Hepatocyte Proliferation and Gene Expression Induced by Triiodothyronine In Vivo and In Vitro

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Subcutaneous injections of hormone triiodothyronine in rats resulted in peak blood levels at 24 hr with return to baseline by 96 hr. The injections stimulated a liver regeneration response that resembled in timing and in magnitude of DNA synthesis (peak, 24 hr) that induced by 40% hepatic resection. The principal proliferation was of hepatocytes. Although there were some temporal differences from the gene expression of transforming growth factor-α (TGF-α), transforming growth factor-β (TGF-β), and c-Ha-ras that are known to follow partial heptectomy, the overall profile of these changes was similar to those after partial resection. The effect was liver specific and could be reproduced three times with no diminution in response in the same animal with injections at 10-day intervals. No response was detected in kidney or intestine. This effect in intact animals contrasted with the minimal ability of triiodothyronine to stimulate hepatocytes in culture. However, when the culture medium was enriched with epidermal growth factor, there was a dose-related response to triiodothyronine. The totality of these experiments provides a preliminary basis for the creation with pharmacological techniques of an in vivo hyperplastic hepatic condition permissive of transcription of new genes, as an alternative to partial heptectomy. Although triiodothyronine was the test agent used, other hepatic growth factors singly or in combination could be candidates for this purpose. (HEPATOLOGY 1994;20:1237-1241.)

The influence of T3 on liver growth control has been demonstrated in rats by reduction in the normal hepatic regeneration response after thyroidectomy (1) and, alternatively, by a proliferative response seen after subcutaneous T3 injections in unaltered animals (2, 3). T3 also has modest hepatotrophic effects (4) that are identified in the Eck fistula screen model by the stimulation of proliferation and the prevention of hepatocyte atrophy (5). In this study, we further characterized the rat liver's response to T3 including the mRNA to T3 expression of the growth-associated genes TGF-α, TGF-β and c-Ha-ras. Our immediate objective was to determine whether the changes in gene expression were comparable to those known to occur after partial heptectomy (6, 7). For the transfection of new genes, a process that is known to be facilitated and prolonged under conditions of cellular proliferation (8), T3 may be an acceptable substitute for partial heptectomy.

MATERIALS AND METHODS

Animals. Male Fischer F344 rats weighing between 180 and 200 gm were obtained from Hilltop Lab Animals, Inc. (Scottsdale, PA). The animals were housed in a temperature- and light-controlled room (21° to 22° C, light from 6 AM to 6 PM) and received food and water ad libitum.

Materials. Type I collagenase (140 to 170 units/mg) was purchased from Worthington Diagnostic Systems (Freehold, NJ). Other materials and sources were formalin (Baxter Health Care Corp., McGaw Park, IL), Eagle's minimal essential medium and fetal calf serum (GIBCO Laboratories, Grand Island, NY), insulin, HEPES, pyruvic acid and T3 (Sigma Chemical Co., St. Louis, MO), EGF (Collaborative Research, Inc., Bedford, MA), [methyl-3H]thymidine (50 to 80 Ci/mmol) (Du Pont—New England Nuclear, Boston, MA), 35-mm tissue culture dishes (Corning, Inc., Corning, NY), gentamycin (Elns-Sinn, Inc., Cherry Hill, NJ), aqueous counting solution (Amersham, Arlington Heights, IL) and monoclonal rabbit Elite anti-human AFP antibodies (Dako Corp., Carpinteria, CA).

Probes Used. cDNA fragments of human c-Ha-ras (cDNA Probe, Oncogene Science, Inc., Cambridge, MA), human TGF-β1 (9) (gift of Dr. A. Derynk, Genentech, Inc., San Francisco, CA) and rat TGF-α (10) (gift of Dr. David C. Lee, University of North Carolina, Chapel Hill, NC) were labeled with [35P]dTCTP by means of random priming (Random Primer Labeling System; BRL, Indianapolis, IN). An oligo for 28S cDNA (Oligo Probe; Oncor, Inc. Gaithersburg, MD) was
labeled with $[^{32}P]dCTP$ by use of T4 kinase (end labeling system; BRL).

**In Vivo Experiments.** Injections (200 µg/100 gm rat body wt) of T3 were made subcutaneously on the backs of the animals in 0.1 ml of 0.01N NaOH–0.9% NaCl solution. Control rats were given similar injections of vehicle solution.

DNA synthesis was determined in the livers, kidneys and small intestines of rats injected intraperitoneally with 50 µCi $[^{3}H]$thymidine/200 gm body wt 1 hr before being killed; the incorporation of $[^{3}H]$thymidine was then determined as described previously (11, 12).

