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I. Effects of a dopamine receptor antagonist on fathead minnow, *Pimephales promelas*, reproduction

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ABSTRACT

Neurotransmitters such as dopamine play an important role in regulating fish reproduction. However, the potential for neuroendocrine active chemicals to disrupt fish reproduction has not been well studied, despite emerging evidence of their discharge into aquatic environments. This study is the first to apply the fathead minnow 21 d reproduction assay developed for the US Endocrine Disruptor Screening Program to evaluate the reproductive toxicity of a model neuroendocrine active chemical, the dopamine 2 receptor antagonist, haloperidol. Continuous exposure to up to 20 μg haloperidol/L had no significant effects on fathead minnow fecundity, secondary sex characteristics, gonad histology, or plasma steroid and vitellogenin concentrations. The only significant effect observed was an increase in gonadotropin-releasing hormone (cGnRH) transcripts in the male brain. Results suggest that non-lethal concentrations of haloperidol do not directly impair fish reproduction. Potential effects of haloperidol on reproductive behaviors and gene expression were examined in a companion study.

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1. Introduction

Reproduction in fish and other vertebrates is predominantly regulated by the hypothalamic–pituitary–gonadal (HPG) axis and a variety of environmental and social cues. Endocrine active chemicals (EACs) that disrupt the normal mechanisms of HPG axis function, or interfere with response to appropriate reproductive cues, have the potential to adversely impact reproductive success. If the impacts are sufficiently widespread, reproductive impairments in individuals can have significant population-level consequences (e.g., Kidd et al., 2007; Miller and Ankley, 2004). Thus, there is a need to characterize the diversity of mechanisms through which xenobiotics might disrupt vertebrate reproduction and develop appropriate techniques to screen chemicals for such activity (e.g., high throughput in vitro bioassays, quantitative structure–activity relationships) and detect their influence in the field (e.g., diagnostic biomarkers).

As part of an on-going research program aimed at systematic characterization of various modes of endocrine disruption using small fish (Ankley et al., 2009), we used a generalized model of the teleost HPG axis (Villeneuve et al., 2007b) to identify likely molecular targets through which xenobiotics could interact with the axis and potentially impair reproduction. This included some relatively well-studied targets such as the estrogen receptor, androgen receptor, and various enzymes involved in steroid biosynthesis. Fathead minnow (*Pimephales promelas*) reproduction assays showed that chemicals that interact with these targets reduce the fecundity of exposed individuals and produce a variety of molecular changes consistent with effects on HPG axis function (e.g., Ankley et al., 2002, 2003, 2005, 2007; Lange et al., 2001; Martinović et al., 2008). Other HPG axis components, such as the gonadotropin-releasing hormone (GnRH) receptors and gonadotropin (GTH) receptors, play key roles in regulating reproduction but were considered unlikely targets for direct modulation by xenobiotics since they bind peptide hormones, which do not have structural analogues that are environmental contaminants. In contrast, various neurotransmitter receptors (e.g., dopamine receptors, gamma amino butyric acid [GABA] receptors) bind...
2. Materials and methods

2.1. Chemical and test organisms

Haloperidol was purchased from Sigma Aldrich (St. Louis, MO, USA; H1512; 99% pure). The reproductively mature adult fathead minnows (6 months old) used in these experiments were obtained from an on-site culture facility at the US EPA Mid-Continent Ecology Division (Duluth, MN, USA). Fish were held at 25 °C under a 16:8 light:dark photoperiod and fed to satiation twice daily with frozen brine shrimp (Artemia). All animals were treated humanely with regard for alleviation of suffering, and all laboratory procedures involving animals were reviewed and approved by an Animal Care and Use Committee in accordance with Animal Welfare Act and Interagency Research Animal Committee guidelines.

