.64	−0.35	0.85	0.87	0.3820
30ppb	0.42	0.64	−0.18	0.69	−0.58	0.5610
Sex(male)	0.60	0.60	−0.39	0.77	0.93	0.3520
Time in middle zone $F_{0,n=93}$						
Intercept	96.50	10.01	76.89	116.11	9.65	<0.001
0.3ppb	−21.00	13.07	−46.61	4.61	−1.61	0.1120
3ppb	−12.81	12.61	−37.53	11.90	−1.02	0.3120
30ppb	−14.89	12.61	−39.60	9.82	−1.18	0.2410

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Table A8.1 – *Continued from previous page*

Sex(male)	−0.96	9.08	−18.75	16.84	−0.11	0.9160
Exploration measures						
Exploration $F_{0,n=93}$						
Intercept	47.06	4.76	37.74	56.39	9.89	<0.001
0.3ppb	15.12	6.21	2.94	27.30	2.43	0.0170
3ppb	4.11	6.00	−7.65	15.86	0.69	0.4953
30ppb	9.48	6.00	−2.27	21.23	1.58	0.1174
Sex(male)	−7.71	4.32	−16.17	0.76	−1.78	0.0778
Activity measures						
Total distance moved in NA $F_{0,n=93}$						
Intercept	2355.00	187.69	1987.14	2722.85	12.55	<0.001
0.3ppb	36.19	245.13	−444.27	516.64	0.15	0.8830
3ppb	192.33	236.53	−271.26	655.92	0.81	0.4183
30ppb	120.66	236.53	−342.93	584.25	0.51	0.6112
Sex(male)	532.19	170.32	198.37	866.02	3.13	0.0024
Mean velocity in NA $F_{0,n=93}$						
Intercept	4.01	0.30	3.42	4.61	13.22	<0.001
0.3ppb	−0.02	0.40	−0.80	0.76	−0.05	0.9581
3ppb	0.30	0.38	−0.45	1.05	0.78	0.4359
30ppb	0.11	0.38	−0.64	0.86	0.30	0.7667
Sex(male)	0.89	0.28	0.35	1.43	3.23	0.0018
Time spent not moving in NA $F_{0,n=93}$						
Intercept	4.80	0.13	4.55	5.05	37.49	<0.001
0.3ppb	0.02	0.17	−0.30	0.35	0.14	0.8874
3ppb	−0.12	0.16	−0.43	0.20	−0.71	0.4775
30ppb	−0.06	0.16	−0.38	0.25	−0.39	0.6985
Sex(male)	−0.34	0.12	−0.57	−0.11	−2.92	0.0045
Total distance moved in NO $F_{0,n=93}$						
Intercept	1126.68	195.18	744.14	1509.22	5.77	<0.001

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Table A8.1 – *Continued from previous page*

	0.3ppb	66.36	254.92	−433.27	566.00	0.26	0.7952
	3ppb	159.16	245.97	−322.94	641.26	0.65	0.5193
	30ppb	−33.49	245.97	−515.58	448.61	−0.14	0.8920
	Sex(male)	631.47	177.12	284.32	978.62	3.57	0.0006
Mean velocity in NO (F0),n=93							
	Intercept	2.02	0.32	1.40	2.64	6.41	<0.001
	0.3ppb	0.03	0.41	−0.78	0.83	0.06	0.9509
	3ppb	0.25	0.40	−0.53	1.03	0.62	0.5349
	30ppb	−0.10	0.40	−0.88	0.68	−0.24	0.8091
	Sex(male)	1.02	0.29	0.46	1.58	3.55	0.0006
Total Distance Moved in MIS (F0),n=93							
	Intercept	1344.41	168.68	1013.80	1675.01	7.97	<0.001
	0.3ppb	−41.13	220.31	−472.93	390.67	−0.19	0.8520
	3ppb	141.26	212.58	−275.39	557.90	0.66	0.5080
	30ppb	−33.92	212.58	−450.57	382.72	−0.16	0.8740
	Sex(male)	700.55	153.07	400.53	1000.57	4.58	<0.001
Mean velocity in MIS (F0),n=93							
	Intercept	2.45	0.25	1.96	2.93	9.81	<0.001
	0.3ppb	0.04	0.33	−0.59	0.68	0.13	0.8936
	3ppb	0.63	0.31	0.02	1.25	2.01	0.0477
	30ppb	0.43	0.31	−0.18	1.05	1.37	0.1730
	Sex(male)	0.97	0.23	0.53	1.41	4.29	<0.001
Aggression measures							
Time spent interacting with mirror (F0),n=93							
	Intercept	216.92	34.03	150.22	283.62	6.37	<0.001
	0.3ppb	2.24	44.45	−84.88	89.36	0.05	0.9600
	3ppb	−46.76	42.89	−130.82	37.30	−1.09	0.2790
	30ppb	−57.37	42.89	−141.43	26.70	−1.34	0.1850
	Sex(male)	34.54	30.88	−26.00	95.07	1.12	0.2660

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Table A8.1 – *Continued from previous page*

Fq interacting with mirror (F0),n=93							
Intercept	34.20	1.17	25.03	45.60	22.85	<0.001	
0.3ppb	1.12	1.21	-1.30	1.63	0.60	0.5490	
3ppb	1.04	1.21	-1.39	1.51	0.23	0.8160	
30ppb	1.18	1.20	-1.22	1.68	0.89	0.3770	
Sex(male)	1.33	1.14	1.03	1.73	2.15	0.0340	
Comparison between assays							
Total Distance Moved between assays							
Intercept	2346.31	156.15	2040.26	2652.35	15.03	<0.001	
Assay(Nobject)	-1218.60	84.10	-1383.43	-1053.77	-14.49	<0.001	
Assay(Mirror)	-994.23	84.10	-1159.06	-829.40	-11.82	<0.001	
Sex(Male)	621.40	134.68	357.44	885.37	4.61	<0.001	
0.3ppb	20.47	193.84	-359.44	400.38	0.11	0.916	
3ppb	164.25	187.03	-202.33	530.83	0.88	0.382	
30ppb	17.75	187.03	-348.83	384.33	0.10	0.925	
Random effects							
Fish ID		310151					
Mean velocity between assays							
Intercept	3.94	0.25	3.45	4.42	15.93	<0.001	
Assay(Nobject)	-1.98	0.13	-2.24	-1.72	-14.89	<0.001	
Assay(Mirror)	-1.34	0.13	-1.61	-1.08	-10.10	<0.001	
Sex(Male)	0.96	0.21	0.54	1.38	4.50	<0.001	
0.3ppb	0.02	0.31	-0.59	0.62	0.05	0.958	
3ppb	0.39	0.30	-0.19	0.97	1.33	0.188	
30ppb	0.15	0.30	-0.43	0.73	0.51	0.614	
Random effects							
Fish ID		0.78					

Table A8.2: Model coefficients of F_1 behavioural measures.

