

LARGE INDUCTION OF TYPE III DEIODINASE (D3) EXPRESSION AFTER PARTIAL HEPATECTOMY IN THE REGENERATING MOUSE AND RAT LIVER.

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Abstract

The deiodinases type 1 (D1) and type 2 (D2) catalyze the activation of T4 to T3, whereas type 3 deiodinase (D3) catalyzes the inactivation of T3 and T4. D3 plays a key role in controlling thyroid hormone bioavailability. It is highly expressed during fetal development, but also in other processes with increased cell proliferation, e.g. in vascular tumors. Since tissue regeneration is dependent on cellular proliferation and is associated with activation of fetal genes, we evaluated deiodinase activities and mRNA expression in rat and mouse liver as well as the local and systemic thyroid hormone status after partial hepatectomy (PH).

We observed that in rats, D3 activity was increased 10-fold at 20 h and 3-fold at 48 h after PH; D3 mRNA expression was increased 3-fold at 20 h. The increase in D3 expression was associated with maximum 2 to 3-fold decreases of serum and liver T3 and T4 levels at 20 to 24 h after PH. In mice, D3 activity was increased 5-fold at 12 h, 8-fold at 24 h, 40-fold at 36 h, 15-fold at 48 h; and 7-fold at 72 h after PH. In correlation with this, D3 mRNA was highest (6-fold increase) and serum T3 and T4 were lowest at 36 h. Furthermore, as a measure for cell proliferation, BrdU incorporation peaked at 20 to 24 h after PH in rats and at 36 h in mice. No significant effect on D1 activity or mRNA expression was found after PH. D2 activity was always undetectable.

In conclusion, we found a large induction of hepatic D3 expression after PH which was correlated with an increased cellular proliferation and decreased serum and liver T3 and T4 levels. Our data suggest that D3 is important in the modulation of thyroid hormone levels in the regenerating liver, in which a decrease in cellular T3 permits an increase in proliferation.

Introduction

A strict regulation of thyroid hormone levels is essential for normal development of different tissues, as well as for the metabolic function of these tissues. The bioavailability of thyroid hormone is regulated by three iodothyronine deiodinases (D1, D2 and D3). D1 and D2 are thyroid hormone activating enzymes that catalyze the conversion of the prohormone thyroxine (T4) to the active hormone 3,3',5-triiodothyronine (T3). D1 is expressed in liver, kidney, thyroid and pituitary, and plays an important role in the production of serum T3 (1). D2 is mainly expressed in brain, pituitary and brown adipose tissue. In tissues such as brain, this enzyme is important for the local activation of T3 (1). D3 is the major inactivating pathway, catalyzing the degradation of T3 to 3,3'-diiodothyronine (T2) and the conversion of T4 to the inactive metabolite reverse T3 (rT3). It is abundantly present in fetal tissues such as liver and brain, and in placenta, pregnant uterus and umbilical arteries and vein (1, 2). The high expression in the fetal compartment suggests that D3 plays an essential role in the regulation of fetal T3 levels, as a mechanism to protect the fetus from excessive exposure to active thyroid hormone. For instance, in the fetal brain, D3 expression is highly regulated in different regions, protecting the brain from excessive T3 until differentiation is required (3).

In contrast to the high D3 activities in the fetus, its activities are normally undetectable in most adult tissues. This is probably why D3 has long been neglected in studies on the regulation of thyroid hormone bioavailability in the adult. However, recent studies have revealed the reexpression of D3 in different pathophysiological conditions, among which cancer, cardiac hypertrophy, myocardial infarction, chronic inflammation and critical illness (4-9). Just like fetal growth, most of these are processes in which proliferation or cell growth take place.

It is generally believed that T3 stimulates the differentiation of cells and decreases their proliferation. Hernandez et al. recently showed that differentiation of precursor cells to adipocytes was associated with a decrease in D3 expression (10), and Dentice et al. demonstrated

that D3 is increased in proliferating keratinocytes (11). Also these findings confirm the concept that in general T3 stimulates cellular differentiation, and suggest a role for D3-mediated low intracellular T3 concentrations in cellular proliferation.

Liver regeneration is also a process dependent on cellular proliferation. Following hepatectomy of 70% of the liver, liver cells switch from a quiescent state to a proliferative state and re-enter the cell cycle (12). In rodents, during the first 4 hours after partial hepatectomy (PH), quiescent hepatocytes enter the cell cycle (G1 phase). This proliferative phase is rather short in rats, but is prolonged in mice, the peak of DNA synthesis being 40-44 h after PH (12, 13). During these processes, many fetal genes, which are not expressed in normal adult liver, are reactivated (12, 14). In the present study we investigated the possible reexpression of D3 in regenerating liver, by determining deiodinase activities and mRNA expression in rat and mouse liver after PH.

Materials and Methods

Materials

[3'-¹²⁵I]T3 and [3',5'-¹²⁵I]T4 were obtained from GE Healthcare (Amersham, UK); and [3',5'-¹²⁵I]rT3 was prepared by radioiodination of 3,3'-T2 as described previously (15). Non-radioactive iodothyronines were purchased from Henning Berlin GmbH (Berlin, Germany); dithiothreitol (DTT) from ICN Biochemicals Inc. (Costa Mesa, CA). 5-Bromo-2'-deoxyuridine (BrdU) and real-time PCR probes and primers were obtained from Sigma (St. Louis, MO), and Eukitt mounting medium from O. Kindler GmbH & Co (Freiburg, Germany).

