

previous results on rats (22). The difference in our findings may depend on the amount of DNA administered to each pig, or due to species specificity of the sialic receptor for sendai viral transfection. However, this gene transduction ratio may be sufficient for the treatment of certain liver diseases including inherited liver enzyme deficiencies, and the application of immunogene therapies. For example, 5% of the normal enzyme activity was reported to be sufficient for the treatment of Crigler Najjar syndrome (37). In addition, the repeated transfections proved to be possible probably because of the low immunogenicity of the HVJ-liposome vector system. Collectively, HVJ-liposome mediated gene transfer using the TVE technique of the liver may be one promising approach in the field of hepatic gene therapy.

The advantages of using HVJ-liposome vector is its low immunogenicity, and its ability to be used for repetitive transfections (29). Using this vector, Bcl2 transfection into the rat liver graft was able to inhibit apoptosis after ischemia reperfusion injury (38, 39), and a FasL transfected liver graft could eliminate the infiltrated cytotoxic lymphocytes efficiently (40). These previous results suggested that the HVJ-liposome vector mediated gene delivery system might therefore be a suitable treatment modality in the transplant setting.

In conclusion, our data shows the potential usefulness of HVJ-liposome as a gene transfer vector for liver directed-gene therapy in a pig model. This method may be applied to either liver transplantation or the treatment of the various liver diseases including inherited liver enzyme deficiencies.

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RESEARCH ARTICLE

# Eradication of hepatocellular carcinoma xenografts by radiolabelled, lipiodol-inducible gene therapy

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The promoter region of the early-growth response-1 (*Egr-1*) gene has been shown to be activated by external radiation, thus making a selective tumoricidal effect possible. A previous experiment showed that the *Egr-1* promoter can be activated by internal radiation using radioisotopes as well as external radiation. Internal radiation using I-131 lipiodol (I-131-Lip) has been established as one of the most useful therapeutic strategies against hepatoma. We herein linked the *Egr-1* promoter to the herpes simplex virus-thymidine kinase (HSV-TK) gene, and investigated its efficacy in hepatoma gene therapy in combination with I-131-Lip. A luciferase assay showed the *Egr-1*-promoter activity to be markedly increased in hepatoma tissue specimens in an I-131-dose-dependent manner, whereas a less than two-fold increase in this activity was observed in other organs. In addition, the radioactivity derived from I-131 was selectively accumulated in the tumor tissue specimens. To examine the efficacy of

*EgrTK/ganciclovir (GCV) gene therapy in vivo*, subcutaneous hepatoma xenografts in nude mice were transfected using a hemagglutinating virus of Japan (HVJ)-liposome vector. Complete tumor regression was observed in all the *EgrTK*-transfected tumors following combination treatment with I-131-Lip and GCV 42 days after treatment without any side effects ( $n=8$ ). In contrast, the tumors continued to grow in all control mice ( $n=10$ ). Furthermore, the serum  $\alpha$ -fetoprotein levels decreased in the combination therapy group, while they increased in the controls. In conclusion, these data indicate that *Egr-1* promoter-based gene therapy combined with internal radiation has a selective effect on hepatoma tumors while also showing an improved *in vivo* efficacy. This combination therapy might, therefore, be an effective human hepatoma gene therapy, even in advanced multiple cases. Gene Therapy (2005) 12, 1633–1639. doi:10.1038/sj.gt.3302531; published online 4 August 2005

**Keywords:** *Egr-1* promoter; hepatoma; HVJ-liposome; internal radiation; mouse model

## Introduction

Hepatocellular carcinoma (HCC) is a common human malignancy with an extremely poor prognosis.<sup>1,2</sup> Gene therapy represents a promising treatment for HCC. Among the potential suicide-gene therapies for cancer, the herpes simplex virus (HSV)-thymidine kinase (TK)/ganciclovir (GCV) system has been widely investigated.<sup>3–5</sup> HSV-TK converts the nontoxic agent GCV into a highly toxic phosphorylated GCV, which acts as a chain terminator of DNA synthesis and an inhibitor of DNA polymerase.<sup>6–8</sup> It is not yet possible to transduce therapeutic genes into all of the target tumor cells. However, one considerable advantage of the HSV-TK system is the 'bystander effect', in which HSV-TK-negative tumor cells are destroyed as a result of their

proximity to any HSV-TK-positive tumor cells that have been exposed to phosphorylated GCV.<sup>9</sup> The tumor-specific activation of the therapeutic gene is crucial in order to minimize the side effects of treatment, and suicide-gene therapy controlled by the  $\alpha$ -fetoprotein (AFP) promoter has proved satisfactory in this respect in both *in vitro* and *in vivo* models.<sup>10–14</sup> However, the level of AFP expression varies among cases, and nonmalignant tissue – particularly, chronic inflammatory tissue – can also express AFP.<sup>2,15</sup> In order to overcome the difficulties of tumor-specific activation, radiation-inducible promoters can provide spatial control of irradiation, and might therefore be useful in the treatment of hepatomas.<sup>16</sup>

We previously reported a novel technique for hepatoma-specific suicide-gene therapy, using a radio-inducible early-growth response-1 (*Egr-1*)-promoter containing six CAAT boxes corresponding to the radiosensitive and serum-responsive elements.<sup>17</sup> The *Egr-1* gene was preferentially expressed in hepatoma cells compared with the surrounding nontumor tissue, and its promoter was significantly activated after radiation exposure. However, this approach might not be suitable for multiple

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hepatomas, because whole-liver irradiation could be fatal to patients: firstly, as a result of direct injury from irradiation, which produces free radicals that cause DNA-strand breaks; and secondly, through the non-specific activation of suicide genes. Recently, the *Egr-1* promoter was shown to be activated not only by external radiation but also by internal radiation from isotopes such as Tc-99m, Ga-67 and I-131.<sup>18</sup> Importantly, I-131-labelled lipiodol (I-131-Lip) concentrates preferentially within hepatomas, where it is retained for significantly longer periods than in non-neoplastic tissues.<sup>19–22</sup> In addition, I-131 generates 606-MeV  $\beta$ -rays that produce a strong effect over short distances, which could therefore enhance the bystander effect through increased intracellular communication.<sup>23</sup>

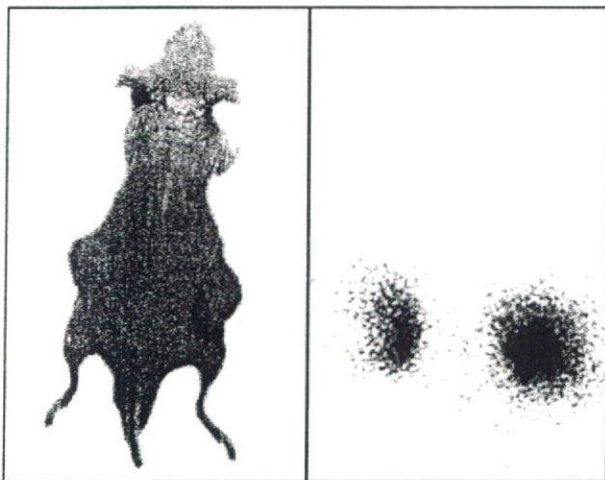
In the present study, we investigated whether combined therapy comprising transduction of the *EgrTK* gene and internal radiation with I-131 was able to reduce tumor development in HCC.

## Results

### Biodistribution of I-131-Lip in tumor-bearing mice analyzed using autoradiography after injection into the tumor xenograft

Autoradiography was carried out after I-131-Lip injection in order to examine whether I-131-Lip accumulated preferentially in the tumor. These mice did not receive iodide-uptake inhibitors such as NaI. As shown in Figure 1, I-131-Lip accumulated preferentially at the tumor site and remained there for up to 7 days, which is consistent with the observed physiological half-life of I-131. In contrast, no I-131-Lip accumulation was detected in any other organs, including the thyroid and gut. A densitometric analysis using Bio-imaging Analyzer System (BAS5000, Fuji) showed the diphasic accumulation of radioisotopes.

Softex roentgenograms revealed the iodide uptake, which was visualized as white dots, only in tumors treated with I-131-Lip. No accumulation in any other organs was observed (data not shown).



**Figure 1** Biodistribution of I-131-lipiodol. I-131 accumulation was detected in mice using autoradiography. I-131-lipiodol (I-131-Lip) accumulated preferentially in the tumor. The iodide uptake was not observed in other organs, such as the thyroid and the gut.

