

Effects of alkylphenols on the biotransformation of diuron and enzymes involved in the synthesis and clearance of sex steroids in juvenile male tilapia (*Oreochromis mossambica*)



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ARTICLE INFO

Article history:

Received 1 June 2016

Received in revised form 18 October 2016

Accepted 19 October 2016

Available online 22 October 2016

Keywords:

Diuron

Alkylphenols

Anti-androgen

Vitellogenin

Biotransformation

Cytochrome P450

ABSTRACT

Previous studies using *in vivo* bioassay guided fractionation indicated that the herbicide diuron (3-(3,4-dichlorophenyl)-1,1-dimethylurea) and alkylphenol (AP)-containing surfactants were detected in fractions of extracts that induced the estrogenic biomarker, vitellogenin in fish exposed to surface water extracts from the United States. However, when the compounds were evaluated individually using *in vivo* estrogenic assays or *in vitro* estrogen receptor assays, estrogenic activity was not observed. Since APs have been shown to alter activity and content of cytochrome P450s (CYP) which convert diuron to potential estrogenic metabolites, the hepatic biotransformation of diuron was measured with and without a 7 day pretreatment of *p*-Octylphenol (OP) and *p*-Nonylphenol (NP) at low (OP 13 ng/L + NP 91 ng/L), and high concentrations (OP 65 ng/L + NP 455 ng/L) in juvenile male Nile tilapia (*Oreochromis niloticus*). Pre-treatment with the OP/NP (AP) mixture caused elevated levels of NADPH-catalyzed formation of 3,4-dichlorophenyl-*N*-methylurea (DCPMU) but not 3,4-dichlorophenylurea (DCPU). Fish were also treated with nominal concentrations of low (40 ng/L) and high (200 ng/L) diuron and each of its three degradation/metabolites: DCPMU, DCPU and 3,4-dichloroaniline (DCA). Additional treatments were conducted with APs and Diuron as a mixture at the low concentrations which mimicked concentrations observed in surface waters. Hepatic vitellogenin (Vtg) mRNA was induced by exposure to the high concentrations of Diuron, as well as DCPMU and DCPU in both concentrations. Brain cytochrome P450 aromatase activity was generally diminished by diuron, its metabolites, and the AP/diuron mixtures. 17 β -Hydroxysteroid dehydrogenase (17 β HSD) levels were also reduced by DCPMU and DCA in the lower concentrations, but not by higher concentrations. While the AP mixture reduced 17 β HSD, the AP/diuron mixture induced testosterone (T) biosynthesis at the single concentration tested. Although CYP3A expression was induced by all diuron metabolites, it was unchanged by the AP mixture. These data indicate that mixtures of AP and diuron enhanced the formation of the metabolite (DCPMU) which induced vitellogenin, and reduced T biosynthetic enzymes (17 β HSD inhibition). Overall, these data showed that APs may have induced the biotransformation of diuron to at least one metabolite, that may disrupt androgen biosynthesis and potentially alter steroid feedback pathways in the central nervous system.

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1. Introduction

Previous studies using *in vivo* bioassay guided fractionation for estrogenic activity in water extracts from the Central Valley and San Francisco Bay Delta in California (USA) identified Diuron (3-(3,4-dichlorophenyl)-1,1-dimethylurea) and alkylphenols in estrogenic fractions of water extracts (Schlenk et al., 2012). Diuron is one of the most widely used herbicides for sugar cane cultivation in

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Brazil (Lourencetti et al., 2008) and a general use herbicide in the United States (Tomlin, 1994). Alkylphenols are derived primarily from alkylphenol ethoxylates which are routinely used as surfactants in a number of applications including pesticide formulations (Xie et al., 2005). When the individual compounds (diuron, octylphenol and *p*-nonylphenol) were administered to primary hepatocytes of rainbow trout (*Oncorhynchus mykiss*) at ambient concentrations, estrogenic activity (induction of vitellogenin mRNA) was not observed (Schlenk et al., 2012). Similarly, when male Japanese medaka (*Oryzias latipes*) were exposed to the individual compounds, hepatic vitellogenin protein was not observed. However, when diuron was combined with several alkylphenol ethoxylates and alkylphenols as well as the pyrethroid, bifenthrin, which was also detected in bioactive fractions, estrogenic activity was observed in the *in vivo* assays, but not the *in vitro* assays (Schlenk et al., 2012).

Diuron has also been shown to be anti-androgenic *in vitro* (Orton et al., 2009), *in vivo* (Pereira et al., 2015) and impair steroidogenesis at relatively high concentrations (15.6–7.3, and 14.5 mg/L, respectively) (Vinggaard et al., 2000). Since responses to the mixtures were only noted *in vivo* during previous studies, the potential targets for diuron and/or the other compounds observed in estrogenically active water extracts are unclear. One possible mechanism may involve the conversion of diuron to other compounds with endocrine activity leading to feminization. In mammals, diuron undergoes biotransformation to 3,4-dichloroaniline (DCA), 3,4-dichlorophenylurea (DCPU) and 3,4-dichlorophenyl-*N*-methylurea (DCPMU) (Tixier et al., 2002; Hodge et al., 1967; Abass et al., 2007). Cytochrome P450 has been shown to be primarily responsible for the demethylation of diuron in mammals to DCPMU and DCPU (Abass et al., 2007), but its role in fish biotransformation is unclear.