Partial hepatectomy model

Wistar rats were obtained from the Hubrecht Laboratory, Utrecht, The Netherlands. Mice on the FVB background were obtained from Harlan, Horst, The Netherlands. The animals were allowed free access to food and water. Mice and rats were anesthetized by buprenorphine and then underwent PH by removing 70% of the liver, or only opening of the abdominal skin in the case of sham-treated control animals. Serum and liver tissues were

collected at PH (t=0) and at different time periods after PH. The same region of the liver was isolated from sham-treated controls. All liver samples were immediately frozen in liquid nitrogen and stored at -80 C until further analysis; serum was stored at -20 C. All procedures performed on the animals were approved by the Committee on Animal Experiments of the Utrecht University (protocols 102648-1 and 102648-2).

Determination of BrdU incorporation

Two hours before ending the experiment animals received an intraperitoneal injection with BrdU in PBS, using 30 µg/gram body weight BrdU for mice or 50 µg/gram body weight BrdU for rats. After the animals were sacrificed liver tissues were fixated in 10% formalin and imbedded in paraffin. Slides were stained overnight with the primary mouse anti-BrdU antibody M0744, clone Bu20a (Dako, Heverlee, Belgium), diluted 1:50 for rats and 1:100 for mice. Subsequently, peroxidase-conjugated rabbit anti-mouse IgG was applied and immunoreactivity was detected using the DakoCytomation Envision+ System, HRP (Dako, Heverlee, Belgium) according to the manufacturer's guidelines. Slides were counterstained with Haematoxylin for 30 seconds and mounted in Eukitt mounting medium. Percentages of BrdU positive cells per slide were calculated after counting 1000 nuclei per animal.

Determination of T4 and T3 concentrations in serum and liver samples

Serum T4 and T3 were measured by Vitros ECI technology (Immunodiagnostic System, Ortho-Clinical Diagnostics, Beerse, Belgium). The iodothyronine levels were determined by highly sensitive and specific radioimmunoassays after extraction and purification of the iodothyronines from the liver, as previously described (16).

Deiodinase activity assays

Tissues were homogenized on ice in 10 volumes 0.1 M phosphate (pH 7.2), 2 mM EDTA, containing 1 mM DTT, using a Polytron (Kinematica, Lucerne, Switzerland). The tissue homogenates were stored at -80 C until further analysis. Protein concentrations were

determined using the method of Bradford (17), using BSA as standard. D1 activities were determined by incubation of 0.1 µM rT3 (including 200,000 cpm [3',5'-¹²⁵I]rT3) for 30 min at 37 C with 10 µg protein/ml tissue homogenate in 0.1 ml 0.1 M phosphate (pH 7.2), 2 mM EDTA, 10 mM DTT (PED10). D2 and D3 activities were determined by incubation of 1 nM [3',5'-¹²⁵I]T4 (200,000 cpm, D2) or 1 nM T3 (including 200,000 cpm [3'-¹²⁵I]T3, D3) for 60 min at 37 C with 5 mg protein/ml tissue homogenate in 0.1 M PED10. Reactions were stopped by the addition of 0.1 ml ice-cold methanol. After centrifugation, 0.1 ml supernatant was mixed with 0.1 ml 0.02 M ammonium acetate (pH 4.0), and 0.1 ml of the mixture was applied to a 4.6 x 250 mm Symmetry C18 column connected to an Alliance HPLC system (Waters, Etten-Leur, The Netherlands), and eluted with a gradient of acetonitrile in 0.02 M ammonium acetate (pH 4.0) at a flow of 1.2 ml/min. The proportion of acetonitrile was increased linearly from 30 to 44% in 10 min. The radioactivity in the eluate was determined using a Radiomatic A-500 flow scintillation detector (Packard, Meriden, CT).

RNA isolation and quantitative RT-PCR

RNA from rat liver was isolated from 25 mg liver tissue using the high pure RNA tissue isolation kit (Roche Diagnostics, Almere, The Netherlands) according to the manufacturer's guidelines. RNA from mouse liver was isolated using Trizol (Invitrogen, Breda, The Netherlands). Subsequently, 10 µg RNA was treated with 10 U DNase (Promega, Leiden, The Netherlands) for 30 min at 37 C, and purified using the RNeasy mini kit RNA cleanup protocol (Qiagen). 500 ng RNA was used for cDNA synthesis using TaqMan RT reagents (Roche Diagnostics). RNA samples were verified to be free from genomic DNA by performing negative control cDNA synthesis reactions of 500 ng total RNA using Taqman RT reagents without reverse transcriptase.

Quantitative real-time PCR was carried out using the ABI PRISM 7700 sequence detection system (Applied Biosystems, Nieuwerkerk aan den IJssel, The Netherlands). Reactions were performed in 25 µl Taqman universal PCR master mix (Roche), containing 20 ng cDNA. mRNA levels are expressed relative to the total

RNA content. Per reaction 200 nM of primers and probe were used for D3, D1 and for the housekeeping gene hypoxanthine-guanine phosphoribosyltransferase (HPRT). Table 1 shows the sequences of the different primers and probes. Real-time PCR reactions were done for 2 min at 50 C and for 10 min at 95 C, followed by 40 cycles of 15 sec at 95 C and for 1 min at 60 C. Cycle at threshold (Ct) values represent the cycle numbers at which probe-derived absorbance reaches the calculated threshold value. Δ Ct values represent the Ct values of the housekeeping gene minus the Ct values of the target gene. Data are expressed as $2^{\Delta Ct}$ or as $2^{\Delta Ct} * 1000$ (relative number of mRNA copies).

Statistics

Results are expressed as means \pm SEM. Significance of differences between means was tested using the Student's t test for unpaired observations of the SPSS 10.1 statistical package (SPSS, Inc., Chicago, IL). *P* values \leq 0.05 were considered significant.

Results

Effects of PH on liver deiodinase expression and BrdU incorporation

We first tested deiodinase activity in rat liver at different time periods after PH (t=0). Figure 1A shows that D3 activity was increased 10-fold at 20 h and 3-fold at 48 h after PH, whereas D1 tended to decrease. D2 activity was absent in all rat liver samples. Next, we determined rat liver deiodinase mRNA levels. D3 mRNA expression tended to increase, whereas D1 mRNA expression tended to decrease at 20 h (Fig. 1B).

