**INTRODUCTION**

Cytochrome P4501A (CYP1A), a drug metabolizing enzyme found in most vertebrates, has been known to be inducible by intrinsic and extrinsic factors. CYP1A reacts specifically to environmental pollutants including polycyclic aromatic hydrocarbons (PAHs) to solubilize them to excrete from the body, and some metabolic intermediates of the environmental pollutants have been proven to be carcinogenic. PAHs induce CYP1A gene after consecutive binding to cytosolic aryl hydrocarbon receptor (AhR) and AhR nuclear translocator (ARNT) in nuclear. On the other hand, vitellogenin (Vg) which is utilized as a nutriment during embryonic period of an oviparous animal is induced in the liver by estrogen (E2) and endocrine disruptors. Therefore the gene expression of CYP1A and Vg have been used as a biomarker for aquatic pollutants (Goksoyr, 1995; Marin & Matozzo, 2004; Sarkar et al., 2006).

**Effect of TBT and PAHs on CYP1A, AhR and Vitellogenin Gene Expression in the Japanese Eel, Anguilla japonica**

**Min Seop Choi¹, Se Ryun Kwon¹, Seong Hee Choi² and Hyuk Chu Kwon¹,⁷**

¹Dept. of Aquatic Life Medical Science, Sun Moon University, Asan 336-708, Korea
²Dept. of Food Science, Sun Moon University, Asan 336-708, Korea

**ABSTRACT**: Gene expressions of cytochrome P4501A (CYP1A), aryl hydrocarbon receptor (AhR) and vitellogenin (Vg) by endocrine disruptors, benzo[a]pyrene (B[a]P) and tributyltin (TBT) were examined in cultured eel hepatocytes which were isolated from eels treated previously with B[a]P (10 mg/kg) or estradiol-17β (20 mg/kg) in vivo, and the relationship between CYP1A, AhR and Vg genes were studied. When the cultured eel hepatocytes were treated with B[a]P (10⁻⁶–10⁻⁵ M) the gene expressions of CYP1A and AhR were enhanced in a concentration-dependent manner. However, when treated with TBT (10⁻⁹–10⁻⁵ M) the gene expressions of CYP1A and AhR were suppressed at high concentrations (10⁻⁶–10⁻⁵ M), while having no effects at low concentrations (10⁻⁹–10⁻⁷ M). Gene expression of Vg was also suppressed by TBT in a concentration-dependent manner in cultured eel hepatocytes which was previously treated in vivo with estradiol-17β.

**Key words**: Eel hepatocytes, B[a]P, TBT, CYP1A, AhR, Vitellogenin
expression of estrogen receptor (ER), Vg and zona radiata of Atlantic salmon (Mortensen & Arukwe, 2007). In the same study the expression of AhR, ARNT and AhR repressor (AhRR) was different as the TBT concentration. In addition, decrease in CYP1A activity was reported in channel catfish and Atlantic salmon (Rice & Roszell, 1998; Mortensen & Arukwe, 2007).

Although numerous studies on the effects of environmental pollutants on the activities of CYP1A and its isomers have been reported, researches on the effects of TBT, one of the most dangerous aquatic pollutants, on the expression of CYP1A, AhR and Vg genes have been limited to a few fish species. Therefore, in the present study the expressions of CYP1A, AhR and Vg genes as affected by TBT treatment were investigated using cultured eel hepatocytes.

**MATERIALS AND METHODS**

1. Animals
Eels (*Anguilla japonica*) weighing 250–300 g were purchased from a local hatchery and kept in plastic tank containing 25°C fresh water. They were not fed during experimental period, and immaturity of genital glands was confirmed by dissection at the end of experiment.

2. Chemicals
Benzo[a]pyrene, estradiol-17β and tributyltin chloride (TBT) were purchased from Sigma Chemical Co (St Louis, MO USA). They were dissolved in DMSO before injection to culture media.

3. Induction of CYP1A and vitellogenin in vivo
For *in vivo* induction of CYP1A and vitellogenin eels were given single intraperitoneal injections of Benzo[a]pyrene (10 mg/kg BW) or estradiol-17β (10 mg/kg BW) dissolved in DMSO. Fish were sampled after 2 days the injection and hepatocytes were isolated as described below.