Each metabolite/degradate weakly binds to the bovine androgen receptor, with relative binding affinities between 0.002 and 0.01% of the affinity of dihydrotestosterone (Orton et al., 2009). DCA, DCPU, and DCPMU all reduced plasma testosterone and 11-keto-testosterone along with gonadal somatic indices, diameter of seminiferous tubules and mean percentages of germ cells in male tilapia (Pereira et al., 2015). However, the mechanism of the anti-androgenic activity was not evaluated.

While the androgen receptor may be a potential target for diuron and its metabolites, a second target may be the hypothalamic-pituitary-gonadal axis which synthesizes and secretes gonadotropic and gonadal steroid hormones, and reproductive process modulators (Agulleiro et al., 2006). These hormones play a key role in fish gametogenesis (Mylonas et al., 2010). In males, for example, gonadal steroids (11-ketotestosterone and testosterone) mediate spermatogonial proliferation and spermiation (Ohta et al., 2007). Since diuron impaired the biosynthesis of these reproductive hormones in fish (Vinggaard et al., 2000; Pereira et al., 2015), the aim of this study was to evaluate the effects of diuron with and without combination of alkylphenols as well as diuron metabolites on two enzymes that are involved in testosterone biosynthesis and metabolism (17 β HSD and aromatase) as well as xenobiotic biotransformation and steroid clearance (CYP3A) in juvenile male tilapia, *Oreochromis mossambicus*.

2. Materials and methods

2.1. Chemicals

All chemicals were 99% pure based on manufacturers information. Diuron and metabolites were ordered from Sigma-Aldrich Chemical Co (St. Louis, Mo).

Table 1

Treatment groups for diuron, metabolite and alkylphenol mixture exposures.

Group number	Treatment
1	Control
2- diuron low	40 ng/L of diuron
3- diuron high	200 ng/L of diuron
4- DCPMU low	40 ng/L of DCPMU
5- DCPMU high	200 ng/L of DCPMU
6- DCPU low	40 ng/L of DCPU
7- DCPU high	200 ng/L of DCPU
8- DCA low	40 ng/L of DCA
9- DCA high	200 ng/L of DCA
10-AP mixture low	13 ng/L of OP + 91 ng/L of NP
11-AP mixture high	65 ng/L of OP + 455 ng/L of NP
12-AP/Diuron mixture	diuron 40 ng/L + 13 ng/L of OP + 91 ng/L of NP

2.2. Test organisms

Juvenile male *Oreochromis mossambicus*, were selected as model organisms because of their worldwide use as a species for aquaculture in the USA and Brazil. Individuals aged one month weighed 78.23 g \pm 3.07 g and had standard lengths of 15.29 cm \pm 0.16 cm. They were obtained from Blue and Beyond Fisheries (Desert Hot Springs, California). Gender was confirmed by microscopic analyses post-mortem. This work was approved by UCR IACUC # 2010-0004.

2.3. Exposure experiments

To a 20L aquarium, 18L of water and one fish was added prior to the addition of chemicals. There were 12 treatment groups, with five fish per group creating five replicates. The temperature was maintained at 28 \pm 2 °C and water was constantly aerated. Fish were acclimated for five days in their respective aquarium before exposure. During the exposure period fish were not fed and the water was changed every two days, to maintain toxicant concentrations and control ammonia concentrations, which were determined throughout the experiment.

The groups of the experiment are described in Table 1. Low concentrations used in this experiment were based upon the measurements observed in surface waters of the San Francisco Bay Delta (Schlenk et al., 2012). High concentrations were arbitrary 5 fold increases.

After seven days of exposure all fish were collected, anesthetized, and tissues (brain and liver) were collected and stored in –80 °C, for analysis.

2.4. Fish microsome isolations

Liver and brain were homogenized using the methods of Lavado et al. (2004) with minor modifications. Gonads were not measured because the tissues did not provide enough biomass. Tissues were homogenized (1:5, w/v) in Phosphate Buffer (100 mM, pH 7.4) containing 100 mM of KCl and 1 mM of EDTA. The homogenized samples were centrifuged at 1500 \times g for 15 min at 4 °C and the resulting supernatant centrifuged again at 12,000 \times g for 12 min at 4 °C. The supernatant from this step was centrifuged again at 100,000 \times g for 60 min to obtain the microsomal fraction as a pellet. The pellet was re-suspended in 1:10 (w/v) in Phosphate Buffer (100 mM, pH 7.4) containing 100 mM of KCl, 1 mM of EDTA and 20% of Glycerol.

For protein quantification, the Bradford (1976) method was used with bovine serum albumin as standard.