We also studied the effects of PH on deiodinase activity and mRNA levels in mice. In this model, D3 activity was increased 5-fold at 12 h, 8-fold at 24 h, 40-fold at 36 h, 15-fold at 48 h, and 7-fold at 72 h after PH compared to t=0, whereas SHAM-treated animals show only a 4-fold increase at 48 h after SHAM surgery (Fig. 2A). D1 activity decreased 2-fold compared to t=0, but was not affected compared to SHAM-treated animals 48 h after SHAM-surgery (Fig. 2A). In all mouse livers D2 activity was absent. Figure 2B shows a 6-fold increase of D3 mRNA at 36 h after PH compared to t=0,

and no significant difference in D1 mRNA levels between time periods after PH.

Figure 3 depicts the levels of BrdU incorporation as a measure of cellular proliferation status at different time periods after PH. In correlation with D3 activity, the BrdU level peaked at 20 to 24 h in rats (data not shown), and at 36 h after PH in mice (Fig. 3).

Effects of PH on systemic and local thyroid hormone levels

Finally, we analyzed local and systemic thyroid hormone levels at PH and at different time periods thereafter. In rats, liver and serum T3 and T4 levels decreased markedly, with maximum 2 to 3-fold decreases at 20 to 24 h after PH (Figs. 4A,B). This correlated well with the peak in D3 activity at 20 h after PH in rats (Fig. 1A). In mice, serum T3 and T4 decreased until minimum levels at 36 h after PH (Fig. 5), coinciding with the peak in D3 activity and BrdU incorporation at 36 h after PH (Figs. 2A,3).

Discussion

D3 is highly expressed in tissues of the fetoplacental unit, and has therefore long been considered to be mainly important in the regulation of the fetal thyroid hormone status. By inactivating the active thyroid hormone T3 and the prohormone T4, D3 is thought to protect developing organs against excessive exposure to thyroid hormone (18). Although D3 is not expressed in most normal adult tissues, recent studies have demonstrated the reexpression of the enzyme in various disease states. For instance, D3 is induced in liver and skeletal muscle of critically ill patients (6), and high levels of D3 are found in vascular tumors (4, 5). Furthermore, in different animal models, D3 was found to be induced in cardiac hypertrophy, myocardial infarction and chronic inflammation (7-9). The finding that in some of these conditions the accelerated degradation of thyroid hormone causes low serum T3 (6) or even severe hypothyroidism (4, 5) indicates the clinical importance of D3 in the regulation of local and systemic thyroid hormone status.

Not only fetal growth but also the above-mentioned pathophysiological conditions are processes in which cellular growth or

proliferation take place. Therefore, we hypothesized that D3-mediated low intracellular T3 levels are needed for cellular proliferation. Interestingly, Hernandez et al. recently showed high D3 expression in proliferating preadipocytes from brown adipose tissue, and a marked decrease in D3 expression when the cells differentiate into mature adipocytes (10). Additionally, by knocking down D3 in basal cell carcinomas Dentice et al. recently proved that D3 promotes the proliferation of malignant keratinocytes by inhibiting the differentiating action of T3 (11). Liver regeneration is also dependent on cellular proliferation and involves induction of various fetal genes (14). Therefore, in the present study we investigated deiodinase expression and local and systemic thyroid hormone status in rat and mouse liver after PH. We found a 10 to 40-fold induction of hepatic D3 activity after PH which was associated with an increased cellular proliferation and decreased local and circulating T3 and T4 levels. Remarkable in this respect is the tendency to a decrease of D1 in rats at 48 h after PH and the 2-fold decrease of D1 in mice at 48 and 72 h after PH. Since D1 is positively regulated by T3 (1), this may indicate that the slight decrease in D1 is secondary to the decrease in the systemic T3 level.

It should be noted that D1 activity was 2-fold decreased and D3 activity was 4-fold increased in SHAM-treated mice compared to $t=0$. Therefore, although D1 activity after PH is 2-fold decreased compared to $t=0$, it is not significantly altered compared to SHAM-treated controls. However, when D3 activity after PH is compared to SHAM-treated controls, it remains highly induced after PH. The mechanisms involved in the decrease of D1 and the increase of D3 activity after SHAM-treatment are unknown. The combined upregulation of D3 and downregulation of D1 by the SHAM operation may be caused by illness (6) and/or inflammation (9). Also the possible influence of the used anesthetics on deiodinase expression should be considered, since it is well known that some anesthetics alter thyroid function. We used the opiate buprenorphine. Baumgartner et al. showed an acute downregulation of D2 and upregulation of D3 in rat brain after administration of the mu-opiate agonist etonitazene (19).

We measured tissue and serum T3 and T4 levels, and found a significant decrease of local and systemic thyroid hormone levels at 20 h after PH in rat and at 36 h after PH in mice, which correlated well with the increase in D3 at these time periods. Tien et al. also studied rT3 levels in mouse liver after PH, and reported on a significant increase in the rT3 level after PH compared to SHAM-treated controls at 36 h after PH, which was correlated to a significant decrease in the T3-regulated genes tyrosine aminotransferase (TAT) and basic transcription element binding protein (BTEB) (20). In agreement with our data, they found 50% decreased D1 at 36 h after PH compared to SHAM-treated controls. However, because of nearly undetectable D3 mRNA expression levels, they did not determine D3 activities (20).

Previous cell culture studies have shown that 12-*O*-tetradecanoylphorbol-13-acetate (TPA), growth factors such as EGF, FGF and TGF β , cyclic AMP, T3, and retinoids are able to induce D3 activity (21-28). However, the molecular mechanisms responsible for the reactivation of D3 in the regenerating liver is still poorly understood. In the proliferating skin hedgehog signalling is involved in the induction of Dio3 expression (11). However, gene expression microarray analyses of regenerating liver after PH do not indicate that the sonic hedgehog/gli protein pathway is involved in this process (29-32).