4. Hepatocyte preparation and culture
Hepatocytes were prepared according to the procedure of Kwon & Mugiya (1994). Eels were anesthetized with 0.1% 2-penoxethanol. The liver was exposed and perfused *in situ* with Ca²⁺-free Ringer solution (120 mM NaCl, 4.7 mM KCl, 1.25 mM KH₂PO₄ and 23 mM NaHCO₃, pH 7.4) *via* the hepatic portal vein for 10 min. The perfusion was followed with the Ringer sloutin containing collagenase (0.35 mg/ml, Type IV, Sigma) at room temperature (ca 25°C) for 25 min. The liver was finally perfused with 30 ml of Ca²⁺-and Mg²⁺-free Ringer solution containing 2 mM EDTA. After perfusion, the liver was freed from the gall bladder and transferred to a beaker containing the Ca²⁺-free Ringer solution (30 ml). It was minced with scissors and pipetted gently. The cell suspension was filtered with a nylon gauze into a 50 ml conical tube and the filtrate was resuspended with Ca²⁺-free Ringer solution and washed three times by centrifugation. Then, the cell precipitate was resuspended in a small volume of the Ringer solution. Cell yield and viability were determined by the Trypan Blue exclusion test. Cells were plated into a 60 mm Petri dish (Falcon) at a density of 3×10⁵ cells/dish. L-15 medium (Sigma) containing 0.2 μM bovine insulin (Sigma), streptomycin (100 μg/ml), and penicillin (70 μg/ml) was used for cell culture. All incubations were carried out in 3 ml of the medium at 25°C under 5% CO₂. The media containing the chemicals were changed every day. All incubations were replicated at least twice.

5. RNA isolation and RT-PCR
Hepatocytes were homogenized with 0.5 ml Trizol reagent (Invitrogen Inc.) on ice and incubated for 15 min at room temperature. After centrifugation (12,000×g, 20 min, 4°C) to remove cell debris, 140 μl of chloroform was added to supernatant. RNA from aqueous phase was precipitated by mixing with isopropanol and incubating samples on ice for 2 hr. After centrifugation (12,000×g, 15 min, 4°C), RNA pellet was washed with 75% ethanol.
and air dried. RNA dissolved in nuclease-free water was determined for concentration and purity using GeneQuant (Biochrom Ltd., Cambridge, England). RNA samples were stored at −80°C before use.

cDNA was synthesized from the isolated total RNA using Oligo(dT)$_{15}$ primer (Promega) and M-MLV reverse transcription (Promega). Primers targeting CYP1A, Vg, AhR and GAPDH genes (Choi, 2012) were used in RT-PCR as shown in Table 1. Constructed cDNA was amplified with Go Taq Green Master Mix (Promega) and 0.2 μM of each PCR primer. PCR procedure included one cycle of 2 min at 95°C, 30 cycles of 40 sec at 95°C, 40 sec at 65°C, 40 sec at 72°C and a final extension step of 7 min at 72°C by using a thermal cycler (Thermocycler, Biometra). The PCR product was separated on an 1% agarose gel (in TBE buffer, 30 g/ml ethidium bromide solution) and visualized using Image Analysis System (Kodak).

6. Statistical analysis

Data were analyzed by student t-test using SPSS version 18.0 for window to compare mean values of experimental group with control (vehicle) (P<0.05).

RESULTS AND DISCUSSION

Changes in gene expressions of CYP1A and AhR by B[a]P (10$^{-6}$ – 10$^{-5}$ M) and TBT (10$^{-9}$ – 10$^{-5}$ M) was investigated using RT-PCR in cultured eel hepatocytes which was treated previously in vivo with B[a]P to induce CYP1A and AhR genes. As shown on Fig. 1, the expression of CYP1A gene by B[a]P was increased in a concentration-dependent manner as compared with DMSO control (p<0.05). Gene expression of CYP1A by B[a]P, which has been reported in olive flounder (Paralichthys olivaceus) and rainbow trout (Oncorhynchus mykiss) (Malmström et al., 2004), was used as a control for studies on changes in CYP1A expression by TBT. When eel hepatocytes were treated with TBT (10$^{-9}$ – 10$^{-5}$ M), decrease in CYP1A was observed at high concentration of TBT (10$^{-6}$ M and 10$^{-5}$ M) while showing no changes at low concentration (10$^{-9}$ – 10$^{-7}$ M) (p<0.05). Changes in the expression of CYP1A by B[a]P and

![Fig. 1. Effects of benzo[a]pyrene (B[a]P) and tributyltin (TBT) on cytochrome p4501A (CYP1A) gene expression in cultured eel hepatocytes.](image-url)

Fig. 1. Effects of benzo[a]pyrene (B[a]P) and tributyltin (TBT) on cytochrome p4501A (CYP1A) gene expression in cultured eel hepatocytes. Hepatocytes which were isolated from eels treated previously with B[a]P (10 mg/kg) in vivo were cultured for 2 days with B[a]P or TBT. Differences in the mRNA level after each treatment were estimated by RT-PCR, and visualized on 1% agarose gels (A). The optical density of each band was quantified in a Bio Image System (Kodak) and normalized to the GAPDH (B). Lanes 1: vehicle (DMSO), 2: B[a]P 10$^{-5}$ M, 3: B[a]P 10$^{-6}$ M, 4: TBT 10$^{-5}$ M, 5: TBT 10$^{-6}$ M, 6: TBT 10$^{-7}$ M, 7: TBT 10$^{-8}$ M, 8: TBT 10$^{-9}$ M. Values are mean±SD (n=3). Asterisk (*) indicates significant difference with respect to the vehicle (control) (p<0.05).
TBT were not observed to be different between 24 hr and 48 hr of incubation (data not shown). These results indicate TBT to act as a suppressor to CYP1A expression in cultured eel hepatocytes. In an *in vivo* study with Atlantic salmon (*Salmo salar*) the expression of CYP1A was suppressed with high concentration of TBT and stimulated with low concentration (Mortensen & Arukwe, 2007). Meanwhile, in the present study with eel hepatocytes the expression of CYP1A was suppressed with high concentration of TBT as in Atlantic salmon, while showing no differences from control at low concentration.