2.5. Diuron biotransformation

Liver microsomal fractions of fish from groups 3 (exposed to 200 ng/L of diuron), 10 (exposed to 13 ng/L of OP + 91 ng/L of NP),

and 11 (exposed to 65 ng/L of OP + 455 ng/L of NP) were incubated for 30 min at 30 °C in 100 mM Tris-HCl buffer (pH 7.4) with 100 μ M of diuron and 7.5 mM of NADPH. An additional 7.5 mM of NADPH was again mixed with the incubation for another 30 min. The reaction was stopped with 250 μ L of ice cold acetonitrile to precipitate proteins and the internal standard (100 μ M caffeine) was added. Recovery rates were 60–70%. Sample was centrifuged at 14,000 \times g for 10 min and the supernatant was collected for HPLC analyses of DCPMU, DCPU and DCA.

HPLC analyses were performed on a SCL-10AVP Shimadzu HPLC System equipped with a 250 \times 4.6 mm Hypersil ODS C18 (5 μ m) reverse-phase column (ThermoFisher Scientific Inc, Waltham, MA) using an absorbance of 250 nm for quantification. The mobile phase an isocratic 50% mixture of acetonitrile and water with a 1 mL/min flow rate. Absorbance of eluates was measured at 254 nm. Quantification of the metabolites was attained through linear extrapolation from standard curves employing 5 concentrations of each compound.

2.6. CYP3A immunoblot analysis

Microsomal CYP3A from liver was determined by Western immunoblot using the method of Lavado et al. (2004) with minor modifications. Microsomes were boiled at 95 °C for 5 min in SDS-PAGE buffer and 10 μ g of protein was separated by electrophoresis using 10% SDS-polyacrylamide gels, first for 30 min in 60 V and after for 80 min in 150 V, and transferred to Nitrocellulose Membranes using a Trans-Blot[®] Semi-Dry Transfer Gel (Bio-Rad Laboratories, Hercules, CA, USA) for 30 min in 15 V. They were probed using a 1:2000 dilution (v/v) of primary rabbit anti-human CYP3A4, incubating overnight at 25 °C. The membrane was rinsed three times with Tris-buffered saline containing 0.2% Tween 20 (v/v) and 0.5% gelatin (w/v). The membranes were incubated for 1 h with Goat Anti-Rabbit IgG (H+L)-AP Conjugate (Bio-Rad, Hercules, CA, USA) and using the Alkaline Phosphatase Conjugate Substrate Kit (Bio-Rad Laboratories) developed. Quantification was carried out with a ChemiDoc[™] XRS+ System (Bio-Rad) using Image Lab Software (Bio-Rad) to measure optical density units/mg protein. Loading was normalized by total protein measurements.

2.7. Vitellogenin analysis

Vitellogenin messenger RNA was measured using the method of Lavado et al. (2013). Total RNA from the liver was extracted using RNeasy[®] Lipid Tissue MiniKit (QIAGEN; Valencia, CA, USA). For cDNA preparation High Capacity cDNA Reverse Transcription Kit (Applied Biosystems[™]; Foster City, CA, USA) was used, and the quality of the extracted RNA was evaluated by spectrophotometry (Nanodrop). Real time quantitative PCR was performed using “iScript[™] One Step” RT-PCR kits with SYBR[®] Green (BioRad; Hercules, CA, USA). Vitellogenin primers (R – 5’GTC CTC CCT GAT CAC ATTA GTT G 3’ and F – 5’CTC AGT TGC TGG AGT ACA GTG 3’) and EF1-alpha primers (R – 5’GCA TAA GCC ATG CCT TGA GTA TAG 3’ and F – 5’CGG TGT CAT CAA GTC CGT TAT C 3’) for housekeeping were used. Analyses were conducted using an iCycler-MyIQ Single Color Real-Time PCR Detection System (Bio-Rad; Hercules, CA, USA) and data analyzed with IQ5 (Bio-Rad; Hercules, CA, USA). Primers were optimized based on annealing temperature, template concentration, and primer concentration. 250 nM of each primer was added to 25 μ L PCR reactions containing SYBR Green RT-PCR Reaction Mix, and 100 ng of cDNA sample. Thermocycling parameters were as follows: 5 mins at 95 °C; 40 cycles of 10 s at 95 °C and 30 s at 57 °C. At the end of each cycle fluorescence data was collected. A melting curve analysis was run between 60 °C and 95 °C following

the amplification reaction. The C(t) was selected to be in the linear phase of amplification.