2.2. Chemical delivery and exposure verification

Solvent-free, saturated aqueous stock solutions of haloperidol were prepared in UV-treated, filtered, Lake Superior water through continuous stirring in a 20 L glass carboy. Stock solution was diluted in Lake Superior water and delivered to 20 L tanks containing 10 L of water, at a continuous flow of 44 ± 1.5 ml/min (mean ± SD) to achieve the appropriate nominal test concentrations. Water samples were collected from the stock solution and exposure tanks over the course of each experiment. Haloperidol concentrations in the water samples were directly analyzed using an Agilent (Wilmington, DE, USA) model 1100 LC-MS equipped with an electrospray interface (API-ES). An aliquot of sample (30 μL) was injected onto a Luna (Phenomenex, Torrance, CA, USA) CN column (2.0 × 50 mm) and eluted isocratically at a flow rate of 0.3 ml/min. The mobile phase consisted 66% acetonitrile, 10 mM ammonium acetate buffer and 0.04% acetic acid. Haloperidol concentrations were determined using mass 376 (SIM positive ion mode) with an external standard method of quantitation. Routine quality assurance analyses (i.e., procedural blanks, spiked matrix, and duplicate samples) were conducted with each day’s sample set. Spike recoveries averaged (± SD) 102 ± 3% and 96 ± 6% for the range finder and reproduction assay, respectively. Average (± SD) agreement among duplicate analyses of the same sample was 97.6 ± 3.1% (n=24). The method detection limit was 0.01 μg haloperidol/L.

2.3. Range-finding experiment

A range-finding experiment was conducted to aid the identification of non-lethal exposure concentrations for use in the reproduction assay. The nominal (target) concentrations for the range-finding experiment were 0, 0.2, 2, 20, and 200 μg haloperidol/L. There was one tank per test concentration and each tank contained three male and three female fathead minnows along with three breeding substrates (10 cm section of PVC pipe cut in half (Ankley et al., 2001)). The fish were exposed for 5 d, during which survival and spawning activity were monitored daily. Haloperidol concentrations in the exposure tanks were measured on days 1, 3, and 5 of the experiment. At the end of the experiment, surviving fish were humanely terminated in a buffered solution of tricaine methanesulfonate (MS-222; Finquel; Argent, Redmond, WA, USA).

2.4. Fathead minnow 21 d reproduction assay

The basic experimental design for the fathead minnow 21 d reproduction assay was described by Ankley et al. (2001) except a paired, rather than group spawning approach was used (e.g., Ankley et al., 2005). Fathead minnows were paired (one male, one female) and placed in tanks at a density of two pairs per tank. Pairs were separated by a water permeable mesh divider and each had its own breeding substrate. The fish were held in the test system, receiving Lake Superior (control) water only for a 14 d acclimation period during which the fecundity and survival of each pair were assessed daily. After 14 d, exposures were initiated with pairs that had spawned successfully during acclimation. Five exposure concentrations (0, 0.02, 0.2, 2, and 20 μg haloperidol/L nominal) were tested. There were four replicate tanks per treatment and two pairs per tank for a total of eight pairs per treatment.

