

Impact of insulin resistance on the progression of chronic liver diseases

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Received August 13, 2008; Accepted September 29, 2008

DOI: 10.3892/ijmm_00000088

Abstract. Recent studies have revealed a close relationship between insulin resistance (IR) and the progression of chronic liver diseases, although relatively little is known regarding the possible mechanisms involved. The aim of this study was to elucidate the impact of IR on the development of liver fibrosis and hepatocarcinogenesis using obese diabetic Otsuka Long-Evans Tokushima Fatty (OLETF) rats. Liver fibrosis development and glutathione-S-transferase placental form (GST-P)-positive pre-neoplastic lesions were both markedly accelerated in OLETF rats, being induced by pig serum and diethylnitrosamine (DEN), respectively. In the fibrosis experiment, α -smooth muscle actin-positive activated hepatic stellate cells (HSCs) also significantly increased in OLETF rats along with augmentation of the hepatic collagen content and transforming growth factor- β_1 . Our *in vitro* study showed that both glucose and insulin stimulated the proliferation of activated HSCs, and the combination treatment exerted an additive effect. In the DEN model, neovascularization, which plays a pivotal role in hepatocarcinogenesis, was up-

regulated in OLETF rats almost in parallel with pre-neoplastic lesion development and a potent angiogenic factor, vascular endothelial growth factor. High glucose and insulin also significantly augmented the *in vitro* neovascularization via extracellular signal-regulated kinase 1/2 phosphorylation. Similar to the effect on the activated HSCs, co-existence of both factors exerted a more potent effect than either single factor. In conclusion, these results indicated that the IR status directly accelerated liver fibrosis development and hepatocarcinogenesis at least partly through the stimulation of activated HSC proliferation and hepatic neovascularization, respectively, in the rat.

Introduction

Hepatocellular carcinoma (HCC) is a malignancy of worldwide significance, and its prevalence is still increasing in Japan, Western Europe and the US (1). HCC commonly develops in patients with liver cirrhosis. Liver cirrhosis is caused by many etiological factors, such as hepatitis B or C (2). Among them, hepatitis virus C (HCV) infection is now the most common underlying cause of these clinical entities, i.e., liver cirrhosis and HCC (3). Recent evidence suggests that HCV infection is associated with an increased risk of type 2 diabetes mellitus (DM) (4-6). Cross-sectional human studies showed that insulin resistance (IR), i.e., co-existence of high glucose and insulin, is a consistent finding in patients with type 2 DM (7). Experimental evidence for the contribution of HCV in the development of IR and DM has been found in the HCV-transgenic mouse model (5). Moreover, several human studies demonstrated that IR is a risk factor of advanced fibrosis in patients with chronic hepatitis C (CHC), and IR itself may contribute to the progression of liver fibrosis in CHC (8-10). Moreover, a similar close interaction was observed in patients with other morbidities, such as non-alcoholic fatty liver diseases (NAFLD) (11). Either high glucose or insulin increased the production and expression of collagen genes in the activated hepatic stellate cells (HSCs), which play a pivotal role in liver fibrosis development *in vitro* (12,13).

In addition to liver fibrogenesis, a close interaction between DM and HCC has been documented. DM is associated with a 2- to 3-fold increase in the risk of HCC regardless of the etiology of chronic liver diseases, including

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Abbreviations: CHC, chronic hepatitis C; DEN, diethylnitrosamine; DM, diabetes mellitus; ERK1/2, extracellular signal-regulated kinase 1/2; ECs, endothelial cells; GST-P, glutathione-S-transferase placental form; HCC, hepatocellular carcinoma; HSCs, hepatic stellate cells; HUVECs, human umbilical vein endothelial cells; IR, insulin resistance; LETO, Long-Evans Tokushima Otsuka rats; NAFLD, non-alcoholic fatty liver diseases; NASH, non-alcoholic steatohepatitis; OLETF, Otsuka Long-Evans Tokushima Fatty rats; QUICKI, Quantitative Insulin Sensitivity Check Index; α -SMA, α -smooth muscle actin; TGF, transforming growth factor; VEGF, vascular endothelial growth factor

Key words: hepatocellular carcinoma, angiogenesis, hepatic stellate cells, glucose, vascular endothelial growth factor

CHC and non-alcoholic steatohepatitis (NASH) (14,15). It is now recognized that angiogenesis plays an important role in many physiological and pathological events (16). Neovascularization and a potent angiogenic factor; namely, the vascular endothelial growth factor (VEGF), play important roles in hepatocarcinogenesis. It has been reported that VEGF expression and hepatic neovascularization exhibit a stepwise increase in pre-neoplastic lesions during hepatocarcinogenesis (17). Several recent investigations have revealed a close relationship between angiogenesis, high glucose and insulin (18,19). However, the possible interaction between IR, angiogenesis and VEGF during hepatocarcinogenesis has not as yet been elucidated.