### *Egr-1*-promoter activity in vivo

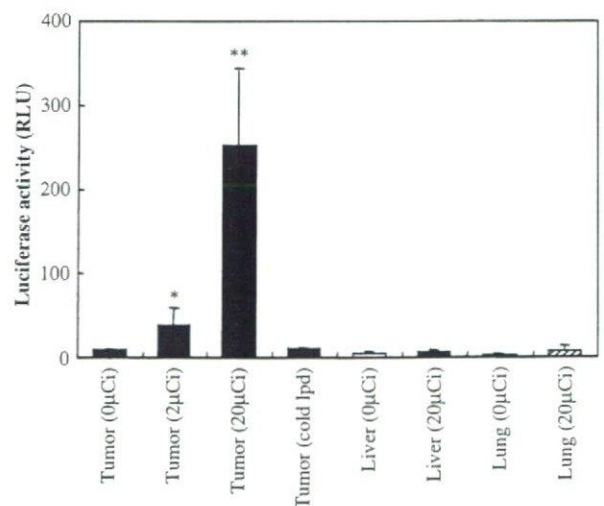
An *in vivo* luciferase assay was performed to examine whether the *Egr-1* promoter could be activated by I-131-Lip in tumor xenografts. *EgrLuc*-HVJ-liposome was introduced into tumors when they reached 8–10 mm in size, along with 0–20  $\mu$ Ci of I-131-Lip, which was injected at 16 h post-transfection. The tumor tissue specimens were excised and examined to determine their luciferase activity 40 h after transfection. Radiation-induced promoter activity was expressed as relative light units, after correction using nonirradiated cell lines. The differences in transfection efficiencies between tumors were negligible. Internal radiation therapy using I-131-Lip significantly increased the *Egr-1*-promoter activity in the hepatoma tissue specimens (4.4-fold at 2  $\mu$ Ci and 28.8-fold at 20  $\mu$ Ci), whereas no increase was observed in the normal liver and lung tissue specimens (Figure 2).

### *Egr-1* protein expression in tumor tissue after I-131-Lip treatment

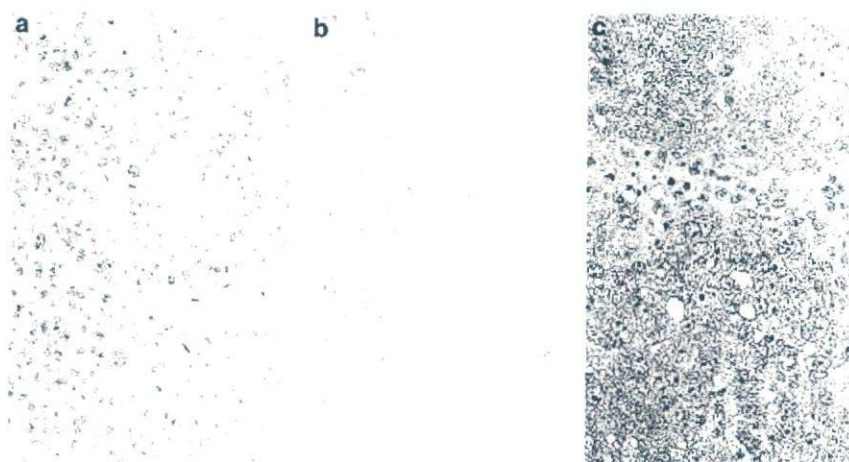
*Egr-1* immunohistochemistry was carried out in tumors after treatment in order to examine the level of *Egr-1* gene expression. Faint staining was detected in the tumor cell clusters of untreated hepatoma tissue (Figure 3a), but not in the absence of the primary antibody (Figure 3b). In contrast, a significant degree of *Egr-1* activation was observed in the tissue specimens treated with I-131-Lip (Figure 3c). These results clearly show that *Egr-1* gene expression can be induced by internal irradiation using I-131, as well as by external radiation exposure.

### *In vivo EgrTK/GCV treatment of tumor xenografts in mice*

The whole-body effective half-life of the therapeutic activity, which was measured daily using a radiation-survey meter, showed an exponential decrease with a mean of 8 days (data not shown). To further investigate



**Figure 2** *Egr-1*-promoter activity in vivo. *Egr-1*-promoter activity was significantly increased by internal irradiation to tumors that were transfected with the *Egr-Luc* gene. However, no increase in the activity was detected in either the liver or lung tissue specimens. \* $P < 0.05$ , \*\* $P < 0.01$  versus control groups.



**Figure 3** *Egr-1* immunohistochemistry of the tumor xenograft. Low levels of *Egr-1* gene expression were observed in nonirradiated hepatoma tissue (a), but not in the absence of the primary antibody (b). Internal irradiation greatly increased *Egr-1* gene expression in hepatoma tissue (c).

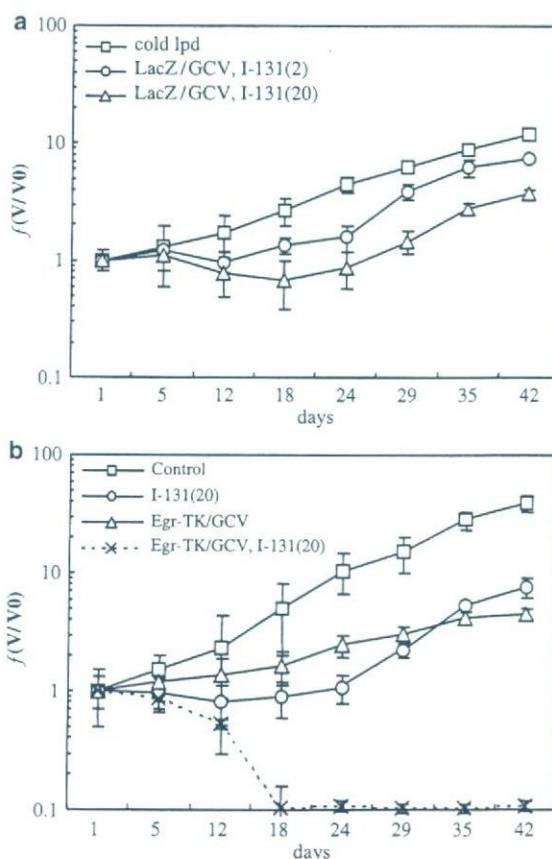
the cytotoxic effects of *EgrTK/GCV* *in vivo*, Huh7 cells ( $1 \times 10^7$ ) were injected subcutaneously into nude mice. Gene therapy was initiated when the tumor volume reached approximately  $200 \text{ mm}^3$ . No spontaneous tumor regression was observed in the untreated tumors during the course of treatment. The data for each treatment group were calculated as a percentage of the initial (day 0) volume, and were plotted as the mean fractional volume  $\pm$  standard error. I-131-Lip alone showed a minor dose-responsive growth-suppressive effect (Figure 4a). In contrast, the *EgrTK*-transfected animals ( $n=8$ ) that received irradiation without GCV exhibited no suppression of the tumor volume compared to the untreated animals ( $n=10$ ). A retardation of tumor growth was seen in the *EgrTK*-transfected group that received GCV without irradiation ( $n=8$ ). Complete tumor regression was observed in all the *EgrTK*-transfected tumors that received I-131-Lip and GCV from 24 days to 42 days after gene therapy (Figure 4b). Thereafter, seven of eight tumors continued to show regression at 80 days post gene therapy, although one remaining tumor slowly regrew.

#### Serum AFP measurement

AFP is commonly used as a diagnostic marker for HCC because of its high sensitivity, and human Huh7 cells produce human AFP. Therefore, in order to estimate the viable volume of implanted hepatoma tumors, the serum AFP levels were measured at the time points indicated in Figure 5. The serum AFP values progressively increased in the untreated controls, whereas a strong suppression was observed in the *EgrTK* plus I-131-Lip group, and these findings were consistent with those from the *in vivo* gene-therapy data.

#### Histological findings

Hematoxylin-eosin staining revealed massive apoptotic and partially necrotic cells with intratumoral lymphocyte-infiltration lesions in the *EgrTK/GCV* plus I-131-Lip group (Figure 6d) compared to the untreated tumor (Figure 6a). In contrast, slight tumor cell damage with necrosis, but without any inflammatory cell infiltration, was observed in the I-131-Lip-injected group (Figure 6c).



**Figure 4** *In vivo* gene-therapy experiments in mice with hepatoma tumor xenografts. The injection of cold lipiodol did not cause tumor retardation. However, slight transient retardation was observed in the I-131-treated group in a radiation dose-dependent manner (a). Minor growth retardation was also observed in the groups treated with *EgrTK/GCV* or I-131-Lip alone. In contrast, complete tumor retardation was observed in the group treated with *EgrTK/GCV* plus I-131-Lip (b).

*EgrTK/GCV* treatment resulted in an increased number of apoptotic tumor cells, but it did not cause any necrotic tumor tissue damage (Figure 6b).