Parameter	Estimate	Variance	Lwr95%CI	Upr95%	Z/t	P
Anxiety measures						
Time in bottom zone NA $F_{1,n=190}$						
Intercept	274.1	65.59	145.54	402.65	4.18	0.003
0.3ppb	134.84	92.05	-45.57	315.25	1.47	0.181
3ppb	65.14	91.92	-115.01	245.30	0.71	0.499
30ppb	69.96	91.98	-110.33	250.24	0.76	0.469
Sex(male)	-86.78	21.19	-128.30	-45.25	-4.10	<0.001
Random effects						
Family		11425				
Time in bottom zone NO $F_{1,n=182}$						
Intercept	319.97	61.36	199.71	440.23	5.22	<0.001
0.3ppb	180.19	85.21	13.18	347.20	2.12	0.068
3ppb	163.90	85.11	-2.92	330.71	1.93	0.092
30ppb	129.86	85.04	-36.82	296.54	1.53	0.167
Sex(male)	-78.44	30.04	-137.31	-19.57	-2.61	0.010
Random effects						
Family		8297				
Boldness measures						
Latency to enter top zone $F_{1,n=190}$						
Intercept	40.80	1.00	40.45	41.26	769.60	<0.001
0.3ppb	6.16	1.76	2.01	18.73	3.20	0.001
3ppb	4.07	1.76	1.34	12.30	2.50	0.013
30ppb	3.71	1.76	1.22	11.25	2.30	0.021
Sex(male)	-2.30	1.00	-2.32	-2.27	-172.80	<0.001
Random effects						
Family		1.85				
Frequency entering top zone $F_{1,n=190}$						

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Table A8.2 – *Continued from previous page*

Intercept	15.85	1.19	11.25	22.42	15.62	<0.001
0.3ppb	-1.35	1.27	-2.14	1.17	-1.27	0.203
3ppb	-1.08	1.26	-1.70	1.46	-0.31	0.755
30ppb	-1.25	1.26	-1.97	1.26	-0.96	0.337
Sex(male)	2.04	1.16	1.51	2.75	4.66	<0.001
Random effects						
Family		0.02				
Fish ID		0.98				
<hr/>						
Time in top zone F ₁ ,n=190						
Intercept	231.82	66.20	102.07	361.56	3.50	0.008
0.3ppb	-126.89	92.92	-309.02	55.24	-1.37	0.209
3ppb	-64.65	92.80	-246.53	117.23	-0.70	0.506
30ppb	-55.04	92.86	-237.05	126.97	-0.59	0.570
Sex(male)	84.21	20.98	43.09	125.32	4.01	<0.001
Random effects						
Family		11693				
<hr/>						
Approach NO F ₁ ,n=182						
Intercept	0.13	0.62	-0.05	-0.28	-3.87	<0.001
0.3ppb	0.63	0.65	-0.34	0.85	0.87	0.386
3ppb	0.59	0.65	-0.30	0.83	0.58	0.565
30ppb	-0.48	0.65	-0.22	0.76	-0.11	0.913
Sex(male)	0.87	0.59	0.76	0.93	5.03	<0.001
Random effects						
Family		0.1831				
<hr/>						
Time in middle zone F ₁ ,n=190						
Intercept	322.58	55.40	214.01	431.15	5.82	<0.001
0.3ppb	104.52	77.75	-47.87	256.90	1.34	0.216
3ppb	54.21	77.64	-97.96	206.38	0.70	0.505
30ppb	43.40	77.69	-108.87	195.68	0.56	0.592
Sex(male)	-69.97	17.75	-104.75	-35.19	-3.94	<0.001

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Table A8.2 – *Continued from previous page*

Random effects							
Family			8166				
Measures of exploration							
Exploration $F_{1,n=190}$							
Intercept	59.55	4.50	50.73	68.38	13.23	<0.001	
0.3ppb	5.80	5.93	-5.82	17.42	0.98	0.329	
3ppb	-1.27	5.86	-12.75	10.21	-0.22	0.828	
30ppb	0.87	5.89	-10.66	12.41	0.15	0.882	
Sex(male)	-12.43	4.26	-20.77	-4.09	-2.92	0.004	
Random effects							
Family			0.0				
Measures of activity							
Total Distance Moved in NA $F_{1,n=190}$							
Intercept	3005.79	213.30	2587.73	3423.85	14.09	<0.001	
0.3ppb	-268.97	284.21	-826.02	288.08	-0.95	0.373	
3ppb	-143.93	281.27	-695.22	407.35	-0.51	0.624	
30ppb	-481.99	282.56	-1035.79	71.82	-1.71	0.129	
Sex(male)	1255.64	185.46	892.14	1619.14	6.77	<0.001	
Random effects							
Family			21379				
Mean velocity in NA $F_{1,n=190}$							
Intercept	5.46	0.31	4.85	6.06	17.64	<0.001	
0.3ppb	-0.04	0.41	-0.84	0.76	-0.10	0.918	
3ppb	-0.15	0.40	-0.94	0.63	-0.38	0.704	
30ppb	-0.83	0.40	-1.62	-0.03	-2.05	0.042	
Sex(male)	1.62	0.29	1.05	2.19	5.54	<0.001	
Random effects							
Family			0.0				
Time spent not moving in NA $F_{1,n=190}$							

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Table A8.2 – *Continued from previous page*

Intercept	4.36	0.17	4.04	4.69	25.96	<0.001
0.3ppb	0.356	0.23	−0.09	0.80	1.56	0.158
3ppb	0.29	0.23	−0.15	0.74	1.29	0.236
30ppb	0.45	0.23	0.01	0.90	1.99	0.083
Sex(male)	−0.55	0.12	−0.79	−0.31	−4.48	< 0.001
Random effects						
Family		0.04				
<hr/>						
Total Distance Moved in NO F ₁ ,n=182						
Intercept	1204.45	195.45	821.37	1587.53	6.16	<0.001
0.3ppb	367.21	259.14	−140.71	875.12	1.42	0.194
3ppb	177.76	258.55	−328.99	684.52	0.69	0.511
30ppb	195.99	257.38	−308.46	700.45	0.76	0.469
Sex(male)	1257.18	173.18	917.76	1596.60	7.26	< 0.001
Random effects						
Family		13634				
<hr/>						
Mean velocity in NO F ₁ ,n=182						
Intercept	2.21	0.36	1.50	2.91	6.12	<0.001
0.3ppb	0.70	0.48	−0.24	1.64	1.46	0.183
3ppb	0.73	0.48	−0.21	1.67	1.52	0.167
30ppb	0.48	0.48	−0.45	1.42	1.01	0.341
Sex(male)	1.86	0.31	1.25	2.47	5.96	< 0.001
Random effects						
Family		0.064				
<hr/>						
Total Distance Moved in MIS F ₁ ,n=190						
Intercept	2205.50	159.50	1892.89	2518.20	13.83	<0.001
0.3ppb	−448.20	210.10	−859.95	−36.53	−2.13	0.034
3ppb	−368.70	207.50	−775.41	37.91	−1.78	0.077
30ppb	−500.10	208.60	−908.81	−91.30	−2.40	0.018
Sex(male)	689.30	150.80	393.78	984.89	4.571	< 0.001
Random effects						

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Table A8.2 – *Continued from previous page*

Family	~0						
Mean velocity in MIS F ₁ ,n=190							
Intercept	4.20	0.29	3.63	4.76	14.51	<0.001	
0.3ppb	-0.14	0.39	-0.90	0.62	-0.36	0.728	
3ppb	-0.20	0.39	-0.95	0.56	-0.51	0.624	
30ppb	-0.64	0.39	-1.39	0.12	-1.64	0.142	
Sex(male)	0.79	0.23	0.33	1.24	3.39	<0.001	
Random effects							
Family	0.07						
Measures of aggression							
Time interacting with mirror F ₁ ,n=190							
Intercept	256.68	25.08	207.53	305.83	10.24	<0.001	
0.3ppb	-110.63	33.02	-175.35	-45.91	-3.35	<0.001	
3ppb	-56.32	32.62	-120.25	7.61	-1.73	0.086	
30ppb	-95.98	32.79	-160.24	-31.72	-2.93	0.004	
Sex(male)	74.42	23.71	27.96	120.89	3.14	0.002	
Random effects							
Family	0.00						
Fq interacting with mirror F ₁ ,n=190							
Intercept	31.62	1.29	19.11	51.94	13.51	<0.001	
0.3ppb	-1.05	1.42	-2.12	1.90	-0.15	0.882	
3ppb	-1.90	1.43	-3.86	1.06	-1.78	0.075	
30ppb	1.05	1.40	-1.92	2.12	0.15	0.884	
Sex(male)	-1.76	1.40	-3.42	1.11	-1.66	0.097	
0.3ppb:Sex(Male)	-1.29	1.63	-2.94	1.79	-0.59	0.555	
3ppb:Sex(Male)	3.75	1.63	1.45	9.68	2.72	0.007	
30ppb:Sex(Male)	1.71	1.64	-1.54	4.48	1.09	0.278	
Random effects							
Family	0.052						
Fish ID	1.29						