In the current study, we examined the *in vivo* role of IR on the progression of chronic liver diseases, i.e., liver fibrosis development and hepatocarcinogenesis, using obese diabetic Otsuka Long-Evans Tokushima Fatty (OLETF) rats, which develop IR after 10-15 weeks of age and non-insulin-dependent DM after 25-30 weeks of age (20). We also performed a set of *in vitro* experiments to elucidate the possible mechanism involved in these processes.

Materials and methods

Animal treatment. Male OLETF and control Long-Evans Tokushima Otsuka (LETO) rats were generously supplied by Otsuka Pharmaceutical Co. (Tokushima, Japan) (20). At the age of 20 weeks, the LETO and OLETF rats were randomly divided into 8 groups and employed for both fibrosis and hepatocarcinogenesis experiments. In the fibrosis experiment, 1.0 ml/kg of pig serum (JRH Biosciences, Inc., St. Lenexa, KS, USA) was intraperitoneally injected twice a week for 4 weeks both in the LETO and OLETF rats, which were designated as Group 1 (G1) and G2, respectively. As a negative control, phosphate buffer saline (PBS) was injected instead in the LETO and OLETF rats (G3 and G4, respectively). In the carcinogenesis experiment, LETO (G5) and OLETF (G6) rats received a single injection of 200 mg/kg of diethylnitrosamine (DEN) (Tokyo Kasei Kogyo Co., Ltd, Tokyo, Japan), partially hepatectomized at week 3 as described previously (21,22), and then sacrificed at the end of week 9. In the carcinogenesis experiment, PBS was injected and partial hepatectomy was performed in the LETO and OLETF rats (G7 and G8, respectively) as a negative control. Each group consisted of 10 rats. At the end of the respective experiment, all rats were anesthetized, the thoracic cavity was opened, and blood samples were withdrawn via cardiac puncture. Several biological markers were assessed by routine laboratory methods. IR was evaluated with the Quantitative Insulin Sensitivity Check Index (QUICKI) as described previously (23). All animal procedures were performed according to standard protocol and in accordance with the standard recommendations for the proper care and use of laboratory animals.

Histological and immunohistochemical staining and quantification. The first section was routinely stained with hematoxylin and eosin for histological examination. Another section was stained with Sirius-red (S-R) to detect fibrosis development. Immunohistochemical staining of enzyme-

altered pre-neoplastic lesions; namely, the placental form of glutathione S-transferase (GST-P) (MBL Co., Ltd., Nagoya, Japan) and α -smooth muscle actin (α -SMA) (Dako, Kyoto, Japan), using formalin-fixed tissue sections, was performed as described previously (24,25). For determination of neovascularization, we employed immunohistochemical detection of CD31 (BD Biosciences, CA, USA), which is widely used as a marker of neovascularization as previously described with frozen sections (26). Briefly, non-specific antibody binding was blocked with serum diluted at 1:50 in PBS, and the endogenous biotin was blocked with 0.1% avidin for 15 min, followed by 0.01% biotin for 15 min. These sections were washed once with PBS after each 15-min incubation period, then reacted with a primary respective antibody for 1 h at room temperature. After washing twice with PBS, sections were incubated with a second biotin-labeled rabbit anti-rat IgG (Novocastra, CA, USA). The endogenous peroxidase was blocked with 0.3% hydrogen peroxide in PBS for 30 min, followed by a 5-min washing with PBS. The sections were then incubated with conjugated streptavidin for 30 min, rinsed again in PBS, and finally incubated with diaminobenzidine for 3 min. We did not count the α -SMA-positive vessels in the portal area, which were assumed to be hepatic arteries. We only included the α -SMA-positive cells in the sinusoidal lining for image analysis. Semi-quantitative analyses of fibrosis development and immunopositive cells of α -SMA, GST-P and CD31 of 7 rats were carried out using Adobe Photoshop software and National Institutes of Health image software as described elsewhere (24-26).