2.8. Liver 17 β -Hydroxysteroid dehydrogenase enzyme activity

Liver microsomal fractions were incubated in 50 mM of Tris-HCl Buffer (pH 7.4) with 0.2 μ M [³H]androstenedione (150,000 dpm), 10 mM MgCl₂ and 300 μ M NADPH in a total volume of 250 μ L, as described in Thibaut and Porte (2004). Samples were incubated for 15 min in 30 °C. To stop reaction 250 μ L of acetonitrile was added and the solution was centrifuged 1500g for 10 min. The supernatant was collected and analyzed in HPLC. HPLC analyzes was performed on a SCL-10AVP Shimadzu HPLC System equipped with a 250 \times 4.6 mm Hypersil ODS C18 (5 μ m) reverse-phase column (ThermoFisher Scientific Inc, Waltham, MA), in a 254 nm of wave-length. Separation of androstenedione and metabolites was performed at 1 mL/min with a mobile phase composed of (A) 75% of water and 25% of acetonitrile, and (B) 25% of water and 75% of acetonitrile. Chromatographic peaks was monitored by on-line radioactivity detection with Radioflow detector LB 509 (Berthold Technologies, BadWildbad, Germany) using Flo-Scint 3 (Packard BioScience, Groningen, The Netherlands) as scintillation cocktail.

2.9. Brain aromatase enzyme activity

Since animals did not have adequate gonadal tissue for aromatase analyses, the brain microsomal fraction was used to measure aromatase activity, using the tritiated-water method as described in Morcillo et al. (1999), with some changes. Each sample was adjusted to 2 mg/ml of protein and was incubated at 25 °C for 30 min with 100 mM Tris-HCl (pH 7.6), 10 μ M unlabeled androstenedione, 4.35 mM NADPH and 0.1 Ci/ μ L [³H]-androstenedione. Assay blanks containing 100 μ L of buffer instead of microsomes were used for every run. The reaction was stopped by placing the tube on ice; organic metabolites and the excess of substrate were immediately eliminated from the aqueous phase by extraction with methylene chloride. Following this, a suspension containing 2.5% (w/v) activated charcoal and 0.25% dextran in Milli-Q water was added. The mixture was centrifuged and the supernatant counted for ³H radioactivity.

2.10. Statistical analyses

Statistical analyses were conducted using Prism5 v5.0a software. Prior to statistical analysis, all data were analyzed to meet the normality and variance assumptions for parametric tests. For normally distributed data, an initial one-way ANOVA was performed to evaluate the differences between treatments. If a P-value less than 0.05 was observed, it was considered statistically significant, and if there was significance, the Tukey’s multiple range test was performed to determine differences between groups. If data did not meet assumptions of the parametric test, a Kruskal–Wallis test and two-tailed multiple comparisons Dunn’s test was used.

3. Results

3.1. Biotransformation

In vitro incubations of diuron with liver microsomes from untreated tilapia catalyzed the NADPH-dependent formation of DCPU, and DCPMU from diuron (Table 1). Fish pretreated with AP significantly increased the rates of DCPMU formation 3-fold after treatment (Table 2).

To evaluate the potential role of CYP3A in the biotransformation of diuron, expression of hepatic CYP3A was measured after treatment with Diuron, its metabolites and the AP mixture. CYP3A was

Table 2
Diuron biotransformation (nmol/min/mg prot.) in liver microsomes of *O. mossambicus* exposed to diuron, octylphenol + nonylphenol (OP+NP), or unexposed (control) for 7 days. Each value represents the mean value of 5 replicates \pm SD.

	Control	Diuron (200 ng/L)	OP+NP (13+91 ng/L)	OP+NP (65+455 ng/L)
DCPMU	0.18 \pm 0.05	0.40 \pm 0.28	0.58 \pm 0.36*	0.31 \pm 0.09
DCPU	0.08 \pm 0.05	0.16 \pm 0.14	0.21 \pm 0.19	0.07 \pm 0.03
DCA	ND	ND	ND	ND

ND = Not detected.

* ($p < 0.05$) compared to the control group.

Table 3
Effects of Diuron, its metabolites/degradates and APs alone and in mixtures on hepatic CYP3A-like expression in *O. mossambicus* after 7 days of exposure. Each value represents the mean \pm SD of 5 replicates.

Compound	Concentration (ng/L)	CYP3A expression
Control	–	1.07 \pm 0.11
Diuron	40	0.65 \pm 0.20
	200	1.60 \pm 0.24
DCPMU	40	0.96 \pm 0.16
	200	4.26 \pm 0.62*
DCPU	40	1.49 \pm 0.22
	200	4.38 \pm 1.08*
DCA	40	1.45 \pm 0.30
	200	2.04 \pm 0.52*
OP+NP	13+91	0.40 \pm 0.09*
	65+455	0.52 \pm 0.15
OP+NP+Diuron	13+91+40	0.86 \pm 0.48

* $p \leq 0.05$.

significantly induced in livers of tilapia following treatment with the diuron metabolites, reduced by AP, and unchanged following pretreatment with AP/diuron (Table 3).

3.2. Estrogenic activity

Vitellogenin expression was statistically higher after seven days of exposure to the high concentrations of Diuron (high concentration), as well as DCPMU and DCPU in both concentrations. Induction was also observed in animals treated with the low concentration of DCA (Fig. 1A). Although trends toward an increase in Vtg were observed with the AP mixture, the only significant difference was observed between diuron treatments and the AP/diuron treatment (ANOVA $p = 0.05$) (Fig. 1B).