Over the course of the exposure, the total number of eggs spawned and the number of fertile eggs produced by each pair were recorded daily. Haloperidol concentrations in the test tanks were quantified approximately every 3 d over the course of the exposure (7/21 d total). Water quality characteristics (mean ± SD) monitored regularly over the course of the study included temperature 25.3 ± 1.0 °C, dissolved oxygen 6.02 ± 0.57 mg/L, pH 7.39 ± 0.08, hardness 45.6 ± 0.5 mg/L as CaCO3, and alkalinity 47.4 ± 0.4 mg/L as CaCO3. After 21 d of exposure, surviving fish were anesthetized in buffered MS-222. Immediately after anesthetization, blood was collected from the caudal vasculature using heparinized microhematocrit tubes. Plasma was separated by centrifugation and stored at −80 °C until extracted and analyzed. Following blood sampling, body mass (sans blood) was measured, secondary sex characteristics (i.e., nuptial tubercles) were evaluated, and gonads, dorsal fatpads (males only), brains, and pituitary glands were removed. Tubercles were scored based on their number and relative size (Jensen et al., 2001; US EPA, 2002). Gonads were weighed and divided into several pieces. A subsample (approximately 10 mg) was used for an ex vivo steroid production assay. The remaining gonad mass was divided into approximately equal thirds (females) or halves (males) with one portion preserved in Daseo-7 for later histological characterization, one portion preserved in RNA later (Sigma R0901), and one portion (ovaries only) snap frozen in liquid nitrogen. Dorsal fatpads were weighed and discarded. Brains and pituitaries, collected separately using fine forceps, were each transferred to pre-weighted tubes containing RNA later. Samples in RNA later were stored at −20 °C until exposed. Snap-frozen samples were stored at −80 °C. All dissection tools were washed with RNaseA Zap (Ambion, Austin, TX, USA) between samples to prevent cross-contamination or degradation by RNases. Gonads preserved for histological analysis were embedded, stained, sectioned and analyzed as described previously (Villeneuve et al., 2008). Ex vivo steroid
production assays were conducted using methods adapted from (McMaster et al., 1995) as described previously (Martinović et al., 2008). Briefly, subsamples of gonad tissue from each fish were transferred to wells of a 48-well plate microplate (Falcon 35–3078, Beckton Dickinson, Franklin Lakes, NJ, USA) containing 500 μl of Medium 199 (M2520; Sigma) supplemented with 0.1 mM isobutylmethylxanthine (IBMX; Sigma I7018) and 1 μg 25-hydroxycholesterol (Sigma)/ml on ice. Samples were incubated at 25 °C overnight (16.5 h), after which media from each well was transferred to a microcentrifuge tube and stored at −20 °C until extracted and analyzed. The mass of each gonad subsample was weighed after collection of the media. Average (+ SD) samples masses were 8.7 ± 6.8 mg for testes and 28.8 ± 13.6 mg for ovaries. Wells containing supplemented medium but no tissue were incubated, sampled, and analyzed along with experimental samples to serve as assay blanks. Steroids were extracted from medium samples (ex vivo) or plasma samples by liquid–liquid extraction with diethyl ether and quantified by radioimmunoassay (Jensen et al., 2001; US EPA, 2002). With the exception of cases where plasma volumes were limiting, both plasma estradiol and plasma testosterone concentrations were quantified. Plasma concentrations of the estrogen-inducible egg yolk precursor protein, vitellogenin (Vtg), were quantified by enzyme-linked immunosorbent assay using a polyclonal antibody to fathead minnow Vtg and purified fathead minnow Vtg as a standard (US EPA, 2002).

Relative abundance of selected mRNA transcripts in ovary, brain, or pituitary samples from the reproduction assay was quantified using quantitative real-time PCR (QPCR). Gonad or brain samples in RNAlater were transferred to TRI Reagent™ (Sigma) and total RNA was extracted from the tissue according to the manufacturer’s protocol. Total RNA was extracted from individual pituitary samples using RNeasy micro kits (Qiagen, Valencia, CA, USA). RNA concentration and quality was evaluated using a Nanodrop® ND-1000 spectrophotometer (Nanodrop Technologies, Wilmington, DE, USA). The 260/280 nm absorbance ratio for all samples was between 1.7 and 2.2. Total RNA samples were diluted to either 10 μg/μl (brains, gonads) or 1 ng/μl (pituitaries) for use in QPCR assays. Relative abundance of mRNA transcripts coding for fathead minnow follicle-stimulating hormone β subunit (FSHβ), luteinizing hormone β subunit (LHβ), and follicle-stimulating hormone receptor (FSHR) was determined using primers, probes, and protocols described previously (Villeneuve et al., 2007a, 2007b). Partial cDNA sequences for a fathead minnow GnRH (c, chicken-type isoform; gnrb2) and D2R were determined using methods similar to those described elsewhere (Villeneuve et al., 2007a, 2007b). Gene-specific primers and probes corresponding to the partial sequences were developed (Table 1) and applied, using the previously described QPCR protocol (Villeneuve et al., 2007a), to measure the relative abundance of cGnRH and D2R transcripts.

Data from the reproduction assay were tested for normality and homogeneity of variance. When data conformed to parametric assumptions, one-way analysis of variance (ANOVA) was used to test for differences across treatments and a post-hoc Duncan’s test was applied to determine which groups differed (p < 0.05). When data did not conform to parametric assumptions, they were either transformed (e.g., log 10) or analyzed using a non-parametric Kruskall–Wallace test (p < 0.05). Ex vivo steroid production data were analyzed by general linear models ANOVA using both haloperidol concentration and the mass of each gonad subsample as independent variables. Pair was treated as the unit of replication for fecundity and fertility. However, statistical conclusions were the same whether each pair or tank was viewed as the unit of replication. Individuals were considered as the unit of replication for all other endpoints, as no tank effects were evident. All statistical analyses were conducted using SAS 9.0 (SAS Institute, Cary, NC, USA).