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Table A8.2 – *Continued from previous page*

Comparison between assays						
Total Distance Moved between assays						
Intercept	2997.98	153.94	2696.26	3299.71	19.48	<0.001
Assay(Nobject)	-1408.50	89.73	-1584.37	-1232.64	-15.70	<0.001
Assay(Mirror)	-1145.07	87.19	-1315.96	-974.19	-13.13	<0.001
Sex(Male)	1072.51	135.49	806.94	1338.07	7.92	<0.001
0.3ppb	-123.83	185.84	-488.08	240.42	-0.67	0.506
3ppb	-122.20	190.01	-494.61	250.22	-0.64	0.521
30ppb	-278.11	190.75	-651.97	95.75	-1.46	0.147
Random effects						
Fish ID: Family ID		626963				
Family ID		0				
Mean velocity between assays						
Intercept	5.32	0.23	4.86	5.78	22.47	<0.001
Assay(Nobject)	-2.43	0.17	-2.77	-2.10	-14.31	<0.001
Assay(Mirror)	-1.60	0.17	-1.92	-1.27	-9.39	<0.001
Sex(Male)	1.42	0.20	1.02	1.81	7.06	<0.001
0.3ppb	0.14	0.28	-0.40	0.68	0.51	0.608
3ppb	0.09	0.28	-0.45	0.64	0.34	0.738
30ppb	-0.36	0.28	-0.91	0.19	-1.28	0.202
Random effects						
Fish ID: Family ID		0.99				
Family ID		~0				

Appendix IX: Behavioural family variation comparisons across treatments

Across a variety of behavioural measures (and across behavioural assays), family variation appeared to significantly influence the F_1 model outputs indicating a genetic component contributed to behavioural phenotypes (Table A9.1). However, it is unclear whether family variation occurred across all treatment groups, or if the atrazine treatment influenced the amount of variation present.

Statistical analysis

To determine if family variance of a particular behavioural measure influenced model outputs, an analysis of variance (ANOVA) was run between a model with the random effect against the same model without the random effect, and a *p-value* was calculated. The amount of family variation is taken from the model outputs with the random factor. In addition, the repeatability at the family level was calculated to provide another measure of genetic influence. Repeatability of each behavioural measure (identified from Table A9.1) was analysed using the ‘*rptR*’ package (Stoffel et al. 2017), with permutations and bootstrap set to 1000. The time spent in the bottom zone of the novel object test, time spent in the bottom zone, top zone and the latency to enter the top zone of the novel arena was analysed under the ‘gaussian datatype’.

To determine the level of within group variation, a test of variance was run. The variance test compares the proportion of two groups and in return gives a variance estimate ratio. The variance estimate ratio (F) is calculated by dividing the variance of one group by another. A proportion where $F < 1$ indicates the numerator group has less variation and inversely the denominator has more. Whereas a $F > 1$ indicates the

opposite; the numerator group has more variation and the denominator group less. As the variance tests are limited to tests along the F distribution, only normally distributed behavioural measures were tested. I compared the proportion of variation across F_1 controls to each of the three treatment groups, and in turn all treatment groups to each other

Results and discussion

There were several behavioural measures that were significantly explained by family level variation. From the novel arena these include; latency to enter the top zone, frequency entering the top zone, time spent in the bottom zone, time spent in the middle zone and the $\log+1(\text{time spent in the top zone})$. From the novel object assay, only the time spent in the bottom half was influenced. And from the mirror test, the frequency entering the mirror zone (summarised in Table A9.1). There was also high variation in the total distance moved in both the novel arena and the novel object assay (but not in the mirror test), but this did not significantly influence model outcomes.

The significant repeatability estimates (Table A9.2) of the behavioural measures identified in Table A9.1, further indicate a genetic component for these behaviours. In particular, the repeatability estimates of the two behavioural measures; time spent in the bottom zone of the novel arena and novel object test are relatively higher compared to the other two measures (latency to enter the top zone and time spent in the top zone of the novel arena test), suggesting a stronger genetic component for these behaviours.

Of the behavioural measures that were normally distributed, there were several instances found in the total distance moved and mean velocity (in all three assays), the standard deviation of exploration and time spent in the middle zone of the novel arena, where the variance ratios were significantly different from one another (see Table A9.3, Table A9.4 and Table A9.5 for raw data).

Table A9.6, Table A9.7 and Table A9.8, shows ranks of which treatment contributed the least, to the most variation for all normally distributed behavioural measures. These tables also summarises the raw data from Table A9.3, Table A9.4 and Table A9.5 to showcase the minimum and maximum ranges of ratios by controls vs treatments (CvT) or treatments vs treatments (TvT).

Overall, the minimum and maximum ranges of the variance ratios show the majority of CvT variation was lower than TvT variation, suggesting at a broad scale, controls exhibited slightly lower variation for some behavioural measures, however in many cases maximum CvT variation was >1 , indicating that lower family variation in the controls is not a consistent pattern. Controls exhibited the lowest variation in the total distance moved (in novel object assay and mirror test) and in the mean velocity (in all three assays), which

is also reinforced by their range of F ratios <1 . Controls exhibited the most variation in the time spent in the bottom half of the novel object assay and time spent interacting with the mirror in the mirror test.

As the time spent in the bottom half of the novel object assay is the only measure that was influenced by family variation, it is possible that less variation in the treatments could be attributed to the atrazine treatments. However, due to the lack of a pattern amongst all other behavioural measures, the lack of a pattern amongst the other treatment levels for this particular behaviour (i.e. the treatment ranks are not ordered from lowest to highest concentration, or vice versa) and the lack of a treatment effect observed (see Chapter 2), suggest that the variation is likely, more attributable to genetic differences. Lastly, I might not have been able to capture differences in the variation produced by families, if they occurred due to the low number of families (three families per treatment) used in this study. More studies with greater numbers of families would be more likely to be able to discern difference if they occur.

In summary, based on this data and for all the behavioural measures, there doesn't appear to be a discernible pattern that would suggest an atrazine treatment effect could influence family level variation, and that difference in at the family level are more likely a result of genetic differences.

Table A9.1: Summary of behavioural models where family level variation influenced model outputs across the novel arena, novel object and mirror test assays. Given are the all models, family variation (variance) and the standard deviation (SD) produced by the inclusion of the random factor (family id), whether or not the model was influenced by family level variation (Model infl.), and the χ^2 test statistic, the degrees of freedom (d.f.) and its p-value.

Novel arena							
Models	Variance	SD	Model infl.	χ^2	d.f.	<i>P-value</i>	
Total distance moved ~ Treatment + Sex	21379	146.2	No	0.0	1	1	
Mean velocity ~ Treatment + Sex	0.0	0.0	No	0.0	1	1	
Exploration ~ Treatment + Sex	0.0	0.0	No	0.0	1	1	
Time spent in bottom zone ~ Treatment + Sex	11425	106.9	Yes	33.6	1	<0.001	
Latency to enter the top zone ~ Treatment + Sex	1.85	1.36	Yes	157.4	1	<0.001	
Frequency entering the top zone ~ Treatment \times Sex	0.98	0.99	No	0.27	1	0.60	
Time spent in the top zone ~ Treatment + Sex	11693	108.1	Yes	35.53	1	<0.001	
Time in the middle zone ~ Treatment + Sex	0.0	0.0	No	0	1	1	
Time spent not moving ~ Treatment + Sex	0.036	0.19	No	0.19	1	0.66	

Novel object							
Models	Variance	SD	Model infl.	χ^2	d.f.	<i>P-value</i>	
Total distance moved ~ Treatment + Sex	13634	116.8	No	0.0	1	1	
Mean velocity ~ Treatment + Sex	0.066	0.25	No	0.0	1	1	
Approached the novel object ~ Treatment + Sex	0.18	0.43	No	0.9	1	0.34	
Time spent in the bottom half ~ Treatment + Sex	8301	91.11	Yes	7.2	1	0.007	