3.3. Steroid biosynthetic enzymes

Aromatase activity in the brain was statistically lower after exposure to the high Diuron treatment. DCPMU in both concentrations, and the low concentration of DCPU also diminished activity (Fig. 2A). Treatment with DCA failed to alter activities. AP treatment with and without the low Diuron concentration also caused significant reductions in aromatase activities (Fig. 2B).

17 β HSD activity was significantly reduced by treatment of fish with the low concentration of DCPMU and DCA (Fig. 3A). Pretreatment with AP also reduced 17 β HSD. However, combinations of diuron with AP led to a non-significant increase in activity (Fig. 3B).

4. Discussion

Previous studies indicated that APs and Diuron were observed in estrogenic fractions that induced vitellogenin in fish treated with concentrations observed in surface waters (Schlenk et al., 2012). Mixtures of APs, diuron and bifenthrin caused induction *in vivo*, but not *in vitro* using primary hepatocyte cells from rainbow trout indicating direct activation of estrogen receptor by any of these

compounds was not a likely mechanism. More recent studies have reported that diuron and its primary metabolites cause reductions of plasma testosterone and 11-ketotestosterone as well as histological alterations of gonads indicating anti-androgenic effects to male fish (Pereira et al., 2015). APs have been shown to modify steroid clearance enzymes and diuron metabolizing enzymes (CYPs) (Lee et al., 1996). Since diuron metabolites were shown to have anti-androgenic effects, the purpose of this study was to determine whether mixtures of AP and diuron affect biotransformation of diuron to the active metabolites, and whether enzymes that are critical in steroid biosynthesis and clearance were impacted.

In tilapia liver microsomes, diuron was readily converted to DCPU, and DCPMU (Table 2). In rats and dogs, DCPU was the predominant metabolite in the urine with small amounts of DCPMU, DCA, 3,4-dichlorophenol, and unchanged diuron detected (Hodge et al., 1967). DCPMU was the predominant metabolite (~35%) formed by NADPH catalysis in mouse liver. Interestingly, when the microsomes were incubated with DCPMU to determine whether additional demethylation occurs only 4% conversion to DCPU was observed (Suzuki and Casida, 1981). Consistent with this observation, liver microsomes from seven mammalian species and recombinant human CYPs, only observed DCPMU formation (Abass et al., 2007). The rank order of formation based on intrinsic clearance (V_{max}/K_m) was dog > monkey > rabbit > mouse > human > minipig > rat. These data indicate biotransformation of diuron in tilapia is qualitatively similar to that of mammals with demethylation of diuron being the predominant enzymatic reaction.

Demethylation of diuron to DCPMU was significantly induced by AP pre-treatment indicating modulation of CYPs in tilapia (Table 2). Evaluation of 12 recombinant human CYPs demonstrated that all were able to catalyze demethylation of diuron (Abass et al., 2007). The highest activities were observed with CYP1A1, CYP1A2, CYP2C19, and CYP2D6. While there are not any known homologous orthologs to CYP2C or CYP2D in fish, CYP1A activities appear to be conserved among vertebrates including fish (Schlenk et al., 2008). However, NP inhibits expression and catalytic activities of CYP1A in rodents (Lee et al., 1996), and most fish (Sturve et al., 2006). In addition, diuron was shown to potently inhibit CYP1A in humans (Abass et al., 2007). Thus, given its lack of induction of CYP1A by NP in fish, induction of diuron metabolism by NP treatment indicates other CYPs may be induced.

NP treatment has been shown to induce CYP3A in rats (Lee et al., 1996) and induce CYP3A at environmentally relevant concentrations (Arukwe et al., 1997). Since diuron and its metabolites caused a significant reduction of plasma androgens (Pereira et al., 2015), it was hypothesized that induction of CYP3A may be a mechanism by which testosterone clearance may be enhanced leading to lower plasma concentrations. However, AP treatment reduced CYP3A expression and when in mixture with diuron, failed to significantly induce vitellogenin expression (Table 3). Lower levels of CYP3A with AP treatment is inconsistent with the enhanced formation of DCPMU by AP suggesting other CYPs not modified by AP may be responsible for the demethylation of diuron. In addition, reduction of CYP content by AP is also inconsistent with dimin-