3. Results

3.1. Toxicity

In the range-finding experiment, 200 μg haloperidol/L was lethal to six of six adult fathead minnows within 3 d. Exposure to 20 μg haloperidol/L or less was not lethal over the 5 d range-finding experiment. Based on these results, 20 μg/L was identified as the maximum target test concentration for the reproduction assay.

3.2. Fathead minnow 21 d reproduction assay

Average (+ SD) haloperidol concentrations measured in water samples collected from the reproduction assay test tanks were 0.017 ± 0.003, 0.204 ± 0.019, 2.27 ± 0.27, and 22.0 ± 2.10 μg/L for the 0.02, 0.2, 2.0, and 20 μg/L nominal treatment groups, respectively. No haloperidol was detected in the control tanks, or in Lake Superior water blanks. There were no significant differences in measured concentrations between replicate tanks within a treatment.

One female from the control group died on the final day of exposure prior to sampling, but there were no mortalities in any of the haloperidol treatments. Cumulative fecundity was greatest for control fish and lowest for fish exposed to 2.0 μg haloperidol/L but there were no statistically significant differences among treatments (Fig. 1), nor were there significant differences in the fertility of spawned eggs (data not shown). Similarly, there were no significant treatment-related differences in fish weight, gonad weight, gonadal somatic index [gonad wt. × body wt. −1 × 100], or male tubercle scores, dorsal pad weight, or dorsal pad index [dorsal pad wt. × body wt. −1 × 100%] (Table 2). Plasma estradiol, testosterone, and vitellogenin concentrations were not

![Fig. 1. Effects of haloperidol on fathead minnow fecundity in a 21 d test. Upper panel shows total cumulative fecundity per treatment group as a function of time. Lower panel shows mean (+ SE) of the total number of eggs spawned per pair (n=8 pairs per treatment) over the 21 d exposure.](image-url)
Among the five gene expression endpoints examined, cGnRH transcript abundance in male brain was the only variable significantly affected by haloperidol exposure (Table 2; Supplementary Figs. S.1 and S.2). Expression of cGnRH mRNA in the brain of male fish exposed to 20 μg haloperidol/L was increased compared to controls (Fig. 2).

Based on incidence and severity, there were no substantial differences in histological findings between exposed males and females and corresponding controls. Fungal elements surrounded by well-circumscribed clusters of macrophages, but no signs of other inflammatory cells or tissue damage, were detected in testes of three males from the 0.02 and 2.0 μg haloperidol/L groups. Additionally, unusual intermingling of yolk granules with cortical alveoli in mature vitellogenic oocytes was noted in the ovaries of three females from 0, 0.2, and 2.0 μg/L groups. However, none of these unusual observations were considered exposure-related. Histologically determined gonad stage (Villeneuve et al., 2007c) was not significantly affected by haloperidol exposure.

4. Discussion

4.1. Reproductive endpoints

This study is the first to apply a fathead minnow assay developed by the US Environmental Protection Agency for its Endocrine Disruptor Screening Program (http://www.epa.gov/endo/) to assess the potential reproductive toxicity of a chemical known to interact directly with a neurotransmitter receptor. Results of this study did not support the hypothesis that exposure to haloperidol would affect the brain mass production significantly by haloperidol treatment in either males or females, nor did ex vivo steroid production vary significantly with treatment (Table 3). Among the five gene expression endpoints examined, cGnRH transcript abundance in male brain was the only variable significantly affected by haloperidol exposure (Fig. 2; Supplementary Figs. S.1 and S.2). Expression of cGnRH mRNA in the brain of male fish exposed to 20 μg haloperidol/L was increased compared to controls (Fig. 2).