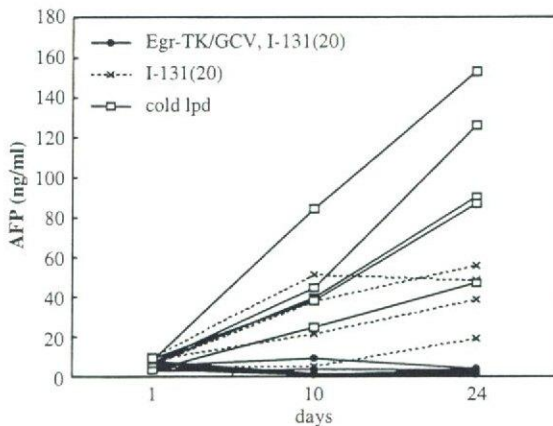
## Discussion

The lymphographic agent lipiodol (also known as iodized oil) has been shown to specifically localize and persist within hepatoma tissue.<sup>24–27</sup> Although lipiodol itself does not seem to have any significant anticancer effects, it allows anticancer drugs to be delivered selectively to tumor tissue when applied as an emulsion.<sup>28</sup> Lipiodol contains sterile iodine 127, so the labelling of lipiodol with radioactive I-131 can be achieved through a simple radioisotopic exchange method.<sup>29</sup> The therapeutic effectiveness is primarily limited to areas in which lipiodol accumulates, as I-131 generates 606-MeV  $\beta$ -rays that produce strong effects over distances of less than 2.0 mm.<sup>30</sup> These findings indicate that I-131-Lip has potential advantages for use

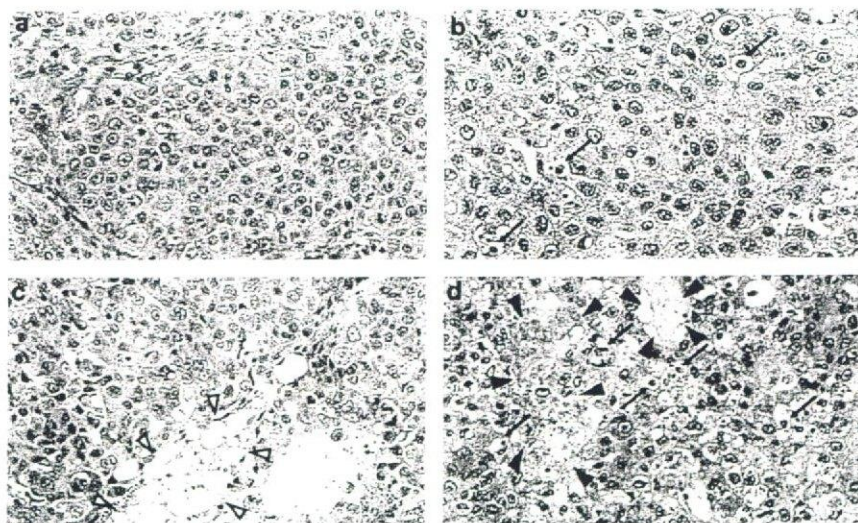
in hepatoma therapy.<sup>31–33</sup> The selective delivery and retention of lipiodol are thought to result from the developed neovasculature, enhanced permeability and the poor reticuloendothelial systems observed in hepatoma tissues. Despite this capacity for the selective accumulation at the tumor site, clinical studies have shown the effectiveness of such I-131-Lip therapy alone to only be temporary, and early HCC recurrence was observed in many cases. In contrast, recent reports have shown that I-131 therapy following a curative resection significantly improved both the disease-free interval and the survival rates.<sup>22,34</sup> Furthermore, no significant side effects – such as liver toxicity, renal disorders, bone-marrow suppression, thyroid dysfunction or respiratory dysfunction – have been reported after I-131-Lip administration.

We previously reported that a total dose of 20 Gy of external ionizing radiation could selectively and effectively activate the *Egr-1* promoter, thus resulting in tumor regression.<sup>17</sup> However, it is not possible to focus specifically on multiple tumor lesions, even when using conformal irradiation. Furthermore, high doses (50 Gy or above) of whole-liver irradiation are known to be toxic to normal livers,<sup>15</sup> and livers with hepatitis or cirrhosis are thought to be even more highly radiosensitive. These factors have clear implications regarding the treatment of HCC.

A recent report demonstrated that the *Egr-1* promoter can be activated by both external and internal radiation, which thus prompted our investigation of I-131-Lip as both an antitumor agent and a molecular switch for radio-inducible promoter *Egr-1*-based gene therapy. Our results revealed a 28.8-fold upregulation of the *Egr-1* promoter in tumors treated with 20  $\mu$ Ci of I-131-Lip compared to the controls, thus resulting in the complete regression of treated tumor xenografts. Immunohistochemical analyses also showed diffuse *Egr-1* activation in the HCC tissue. Clinical trials have used between 50 and 70 mCi of I-131-Lip administered via a tumor-supplying



**Figure 5** Measurements of serum AFP in tumor-bearing mice during treatment. The serum AFP values significantly decreased after treatment in the *EgrTK/GCV* plus I-131-Lip group, which was consistent with the data on tumor size. In addition, a slight decrease in serum AFP levels was observed in the group treated with I-131-Lip alone.



**Figure 6** Histological findings in tumor tissue after treatment. Hematoxylin-eosin staining revealed apoptotic shrinkage in tumor cells (arrows) in the *EgrTK/GCV* group after treatment (b) compared to the normal control (a). In addition, severe steatotic changes (open triangle) were observed in the cytoplasm of the tumor cells of the I-131-Lip group (c). Both apoptotic cells (arrow) and necrotic area (arrow heads) were found in the group treated with *EgrTK/GCV* plus I-131-Lip (d), which supported the growth-suppressive effect of the combination therapy reported in the *in vivo* gene-therapy experiments.

artery. The local radioactivity of the 20  $\mu$ Ci dose is therefore estimated to be 10 times lower than that used in clinical I-131-Lip therapy. The selective uptake of I-131-Lip into tumor cells was also confirmed by autoradiography.

Many different genes can be activated after irradiation, including *c-fos*, *c-jun*, *TNF- $\alpha$*  and *NF- $\kappa$ B*. However, the advantage of the *Egr-1* promoter is that it is tightly regulated by irradiation and its basal activity is relatively low in nonirradiated tumors. Indeed, our *in vivo* luciferase assay data showed the leakage of the transduced suicide gene to be extremely low, thus suggesting that I-131-Lip could efficiently regulate therapeutic gene expression. Although, we have not examined whether restimulation of the *Egr-1* promoter is possible after multiple treatments with internal radiation, our *in vivo* gene-therapy data suggested that *HSV-TK* gene expression might be prolonged. We have previously shown that the *Egr-1* promoter was relatively hepatoma tissue specific.<sup>17</sup> More interestingly, the *Egr-1* gene has recently been shown to be overexpressed in prostate cancer and to be closely associated with its carcinogenesis.<sup>35</sup>

Regarding the optimal method for administering I-131-Lip to patients, arterial chemoembolization is generally performed to treat multiple HCC. However, repeated embolization is limited by ischemic damage to normal livers and/or occlusion of the main feeding artery. Importantly, I-131-Lip therapy without embolization can be repeated in patients with hepatoma, thus making internal radiation-controlled suicide-gene therapy suitable for use even in regions of tumor recurrence.

Regarding future human clinical trials, a subcutaneous HCC implantation model may not be an ideal situation. If adequate animal models are developed, further studies will be necessary to establish an adequate treatment method for intrahepatic tumors with this strategy with an ultimate goal of future clinical application.

Up to now, adenoviruses have proven to be the most efficient gene-transfer vectors. However, the clinical findings from gene therapies using these vectors directed at the liver have also revealed fatal side effects that occur as a result of systemic inflammatory responses.<sup>36</sup> Safer gene-transfer methods, such as nonviral vectors, must therefore be considered for liver-related gene therapy. In the present study, we used an HVJ-liposome vector for the transfer of genes, which proved to be efficient, safe and simple to use *in vivo*. This vector was constructed from inactivated HVJ envelope proteins and anionic liposomes. It therefore had a low immunogenicity, which allowed for repeated administrations without any significant tissue damage.<sup>37-39</sup> The gene-transfer efficiency might thus be further improved by using other lipid formations or cointroduced proteins. Single applications of HVJ-liposome-mediated gene transfer are transient, as the transferred gene remains episomal in the nucleus, similar to adenoviruses and other physical methods. However, repeated gene transfer might result in a long-term gene expression. Moreover, previous studies have detected no obvious functional or histological liver damage after gene transfer using the HVJ-liposome vector.<sup>39</sup> These preliminary results suggest that the HVJ-liposome method has several advantages compared with other gene-delivery systems for the *in situ* transduction of cancer cells, even though the vector itself does not have any specificity for tumor cells.

In conclusion, radio-inducible suicide-gene therapy in combination with I-131-Lip treatment was found to enhance the efficacy of gene therapy for hepatomas. This approach might therefore represent a potentially effective treatment modality for clinically unresectable multiple HCC.

## Materials and methods

### Plasmid construction

EgrLuc and EgrTK plasmids were constructed according to the method reported previously.<sup>17</sup> Briefly, a region of the murine *Egr-1* promoter (-425 to +0) was amplified using the polymerase chain reaction with specific primers from the pE425 plasmid,<sup>40</sup> under the kind guidance of Dr DW Kufe of the Dana-Farber Cancer Institute, Harvard Medical School, USA. The polymerase chain reaction fragment was inserted upstream of firefly luciferase or the *HSV-TK* gene, producing the EgrLuc or EgrTK plasmid, respectively.