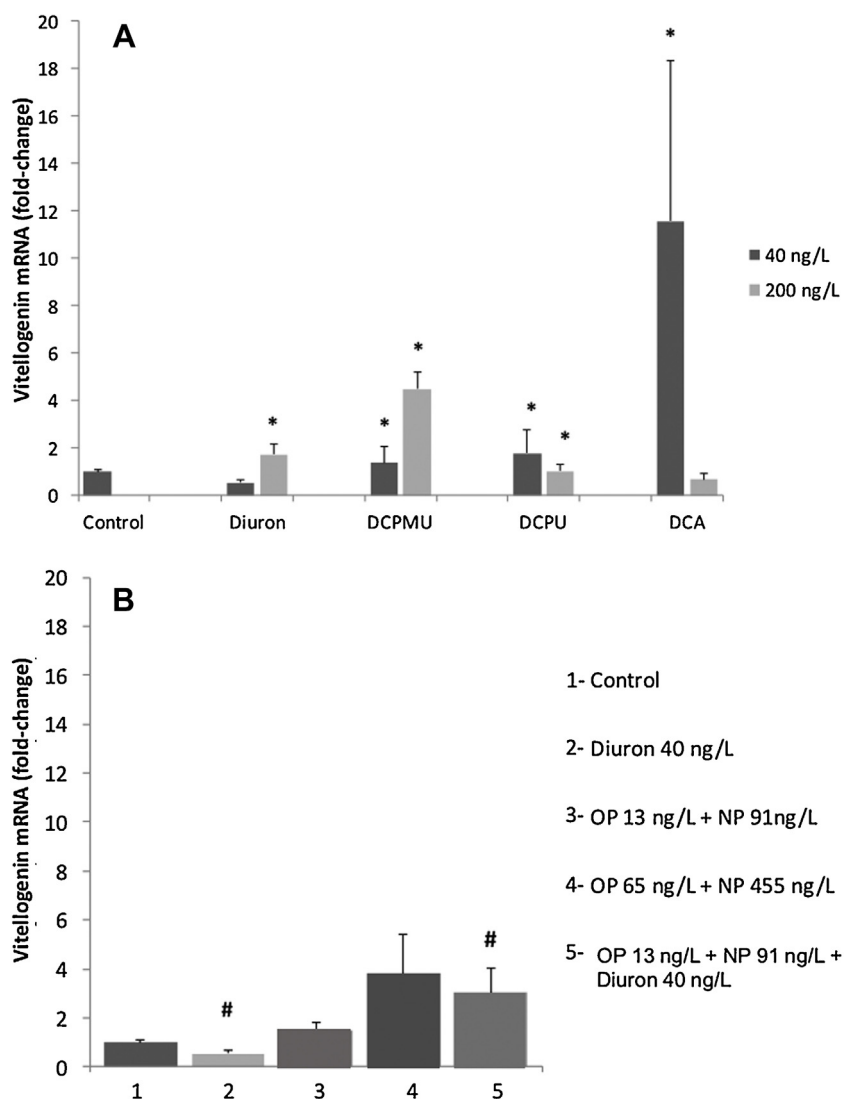


Fig. 1. Vitellogenin mRNA expression in liver of *O. mossambicus* exposed to 40 and 200 ng/L of Diuron, DCPMU, DCPU and DCA for 7 days (A) and mixtures diuron 40 ng/L, octylphenol (OP) 13 ng/L + nonylphenol (NP) 91 ng/L, OP 65 ng/L + NP 455 ng/L, and OP 13 ng/L + NP 91 ng/L + Diuron 40 ng/L for 7 days. (B). Each value represents the mean \pm SD of five replicates. * represents significant difference compared to control; # represents significant difference between treatments ($p < 0.05$).

ished testosterone concentrations caused by diuron observed in earlier tilapia studies (Pereira et al., 2015). CYP3A is the initial clearance enzyme for testosterone in vertebrates (Schlenk et al., 2008). Hydroxylation at the 6 β and 16 β position is followed by Phase II conjugation through sulfation or glucuronidation (Schlenk et al., 2008).

Although AP/Diuron modestly induced vitellogenin transcription relative to diuron alone, demethylated metabolites and the high concentration of diuron induced vitellogenin mRNA at equal efficacies and in the case of DCPMU, a concentration dependent manner (Fig. 1A). None of the compounds are estrogen receptor agonists, but diuron has relatively weak anti-estrogenic and anti-androgenic binding activities (Orton et al., 2009). Consistent with these effects, earlier studies in Japanese medaka (*Oryzias latipes*) showed that exposure to 41 ng/L of diuron alone failed to induce vitellogenin (Schlenk et al., 2012). Unfortunately, higher concentrations such as the 200 ng/L in this study, were not tested in medaka. The similarity in responses between the two species at the 40 ng/L concentration is significant since the current study utilized qPCR assessment of vitellogenin transcription in the liver, whereas the medaka measurements utilized enzyme linked immunosorbent

assays of hepatic vitellogenin protein. Since vitellogenin is transcriptionally regulated by estrogens, measurements of mRNA have been shown to be more sensitive in short duration assessments, while measurements of protein have a longer half-life and may be more robust during chronic exposures (Schultz et al., 2001). Responses in two species to the same compound indicate vitellogenin mRNA and/or protein in male or sexually immature animals provide consistent indications of estrogenic activity.