As for the TGF β /Smad signalling pathway, TGF β 1 expression does increase after PH, with a peak level 24-48 h after operation (33-35). However, this increase in TGF β is accompanied by a downregulation of the TGF β receptors and by an increase of the transcriptional repressors SnoN and Ski (33-35), and TGF β does not increase but suppresses hepatocyte proliferation through direct effects on the hepatocytes (36).

Another mechanism involved in the induction of D3 after PH might be cellular hypoxia. Interestingly, Simonides et al. recently showed that hypoxia induces D3 expression via a hypoxia inducible factor (HIF)-dependent pathway (37). Since PH is well recognized as a hypoxic-ischemic injury, and HIF-1 α is induced after PH (38), the HIF-dependent pathway may well contribute to the induction of D3 after PH at the transcriptional level.

Regulation of D3 expression is complex, because the *Dio3* gene is imprinted, with preferential expression from the paternal allele (28). The gene is located in an imprinted region on human chromosome 14, mouse chromosome 12 and rat chromosome 6. One of the other imprinted genes in this region is called *Dio3os*, and is transcribed from the opposite DNA strand compared to *Dio3* (28). Interestingly, Hernandez et al. demonstrated an inverse correlation of the paternally imprinted *Dio3* gene with the maternally imprinted *Dio3os* gene (10). This *Dio3os* gene partially overlaps the coding region of the *Dio3* gene, and may thus interfere with the translation of the Dio3 transcript.

In summary, we found a large induction of D3 expression in regenerating liver which was correlated with an increased cellular proliferation and decreased local and systemic T3 and T4 levels. Future research should elucidate the roles of hypoxia and *Dio3os* in the regulation of Dio3 transcription and translation. Our findings support the concept that D3-mediated low cellular T3 is needed for cellular proliferation. Future studies, e.g. using conditional D3 knock-out mice, should further unravel the role of D3 in this process.

Acknowledgements

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Legends

- Fig. 1. Effect of partial hepatectomy in rats on D1 and D3 activity (A) and mRNA level relative to HPRT (B). Reaction conditions for the D3 activity assay were 1 nM ^{125}I -labeled T3, 5 mg protein/ml tissue homogenate, and 1 h incubation. Reaction conditions for the D1 assay were 100 nM ^{125}I -labeled rT3, 10 μg protein/ml tissue homogenate, and 1 h incubation. Results are the means \pm SEM (n=2-4). Significance of differences is indicated as follows: ***: $P<0.001$ vs. control (0).
- Fig. 2. Effect of partial hepatectomy in mice on D1 and D3 activity (A) and mRNA level relative to HPRT (B). Reaction conditions for the D3 activity assay were 1 nM ^{125}I -labeled T3, 5 mg protein/ml tissue homogenate, and 1 h incubation. Reaction conditions for the D1 activity assay were 100 nM ^{125}I -labeled rT3, 10 μg protein/ml tissue homogenate, and 1 h incubation. Results are the means \pm SEM (n=3). Significance of differences is indicated as follows: ***: $P<0.001$ vs. control (0); **: $P<0.01$ vs. control (0); *: $P<0.05$ vs. control (0); ++: $P<0.01$ vs. control (48 SHAM); +: $P<0.05$ vs. control (48 SHAM).
- Fig. 3. Effect of partial hepatectomy in mice on BrdU incorporation in mice. Results are the means \pm SEM (n=3). Significance of differences is indicated as follows: ***: $P<0.001$ vs. control (0).
- Fig. 4. Effect of partial hepatectomy on liver T3 and T4 levels in rats (A) or on serum T3 and T4 levels in rats (B). Results are means \pm SEM (n=2-5). Significance of differences is indicated as follows: ***: $P<0.01$ vs. control; **: $P<0.01$ vs. control. (0); *: $P<0.05$ vs. control (0).
- Fig. 5. Effect of partial hepatectomy on serum T3 and T4 levels in mice. Results are the means \pm SEM (n=3-5). Significance of differences is indicated as follows: **: $P<0.01$ vs. control. (0); *: $P<0.05$ vs. control (0).

Table 1. Oligonucleotide primers used for real-time PCR.

Gene	Sense primer	Antisense primer	Probe
D3	5'-TTCATGGCGCGGATGAG-3'	5'-GATGATAAGGAAGTCAACGTCGC-3'	5'-FAM-TTCCAGCGCCTGGTCACCAAGTACC-TAMRA-3'
D1	5'-ATTGACCAGTTCAAGAGACTCGTAG-3'	5'-CCACGTTGTTCTTAAAAGCCCA-3'	5'-FAM-TCATTTACATTGAAGAAGCTCACGCCACAGA-3'
HPRT	rat 5'-TATCAGACTGAAGAGCTACTGTAATGACC-3' mouse 5'-TATCAGACTGAAGAGCTACTGTAATGATC-3'	5'-TTACCAGTGTCAATTATATCTTCAACAATC-3'	5'-FAM-TGAGAGATCATCTCCACCAATAACTTTTATGTCCC-TAMRA-3'