### Cell lines

The human hepatoma cell line Huh7, which was kindly provided by Dr H Nakabayashi of the Department of Biochemistry, Hokkaido University School of Medicine, Japan, was cultured in RPMI 1640 medium (Gibco, Long Island, NY, USA) supplemented with 5% fetal bovine serum, 100 U/ml penicillin and 100  $\mu$ g/ml streptomycin.

### Preparation of the HVJ-liposome vector

The HVJ-liposome vector was prepared according to the method described previously. Briefly, phosphatidylserine, phosphatidylcholine and cholesterol were mixed at a weight ratio of 1:4.8:2. The lipid mixture (10 mg) was dried by the removal of chloroform, and then hydrated in 200  $\mu$ l balanced salt solution containing DNA-high-mobility group I complex (200:64  $\mu$ g), which had been incubated previously at 20°C for 1 h. Liposomes were prepared by shaking and sonication. The liposome suspension (10 mg lipids) was mixed with HVJ (30 000 HAU) in a total volume of 1 ml balanced salt solution. The mixture was incubated at 4°C for 10 min, and then for 1 h with shaking at 37°C. Free HVJ was removed from the HVJ-liposome solution using sucrose density gradient centrifugation, and the second layer of sucrose containing the HVJ-liposome vector was collected.

### Labelling of lipiodol with I-131

I-131-labelled lipiodol, which was produced by replacing the iodine residue of lipiodol with I-131-C104 under aseptic conditions, was supplied by The First Radioisotope Co. Ltd (Tokyo, Japan). The labelling index was consistently more than 99% and the radioactivity of the I-131-labelled lipiodol was 131 MBq/ml.

### In vivo luciferase assay

To correct for transfection efficiencies, the cytomegalovirus-*renilla* (sea pansy) luciferase vector was cotransfected in each experiment. EgrLuc (196  $\mu$ g) and cytomegalovirus-Luc (4  $\mu$ g) plasmids were encapsulated in the HVJ-liposome and transfected into established tumors in nude mice. Based on the findings of previous reports, we administered 2-20  $\mu$ Ci of I-131-labelled

lipiodol to each animal in a total volume of 100  $\mu$ l, diluted with cold lipiodol 24 h after transfection.<sup>41</sup> Subsequently, 48 and 72 h after transfection, the tumors, livers and lungs of the animals were excised and immersed in 1200  $\mu$ l of  $3 \times$  reporter lysis buffer, and then were cut with sterile scissors. The mixture was transferred to a 15 ml falcon tube and homogenized using a sonicator for 5–10 min, followed by centrifugation at 3000 r.p.m. for 5 min. The supernatant was then transferred into a 1.5 ml Eppendorf tube. Aliquots were assayed for luciferase activity in a Lumat LB9501 luminometer (Berthold Systems, Aliquippa, PA, USA) using 20  $\mu$ l of supernatant and 100  $\mu$ l of reconstituted luciferase-assay reagent (Promega, Madison, WI, USA). The light units produced were measured for 20 s, 3–5 min after mixing. The reagents used in this step were obtained from the Dual-Luciferase<sup>®</sup> Reporter Assay System (Promega).

**In vivo imaging and quantitation of accumulated radionuclides in tumor xenografts in Balb/c nu/nu mice**  
Autoradiography was carried out in order to visualize I-131 accumulation in the tumors. The mice were placed under general anesthesia by the intraperitoneal injection of a sterile 2.5% solution of 2-2-2-tribromoethanol (Aldrich chemicals, Milwaukee, WI, USA). The anaesthetized animals were then placed on an imaging plate (Fuji, Tokyo, Japan) in a supine position in a dark room for 5 min, followed by a densitometric analysis using a Fujix Bioimaging Analyzer BAS2000 (Fuji).

An *in vivo* imaging study was also performed in order to examine the distribution of I-131 in established tumors in nude mice. After 3–5 weeks, when the tumor diameter had reached 8–10 mm, 200  $\mu$ l of *EgrLuc*-HVJ-liposome solution was injected into the tumor.

#### *In vivo gene therapy*

Female nude mice (Charles-River Japan, Tokyo, Japan) aged 4–6 weeks were injected subcutaneously on both flanks with 200  $\mu$ l of an Huh7 cell suspension ( $1 \times 10^7$  cells) in PBS. A transduction efficiency to the subcutaneous tumor of more than 50% was attained using the HVJ-liposome method, as reported previously.<sup>17</sup> After 3–5 weeks, when the tumor diameter had reached 8–10 mm, 200  $\mu$ l of the *EgrTK*-HVJ-liposome was injected (on days 1 and 8). For the *in vivo* luciferase assay experiments, 0–20  $\mu$ Ci I-131-lipiodol was injected into the tumor xenografts on day 2. The doses were selected on the basis of the clinical data and our pilot study. GCV administration (20 mg/kg per day injected intraperitoneally) was initiated 3 h after transfection and then continued for 2 weeks.

The mice were divided into four groups as follows: untreated animals (Group A;  $n=10$ ); those receiving *EgrTK*-vector transfection and irradiation (Group B;  $n=8$ ); those receiving *EgrTK*-vector transfection and GCV (Group C;  $n=8$ ) and those receiving *EgrTK*-vector transfection, irradiation and GCV (Group D;  $n=8$ ). None of the mice showed signs of wasting or any other indications of toxicity. The tumor volume was calculated using the formula  $1/2 \times (ab^2)$ , where 'a' represents the longer diameter and 'b' represents the shorter diameter. The care and treatment of the animals were in accor-

dance with the guidelines of the Nagasaki University Institutional Ethics Committee, Japan.

#### *Measurement of serum AFP*

In parallel with the measurement of tumor size, blood samples were collected from representative mice at various intervals by retro-orbital bleeding. Serum AFP levels were analyzed using a radioimmunoassay.<sup>42</sup>

#### *Hematoxylin–eosin staining*

To evaluate the mechanism causing the synergistic effects of *HSV-TK* gene therapy and internal radiation by I-131-Lip, samples of subcutaneous tumors were carefully dissected and removed at various intervals during therapy. The tumor tissues were fixed with 10% neutral formalin, embedded in paraffin and histologically examined.

#### *Egr-1 immunohistochemistry*

Immunohistochemical testing was performed in the hepatoma xenografts in order to evaluate the activation of the *Egr-1* gene after internal radiation. The samples were fixed with 4% paraformaldehyde and embedded in paraffin. The tissues were then cut into 3  $\mu$ m sections, deparaffinized in xylene and rehydrated. The sections were preincubated with normal bovine serum to prevent nonspecific binding, and then were incubated overnight at 4°C with an optimal dilution (5  $\mu$ g/ml) of the primary mouse monoclonal antibody against human *Egr-1* (Oncogene Science, Uniondale, NY, USA). The slides were sequentially incubated with the secondary antibody and the reaction products were visualized by DAB using a SIMPLE STAIN MAX-PO (MULTI) kit (Nichirei, Tokyo, Japan). Negative control samples were prepared by replacing the primary antibody with nonimmune murine serum.

#### *Statistical analysis*

All data are presented as the mean  $\pm$  standard deviation. Statistical analyses were performed using the Mann–Whitney *U*-test. Probability (*P*) values  $<0.05$  were considered to be statistically significant.

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## Review Article

# Liver Repopulation: A New Concept of Hepatocyte Transplantation

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### Abstract

Hepatocyte transplantation has been recognized as an alternative strategy for organ transplantation because the supply of donor livers is limited. However, in conventional hepatocyte transplantation, only 1%–10% of the liver replaced with transplanted hepatocytes. Recently a novel concept termed “liver repopulation” has been established, where the whole recipient liver can be replaced by a small number of donor hepatocytes. To induce liver repopulation, growth advantage of the donor hepatocytes against the host liver seems to be required according to the data of previous studies. Additionally, various cell sources, including bone marrow cells and other stem cells, could potentially be used as donor cells for liver repopulation. In this article, we discuss recent progress and future perspectives of this emerging technology.

**Key words** Hepatocyte transplantation · Liver repopulation

### Introduction

Many patients with inherited liver disorders and liver failure have been successfully treated by orthotopic liver transplantation (OLT). However, this procedure is associated with several problems; primarily, donor scarcity, the lifelong use of immunosuppressive drugs, and extraordinary expenses. Thus, much interest is now focused on hepatocyte transplantation as part of regenerative medicine. The advantages of hepatocyte transplantation include:

1. It is less invasive
2. It costs much less
3. Cells from a single donor can be used for multiple recipients
4. Cells can be cryopreserved
5. Autologous hepatocytes can be isolated from the small piece of resected tissue and genetically modified as ex vivo gene therapy

The wider use of hepatocytes to treat liver disease will not be possible until hepatocytes for cell transplantation become more readily available. Liver tissue obtained from patients without any liver disease, but whose liver is unsuitable for OLT, is the usual source of hepatocytes. However, the availability of viable hepatocytes is limited, and as yet there are very few medical centers performing hepatocyte transplantation worldwide. To address this issue, strategies to amplify the transplanted hepatocytes are being vigorously investigated. In particular, the new concept of “liver repopulation” has great potential for future clinical trials of hepatocyte transplantation.