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### Table 2

<table>
<thead>
<tr>
<th>Endpoint</th>
<th>Sex</th>
<th>Haloperidol concentration (nominal; μg/L)</th>
<th>0</th>
<th>0.02</th>
<th>0.2</th>
<th>2.0</th>
<th>20</th>
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<tbody>
<tr>
<td>Mass (g)</td>
<td>M</td>
<td>3.90 ± 0.65</td>
<td>3.95 ± 0.82</td>
<td>4.05 ± 0.82</td>
<td>3.28 ± 0.76</td>
<td>3.46 ± 0.67</td>
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<tr>
<td></td>
<td>F</td>
<td>1.52 ± 0.24</td>
<td>1.48 ± 0.24</td>
<td>1.57 ± 0.24</td>
<td>1.43 ± 0.27</td>
<td>1.51 ± 0.25</td>
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<tr>
<td>Gonad mass (mg)</td>
<td>M</td>
<td>48 ± 13</td>
<td>45 ± 15</td>
<td>57 ± 24</td>
<td>58 ± 44</td>
<td>37 ± 14</td>
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<tr>
<td></td>
<td>F</td>
<td>183 ± 52</td>
<td>173 ± 101</td>
<td>192 ± 58</td>
<td>161 ± 75</td>
<td>189 ± 63</td>
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<tr>
<td>GSI (%)</td>
<td>M</td>
<td>1.25 ± 0.26</td>
<td>1.12 ± 0.31</td>
<td>1.41 ± 0.60</td>
<td>1.82 ± 1.61</td>
<td>1.02 ± 0.24</td>
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<td></td>
<td>F</td>
<td>121 ± 2.5</td>
<td>112 ± 4.3</td>
<td>125.5 ± 3.7</td>
<td>109 ± 3.2</td>
<td>123 ± 3.1</td>
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<tr>
<td>Dorsal pad mass (mg)</td>
<td>M</td>
<td>140 ± 71</td>
<td>163 ± 69</td>
<td>114.3 ± 32</td>
<td>78 ± 36</td>
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<td></td>
<td>F</td>
<td>3.6 ± 1.5</td>
<td>4.1 ± 1.4</td>
<td>2.8 ± 0.7</td>
<td>2.6 ± 1.6</td>
<td>3.6 ± 2.2</td>
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<tr>
<td>Tubercle score</td>
<td>M</td>
<td>33.1 ± 6.4</td>
<td>29.6 ± 5.6</td>
<td>31.9 ± 6.2</td>
<td>34.0 ± 5.1</td>
<td>32.7 ± 8.5</td>
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</tbody>
</table>

a Males only.
b Whole body wet weight measured immediately following blood sampling.

### Table 3

<table>
<thead>
<tr>
<th>Endpoint</th>
<th>Sex</th>
<th>Haloperidol concentration (nominal; μg/L)</th>
<th>0</th>
<th>0.02</th>
<th>0.2</th>
<th>2.0</th>
<th>20</th>
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</thead>
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<td>Ex vivo T (ng/ml)</td>
<td>M</td>
<td>2.61 ± 1.24; 8</td>
<td>2.24 ± 0.90; 8</td>
<td>2.28 ± 1.24; 8</td>
<td>1.97 ± 1.56; 8</td>
<td>2.75 ± 1.03; 8</td>
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<td>0.47 ± 0.39; 7</td>
<td>0.23 ± 0.34; 8</td>
<td>0.23 ± 0.21; 8</td>
<td>0.37 ± 0.35; 8</td>
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<td>Ex vivo E2 (ng/ml)</td>
<td>M</td>
<td>&lt; 0.05b; 8</td>
<td>&lt; 0.05b</td>
<td>&lt; 0.05; 8</td>
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<td>2.55 ± 0.69; 7</td>
<td>2.63 ± 0.99; 8</td>
<td>2.03 ± 0.35; 8</td>
<td>2.20 ± 0.61; 8</td>
<td>1.82 ± 0.74; 8</td>
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</tr>
<tr>
<td>Plasma T (ng/ml)</td>
<td>M</td>
<td>13.8 ± 2.8; 8</td>
<td>11.1 ± 5.1; 8</td>
<td>12.7 ± 4.2; 8</td>
<td>14.1 ± 6.6; 8</td>
<td>16.2 ± 4.4; 8</td>
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<td>F</td>
<td>4.03 ± 3.60; 5</td>
<td>4.63 ± 3.62; 7</td>
<td>4.33 ± 2.95; 6</td>
<td>7.20 ± 4.75; 6</td>
<td>6.98 ± 3.88; 6</td>
<td></td>
</tr>
<tr>
<td>Plasma E2 (ng/ml)</td>
<td>M</td>
<td>0.43 ± 0.28; 8</td>
<td>0.43 ± 0.33; 8</td>
<td>0.36 ± 0.30; 8</td>
<td>0.43 ± 0.23; 8</td>
<td>0.37 ± 0.37; 8</td>
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<tr>
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<td>6.99 ± 3.55; 7</td>
<td>10.0 ± 3.22; 7</td>
<td>9.63 ± 3.61; 8</td>
<td>7.73 ± 4.80; 7</td>
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<tr>
<td>Plasma Vtg (mg/ml)</td>
<td>M</td>
<td>1.58 ± 1.30; 8</td>
<td>1.38 ± 1.15; 8</td>
<td>1.66 ± 1.54; 8</td>
<td>2.63 ± 3.32; 7</td>
<td>3.15 ± 3.76; 8</td>
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<tr>
<td>(mg/ml)</td>
<td>F</td>
<td>18.0 ± 9.54; 7</td>
<td>13.5 ± 6.55; 8</td>
<td>14.0 ± 5.60; 8</td>
<td>16.9 ± 12.2; 8</td>
<td>17.5 ± 9.00; 8</td>
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</tr>
</tbody>
</table>