Proteins were determined by the method of Lowry et al. (13). Serum levels of T3 were determined by means of RIA (Ciba Corning MAGIC T3 commercial kit; Ciba Corning Diagnostic Labs, Burlingame, CA). Serum T3 and TGF-β were determined by use of RIA (Ciba Corning Diagnostic Labs, Burlingame, CA).

In Vitro Experiments. Hepatocytes from the livers of unaltered 7-wk-old male rats (180 to 200 gm) were isolated by means of a modification (16) of the in situ two-step collagenase perfusion technique (17, 18). Hepatocytes were dispersed and washed twice with cold Ca++-free perfusion buffer and resuspended in serum-free Williams E medium supplemented with EGF on fibronectin-coated dishes. Viability was determined by means of trypan blue exclusion, and only preparations with viability greater than 90% at the outset were used. Cell number was determined with a hemocytometer. The cells were plated at a cell density of 3 x $10^5$35 mm in Corning tissue-culture dishes containing 1.5 ml medium and maintained at 37° C in a 5% CO2 atmosphere. After a 3-hr attachment period, the medium was aspirated, and 1.5 ml of Williams E medium with or without EGF, 10 ng/ml, was added. T3 and TGF-β, when used, were at the concentrations reported in Table 1.

To determine *in vitro* DNA synthesis, we added 3 µCi $[^{3}H]$thymidine to each well and maintained it from 48 to 72 hr of the culture period. When the cells were harvested, DNA content was determined by use of the microfluorometric method of Setaro and Morley (19), and DNA synthesis was measured as we have previously reported (16).

**Table 1. Effects of different concentrations of T3 on hepatocyte proliferation in medium enriched with EGF and TGF-β**

<table>
<thead>
<tr>
<th>Medium additions (ng/ml)</th>
<th>$[^{3}H]$Thymidine incorporation (cpm x 10(^{-3}/\mu g) proteins)</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>102 ± 61*</td>
</tr>
<tr>
<td>T3 (1 ng)</td>
<td>124 ± 70</td>
</tr>
<tr>
<td>EGF + T3 (0.01 ng)</td>
<td>1,109 ± 98</td>
</tr>
<tr>
<td>EGF + T3 (0.1 ng)</td>
<td>1,480 ± 129*</td>
</tr>
<tr>
<td>EGF + T3 (1 ng)</td>
<td>1,920 ± 131*</td>
</tr>
<tr>
<td>EGF + T3 (10 ng)</td>
<td>1,811 ± 191*</td>
</tr>
<tr>
<td>EGF + T3 (1 ng) + TGF-β (1 ng)</td>
<td>304 ± 148</td>
</tr>
</tbody>
</table>

*Data expressed as mean ± S.D.

**Table 2. DNA synthesis in kidney and small intestine from control (vehicle) and T3-treated rats**

<table>
<thead>
<tr>
<th>Organ</th>
<th>Vehicle (24 hr)</th>
<th>T3 24 hr</th>
<th>T3 48 hr</th>
</tr>
</thead>
<tbody>
<tr>
<td>Kidney</td>
<td>2.6 ± 0.4*</td>
<td>2.5 ± 0.4</td>
<td>2.4 ± 0.3</td>
</tr>
<tr>
<td>Small intestine</td>
<td>13.2 ± 2.2</td>
<td>15.7 ± 0.8</td>
<td>13.7 ± 2.5</td>
</tr>
</tbody>
</table>

*Data expressed as mean ± S.D.

**Gene Expression Studies.** Total cellular RNA from the rat livers was extracted by use of RNAzol (Biotex, Houston, TX). Total RNA (20 µg) was subjected to electrophoresis in a 1% agarose gel in 10 mmol/L NaPO4 with constant recirculation. The fractionated RNA was transferred to a Zetabind nylon membrane (Whatman, Maidstone, UK) overnight in 20 × standard saline citrate. After transfer, the blot was fixed with UV light (short wave, 254 nm). cDNA probes were labeled with $[^{3}P]$ with a random-primed labeling kit. Prehybridization, $[^{3}P]$-labeled denatured probes in Church buffer were added at 70° C for hybridization (20). After hybridization, the membranes were washed twice in buffer A (1% BSA, 5% SDS, 0.5 mmol/L sodium phosphate, 1 mmol/L EDTA). For hybridization, $[^{3}P]$-labeled denatured probes in Church buffer were added at 70° C for hybridization (20). After hybridization, the membranes were washed twice in buffer A (1% BSA, 5% SDS, 0.5 mmol/L sodium phosphate and 1 mmol/L EDTA) at 70° C for 20 min and then washed in buffer B (1% SDS, 40 mmol/L sodium phosphate and 1 mmol/L EDTA) at 70° C for 20 min (four times).