### Hepatocyte Transplantation for Liver Failure

Experimental studies have shown that acute liver failure induced by the administration of liver toxic drugs such as D-galactosamine or by 90% hepatectomy can be treated by hepatocyte transplantation in various animals.<sup>1,2</sup> In rats with hepatic encephalopathy caused by an end-to-side portacaval shunt, which reflects fulminant hepatic failure (FHF), the intrasplenic transplantation of hepatocytes resulted in improved survival with recovery from hepatic coma.<sup>3</sup> Hepatocyte transplantation in rats with chronic decompensated liver cirrhosis stabilized the total bilirubin level and prothrombin time, and improved serum albumin and encephalopathy, thereby increasing survival.<sup>4–6</sup>

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**Table 1.** Clinical trial of hepatocyte transplantation for liver failure

First author <sup>Ref.</sup> (year)	Target disease	No. of cells transplanted	Site of transplantation	Outcome
Mito <sup>7</sup> (1992)	LC, CH 10 cases	1–60 × 10 <sup>7</sup>	Spleen	One patient recovered Clinically applicable
Habibullah <sup>8</sup> (1994)	FHF 7 cases	6 × 10 <sup>7</sup> /kg BW	Abdominal cavity	Survival and encephalopathy improved
Strom <sup>25</sup> (1997)	FHF 5 cases	2.8–29 × 10 <sup>7</sup>	Spleen	Ammonemia improved and 3 patients were bridged to OLT
Bilir <sup>26</sup> (2000)	FHF 5 cases	1–3.9 × 10 <sup>10</sup>	Spleen	Ammonemia improved Brain edema improved

LC, liver cirrhosis; CH, chronic hepatitis; FHF, fulminant hepatic failure; OLT, orthotopic liver transplantation; BW, body weight

The transplantation of hepatocytes in patients with acute liver failure may replace lost hepatocytes and serve as a bridge to OLT, or extend survival long enough to allow recovery and regeneration of the injured liver. Table 1 summarizes the clinical trials of hepatocyte transplantation against liver failure which have been performed to date. The first clinical trial was performed in 1991 by Mito et al.,<sup>7</sup> who transplanted autologous hepatocytes into the spleen of ten patients with liver cirrhosis or chronic hepatitis. The transplanted hepatocytes could be seen in the spleens for several months. Although the results were not conclusive, probably because of the insufficient number of hepatocytes isolated, it is noteworthy that hepatocyte clusters were found in the spleens. This pioneering study paved the way for subsequent clinical trials of hepatocyte transplantation. In 1994, Habibullah et al.<sup>8</sup> reported injecting 5 × 10<sup>9</sup> human fetal hepatocytes into the peritoneal cavity of patients with FHF, which decreased the plasma levels of ammonia and serum bilirubin and improved survival slightly. So far, hepatocyte transplantation has only been performed clinically as a "bridge to OLT" for FHF or chronic liver failure.

#### Hepatocyte Transplantation for Inherited Metabolic Diseases

Hepatocyte transplantation does not require replacement of the entire organ for many liver diseases, particularly the single gene defect-derived liver disorders listed in Table 2. Patients with ornithine transcarbamylase (OTC) deficiency,  $\alpha$ 1-antitrypsin deficiency,

and Crigler–Najjar syndrome type 1 have all been treated with hepatocyte transplantation. One infant with OTC deficiency received more than 4 × 10<sup>9</sup> hepatocytes, which resulted in temporary relief of hyperammonemia and protein intolerance.<sup>9</sup> The metabolic stability achieved was lost 11 days after transplantation, presumably because the transplanted cells were rejected as a result of inappropriate immunosuppression. However, this infant underwent successful OLT at 6 months of age.

The long-term therapeutic efficacy of hepatocyte transplantation was demonstrated in a patient with Crigler–Najjar syndrome type 1, who received 7.5 × 10<sup>9</sup> hepatocytes via a percutaneous portal vein catheter.<sup>10</sup> High-pressure liquid chromatography analysis of bile samples showed the excretion of conjugated bilirubin, reflecting the presence of functioning hepatocytes in the host liver. The serum bilirubin levels declined remarkably, allowing the time of phototherapy to be reduced from 12 h/day to 6 h/day.

Hereditary hypercholesterolemia is a fatal disease caused by the deficiency of a single-gene low-density lipoprotein (LDL) receptor. Chowdhury et al.<sup>11</sup> developed a method of harvesting primary hepatocytes from Watanabe rabbits, a model of LDL-receptor deficiency, and transducing them with a recombinant retrovirus expressing the LDL receptor gene. The transduced hepatocytes were then transplanted back into the rabbits. Long-term transgene expression and a 20%–30% reduction in serum cholesterol levels were confirmed during several months of observation. A clinical trial in patients with familial hypercholesterolemia resulted in a slight reduction of serum LDL cholesterol levels, but because the liver cell mass replaced was estimated to be

**Table 2.** Clinical trials of hepatocyte transplantation for inherited liver enzyme deficiencies

First author <sup>Ref.</sup> (year)	Diseases	No. of cells transplanted	Site of transplantation	Outcome
Grossman <sup>27</sup> (1995)	LDL receptor deficiency 5 cases	$1 \times 10^8/\text{kg BW}$	Portal vein	Cholesterol value dropped
Strom <sup>25</sup> (1997)	$\alpha$ 1-AT deficiency	$2.2 \times 10^7$	Splenic artery	Bridged to OLT 3 days later
Fox <sup>10</sup> (1998)	Crigler-Najjar syndrome type I	$7.5 \times 10^9$	Portal vein	50%–60% decrease in total bilirubin
Muraca <sup>28</sup> (2002)	Glycogen storage disease type Ia	$2 \times 10^9$	Portal vein	Food intake was normalized for 9 months
Horslen <sup>9</sup> (2003)	OTC deficiency	$4 \times 10^9$	Portal vein	Temporary relief of hyperammonemia and protein intolerance
Sokal <sup>29</sup> (2003)	Refsum's disease	$2 \times 10^6$	Portal vein	Total bile acids and abnormal DHCA decreased for 1 year

LDL, low-density lipoprotein; OTC, ornithine transcarbamylase;  $\alpha$ 1-AT,  $\alpha$ 1-antitrypsin; BW, body weight; OLT, orthotopic liver transplantation; DHCA, dihydroxycoprostanic acid

less than 1%, the serum cholesterol reduction was not therapeutically beneficial.<sup>12</sup>

### Limitations of Hepatocyte Transplantation

Although hepatocyte transplantation has been performed safely, sometimes with long-term benefits, in humans, its application is still limited. Because of the critical shortage of donor organs, there are insufficient isolated hepatocytes for hepatocyte transplantation. Furthermore, transplanted hepatocytes cannot proliferate in the host liver without adequate growth stimulus. Unfortunately, the use of partial hepatectomy or cell growth factor treatment to stimulate liver growth and increase the proportion of transplanted hepatocytes in the liver has generally failed to show a significant benefit. Because the host hepatocytes divide during liver regeneration after hepatectomy, both the host and the transplanted hepatocytes should respond similarly to this regenerative stimulus, but repeated cell transplantations have been performed without success.<sup>13</sup> Thus, interest is now focused on finding a way to provide selective growth pressure to the transplanted hepatocytes.

### A New Concept: Repopulation of Transplanted Hepatocytes

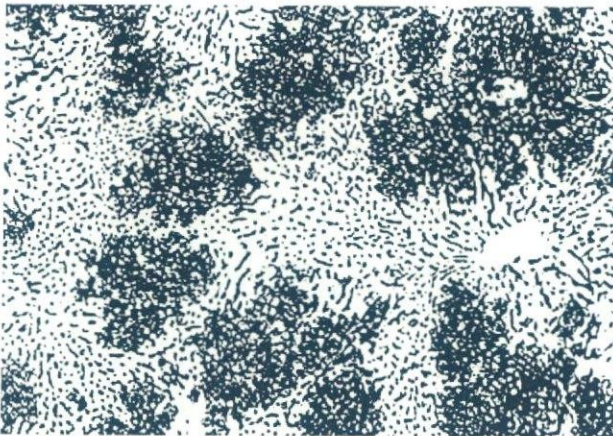
Data from clinical trials suggest that the efficacy of hepatocyte transplantation has been hampered by the fact that only a small percentage of the host hepatocytes

can be replaced with transplanted hepatocytes. Experimental studies have shown that mature hepatocytes have the potential to repopulate diseased livers *in vivo* under regenerative stimulus directed at the transplanted cells. In other words, replacement of the host liver cells with transplanted hepatocytes requires a combination of proliferative stimuli to the hepatocytes and suppression of the division of host hepatocytes. An example of this concept has been found in urokinase plasminogen activator transgenic mice (uPA), in which the life span of the host hepatocytes is restricted, resulting in preferential proliferation of the transplanted normal hepatocytes until the whole liver is replaced.<sup>14</sup> Similar data were obtained by using fumarylacetoacetate hydrolase (FAH)-deficient mice, a model of hereditary tyrosinemia.<sup>15</sup> To popularize the concept of "liver repopulation," chemical and physical devices to inhibit the division of host hepatocytes are being explored. Using one approach, a recipient was treated with a DNA-kilating alkaloid, retrorsine, followed by partial hepatectomy and hepatocyte transplantation.<sup>16</sup> This resulted in the transplanted cells progressively repopulating the host liver. Instead of retrorsine, we used hepatic irradiation to arrest the cell cycle of the host hepatocytes, which resulted in near total liver replacement by the transplanted hepatocytes<sup>17,18</sup> (Fig. 1). We also showed that partial hepatectomy can be replaced by adenovirus-mediated Fas ligand gene transfer, which induces strong mitotic stimuli by causing apoptotic cell killing of endogenous hepatocytes. Table 3 lists the various methods used to achieve massive liver repopulation.