Hepatic collagen, transforming growth factor- β_1 and vascular endothelial growth factor expression. After equalization of the protein content, the hepatic collagen content was determined using a Sircol Soluble Collagen Assay Kit (Biocolor Ltd., New Town Abbey, Northern Ireland) in 200 mg of frozen liver samples according to the manufacturer's instructions. TGF- β_1 and VEGF in the liver were determined using an enzyme-linked immunosorbent assay (ELISA) kit (R&D Systems, Tokyo, Japan), according to the manufacturer's instructions.

***In vitro* proliferation assays of hepatocellular carcinoma.** The primary HSCs were isolated from the liver of OLETF rats, as described previously (27) with a minor modification. The cell viability was >95% as determined by the Trypan blue exclusion test. Freshly isolated HSCs were plated at a density of 5×10^5 cells/ml on uncoated plastic dishes. After a 5-day culture, HSCs became myofibroblast-like with reduced lipid vesicles and increased immunoreactive α -SMA, and at 7 days after planting, all cells were well-spread and α -SMA positive (28). The effects of D-(+)-glucose (Nacalai Tesque, Kyoto, Japan) and human recombinant insulin (Gibco, Tokyo, Japan) on the proliferation of HSCs were detected by bromodeoxyuridine (BrdU) incorporation using a Cell Proliferation ELISA, BrdU kit (Roche Applied Science, Mannheim, Germany) according to the manufacturer's instructions.

***In vitro* angiogenesis assay and extracellular signal-regulated kinase 1/2 phosphorylation in endothelial cells (ECs).** *In vitro* angiogenesis was assessed as the formation of capillary-like structures of human umbilical vein endothelial cells

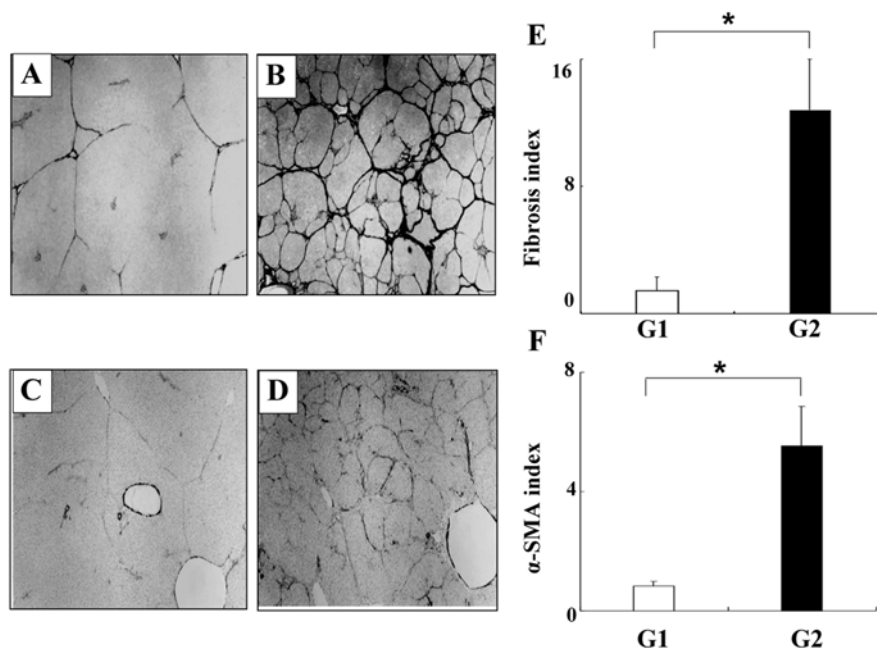


Figure 1. Representative microphotographs of liver fibrosis development and HSC activation. Pig serum resulted in marked fibrosis development in the (B) OLETG G2 rats whereas little fibrosis development was observed in the (A) LETO G1 rats (Sirius-red staining; original magnification, $\times 40$). Similar to the findings of fibrosis development, the α -SMA-immunopositive-activated HSCs significantly increased in the liver of the (D) OLETG G2 rats as compared with the (C) LETO G1 rats (original magnification, $\times 40$). (E) Densitometric analysis showed that the fibrosis area almost corresponded to the histological findings. (F) Along with the histological findings, image analysis showed that the α -SMA-positive lesions significantly increased in the G2 rats (as compared with the G1 rats) at a similar magnitude to the liver fibrosis development. The data represent the mean \pm SD ($n=10$). Statistically significant difference between the indicated groups ($p<0.01$).