Fig. 1A

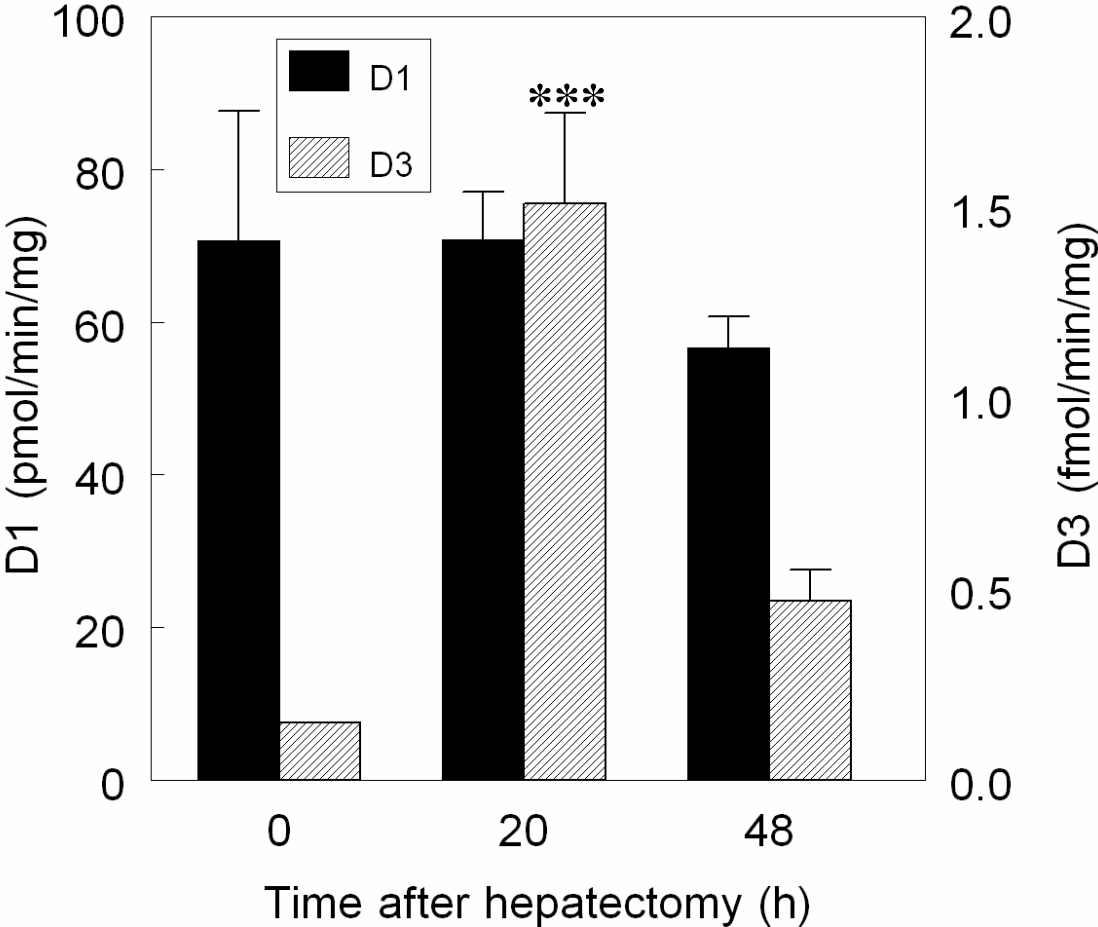


Fig. 1B

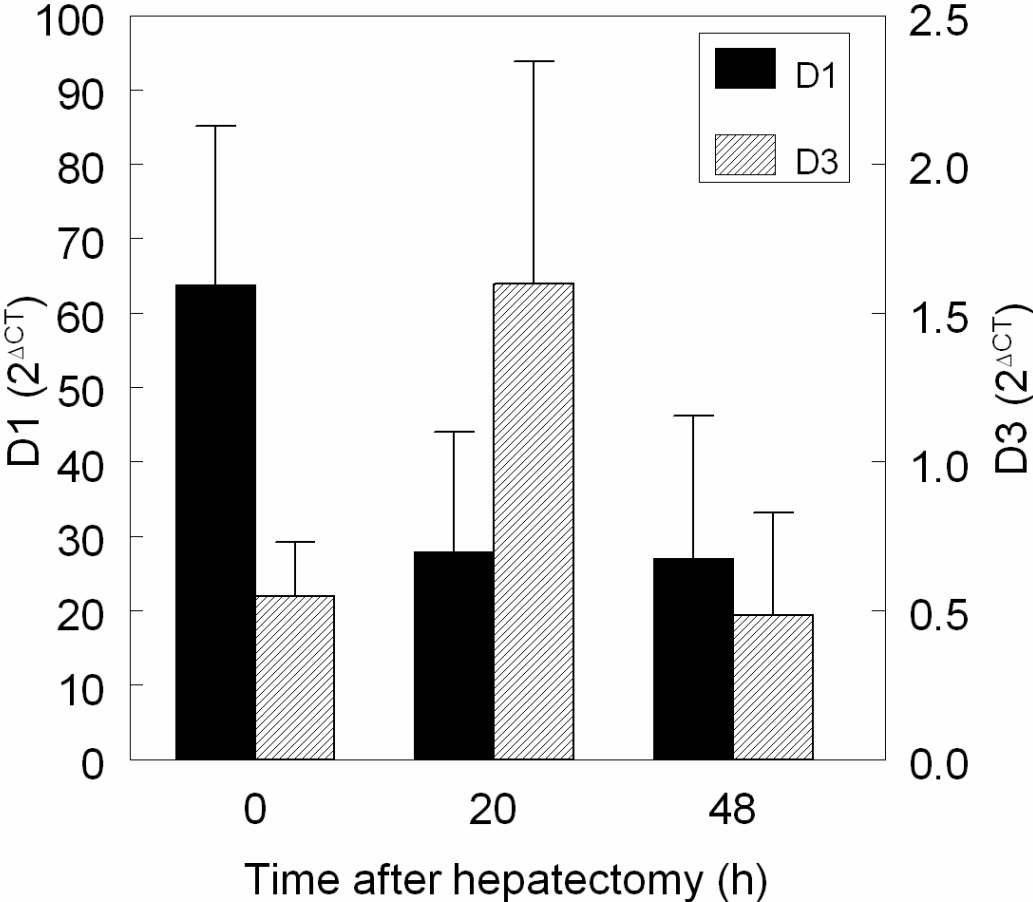