**Table 3.** Repopulation of transplanted hepatocytes in experimental animals

First author <sup>Ref.</sup> (year)	Recipient animal	Preconditioning of the host	Character of transplanted hepatocytes
Moscioni <sup>30</sup> (1996)	NAR rat	PBL	Normal
Ilan <sup>31</sup> (1997)	Gunn rat	PBL	Normal
Laconi <sup>16</sup> (1998)	DPPIV <sup>-/-</sup> rat	PH + retrorsine	DPPIV <sup>+/+</sup>
Mignon <sup>32</sup> (1998)	Normal mouse	Fas antibody treatment (Jo2)	Bcl2 transduced
Overturf <sup>25</sup> (1998)	FAH deficient mouse	NTBC administration	FAH transduced
Guha <sup>33</sup> (1999)	DPPIV <sup>-/-</sup> rat	PH + RT	DPPIV <sup>+/+</sup>
Oren <sup>34</sup> (1999)	DPPIV <sup>-/-</sup> rat	T3 and retrorsine	DPPIV <sup>+/+</sup>
Laconi <sup>16</sup> (2001)	DPPIV <sup>-/-</sup> rat	Retrorsine alone	DPPIV <sup>+/+</sup>
Guha <sup>17</sup> (2002)	Gunn rat	PH + RT	Normal
Guo <sup>35</sup> (2002)	C57BL/6J mouse	CCl <sub>4</sub> , retrorsine	X-gal transduced
Malhi <sup>36</sup> (2002)	DPPIV <sup>-/-</sup> rat	IR injury; RT	DPPIV <sup>+/+</sup>
Mallet <sup>37</sup> (2002)	Normal rat	Fas antibody treatment (Jo2)	Bcl2 transduced
Mitchell <sup>38</sup> (2002)	Normal rat	Fas antibody treatment (Jo2)	BclXL transduced
Takahashi <sup>18</sup> (2003)	Gunn rat	Fas-Ligand gene transfer + RT	Normal
Yuan <sup>39</sup> (2003)	DPPIV, Rag2 <sup>-/-</sup> rat	CCl <sub>4</sub>	p27 <sup>-/-</sup>
Suzuki <sup>40</sup> (2004)	Normal rat	PH + retrorsine	c-Met <sup>+</sup> CD45 <sup>-</sup> sorted

PH, partial hepatectomy; DPPIV, dipeptidyl-peptidase IV; RT, radiation treatment; IR, ischemia-reperfusion injury; FAH, fumarylacetoacetate hydrolase; NTBC, 2-(2-nitro-4-trifluoromethylbenzoyl)-1,3-cyclohexanedione; PBL, portal branch ligation; SD, Sprague-Dawley



**Fig. 1.** Hepatocyte repopulation in F344 rats that received hepatic irradiation. Dipeptidyl-peptidase IV (DPPIV) +F344 hepatocytes (stained red) were transplanted intrasplenically into congenic, DPPIV-deficient F344 rats subjected to whole liver irradiation (50 Gy)

### Novel Sources of Hepatocytes

The success of hepatocellular transplantation depends greatly on the eventual number of viable hepatocytes transplanted. Autologous hepatocytes are an ideal source because immunosuppression is not required; however, realistically, when cells are obtained from allogenic donors, immunosuppressant therapy is needed. To address this issue, a novel source of hepatocytes should be investigated. Immortalizing isolated hepatocytes is one approach.<sup>19,20</sup> Although successful results have been reported, complete suppression of

tumorigenic potential is a major concern when transplanting these immortalized cells to humans. Xenogenic cells such as porcine hepatocytes have also been considered as an unlimited source of mature hepatocytes. These cells were originally used as a bioreactor in artificial liver support systems with encouraging results,<sup>21,22</sup> but reports of porcine endogenous retrovirus, which can infect human cells, halted the use of porcine hepatocytes.<sup>23</sup> Stem/progenitor cells are the most promising source of hepatocytes. Recent advances in stem cell biology have changed the dogma concerning lineage relationships. Hepatocytes derived from bone marrow cells were proven capable of repopulating the host liver in a murine model of tyrosinemia, although it is not clear what signals are implicated in the differentiation of the bone marrow cells into mature hepatocytes.<sup>24</sup>

### Future Perspectives

The idea that isolated cells can be transplanted into the body and directed to repair damaged organs is gaining much attention. As described in this review, hepatocyte transplantation is technically feasible and clinically beneficial; however, the number of transplanted hepatocytes that survive in the host liver limits the efficiency of hepatocyte transplantation. We think that liver repopulation technology will enable the expansion of a few transplanted hepatocytes, which can be obtained from small pieces of liver from the patient (autologous) or healthy donors (allogenic). In our mind, preparative liver irradiation of the host liver can be translated in the clinical settings as follows. A partial hepatectomy is performed in patients with liver dis-

eases, and hepatocytes are isolated from the resected liver tissue. Then, after irradiation to the remnant liver, the isolated hepatocytes (with or without genetic modification) are transplanted back into the patient. If the host liver is not suitable for cell isolation or transplantation, the allogenic hepatocytes obtained from the healthy donors can be used instead, although this requires immunosuppression.

In conclusion, hepatocyte transplantation to treat human liver disorders is still in the early stages of development. Further research and clinical trials are needed, but hepatocyte transplantation shows the promise of becoming a therapeutic option for patients with end-stage liver diseases, particularly those awaiting OLT.

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# 成人生体肝移植におけるネオーラル投与の実際

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*Optimal application of Neoral in adult living donor liver transplantation*

**key words** : 成人生体肝移植, ネオーラル, AUC

近年, 生体肝移植において, 免疫抑制剤の血中濃度が脳死肝移植に比して相対的に高値を示すこと<sup>1)</sup>, タクロリムス(FK506)の至適投与量がグラフト重量に規定されること<sup>2)</sup>などが報告され, 免疫抑制剤の薬物動態が生体肝移植と脳死肝移植とで異なることが示されている。

そこで, 生体肝グラフトの再生がシクロsporin(ネオーラル)の代謝に及ぼす影響について検討し, 生体肝移植における therapeutic drug monitoring(TDM)に基づいたネオーラルの至適使用法の確立を目指した<sup>3)</sup>。

## 対象と方法

2001年4月から現在までネオーラル経口投与をベースとする免疫抑制療法を施行し, 薬物動態を解析できた10例の成人生体肝移植症例を対象

とした。

術前日, 術後3, 7, 21病日(POD)に経時的(ネオーラル投与後1, 2, 4, 6, 12時間後)に末梢静脈血および門脈血を採取し, area under the time concentration curve(AUC)を測定し, 7POD, 30PODに施行したCT volumetryによりグラフト肝容量を測定した。

免疫抑制はネオーラルとステロイドの2剤で行い, ネオーラル投与量は術前日5 mg/kg/dayで開始し, 術後4週間はトラフ値200~300 ng/mLを目標とした。

また, Level/Dose比(L/D比) = トラフ値/投与量,  $\Delta$  AUC = 門脈血 AUC - 末梢血 AUC, 肝除去率 =  $\Delta$  AUC/門脈血 AUC, などについても検討した。

## 結果

### (1) 周術期のAUC

術前はいずれの症例においてもC<sub>2</sub>がピーク

表1 ネオーラル使用例の臨床成績: FK506との対比

(Mean follow up)	Neoral-based (24.3 months)	FK506-based (41.5 months)
Total cases (Male/Female)	10 (6/4)	15 (5/10)
Acute rejection	3 (30.0%)	6 (40.0%)
Infection	2 (20.0%) CMV enterocolitis, HCV acute hepatitis	3 (20%) CMV enterocolitis, sepsis, pulmonary aspergillosis
Side effects (others)	2 (20.0%) seizure, renal failure	2 (13.3%) renal failure
Graft or patient loss	1 (10.0%) PSC recurrence (reALDLT)	2 (13.3%) idiopathic cardiac failure, pulmonary aspergillosis