Mirror test							
Models	Variance	SD	Model infl.	χ^2	d.f.	<i>P-value</i>	
Total distance moved ~ Treatment + Sex	~0	~0	No	0	1	1	
Mean velocity ~ Treatment + Sex	0.07	0.27	No	0	1	1	
Time spent interacting with the Mirror ~ Treatment + Sex	~0	~0	No	0	1	1	
Frequency entering mirror zone ~ Treatment \times Sex	1.29	1.14	No	2.46	1	0.12	

Table A9.2: Family level repeatability estimates (R) for behavioural traits identified as having a genetic influence in Table A9.1; i.e. time spent in the bottom zone, latency to enter the top zone and time spent in the top zone of the novel arena, as well as the time spent in the bottom zone of the novel object test. Lower and upper confidence intervals (CI), standard error (SE) and P values are reported

	Behaviour	R	Lower 95% CI	Upper 95% CI	SE	P-value
	Time spent in the bottom zone of the novel arena	0.33	0.11	0.52	0.11	7.68E-12
	Latency to enter the top zone of the novel arena	0.12	0.00	0.26	0.07	1.74E-03
	Time spent in the top zone of the novel arena	0.34	0.12	0.54	0.11	5.59E-12
	Time spent in the bottom zone of the novel object test	0.22	0.05	0.40	0.09	3.58E-07

Table A9.3: Relative family variation during the novel arena test.

Total distance moved												
	0.3ppb				3ppb				30ppb			
Treatment	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value
control	0.67	0.37	1.20	0.17	0.96	0.54	1.71	0.89	1.21	0.68	2.16	0.52
0.3ppb					1.44	0.80	2.58	0.22	1.81	1.0	3.26	0.05
3ppb									1.26	0.70	2.25	0.43
30ppb												

Mean velocity												
	0.3ppb				3ppb				30ppb			
Treatment	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value
control	0.59	0.33	1.06	0.08	0.88	0.50	1.57	0.67	0.9	0.51	1.61	0.73
0.3ppb					1.49	0.83	2.67	0.18	1.52	0.85	2.75	0.16
3ppb									1.03	0.57	1.83	0.93
30ppb												

Exploration												
	0.3ppb				3ppb				30ppb			
Treatment	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value
control	0.83	0.46	1.48	0.53	1.80	1.01	3.20	0.05	2.08	1.16	3.70	0.001
0.3ppb					2.17	1.21	3.91	0.01	2.50	1.39	4.51	0.00
3ppb									1.15	0.64	2.06	0.63
30ppb												

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Table A9.3 – *Continued from previous page*

Time spent in the bottom zone												
	0.3ppb				3ppb				30ppb			
Treatment	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value
control	1.16	0.65	2.07	0.62	0.96	0.54	1.71	0.90	1.04	0.58	1.85	0.90
0.3ppb					0.83	0.46	1.50	0.53	0.90	0.50	1.62	0.71
3ppb									1.08	0.60	1.93	0.80
30ppb												

Time spent in the middle zone												
	0.3ppb				3ppb				30ppb			
Treatment	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value
control	1.37	0.76	2.44	0.29	1.42	0.80	2.52	0.23	2.15	1.20	3.83	0.01
0.3ppb					1.04	0.58	1.86	0.90	1.57	0.87	2.83	0.13
3ppb									1.51	0.84	2.71	0.16
30ppb												

log+1(Time spent in the top zone)												
	0.3ppb				3ppb				30ppb			
Treatment	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value	F	L. CI	U. CI	P-value
control	0.80	0.44	1.42	0.44	1.05	0.59	1.87	0.86	1.08	0.60	1.92	0.79
0.3ppb					1.33	0.74	2.38	0.34	1.36	0.75	2.45	0.30
3ppb									1.02	0.57	1.83	0.94
30ppb												

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Table A9.6: Ranked family variation produced across treatments for measures during the novel arena test. Included is the relative minimum and maximum variation when controls are compared to the three treatment levels (0.3ppb, 3ppb and 30ppb) and when the treatment levels are compared with each other.

Total distance moved	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.67	1.21	30ppb	Control	3ppb	0.3ppb
Treatment vs treatment variation	1.26	1.81				
Mean velocity	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.59	0.90	Control	3ppb	30ppb	0.3ppb
Treatment vs treatment variation	1.03	1.52				
Standard deviation of exploration	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.83	2.08	30ppb	3ppb	Control	0.3ppb
Treatment vs treatment variation	1.15	2.50				
Time spent in the bottom zone	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.96	1.16	0.3ppb	30ppb	Control	3ppb
Treatment vs treatment variation	0.83	1.08				
log+1(Time spent in the top zone)	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.80	1.08	30ppb	3ppb	Control	0.3ppb
Treatment vs treatment variation	1.02	1.36				
Time spent in the middle zone	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	1.37	2.15	30ppb	3ppb	Control	0.3ppb
Treatment vs treatment variation	1.04	1.57				
log+1(Time spent not moving)	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.83	1.06	30ppb	Control	3ppb	0.3ppb
Treatment vs treatment variation	1.10	1.27				

Table A9.7: Ranked family variation produced across treatments for measures during the novel object test. Included is the relative minimum and maximum variation when controls are compared to the three treatment levels (0.3ppb, 3ppb and 30ppb) and when the treatment levels are compared with each other.

Total distance moved	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.41	0.70	Control	30ppb	3ppb	0.3ppb
Treatment vs treatment variation	1.16	1.71				

Mean velocity	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.39	0.66	Control	30ppb	3ppb	0.3ppb
Treatment vs treatment variation	1.01	1.68				

Time spent in the bottom half	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	1.34	1.60	0.3ppb	30ppb	3ppb	Control
Treatment vs treatment variation	0.84	1.22				

Table A9.8: Ranked family variation produced across treatments for measures during the mirror test. Included is the relative minimum and maximum variation when controls are compared to the three treatment levels (0.3ppb, 3ppb and 30ppb) and when the treatment levels are compared with each other.

Total distance moved	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.58	0.82	Control	30ppb	3ppb	0.3ppb
Treatment vs treatment variation	1.14	1.41				

Mean velocity	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	0.33	0.80	Control	3ppb	30ppb	0.3ppb
Treatment vs treatment variation	0.79	2.46				

Time spent interacting with the mirror	Range		Ranked treatment variation			
	Min	Max	Least	2 nd least	2 nd most	Most
Control vs treatment variation	1.08	1.75	30ppb	0.3ppb	3ppb	Control
Treatment vs treatment variation	0.79	1.62				