Table I. Characteristic features of the LETO and OLETG rats.

Group	LETO	OLETF
Body weight (g)	515.0 \pm 20.8	671.3 \pm 35.2 ^a
Liver weight (g)	14.0 \pm 1.6	25.8 \pm 4.3 ^a
Ratio (liver to body)	0.027 \pm 0.004	0.038 \pm 0.007 ^a
Albumin (g/dl)	3.9 \pm 0.3	3.7 \pm 0.5
T-Bil (mg/dl)	0.06 \pm 0.01	0.14 \pm 0.08
ALT (IU/l)	37.5 \pm 1.7	98.3 \pm 18.4 ^a
Glucose (mg/dl)	178.5 \pm 17.3	271.3 \pm 41.0 ^a
Insulin (nM/ml)	50.6 \pm 7.1	128.2 \pm 10.4 ^a
QUICKI	0.244 \pm 0.009	0.206 \pm 0.003 ^a

Values are the means \pm SD. Statistically different vs. the LETO rats ($^aP<0.01$).

(HUVECs) co-cultured with human diploid fibroblasts as described previously (29). The experimental procedure followed the instructions provided with the angiogenesis kit (Kurabo, Tokyo, Japan). Computer-assisted quantitation of tubule formation was performed as in the *in vivo* assay. The effect of glucose and insulin on extracellular signal-regulated kinase 1/2 (ERK1/2) phosphorylation in HUVECs was evaluated by the Cellular Activation of Signaling ELISA (CASE) kit (ActiveMotif, Tokyo, Japan), according to the manufacturer's protocol.

Statistical analysis. To assess the statistical significance of the inter-group differences in the quantitative data, Bonferroni's multiple comparison test was used after one-way ANOVA. This was followed by Barlett's test to determine the homology of variance.

Results

General findings in the OLETG and LETO rats. The characteristic features of the OLETG and LETO rats at the age of 20 weeks are shown in Table I. The final body weights and relative liver weights of the OLETG rats were both higher than those of the LETO rats. Regarding the serological data, the serum alanine aminotransferase (ALT) was significantly higher in the OLETG rats than that in the LETO rats, whereas no significant differences in the levels of albumin and total bilirubin were observed. Considering the IR status, the serum levels of glucose and insulin were significantly higher in the OLETG than those in the LETO rats. The value of QUICKI in the OLETG rats was significantly lower than that in the LETO rats, confirming the IR status in the OLETG rats as described previously (20).

Liver fibrosis development in the pig serum-treated rats. Histological examination revealed that 4-week treatment with pig serum resulted in a marked fibrosis development in the OLETG rats (G2) whereas little fibrosis development was observed in the LETO rats (G1) (Fig. 1B and A, respectively). We next carried out the α -SMA immunohistochemical analysis to examine the effects of IR status on HSC activation. The

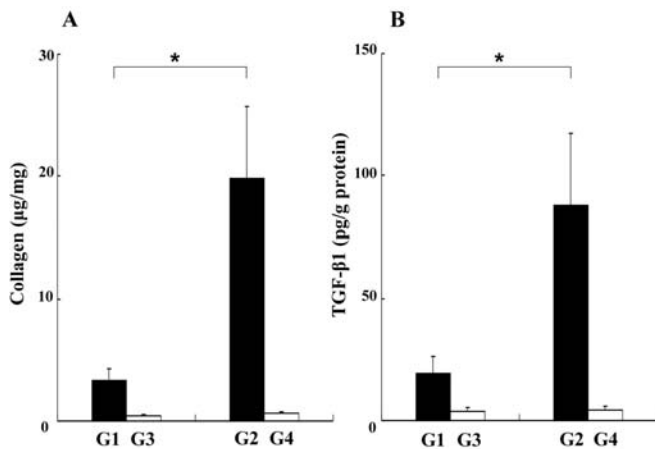


Figure 2. The effect of IR status on the hepatic collagen content (A) and TGF- β_1 expression (B) in the liver of the LETO (G1) and OLETF (G2) rats. Similar to the fibrotic area, hepatic collagen and TGF- β_1 significantly increased in the G2 rats as compared with the G1 rats. The magnitudes of these indices were almost similar to that of α -SMA expression. No increase in hepatic collagen and TGF- β_1 was observed in the liver of the PBS-treated LETO and OLETF rats (G3 and G4, respectively). The data represent the mean \pm SD (n=10). Statistically significant difference between the indicated groups ($^*p<0.01$).