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値で、 $C_2$ 、AUCは脳死肝移植で報告されているデータとほぼ同様であった( $C_2 = 1,013 \pm 216$  ng/mL,  $AUC_{0-12} = 6,952 \pm 1,143$  ng·hr/mL)。一方、3PODではピークが明らかでない症例や遅れる症例が多く、 $C_{max}$   $484 \pm 263$  ng/mLと低かった。7PODでは術前のAUCに戻る傾向であったが、脳死肝移植での報告<sup>4)</sup>とくらべて $C_0$ 、 $C_2$ 、AUCともに低値であった。

術後1週間以内の $AUC_{0-12}$ は $C_2$ のみならず $C_0$ ともよく相関し( $r = 0.773$ ,  $p = 0.0001$ )、トラフ値でモニターしても支障のないことが示唆された。

### (2) 肝再生に伴うシクロスポリンの代謝変動

シクロスポリンの肝除去率は7PODでは3PODにくらべ有意に増加し、グラフト容積の増加率と相関したが、プロトロンビン時間やASTなどの肝機能とは相関しなかった。30PODのL/D比はグラフト容積/標準肝容積(GV/SLV)と有意な負の相関( $r = -0.699$ ,  $p < 0.05$ )を示し、肝再生に伴うグラフト容積の増大がシクロスポリンの代謝亢進に関与することが示唆された。

生体肝移植では、グラフトサイズがシクロスポリンの代謝に影響を与えることを考慮してネオー

ラルの投与量を加減する必要がある。

### (3) 臨床成績

ネオーラルを*de novo*で使用した症例の成績は表1のごとくで、FK506をベースにした症例と拒絶率、合併症率、生存率などに差はなく、良好であった。

### おわりに

ネオーラルは、生体肝移植においても安全・有効に使用できたが、脳死肝移植にくらべて吸収・代謝に違いがあり、生体肝移植独自の投与プロトコルが必要である。

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# Identification of Novel HCV Subgenome Replicating Persistently in Chronic Active Hepatitis C Patients

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In an effort to clarify the life cycle of HCV, the HCV genome in liver biopsies taken from chronic active hepatitis C patients undergoing interferon treatment was investigated. Molecular cloning by long distance reverse-transcription polymerase chain reaction (RT-PCR) revealed that the HCV genome in two patients with high viral loads in the liver had in-frame deletions of approximately 2 kb between E1 and NS2, which encode the E1–NS2 fusion protein and six other HCV proteins: core, NS3, NS4A, NS4B, NS5A, and NS5B. Among the remaining 21 chronic active hepatitis C patients, these types of deletion were found in another two patients and in two hepatocellular carcinoma patients. Out-of-frame deletions in the structural region were isolated from the other five patients, but the dominant RT-PCR products were non-truncated genomes. Retrospective analysis of a series of serum samples taken from a patient carrying the subgenome with the in-frame deletion revealed that both the subgenome and the full genome persisted through the 2-year period of investigation, with the subgenome being predominant during this period. Sequence analysis of the isolated cDNA suggested that both the subgenome and the full genome evolved independently. Western blotting analysis of HCV proteins from the HCV subgenome indicated that they were processed in the same way as those from the full genome. HCV subgenomes thus appear to be involved in the HCV life cycle. *J. Med. Virol.* 77:399–413, 2005. © 2005 Wiley-Liss, Inc.

**KEY WORDS:** HCV; deletion; replication; biopsy

## INTRODUCTION

Hepatitis C virus (HCV) is primarily transmitted via blood and blood-derived materials [Alter et al., 1989]

and often causes chronic hepatic diseases that progressively worsen to chronic active hepatitis, cirrhosis, and finally to hepatocellular carcinoma (HCC) [Kiyosawa et al., 1990, 1994, 2004; Alter and Seeff, 2000]. Interferon (IFN) and interferon with ribavirin treatment are effective in eradicating HCV from patients [Iino et al., 1994; McHutchison and Fried, 2003], improving liver histological findings, and in prolonging life in patients with hepatitis C [Yoshida et al., 1999; Kasahara et al., 2004]; however, their efficacy is limited.

HCV was first identified as cDNA clones, and was characterized molecularly using cDNA isolates [Choo et al., 1989; Kuo et al., 1989]. The HCV genome is single-stranded RNA of about 9,600 nucleotides with an untranslated region (UTR) at each end, and encodes a polyprotein of about 3,010 amino acids [Choo et al., 1989; Kato et al., 1990], which is processed into 10 proteins by a host peptidase and two HCV proteases [Hijikata et al., 1991, 1993; Grakoui et al., 1993a]; Core, E1, and E2 are structural proteins for virion formation, and NS3, NS4A, NS4B, NS5A, and NS5B are components of the replication machinery for the RNA genome [Houghton et al., 1994]. However, isolation of virion particles has been difficult owing to a lack of in vitro culture systems for HCV.

HCV replication in chimpanzee following intrahepatic injection of an RNA transcript from HCV genomic cDNA proved that a molecular clone could represent a functional HCV genome [Yanagi et al., 1997]. Lohmann

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et al. [1999] established a dicistronic subgenomic RNA that replicates in a hepatoma cell line (Huh7) and consists of the HCV NS protein coding region, the 5'- and 3'-UTR of HCV and a selective marker gene. HCV subgenomic RNA replicon systems are vital to the study of the mechanisms of HCV RNA replication, but there remain problems with regard to viral replication. Information obtained from liver biopsies of hepatitis C patients thus remains important in clarifying the life cycle of HCV.

Histological grading for diagnosis [Perrillo, 1997], immunohistochemical analysis, immuno-staining [Infantolino et al., 1990; Hiramatsu et al., 1992], electron microscopic analysis [Fagan et al., 1992], in-situ hybridization and in-situ reverse transcription polymerase chain reaction (RT-PCR) [Lau et al., 1996; Dries et al., 1999], and quantitation of HCV RNA in liver biopsy specimens [Sakamoto et al., 1994; Nuovo et al., 2002] have all been used to demonstrate HCV replication in liver. However, molecularly characterized data have been limited; HCV RNA isolated from liver was found to be equivalent in size to the well-characterized RNA seen in circulating HCV [Nielsen et al., 2004], thus confirming HCV replication in liver.

In order to obtain data to elucidate the nature of HCV in liver, viral loads and the structure of the HCV genome in patient liver biopsy specimens were examined. A highly sensitive ELISA for quantitation of the HCV core antigen [Aoyagi et al., 1999; Tanaka et al., 2000] and a quantitative RT-PCR system were applied to estimate viral loads in serum and liver biopsies. For structural analysis of the whole genome in specimens, molecular clones were used. Surprisingly, it was found that novel HCV subgenomes were predominant in several patients and, in one patient, these persisted for several years. The nature of these subgenomes are described and discussed in this paper.

## MATERIALS AND METHODS

### Samples and Antibodies

Serum and liver biopsy specimens were taken from patients undergoing IFN- $\alpha$  treatment at Shinshu University Hospital. Informed consent was obtained from all patients from whom samples were taken. A 7.2 mega-unit dose of IFN- $\alpha$  was administered daily for 2 weeks, followed by three times per week for 22 weeks. Serum samples were collected 1 or 2 months before administration of IFN- $\alpha$ . A series of serum specimens was collected from one patient from before IFN treatment until 1 year after treatment. Liver biopsy specimens from two HCC patients were obtained from surgically removed cancerous liver tissues.

An anti-core monoclonal antibody (5E3) has been described previously [Kashiwakuma et al., 1996]. Anti-E1 monoclonal antibody was raised against recombinant E1 and E1/E2 proteins expressed in Sf-9 cells infected with recombinant baculoviruses (Yamaguchi unpublished). Anti-NS3 rabbit polyclonal antibody was purified from the serum of rabbits immunized with the

recombinant NS3 antigen expressed in *E. coli* [Saito et al., 1992]. Anti-mouse and rabbit immunoglobulin antibody conjugated with HRP were purchased from Kirkegaard & Perry Laboratories, Inc. (Gaithersburg, MD) and Bio-Rad Laboratories (Tokyo, Japan), respectively. All primers were purchased from Sigma Genosys (Tokyo, Japan) and Texas Genomics Japan (Tokyo, Japan), and sequences are available on request.

### Quantitation of HCV Core Antigen

Quantities of HCV core antigen were measured by EIA as described previously [Aoyagi et al., 1999; Kato et al., 2003]. The concentration of core antigen was expressed in fmol/L, and the cut-off value of the assay was set at 7.5 fmol/L. For quantitation of the core antigen in liver, extracts were diluted to 100  $\mu$ g of liver protein per milliliter with negative control serum before pretreatment of the samples. Samples were heated at 56°C for 30 min with pretreatment solution containing SDS, CHAPS, and Triton X-100, and were then added to wells pre-coated with anti-HCV core antibodies, and reaction buffer was used to fill the wells. Captured core antigen after 1-hr incubation was reacted for 30 min with anti-HCV antibodies conjugated with horseradish peroxidase after stringent washing. Bound enzyme activities were measured using a Fusion plate reader (PerkinElmer, Tokyo, Japan) with a chemiluminescent reagent (SuperSinal Pico ELISA, Pierce, Rockford, IL).