Appendix X: Phenotype selection for dissections

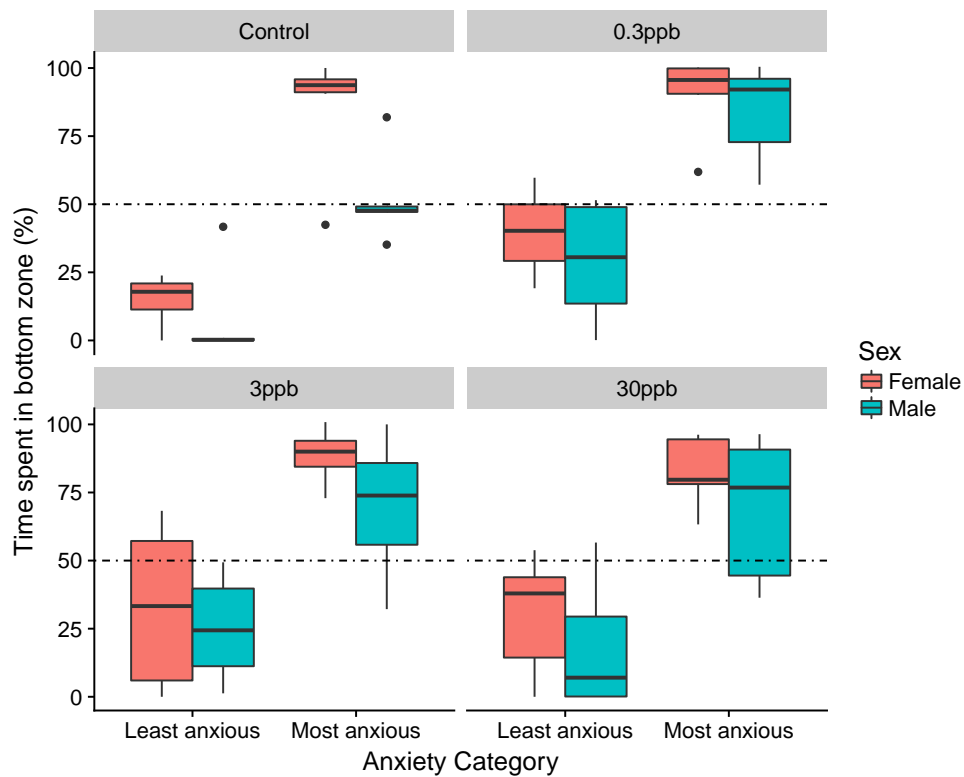


Figure A10.1: Anxiety phenotype (time spent in the bottom zone) of the least and most anxious fish across treatments and sex, for further selection for qPCR analysis. Sample sizes were: Controls n=10-12; 0.3ppb n=12; 3ppb n=10-14; 30ppb n=11 per sex.

Appendix XI: Correlation of anxiety and aggression phenotypes

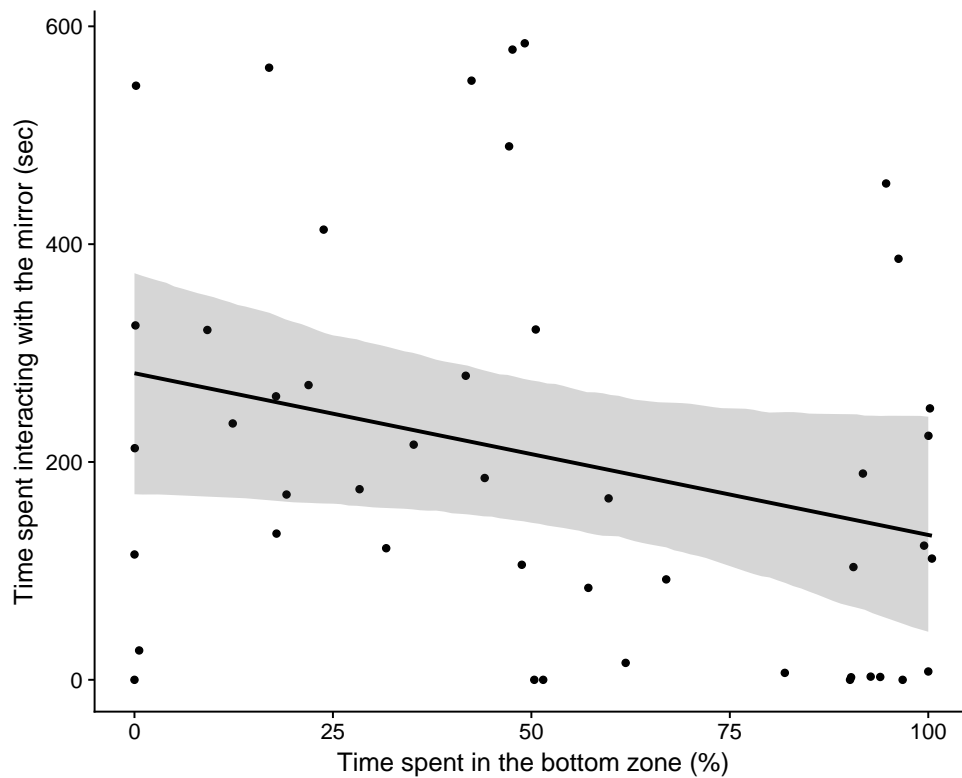


Figure A11.1: Correlation of anxiety and aggression phenotypes for qPCR, (Time spent in the bottom zone Est. -1.29 ± 0.74 ; $-2.75, 0.17$ CI; T -value = -1.74 ; $P = 0.090$).

Appendix XII: Efficiency of qPCR

runs

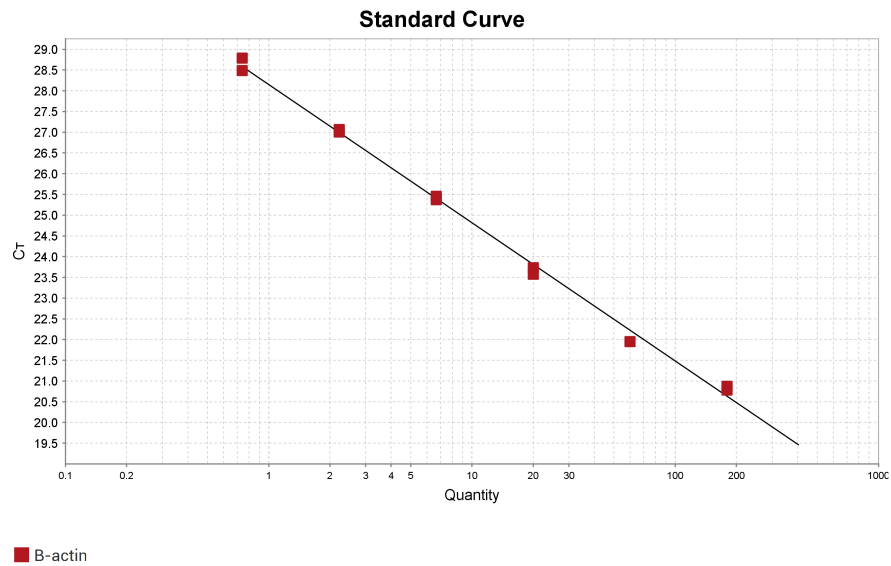


Figure A11.2: Efficiency of *b-actin*.

Appendix XIII: Validation of *b-actin* as a reference gene

Statistical analysis

To validate mRNA of *b-actin* as a suitable reference gene, I ran several linear mixed effects models in R (version 3.5.0; Team 2018), using the package ‘*lme4*’ (Bates et al. 2015). The behavioural measures (anxiety phenotype [least and most anxious]; aggression phenotype [least and most aggressive]; time spent in the bottom zone and time spent interacting with the mirror (both continuous behavioural measurements were further centred), treatment (control and 0.3ppb) group and sex (male and female) were used as predictors. *b-actin* was log transformed to improve normality and was used as the response variable. Family id was used as a random effect to account for variation in genetics. Confidence intervals were calculated using the ‘*confint*’ function, p-values were calculated using the ‘*lmerTest*’ package (Kuznetsova et al. 2017). The full model output is listed in the Table A13.1 and Table A13.2 below.

Table A13.1: Model outputs of *b-actin* mRNA transcripts for the least and most anxious/aggressive phenotypes. For all models, results are presented with estimates, lower and upper 95% confidence intervals (Lwr95%CI and Upr95%CI), the *t* value and the *p* value.

Parameter	Estimate	Variance	Lwr95%CI	Upr95%	<i>T</i>	<i>P</i>
Anxious phenotypes to <i>b-actin</i> mRNA						
Intercept	4.31	0.37	3.59	5.04	11.59	<0.001
Anxious(most)	0.14	0.24	-0.32	0.60	0.60	0.554
Treatment(0.3ppb)	0.02	0.47	-0.90	0.95	0.05	0.963
Sex(male)	-0.16	0.24	-0.62	0.31	-0.67	0.510
Random effects						
Family		0.25				
Aggressive phenotypes to <i>b-actin</i> mRNA						
Intercept	4.52	0.37	3.79	5.25	12.17	<0.001
Aggressive(most)	-0.27	0.23	-0.73	0.18	-1.174	0.248
Treatment(0.3ppb)	0.02	0.47	-0.90	0.95	0.05	0.963
Sex(male)	-0.16	0.23	-0.62	0.30	-0.67	0.506

Table A13.2: Model outputs of *b-actin* mRNA transcripts for anxiety and aggression levels. For all models, results are presented with estimates, lower and upper 95% confidence intervals (Lwr95%CI and Upr95%CI), the *t* value and the *p* value.