α -SMA-immunopositive-activated HSCs were more markedly increased in G2 than in G1 (Fig. 1D and C, respectively). The densitometric analysis confirmed that the activated HSCs increased significantly at a similar magnitude to the liver fibrosis development (Fig. 1E and F, respectively). The hepatic collagen content and TGF- β_1 expression also increased in G2 as compared with G1 (Fig. 2A and B, respectively). No fibrosis development was found in the control groups of the OLETF and LETO rats, which received PBS (G3 and G4, respectively).

Effects of high glucose and insulin on the proliferation of HSCs in vitro. To evaluate whether the IR status, i.e., high glucose and insulin, directly affected the activated HSCs, we examined the effects of high glucose and insulin on the proliferation of the activated HSCs *in vitro*. We measured BrdU incorporation of HSCs incubated with glucose and insulin at various concentrations (100, 200, 400, 600 mg/dl and 1, 10, 100, 1000 nM, respectively) for 24 h. As shown in Fig. 3A and B, glucose and insulin treatment stimulated the proliferation of the activated HSCs in a dose-dependent manner. We next examined the combined effect of glucose and insulin on the proliferation of the activated HSCs. Either single treatment with glucose (271.3 mg/dl) or insulin (128.2 nM), which almost corresponded to the serum levels in the OLETF rats, stimulated the proliferation of the activated HSCs, and co-existence of both conditions exerted a more potent effect as compared with either single condition (Fig. 3C). We also observed that neither serum levels of insulin nor glucose in the OLETF rats promoted proliferation of quiescent primary HSCs (data not shown).

Tumor development in DEN-treated rat liver. In the hepatocarcinogenesis model, the GST-P-positive pre-neoplastic lesions in the OLETF rats (G6) were generally larger than those in the LETO rats (G5) (Fig. 4B and A, respectively). As shown in Fig. 4C, the mean size of the pre-neoplastic lesions in the G6 rats was significantly larger than that in the G5 rats. On the other hand, no marked differences were observed in the number of the GST-P-positive lesions between the G5 and G6 rats (Fig. 4D). No GST-P-positive lesions developed in the PBS-treated OLETF and LETO rats (G7 and G8, respectively) (data not shown).

Effects of high glucose and insulin on neovascularization. Since neovascularization has been shown to play an important

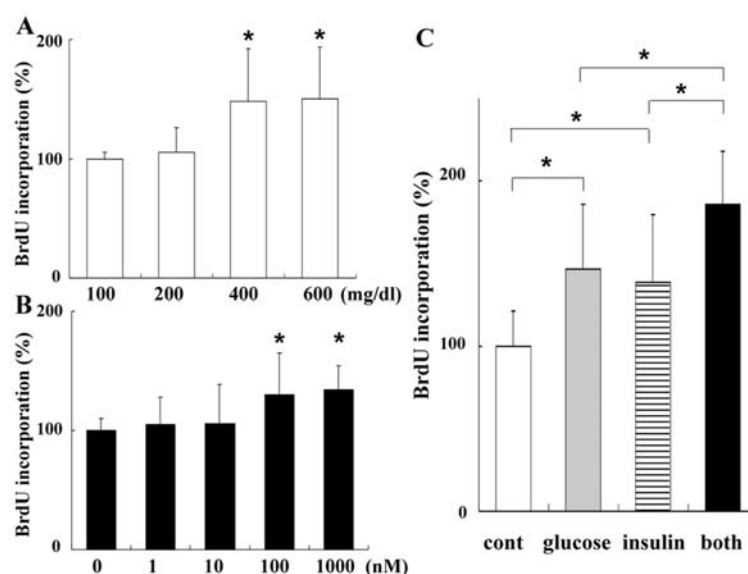


Figure 3. The effects of glucose (A) and insulin (B) on the proliferation of activated HSCs *in vitro*. Proliferation was measured in the presence of glucose (A) and insulin (B) at various concentrations (100, 200, 400, 600 mg/dl and 1, 10, 100, 1000 nM, respectively). Both glucose and insulin treatment stimulated the proliferation of activated HSCs in a dose-dependent manner. (C) The effect of co-existence of glucose and insulin on the proliferation of activated HSCs *in vitro*. The combination treatment of glucose and insulin significantly stimulated the proliferation of activated HSCs as compared with either single treatment. The concentrations of glucose and insulin were 271.3 mg/dl and 128.2 nM, respectively, which almost corresponded to the serum levels in the OLETF rats. The data represent the mean \pm SD (n=12). Statistically significant difference between the indicated groups ($^*p<0.01$).