### Real-Time PCR Assay for HCV RNA

HCV RNA was recovered from samples by using the QIAamp viral RNA kit (QIAGEN K.K., Tokyo, Japan) according to the manufacturer's instructions. HCV RNA was reverse-transcribed and amplified using QuantiTect One-Step RT-PCR kit (QIAGEN) with primers. For quantitation of the 5'-UTR, the forward primer, chiba-s (5'-TAGTGGTCTGCGGAACCGGT-3'), and reverse primer, chiba-as (5'-TGCACGGTCTACGAGACCT-3'), yielded fragments corresponding to nucleotides 141–339 of HCV RNA. In the case of the E2 region, HC1986S (5'-TGGTTCGGCTGYACATGGATGAA-3') and HC2199AS (5'-GGRTAGTGCCARAGCCTGTATGGGTA-3') primers were used. Reactions were performed with a LightCycler system (Roche Diagnostics K.K., Tokyo, Japan), and fluorescence by SYBR green was monitored after each elongation reaction for real-time monitoring of DNA products during PCR. The amount of HCV RNA was calculated according to the calibration curve produced with serial dilutions of standard RNA synthesized by T7 RNA polymerase (Ambion, Inc., Austin, TX) from plasmids carrying the HCV cDNA isolate (genotype 1b). To examine the specificity of PCR, the melting point of DNA products was analyzed by melting curve analysis using LCDA software (Roche Diagnostics).

### Cloning and Analysis of HCV cDNA

HCV cDNA was amplified by long distance RT-PCR (LD-RT-PCR) as described previously [Tellier et al., 1996;

TABLE I. Viral Data of Patients

Patient No.	HCV genotype	Viral loads				PCR primer sets for positive results		
		Serum		Liver biopsy		Non-truncated genome	Truncated genome	Test primer sets
		Core antigen (fmol/L)	HCV RNA (copies/ml)	Core antigen (fmol/g protein)	HCV RNA (copies/g protein)			
368	1b	17,108.5	$1.73 \times 10^5$	5,462.4	$4.71 \times 10^8$	—	j	j
207	1b	12,695.2	$1.44 \times 10^5$	30,792.3	$1.43 \times 10^{10}$	NT	NT	NT
204	1b	5,082.4	$5.74 \times 10^4$	8,779.7	$2.22 \times 10^9$	i	—	i
274	1b	1,034.4	$4.24 \times 10^3$	2,651.7	$3.56 \times 10^7$	a, b, c, d	e	a-e
193	1b	988.8	$3.09 \times 10^4$	14,519.9	$1.07 \times 10^9$	a, b, c, d, e	—	a-e
331	1b	922.2	$2.03 \times 10^3$	2,387.1	$2.84 \times 10^8$	a, c, d	b, e	a-e
325	1b	623.5	$3.82 \times 10^3$	10,127.9	$7.28 \times 10^7$	a, b, c, d, e	d	a-e
288	1b	254.5	$1.00 \times 10^1$	4,037.9	$9.50 \times 10^6$	a, b, c, d, e	d, e	a-e
299	1b	166.6	$1.14 \times 10^3$	1,287.8	$5.35 \times 10^7$	c, d	—	a-e
295	1b	1.0	$5.11 \times 10^1$	261.5	$2.62 \times 10^7$	a, b, c, d, e	b	a-e
171	1b	1,077.3	$6.42 \times 10^3$	3,781.8	$6.91 \times 10^6$	c, d	b	a-e
257	1b	12.7	$1.06 \times 10^2$	568.5	$2.78 \times 10^7$	d	—	a-e
372	1b	723.7	$2.28 \times 10^4$	1,784.1	$3.35 \times 10^8$	a, b, c, d, e	—	a-e
373	1b	597.0	$8.31 \times 10^3$	33,919.0	$2.65 \times 10^9$	—	a, c, d	a-e
248	2a	209.3	$2.58 \times 10^2$	4,417.1	$3.70 \times 10^8$	—	—	a-e
235	2a	3,616.2	$3.66 \times 10^2$	7,462.1	$1.55 \times 10^9$	c	—	a-e
203	2b	95.1	$1.46 \times 10^2$	5,590.9	$1.82 \times 10^9$	—	b, d	a-e
178	2b	34.5	$8.08 \times 10^1$	609.1	$4.51 \times 10^7$	—	—	a-e
297	2	3,112.7	$8.35 \times 10^3$	2,883.6	$1.14 \times 10^8$	—	—	a-e
298	2a	180.0	$8.09 \times 10^2$	3,015.0	$1.76 \times 10^9$	b	—	a-e
305	2a	173.6	$1.12 \times 10^3$	1,782.8	$5.96 \times 10^7$	b	—	a-e
201	2a	127.6	$2.40 \times 10^3$	497.6	$1.87 \times 10^7$	—	—	a-e
357	2	227.2	$3.11 \times 10^3$	321.9	$2.29 \times 10^7$	—	—	a-e

NT: not tested.

Yanagi et al., 1998]. HCV cDNA was synthesized using Superscript II reverse transcriptase (Invitrogen K.K., Tokyo, Japan) with HC1b9405R primer (5'-GCCTA-TTGGCCTGGAGTGTTTAGCTC-3'). After RNase H

(Invitrogen) treatment at 37°C, a cDNA mixture was subjected to PCR with KlenTaq DNA polymerase (BD Biosciences Clontech, Tokyo, Japan), HClong A1 primer (5'-GCCAGCCCCCTGATGGGGGCGACA-

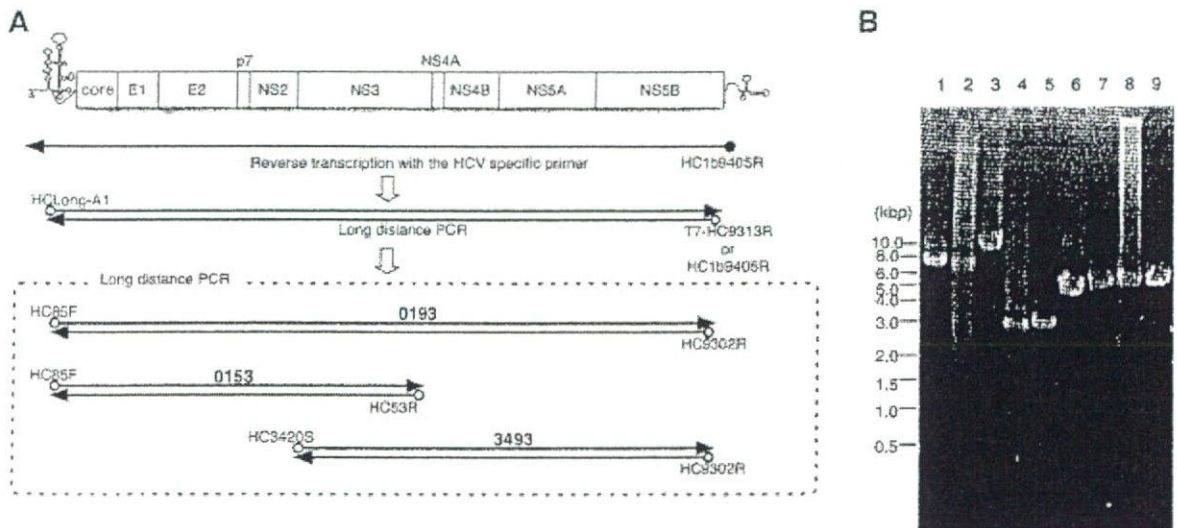


Fig. 1. Molecular cloning of HCV genome by long distance reverse-transcription PCR. A: Schematic view of HCV RNA is shown at the top of the figure. HCV cDNA, which was synthesized from total RNA from liver using reverse transcriptase with HCV-specific primers for the 3'-UTR, was amplified by nested PCR with HCV-specific primers. The longest LD-RT-PCR product, 0193, covered 99% (amino acids 1-2,987) of the HCV polyprotein coding sequence of genotype 1b HCV (length: 3,011 amino acids). Fragments obtained by LD-RT-PCR using HC85F

and HC9302R, HC85F, and HC53R; and HC3420S and HC9302R were designated 0193, 0153, and 3493, respectively. After agarose gel electrophoresis, LD-RT-PCR products from liver biopsy samples were stained with ethidium bromide. B: Lanes 1-3, lanes 4-6, and lanes 7-9 represent 0193, 0153, and 3493 fragments from Patient 207 (lanes 1, 4, and 7), 373 (lanes 2, 5, and 8), and control HCV cDNA (lanes 3, 6, and 9), respectively. The positions of markers are indicated at the left side of the image.