Effects of B[a]P and TBT on gene expression of AhR are shown on Fig. 2. Gene expression of AhR was increased with B[a]P. As CYP1A has been known to be induced by xenobiotics such as PAHs including B[a]P which binds to the AhR and thereby induce the CYP1A (Hahn, 1998), in the present study with eel hepatocytes the gene expressions of both CYP1A and AhR were shown to be increased by B[a]P. The gene expression of AhR was significantly suppressed (p<0.05) by TBT at 10^{-6} M and 10^{-5} M, but at 10^{-9}–10^{-7} M it did not show any differences from control group. The expression of AhR at 24 hr was not different from 48 hr of incubation (data not shown). In an *in vivo* study with Atlantic salmon, although hepatic mRNA expression of AhR was not significantly changed by treatment of TBT, it was higher in fishes treated with low concentration of TBT (50 ul/L) than in fishes treated with high concentration of TBT (250 ul/L) (Mortensen & Arukwe, 2007). The results on the control of AhR gene expression by TBT in the present study with eel hepatocytes are not in accord with the *in vivo* study with Atlantic salmon. Further studies in other fish species are needed to prove whether the discordance was caused by *in vivo* and *in vitro* difference or species difference.

Effects of TBT on vitellogenin (Vg) gene expression were studied in cultured eel hepatocytes. Hepatocytes which were isolated from eels treated previously with B[a]P (10 mg/kg) *in vivo* were cultured for 2 days with B[a]P or TBT. Differences in the mRNA level after each treatment were estimated by RT-PCR, and visualized on 1% agarose gels (A). The optical density of each band was quantified in a Bio Image System (Kodak) and normalized to the GAPDH (B). Lanes 1: vehicle (DMSO), 2: B[a]P 10^{-5} M, 3: B[a]P 10^{-6} M, 4: TBT 10^{-5} M, 5: TBT 10^{-6} M, 6: TBT 10^{-7} M, 7: TBT 10^{-8} M, 8: TBT 10^{-9} M. Values are mean±SD (n=3). Asterisk (*) indicates significant difference with respect to the vehicle (control) (p<0.05).

Estradiol-17β (E_{2}) *in vivo* were cultured for 4 days with TBT. Differences in the Vg mRNA level after TBT treatment were estimated by RT-PCR (Fig. 3). At both 2 and 4 days of culture the expression of Vg mRNA in hepatocytes treated with 10^{-7}–10^{-5} M TBT was significantly decreased compared with control cells, while when treated with low concentration (10^{-9}–10^{-8} M) no difference in Vg expression was observed. Endocrine disruptors acting like E2 have been known to increase Vg gene expression (Jung et al., 2012). However, despite the TBT is classified as an endocrine disruptor it suppressed the expression of Vg gene in the present study. The suppression of Vg gene expression by TBT in this study is coincide well with the *in vivo* result.
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Fig. 3. Effects of tributyltin (TBT) on vitellogenin (Vg) gene expression in cultured eel hepatocytes. Hepatocytes which were isolated from eels treated previously with estradiol-17β (20 mg/kg) in vivo were cultured for 2 days with TBT. Differences in the mRNA level after TBT treatment were estimated by RT-PCR, and visualized on 1% agarose gels (A). The optical density of each band was quantified in a Bio Image System (Kodak) and normalized to the GAPDH (B). Lanes 1: vehicle (DMSO), 2-6: TBT 10^{-5} M – 10^{-7} M. Values are mean±SD (n=3). Asterisk (*) indicates significant difference with respect to the vehicle (control) (p<0.05).

with Atlantic salmon (Mortensen & Arukwe, 2007). Although Vg gene has been known to be expressed through activation of estrogen receptor (ER) in the presence of E2, activation of ER with TBT has been shown to be different between ER isomers. The expression of ERα was elevated, while ERβ decreased (Mortensen & Arukwe, 2007). Therefore, whether the suppression of Vg expression with TBT was through ER is not clear. As the present study with eel hepatocytes did not prove the point, more indept studies on the expression of Vg and ER are needed.

In summary, the gene expressions of both AhR and CYP1A were shown to be suppressed by TBT, and the suppression of CYP1A was thought to be through AhR. Expression of Vg gene was also observed to be suppressed by TBT. Therefore, as in the present results, the expressions of CYP1A, AhR and Vg in eel hepatocytes would be suggested to be valuable tools for detecting and studying endocrine disruptors and environmental pollutants such as PAHs, PHHs and PCB.

ACKNOWLEDGEMENT

This work was supported by the Sun Moon University Research Grant of 2010.

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(Received 4 September 2012, Received in revised form 27 September 2012, Accepted 28 September 2012)