Parameter	Estimate	Variance	Lwr95%CI	Upr95%	<i>T</i>	<i>P</i>
Time spent in the bottom zone						
Intercept	4.40	0.35	3.72	5.08	12.68	<0.001
Centred bottom time	0.00	0.00	-0.00	0.01	1.102	0.277
Treatment(0.3ppb)	-0.06	0.47	-0.98	0.87	-0.12	0.911
Sex(male)	-0.10	0.24	-0.58	0.37	-0.43	0.668
Time spent interacting with mirror						
Intercept	4.42	0.36	3.72	5.12	12.340	<0.001
Centred mirror time	-0.00	0.00	-0.00	0.00	-0.74	0.463
Treatment(0.3ppb)	-0.061	0.49	-1.02	0.90	-0.13	0.906
Sex(male)	-0.13	0.24	-0.60	0.34	-0.54	0.595

Appendix XIV: Model selection using AIC_C

Table A14.1: Ranked models by small-sample-size-corrected Akaike Information Criterion (AIC_C) for the inclusion or exclusion of interaction terms in models describing changes in mRNA of the genes: *slc6a4b*, *htr1B* or time in the bottom zone (Tbz) by treatment:sex interactions, treatment:behaviour interactions or mRNA:treatment interactions. The top five ranked models per each AIC_C analysis are shown. Model number indicates the order the model was generated. A + indicates the presence of the main predictor or the interaction term: sex, treatment (Trt), sex:treatment (Sex:Trt), behavioural phenotype (Beh), behaviour phenotype:treatment (Beh:Trt) and *htr1Aa* mRNA:treatment (H1A:Trt). Presence of estimate indicates the inclusion of the continuous main predictor *htr1Aa* mRNA (H1A). Also included are the intercepts (int), degree of freedom (df), log-Likelihood (logLik), AIC_C, change in AIC_C value (Δ) and the AIC_C probability for each model (weight). Models in bold indicate models with $\Delta\text{AIC}_C \leq 6$.

Model number	Int	Sex	Trt	Sex:Trt	df	logLik	AIC _C	Δ	weight
<i>slc6a4b</i> mRNA~Treatment*Sex									
1	1.062				3	-20.175	46.9	0	0.705
3	1.143			+	4	-20.445	49.9	2.94	0.162
2	1.117	+			4	-20.908	50.8	3.87	0.102
4	1.192	+	+		5	-21.244	54	7.07	0.021
8	1.272	+	+	+	6	-20.614	55.4	8.46	0.01
Model number	Int	Beh	Trt	Beh:Trt	df	logLik	AIC _C	Δ	weight
<i>htr1B</i> mRNA~Anxiety*Trt									
1	0.9693				3	22.221	-37.90	0.0	0.848
3	1.008		+		4	21.199	-33.4	4.45	0.092
2	0.993	+			4	20.674	-32.4	5.5	0.054
4	1.032	+	+		5	19.651	-27.8	10.07	0.006
8	1.063	+	+	+	6	19.274	-24.4	13.48	0.001
<i>slc6a4b</i> mRNA~Aggression*Trt									
1	1.062				3	-20.175	46.9	0	0.624
2	1.146	+			4	-20.251	49.5	2.56	0.174
3	1.143		+		4	-20.445	49.9	2.94	0.143
4	1.227	+	+		5	-20.501	52.5	5.58	0.038
8	1.315	+	+	+	6	-19.799	53.8	6.83	0.021
Model number	Int	H1A	Trt	H1A:Trt	df	logLik	AIC _C	Δ	weight
Tbz~ <i>htr1Aa</i> mRNA*Trt									
8	40.14	17.28	+	+	6	-212.15	438.4	0	0.972
4	40.79	8.41	+		5	-217.12	445.7	7.3	0.025
3	41.41		+		4	-220.82	450.6	12.17	0.002
2	51.58	7.864			4	-222.18	453.3	14.89	0.001
1	51.63				3	-225.73	458	19.58	0

Appendix XV: Model outputs for relative mRNA level data

Table A15.1: Model outputs of F₁ gene expression data.

Parameter	Estimate	Variance	Lwr95%CI	Upr95%	Z/t	P
<i>slc6a4a</i> expression						
Intercept	0.59	0.11	0.37	0.82	5.81	5.188
Treatment(0.3ppb)	-0.03	0.15	-0.32	0.26	-0.20	0.851
Sex(male)	-0.04	0.10	-0.23	0.15	-0.42	0.681
Random effects						
Family	0.11					
<i>slc6a4b</i> expression						
Intercept	1.08	0.07	0.94	1.22	15.47	<0.001
Treatment(0.3ppb)	-0.15	0.11	-0.37	0.06	1.42	0.157
Sex(male)	-0.12	0.11	-0.33	0.10	1.05	0.294
Random effects						
Family	0.0					
<i>htr1Aa</i> expression						
Intercept	-0.33	0.26	-0.84	0.18	-1.27	0.230
Treatment(0.3ppb)	-0.64	0.37	-1.36	0.07	-1.76	0.109
Sex(male)	-0.71	0.35	-1.39	-0.02	-2.03	0.049
Trt:Sex	1.05	0.48	0.11	2.00	2.19	0.035

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Table A15.1 – *Continued from previous page*

Random effects						
Family	0.04					
<i>htr1b</i> expression						
Intercept	1.03	0.09	0.86	1.19	12.02	<0.001
Treatment(0.3ppb)	−0.08	0.12	−0.31	0.16	−0.64	0.558
Sex(male)	−0.04	0.04	−0.11	0.03	−1.04	0.307
Random effects						
Family	0.02					
<i>htr2b</i> expression						
Intercept	0.66	0.10	0.46	0.86	6.43	<0.001
Treatment(0.3ppb)	0.04	0.13	−0.23	0.30	0.27	0.800
Sex(male)	0.00	0.08	−0.16	0.16	−0.04	0.970
Random effects						
Family	0.0					

Table A15.2: Model outputs of the least/most anxious and aggressive phenotypes and relative mRNA levels of candidate genes *slc6a4a*; *slc6a4b*; *htr1Aa*; *htr1B* and *htr2B* of F₁ offspring from control and 0.3ppb atrazine treated males normalised to *b-actin* (from whole brain samples). For all models, results are presented with estimates, lower and upper 95% confidence intervals (Lwr95%CI and Upr95%CI), the *t* value and the *p* value.

Parameter	Estimate	Variance	Lwr95%CI	Upr95%	Z/t	P
Anxiety phenotype						
log <i>slc6a4a</i>						
Intercept	-0.39	0.24	-0.87	0.08	-1.62	0.15
Anxious(most)	-0.07	0.24	-0.53	0.40	-0.29	0.77
Treatment(0.3ppb)	-0.08	0.29	-0.66	0.50	-0.27	0.80
Random effects						
Family		0.05				
<i>slc6a4b</i>						
Intercept	1.14	0.09	0.96	1.32	12.12	<0.001
Anxious(most)	0.01	0.11	-0.20	0.22	0.06	0.955
Treatment(0.3ppb)	-0.15	0.11	-0.36	0.06	-1.44	0.157
Random effects						
Family		0				
log <i>htr1Aa</i>						
Intercept	-0.76	0.25	-1.25	-0.27	-3.048	0.018
Anxious(most)	0.23	0.25	-0.26	0.71	0.91	0.37
Treatment(0.3ppb)	-0.15	0.30	-0.74	0.44	-0.50	0.645
Random effects						
Family		0.044				
<i>htr1B</i>						
(Intercept)	0.97	0.06	0.85	1.09	15.84	<0.001
Anxious(most)	-0.05	0.04	0.04	0.03	1.26	0.207
Treatment(0.3ppb)	-0.08	0.12	-0.31	0.16	0.64	0.521

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Table A15.2 – *Continued from previous page*