The washed, air-dried membranes were subjected to autoradiography at −70° C. To check for equivalent transfer of RNAs from agarose gel to membrane, we stained the gel with ethidium bromide after capillary transfer: Little RNA was found to remain in any lane. Before probe hybridization, the membranes were checked under UV light; similar amounts of RNA were found to have been transferred to each lane. This was always confirmed with separate probing with cDNA for 28S RNA. The probes used were described. The products were subjected to electrophoresis on a 2% agarose gel with molecular-weight markers. The determination of TGF-α, TGF-β, and Ha-ras was chosen because the behavior of those genes has been completely established during the course of liver regeneration after partial hepatectomy (7, 9, 10).

**Statistical Analysis.** Data are reported as mean ± S.D. The Student one-tailed t test was used for statistical analysis of the data. Any p value less than 0.05 was considered significant.

**RESULTS**

**In Vivo Experiments.** The serum concentration of T3 rose from control of 0.7 ± 0.3 ng/ml to 38.9 ± 5.3 ng/ml after 24 hr and had returned to control values by 4 days. Repeat injections 10 and 20 days later caused 24-hr peak levels of 28.2 ± 6.5 and 46.3 ± 7.4 ng/ml, respectively, with return to control by 96 hr.

A peak response in DNA synthesis and mitoses was found at 24 hr followed by a lower second peak at 48 hr. Return to baseline required 96 hr (Fig. 1). When a second injection of T3 was given to the same animals 10 and 20 days after the first one, the same response was
obtained on all three occasions (Fig. 2). On histological and immunohistochemical examination, the mitotic figures were found predominantly in the AFP-negative hepatocytes.

[3H]thymidine incorporation in the kidneys and small intestine was not affected by T₃ administration (Table 2).

**In Vitro Experiments.** T₃ alone had a minimal stimulatory effect but caused a dose-related amplification of the DNA synthesis response to the primary hepatocyte mitogen EGF (Table 1).

**Gene Expression.** T₃ induced distinct peaks of TGF-α expression, the principal one at 24 hr (Fig. 3) with a subsequent return almost to baseline. The second peak of increased expression appeared at 72 hr after T₃ injection. The expression of c-Ha-ras was increased from 24 hr to 120 hr (Fig. 4). The peak expression of TGF-β was from 72 hr to 120 hr (Fig. 4). The mRNA of 28S, a nongrowth gene, was used as an internal standard in all experiments to which other RNAs could be compared (Figs. 3 and 4).

**DISCUSSION**

The simplicity of T₃ injection for induction of hepatic proliferation in vivo suggests that this method might be used to facilitate gene incorporation into the hepatocyte DNA under experimental conditions. Because partial hepatectomy has been most commonly used in intact
animals to provide the hyperplasia necessary for efficient and prolonged gene transfection (8, 21-23), comparisons of the two techniques are in order.

In most respects, the response to the T₃ doses used was similar to that following a moderate hepatic resection, with a peak response at 24 hr and return to baseline by 4 days. The secondary peak of DNA synthesis, at 48 hr, was not attributable to delayed ab-
sorption of the subcutaneously injected \( T_3 \), in view of the \( T_3 \) serum levels, which were maximal at day 1 and back to baseline by day 4 in association with proliferation of nonparenchymal cells, which are well known to have a delayed time of proliferation (24). A secondary small burst of DNA synthesis also has been seen after partial hepatectomy (25) and may represent in part a temporarily dissociated bile duct cell response (26-27). However, it was clear from the histopathological studies that the principal proliferation after \( T_3 \) was that of hepatocytes. Of considerable practical importance in the gene therapy context was the induction of undiminished proliferative response at 10-day intervals.

The changes in liver mRNA of hepatic growth-associated genes also were identical after \( T_3 \) injection to those known to occur after partial hepatectomy, including the expression of TGF-\( \alpha \) at 24 hr, TGF-\( \beta \) at 72 to 120 hr and c-Ha-ras from 24 to 120 hr. Minor deviations from the orderly sequence after partial hepatectomy included a delayed secondary peak of TGF-\( \alpha \) mRNA at 72 hr and more prolonged expression of c-Ha-ras.

These experiments support the feasibility of using a pharmacological approach as an alternative to hepatic resection for creation of a hyperplastic microenvironment conducive to gene insertion. For this purpose, \( T_3 \) is only one of several candidates—including hormones, growth factors and some of the most potent T-cell-directed immunosuppressive agents (4). The need to screen for such substances for their hepatic growth effect in intact animals as opposed to tissue culture conditions was emphasized before (4) and exemplified in the in vitro experiments of this report, in which \( T_3 \) alone was almost inert. The dose-related effect of \( T_3 \) was detected in tissue culture only when it was added to a medium enriched with EGF. This finding is a simple demonstration of the complexity of the process of liver regeneration in vivo, which is a multifactorial phenomenon that includes several growth factors, hormones and cellular matrix components, along with complicated cell-to-cell interactions.

REFERENCES