a ng/ml of incubation medium, following 16.5 h incubation.
b Method detection limit for the E2 radioimmunoassay was 0.05 ng/ml. Testes samples generally did not produce detectable E2 concentrations over a 16.5 h incubation period.
c ng/ml plasma.
d Note, male and female Vtg concentrations are presented using different units as indicated.
fecundity of fathead minnows exposed over a 21 d period. The average productivity of fish exposed to 2.0 µg/L was less than that of other groups, but the difference was not statistically significant, nor was there a clear concentration-dependent trend. Similarly, there was no evidence that exposure to a D2R antagonist enhanced spawning activity as had been reported for grey mullet (Aizen et al., 2005). The general lack of significant effects on reproductive parameters including secondary sex characteristics, gonad histology, and plasma steroid and Vtg concentrations was consistent with the lack of effect on fecundity or fertility.

4.2. Gene expression

Supervised analysis of the effects of haloperidol exposure on the abundance of mRNA transcripts coding for five different proteins was conducted by QPCR. Examination of D2R transcripts in brain did not provide any evidence to suggest that receptor expression had increased to offset the antagonistic effects of the chemical. Similarly, the lack of significant effects on the expression of GtH β subunits in the pituitary and FSHR in the testes did not provide any indication of drastic shifts in GtH production or signaling, although we note that the regulatory effect of dopamine is thought to be at the level of GtH release, rather than transcription (Trudeau, 1997). Among the gene expression endpoints examined in the reproduction assay, the only suggestion that haloperidol may have been affecting the HPG axis of the exposed fish was a significant increase in cGnRH transcripts in the brain of males exposed to 20 µg haloperidol/L. This increase is broadly consistent with the idea that haloperidol exposure would release dopaminergic inhibition of GnRH synthesis and/or secretion. However, there was little corroborating evidence supporting that hypothesized effect. The same response could not be examined in the brains of females due to a technical mishap during RNA extraction. There were no significant changes in the abundance of LHβ or FSHβ transcripts in the pituitary tissue of exposed males or females. Finally, methods for measuring circulating peptide hormone concentrations (e.g., GnRH, GtH) are not currently available for the fathead minnow. Ultimately, whether haloperidol exposure modulated GnRH expression and/or secretion or not, results of the reproduction study did not indicate that haloperidol acted as a direct reproductive toxicant.

5. Conclusions

Results of our study did not support the hypothesis that exposure to the D2R antagonist, haloperidol, would directly disrupt fish reproduction. The potential for D2R antagonism to interfere with behavior or other aspects of transcriptional signaling in reproductive tissues was examined in a companion study (Villeneuve et al., 2009).

Acknowledgments

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Appendix A. Supporting information

Supplementary data associated with this article can be found in the online version at doi:10.1016/j.ecoenv.2009.09.007.

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