Random effects	0.02 Family					
<i>htr2B</i> Intercept	0.77	0.11	0.55	0.98	6.89	<0.001
Anxious(most)	−0.22	0.11	−0.44	0.01	−1.89	0.067
Treatment(0.3ppb)	−0.12	0.16	−0.42	0.19	−0.76	0.474
Anxious(most):Trt	0.31	0.16	0.00	0.62	1.94	0.059
Aggressive phenotype						
log <i>slc6a4a</i>						
Intercept	−0.25	0.24	−0.72	0.23	−1.02	0.339
Aggressive(most)	−0.36	0.23	−0.82	0.09	−1.58	0.122
Treatment(0.3ppb)	−0.08	0.29	−0.66	0.50	−0.27	0.800
Random effects						
Family		0.05				
log <i>slc6a4b</i>						
Intercept	1.10	0.08	0.94	1.25	13.70	<0.001
Aggressive(most)	−0.15	0.11	−0.37	0.06	1.42	0.155
Treatment(0.3ppb)	−0.17	0.11	−0.38	0.04	1.55	0.121
Random effects						
Family		0.02				
log <i>htr1Aa</i>						
Intercept	−0.53	0.25	0.53	0.94	−2.13	0.07
Most Aggressive	−0.23	0.25	−0.31	0.00	−0.94	0.36
AZT0.3	−0.15	0.30	−0.23	0.30	−0.50	0.64
Random effects						
Family		0.04				
log <i>htr1B</i>						
Intercept	1.01	0.09	0.84	1.18	11.84	0.00
Aggressive(most)	0.00	0.04	−0.07	0.07	−0.02	0.99
Treatment(0.3ppb)	−0.08	0.12	−0.31	0.15	−0.66	0.54
Random effects						
Family		0.02				

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Table A15.2 – *Continued from previous page*

<i>htr2B</i>						
Intercept	0.73	0.10	0.53	0.94	7.13	0.00
Aggressive(most)	-0.15	0.08	-0.31	0.00	-1.97	0.06
Treatment(0.3ppb)	0.04	0.13	-0.23	0.30	0.27	0.80
Random effects						
Family		0.02				

Table A15.3: Model outputs of anxiety (time spent in the bottom zone), aggression (time spent interacting with the mirror) and relative mRNA levels of candidate genes *slc6a4a*; *slc6a4b*; *htr1Aa*; *htr1B* and *htr2B* of F₁ offspring from control and 0.3ppb atrazine treated males normalised to *b-actin* (from whole brain samples). For all models, results are presented with estimates, lower and upper 95% confidence intervals (Lwr95%CI and Upr95%CI), the *t* value and the *p* value.

Parameter	Estimate	Variance	Lwr95%CI	Upr95%	<i>T</i>	<i>P</i>
Anxiety						
Time spent in the bottom zone						
Intercept	40.92	7.18	26.85	54.98	5.70	<0.001
c.log <i>slc6a4a</i>	11.94	8.94	-5.57	29.46	1.34	0.189
Treatment(0.3ppb)	19.64	9.93	0.17	39.11	1.98	0.055
c.log <i>slc6a4a</i> :Treatment(0.3ppb)	-22.27	12.43	-46.64	2.09	-1.79	0.080
Random effects						
Family		0				
Intercept	40.62	7.40	26.12	55.13	5.49	<0.001
c. <i>slc6a4b</i>	9.77	14.41	-18.48	38.02	0.68	0.50
Treatment(0.3ppb)	21.05	10.36	0.74	41.36	2.03	0.05
Random effects						
Family		0				
Intercept	40.14	7.02	26.37	53.90	5.72	<0.001
c.log <i>htr1Aa</i>	17.28	7.58	2.42	32.13	2.28	0.028
Treatment(0.3ppb)	20.56	9.73	1.49	39.62	2.11	0.041
c.log <i>htr1Aa</i> :Treatment(0.3ppb)	-21.08	11.69	-43.98	1.82	-1.80	0.078
Random effects						
Family		0				
Intercept	42.24	7.78	27.00	57.48	5.43	0.009

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Table A15.3 – *Continued from previous page*

<i>c.htr1B</i>	−17.12	31.18	−78.23	43.99	−0.55	0.593
Treatment(0.3ppb)	18.01	10.94	−3.43	39.45	1.65	0.191
Random effects						
Family		15.89				
Intercept	41.32	7.35	26.91	55.72	5.62	<0.001
<i>c.htr2B</i>	−4.73	17.55	−39.13	29.68	−0.27	0.789
Treatment(0.3ppb)	19.72	10.19	−0.25	39.69	1.94	0.060
Random effects						
Family		0				
Aggression						
Time spent in the mirror zone						
Intercept	282.55	35.56	212.86	352.23	7.95	<0.001
<i>c. log slc6a4a</i>	−42.57	30.80	−102.94	17.80	−1.38	0.17
Treatment(0.3ppb)	−148.91	49.25	−245.44	−52.37	−3.02	<0.001
Random effects						
Family		0.00				
Intercept	297.64	37.03	225.07	370.21	8.04	0.001
<i>c. slc6a4b</i>	−196.40	64.83	−323.46	−69.34	−3.03	<0.001
Treatment(0.3ppb)	−176.89	51.91	−278.63	−75.15	−3.41	0.03
Random effects						
Family		810.3				
Intercept	282.03	36.25	210.99	353.07	7.78	<0.001
<i>c.log htr1Aa</i>	−16.87	29.81	−75.30	41.56	−0.57	0.574
Treatment(0.3ppb)	−147.92	50.26	−246.44	−49.40	−2.94	0.005
Random effects						
Family		0				
Intercept	299.61	34.53	231.93	367.29	8.68	<0.001
<i>c. htr1B</i>	−420.75	139.49	−694.14	−147.35	−3.02	0.01
Treatment(0.3ppb)	−181.50	48.56	−276.67	−86.33	−3.74	0.03

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Table A15.3 – *Continued from previous page*

Random effects							
Family		211					
Intercept	280.05	39.97	201.71	358.39	7.01	<0.001	
c. <i>htr2B</i>	-95.49	87.04	-266.09	75.11	-1.10	0.28	
Treatment(0.3ppb)	-143.06	55.60	-252.03	-34.10	-2.57	0.08	
Random effects							
Family		984.1					

Table A15.4: Cross validation model outputs of *slc6a4a* and *htr1Aa* mRNA for interactive anxiety:treatment differences. For all models, results are presented with estimates, lower and upper 95% confidence intervals (Lwr95%CI and Upr95%CI), the *t* value and the *p* value.

Parameter	Estimate	Variance	Lwr95%CI	Upr95%	T	P
Controls						
(Intercept)	41.41	7.61	26.50	56.32	5.44	<0.001
c.log. <i>slc6a4a</i>	11.94	9.49	-6.65	30.54	1.26	0.223
Treatment						
(Intercept)	60.95	6.45	48.30	73.60	9.45	<0.001
c.log. <i>slc6a4a</i>	-10.33	8.13	-26.26	5.60	-1.27	0.217
Controls						
(Intercept)	41.41	7.05	27.59	55.23	5.87	<0.001
c.log. <i>htr1Aa</i>	17.28	7.64	2.31	32.24	2.26	0.0349
Treatment						
(Intercept)	60.95	6.66	47.90	74.00	9.16	<0.001
c.log. <i>htr1Aa</i>	-3.80	8.84	-21.12	13.51	-0.43	0.671

Appendix XVI: Additional measures of relative mRNA level data

Methods

Due to the least and most anxious fish being selected, additional analyses were carried out on the behavioural phenotype defined as a categorical variable. The least and most aggressive fish were selected based on highest and lowest time spent interacting with the mirror from their family group, per sex.

Statistical analysis

To investigate associations between mRNA transcripts of candidate genes and behaviour, LMMs were conducted with candidate genes as response variables with treatment, behavioural phenotype (fish categorised as either the least or most anxious/ aggressive) and an interaction term as predictors. Sex as a main effect was non-significant and thus removed from further analyses between behavioural phenotypes.