The mechanisms of vitellogenin induction by diuron and its demethylated metabolites are unclear, but may be related to alterations of circulating steroid hormones. Vitellogenin expression is directly controlled by E2, which is typically synthesized *via* CYP19 catalyzed aromatization of testosterone in the gonad. In contrast to the single form found in mammals, two genes of CYP19 have been found in teleosts; one of which is expressed mainly in the brain (CYP19a1b) and the second mainly in the gonad (CYP19a1a) (see Diotel et al., 2010 for review). While the gonadal forms are thought to be more involved in general reproduction, the brain forms appear to control many other neuroendocrine responses such as behavior. Brain aromatase activity was reduced by nearly all diuron treatments in tilapia (Fig. 2). Reductions in brain aromatase

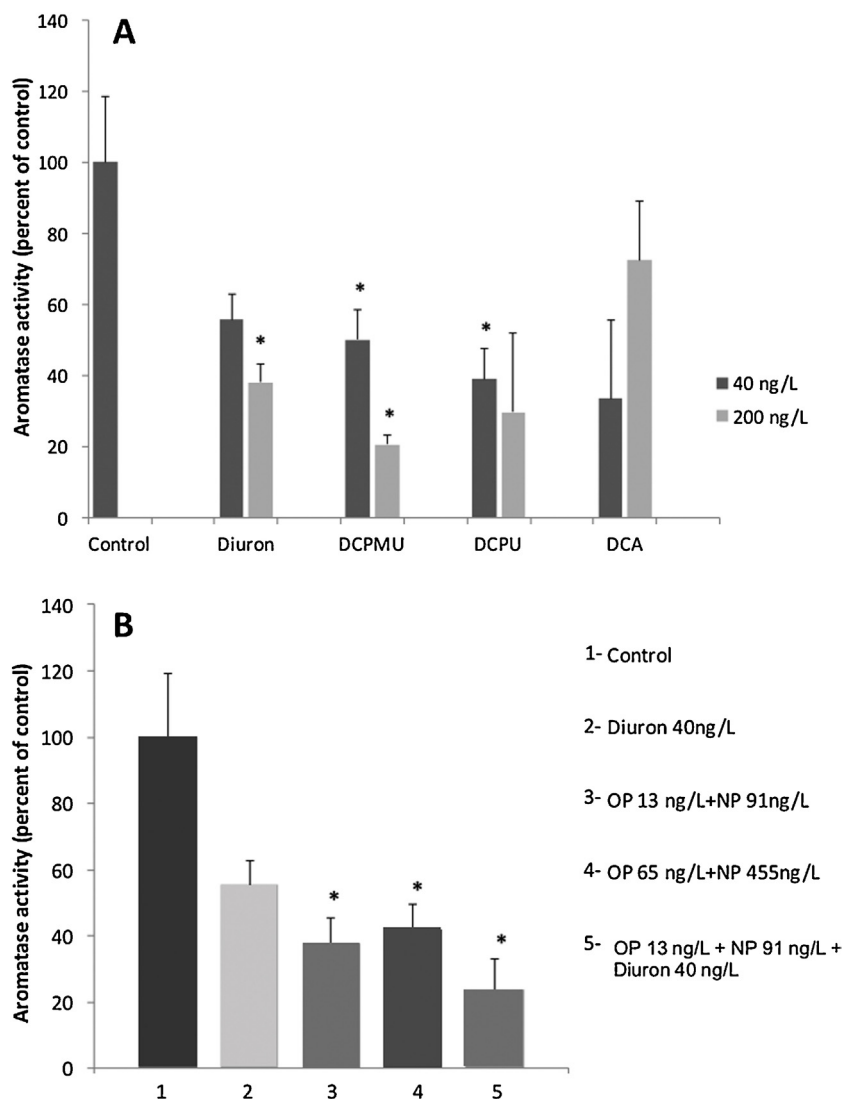


Fig. 2. Aromatase activity in brain of *O. mossambicus* exposed to 40 and 200 ng/L of Diuron, DCPMU, DCPU and DCA for 7 days (A) and mixtures diuron 40 ng/L, octylphenol (OP) 13 ng/L + nonylphenol (NP) 91 ng/L, OP 65 ng/L + NP 455 ng/L, and OP 13 ng/L + NP 91 ng/L + Diuron 40 ng/L for 7 days (B). Each value represents the mean \pm SD of five replicates * ($p < 0.05$).

may have a number of significant neuroendocrine outcomes including altered release of neuronal E2 or other hypothalamic pituitary signals. Several studies have shown that diuron impaired steroidogenesis in various organisms (Cardone et al., 2008; Kojima et al., 2004) including fish (Vinggaard et al., 2000; Pereira et al., 2015). CYP19a1b is up-regulated *in vivo* by certain androgens, notably testosterone (Diotel et al., 2010). Although putative androgen-responsive elements were identified in the promoter region of cyp19a1b, transfection experiments showed that neither androgen receptor nor the potential response element were required for the stimulation of cyp19a1b by testosterone (Diotel et al., 2010). Consequently, the potential for diminished levels of testosterone biosynthesis caused by diuron or its metabolites may be consistent with reductions in brain aromatase in tilapia.

The fact that non-aromatized androgens (11-KT) were also diminished following treatment with diuron and its metabolites suggests that steroid biosynthesis upstream of androstenedione may also be a potential target (Pereira et al., 2015). Consistent with vitellogenin expression, DCPMU also inhibited 17 β HSD (Fig. 3). The failure of diuron alone to inhibit 17 β HSD is consistent with earlier studies that showed 17 β HSD was unaltered by diuron in

fish ovarian microsomes (Thibaut and Porte, 2004). Consequently, conversion of diuron to demethylated metabolites may be a key factor in the initiation of anti-androgenic activity. Additional studies focusing on estrogen levels, gonadal aromatase (in older animals), as well as steroid biosynthetic enzymes upstream of 17 β HSD or aromatase are needed to better understand this potential pathway.