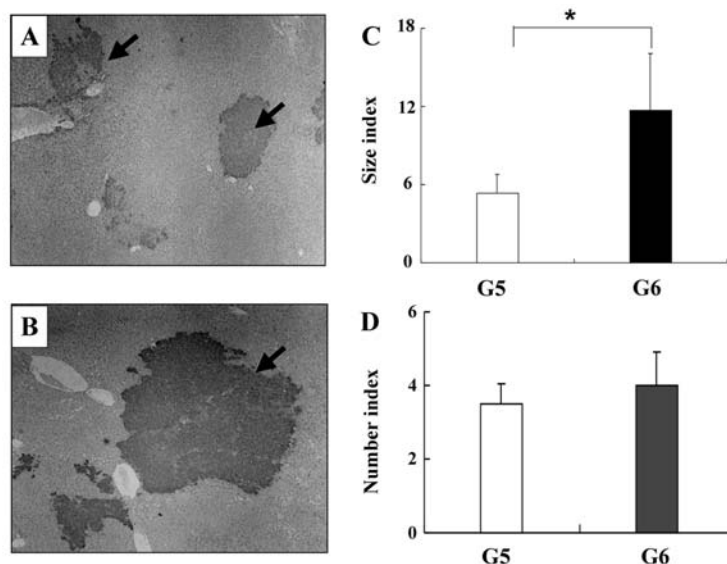


Figure 4. Representative microphotographs and indices of the GST-P-positive pre-neoplastic lesions in the liver. (A) In the DEN-treated LETO (G5) rats, GST-P-positive lesions moderately developed. (B) A marked development of pre-neoplastic lesions was observed in the DEN-treated OLETF (G6) rats as compared with the LETO rats. The arrows indicate the GST-P-positive pre-neoplastic lesions (original magnification, x40). (C and D) The mean size of the pre-neoplastic lesions in the G6 rats was significantly larger than that in the G5 rats, whereas no marked differences were observed in the number of GST-P-positive lesions between the G5 and G6 rats. The data represent the mean \pm SD (n=10). Statistically significant difference between the indicated groups (* p <0.01).

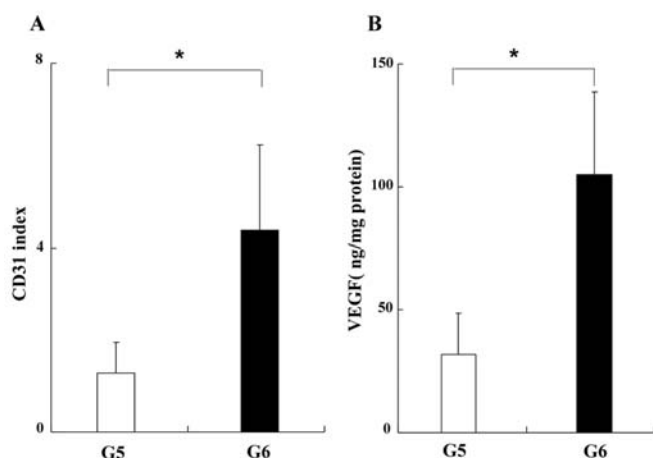


Figure 5. The effects of the IR status on the development of neovascularization and VEGF in the liver. Similar to the development of GST-P-positive lesions, hepatic neovascularization, which was evaluated with immunohistochemical analysis with CD31 (A) and VEGF expression (B) were significantly higher in the OLETF (G6) than in the LETO (G5) rats. The expression levels of neovascularization and VEGF in G6 rats were similar to those during the development of GST-P-positive pre-neoplastic lesions. MV, microvessel. The data represent the mean \pm SD (n=10). Statistically significant difference between the indicated groups (* p <0.01).

role in the hepatocarcinogenesis, we investigated the possible involvement of neovascularization in the current study. We first examined the effect of IR on neovascularization in the liver to elucidate whether or not the augmented effect of IR on GST-P-positive lesions is associated with alteration of angiogenesis. We observed that the number of CD31-immunopositive neo-vessels was much higher in the liver of the OLETF (G6) than in the LETO (G5) rats (Fig. 5A). These

neo-vessels were mainly localized in the pre-neoplastic lesions. We also performed an immunohistochemical analysis with another marker of neovascularization (CD105, endoglin), and the results were similar to those of CD31 (data not shown). Hepatic VEGF expression was also significantly higher in the G6 than in the G5 rats (Fig. 5B). The magnitude of VEGF augmentation was of a similar level to the development of neovascularization.