Fig. 2A

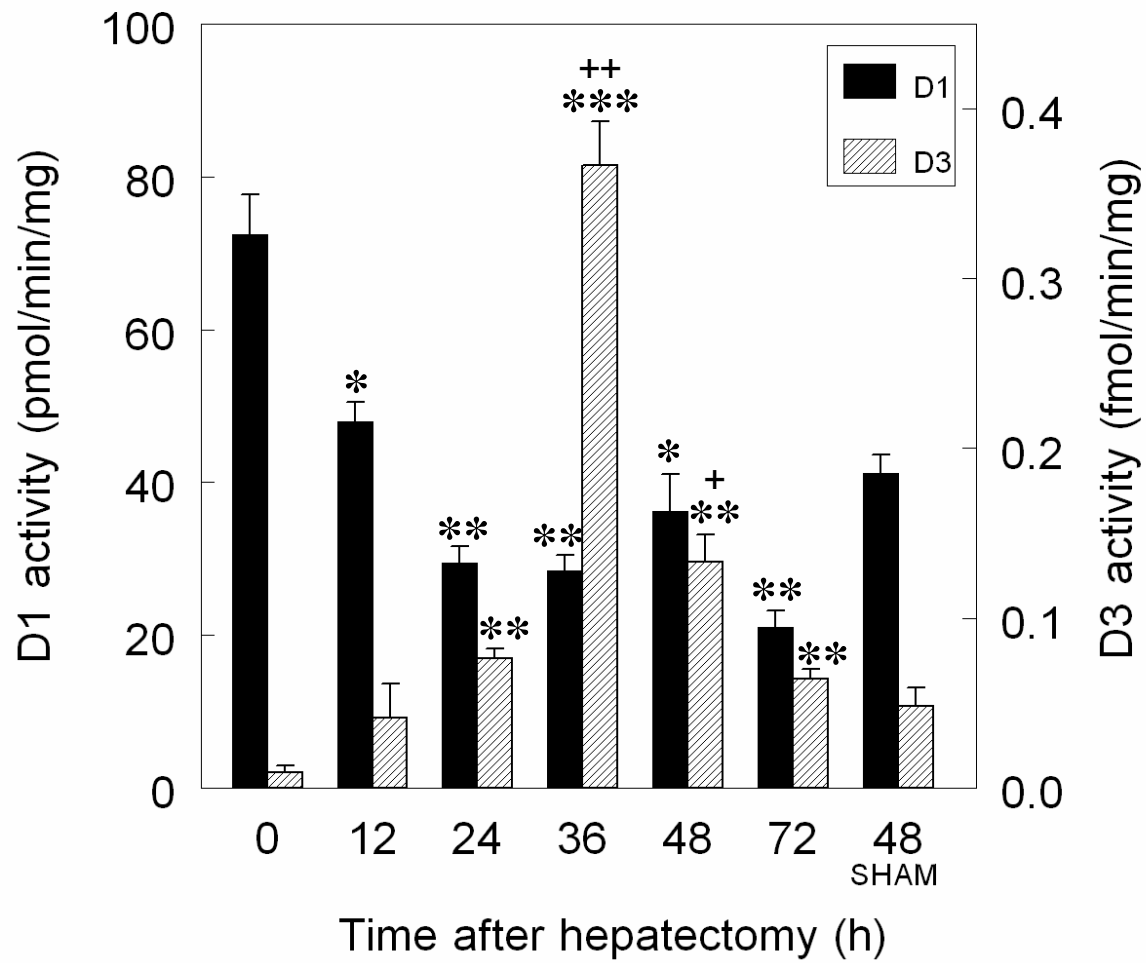


Fig. 2B

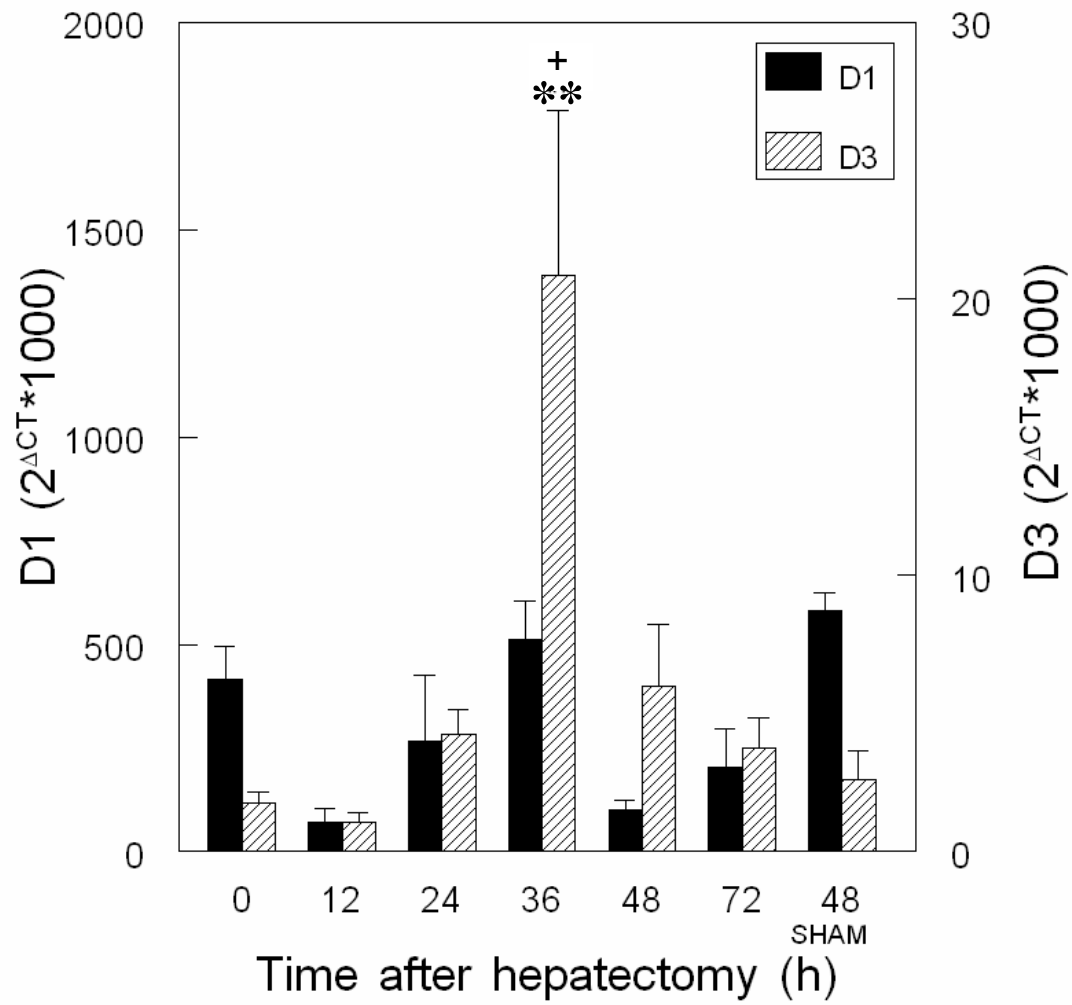