5. Conclusion

Combined exposure to AP/Diuron caused a modest vitellogenin induction compared to diuron alone. Evaluation of targets that involve androgen transformation (CYP19) and clearance (CYP3A) suggest that enhanced biotransformation to metabolites with equivalent activity by AP, resulted in feminization (Vtg mRNA) possibly through anti-androgenization (reduced Testosterone formation *in vitro*). The relationships between anti-androgenization and feminization further suggest that impacts on brain aromatase as well as other steroid biosynthetic pathways may provide additional information regarding potential mechanisms for the estrogenic effects observed with diuron.

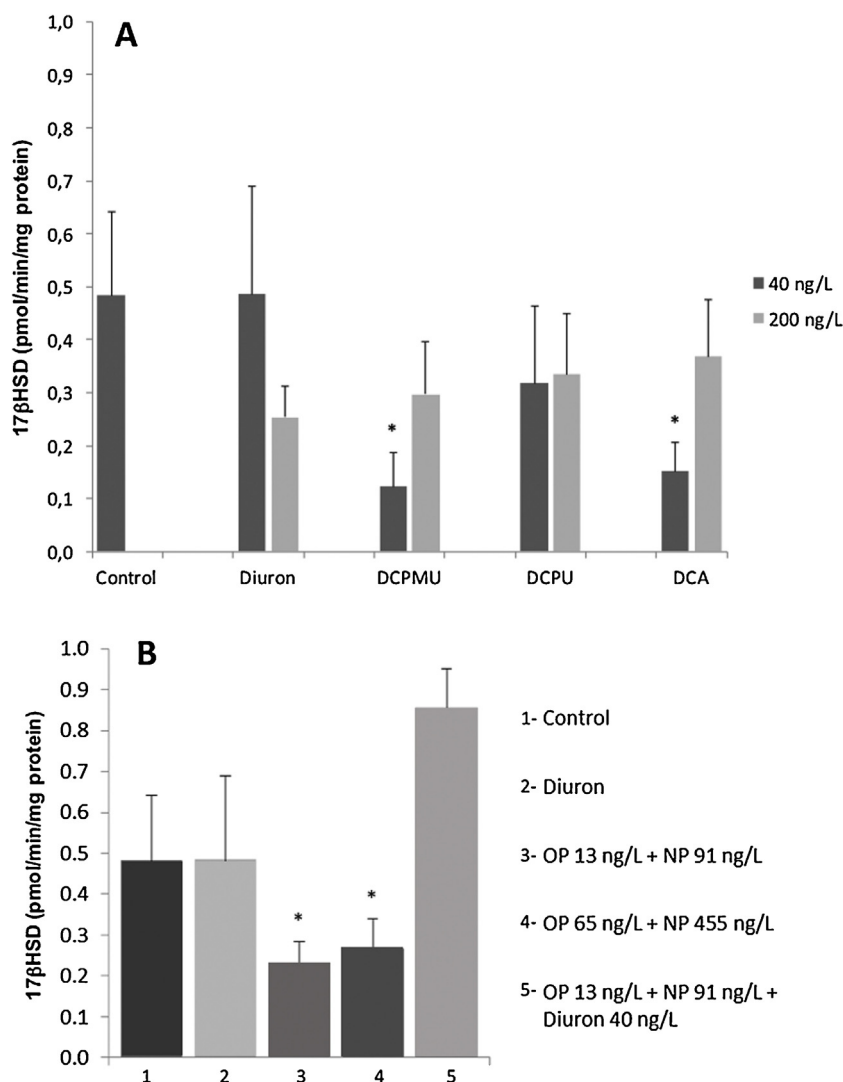


Fig. 3. Activity of liver 17βHSD in *O. mossambicus* exposed to 40 and 200 ng/L of Diuron, DCPMU, DCPU and DCA for 7 days (A) and mixtures diuron 40 ng/L, octylphenol (OP) 13 ng/L + nonylphenol (NP) 91 ng/L, OP 65 ng/L + NP 455 ng/L, and OP 13 ng/L + NP 91 ng/L + Diuron 40 ng/L for 7 days (B). Each value represents the mean ± SD of five replicates (* $p < 0.05$).

Acknowledgements

The authors wish to thank Ramon Lavado for helpful discussions for enzymatic assays. This project was supported by a grant to EAA (2011/52061-8) and graduate fellowships to AAF (2014/18825-9) from FAPESP and CAPES (3000-13-3) of the Brazilian Government. Funding was also provided to EAA through CNPq (307603/2014-8) and to DS through CNPq (401884/2012-0) and the UCR, Agricultural Experiment Station Resource Allocation Support Program.

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