Results

Anxiety

No significant relationship was observed for anxiety phenotype and *slc6a4a* mRNA transcript (Figure A16.1 A), nor were there any significant differences observed for *slc6a4b* and *htr1B* mRNA transcripts (after model averaging) between the least anxious and most anxious fish from either treatment (Figure A16.1 B and D respectively). On average, more anxious control fish showed an up-regulation of *htr1Aa* with the pattern reduced in atrazine treated fish. However no significant relationships were observed (Figure A16.1 C). More anxious control fish showed a marginally non-significant down-regulation of *htr2B* mRNA (Est. -0.22

[−0.44, 0.01 CI]). Conversely, fish from atrazine treated fathers exhibited the opposite pattern, but this interactive effect was also marginally non-significant (Est. 0.31 [0.0, 0.62 CI]; Figure A16.1 E).

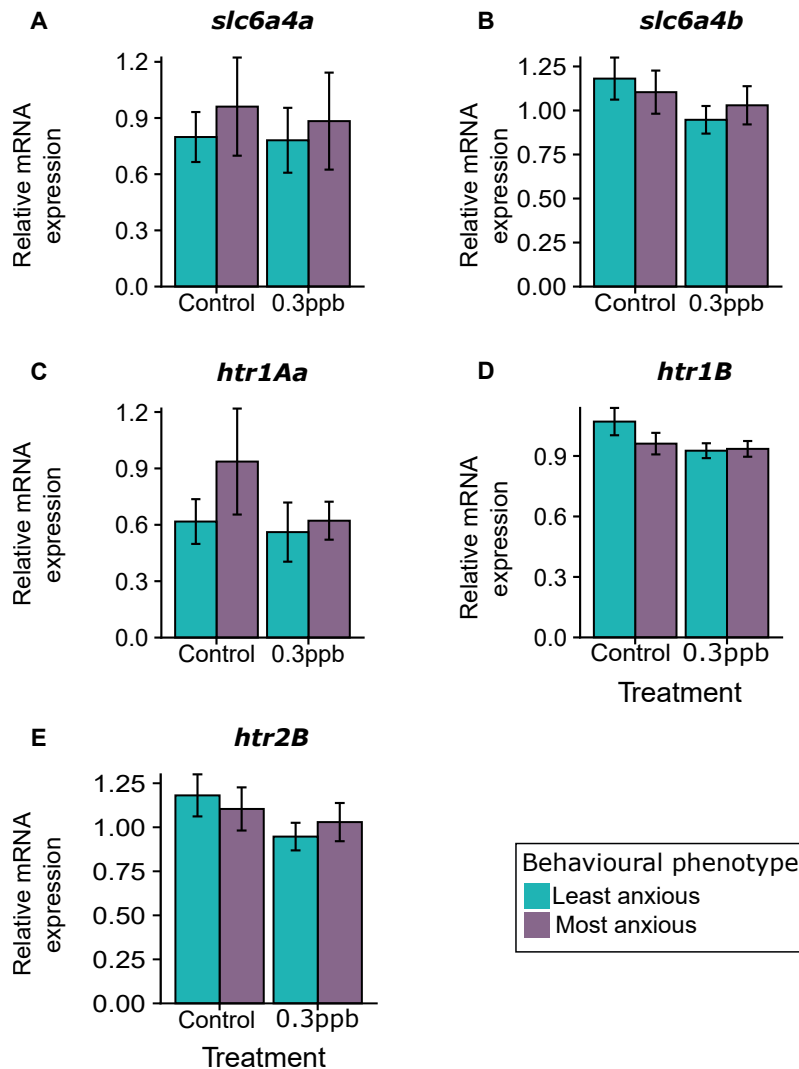


Figure A16.1: Relationship between anxiety phenotype (fish categorised as the least and most anxious) and relative mRNA levels of the candidate genes (A) *slc6a4a*; (B) *slc6a4b*; (C) *htr1Aa*; (D) *htr1B* and (E) *htr2B* of F₁ offspring from control and 0.3ppb atrazine treated males normalised to *b-actin* (from whole brain samples). Sample sizes for control offspring were n=22 (males=10; females=12; progeny per family=6-8; families n=3), and for 0.3ppb offspring were n=24 (males=12; females=12; progeny per family=8; families n=3).

Aggression

There was no statistically significant relationship in the mRNA transcripts of *slc6a4a*, *slc6a4b* (after model averaging), *htr1Aa* and *htr1B* in fish categorised as either the most or least aggressive (Figure A16.2 A; B; C; D respectively). However, more aggressive fish showed a non-significant down regulation in *htr2B* mRNA (Est. -0.15 [$0.0, -0.94$ CI]) regardless of treatment (Figure A16.2 E).

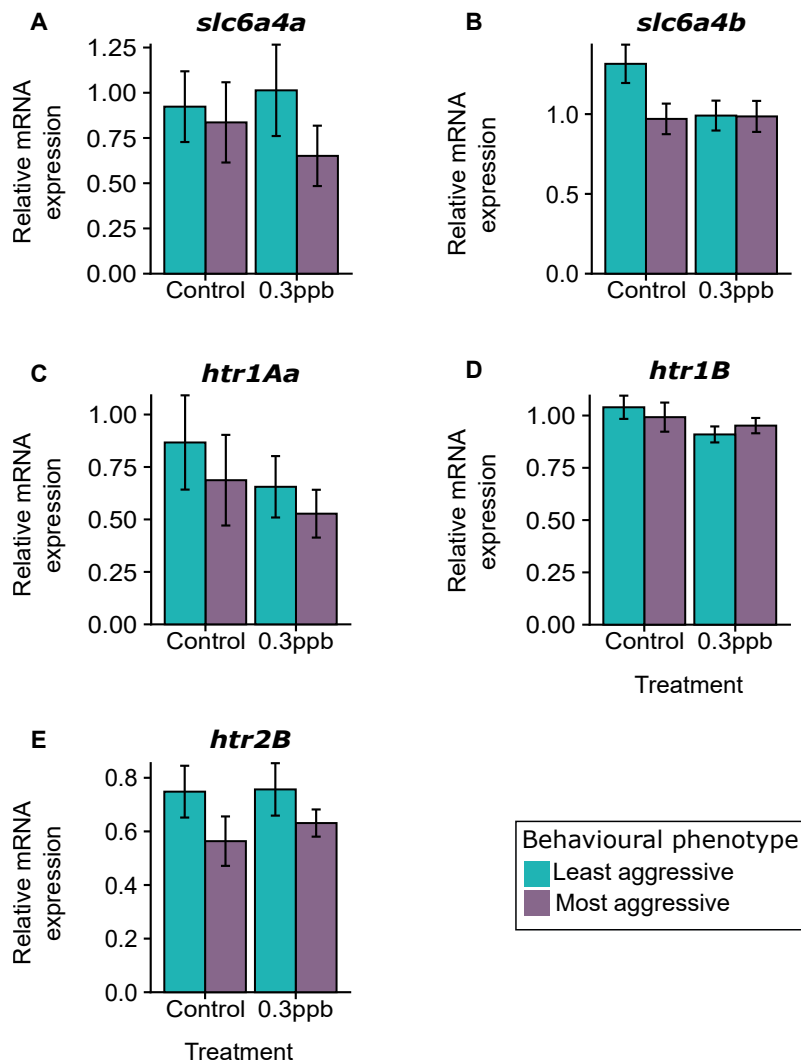


Figure A16.2: Relationship between aggression phenotype (fish categorised as the least and most aggressive) and relative mRNA levels of the candidate genes (A) *slc6a4a*; (B) *slc6a4b*; (C) *htr1Aa*; (D) *htr1B* and (E) *htr2B* of F₁ offspring from control and 0.3ppb atrazine treated males normalized to *b-actin* (from whole brain samples). Sample sizes for control offspring were n=22 (males=10; females=12; progeny per family=6-8; families n=3), and for 0.3ppb offspring were n=24 (males=12; females=12; progeny per family=8; families n=3).