Fig. 3

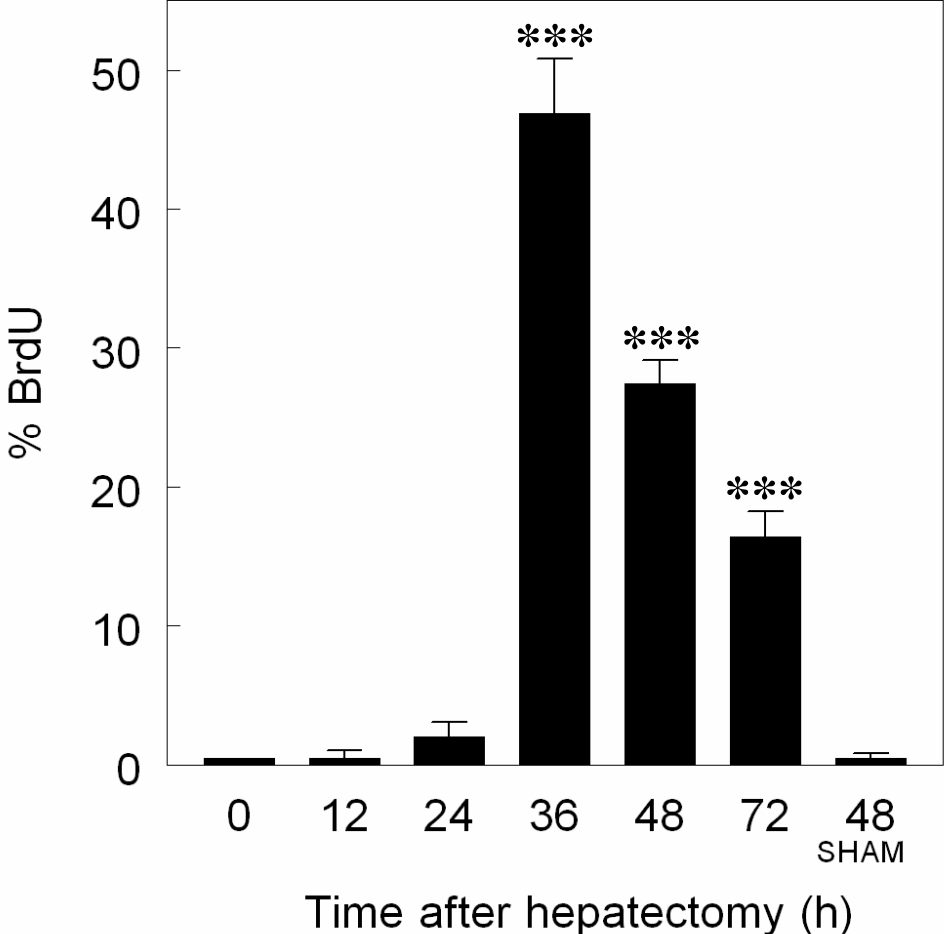


Fig. 4A

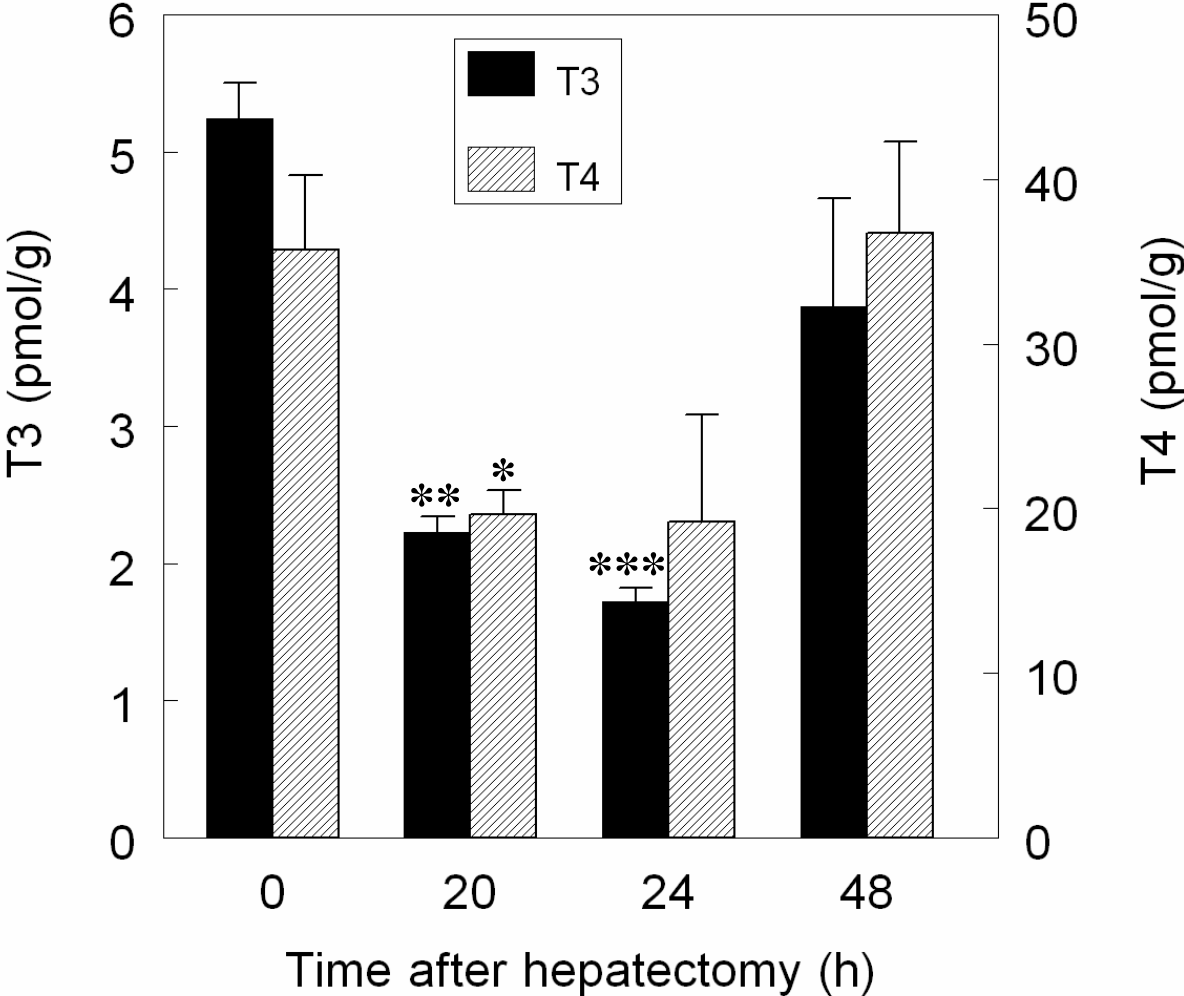


Fig. 4B

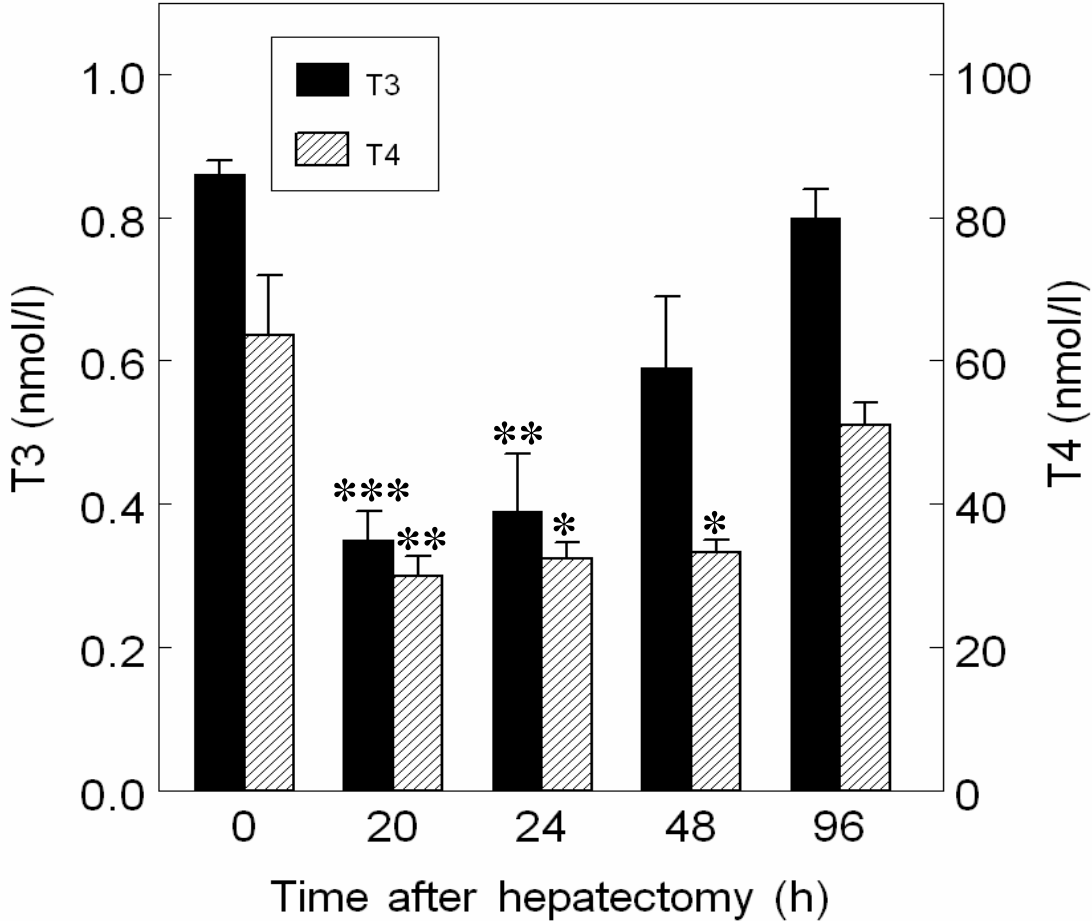


Fig. 5