To elucidate the direct interaction between high glucose, insulin and angiogenesis, we next performed a set of *in vitro* experiments. Either high glucose or insulin treatment stimulated EC tubular formation, and the combination treatment with glucose and insulin exerted a more potent angiogenic activity as compared with the single condition. These promoting effects were suppressed by treatment with VEGF-neutralizing monoclonal antibody (VEGF-mAb) (Fig. 6). Since high glucose and insulin reportedly stimulate tubular formation of ECs via ERK 1/2 (18), we also examined the effect of glucose and insulin at doses corresponding to the serum levels in the OLETF rats. As shown in Fig. 7, ERK1/2 phosphorylation significantly increased in either glucose- or insulin-treated ECs, and the combination treatment of glucose and insulin exerted more ERK1/2 phosphorylation as compared with either single treatment. Similar to tubular formation, this augmentation was almost attenuated by treatment with VEGF mAb.

Discussion

Liver cirrhosis and HCC are known as terminal clinical manifestations in chronic liver diseases regardless of the etiology (1,2,30). Recent studies have revealed a possible interaction between these entities and IR; i.e., co-existence of high glucose and insulin. In the current study, we found that

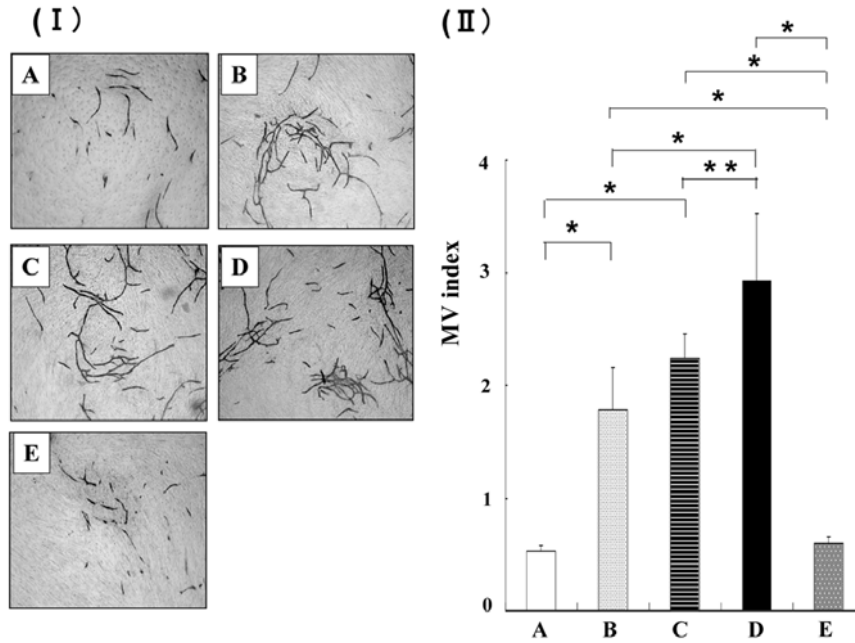


Figure 6. (I) Representative features and index of *in vitro* EC tubular formation following treatment with glucose and/or insulin. (A) PBS-treated control group. (B and C) Glucose (271.3 mg/dl)- and insulin (128.2 nM)-treated groups, respectively. (D) Combination-treated group. Treatment with either glucose or insulin increased the EC tubule formation as compared with the PBS-treated control group. The combination treatment with glucose and insulin caused a further augmentation as compared with either single agent treatment. (E) These promoting effects were mostly attenuated by treatment with VEGF-neutralizing monoclonal antibody (VEGF mAb). (II) Semi-quantitative analysis with image analyzer system confirmed the above-mentioned findings. MV, microvessel. The data are expressed as the mean \pm SD (n=12). Statistically significant differences between the indicated groups (* p <0.01 and ** p <0.05, respectively).

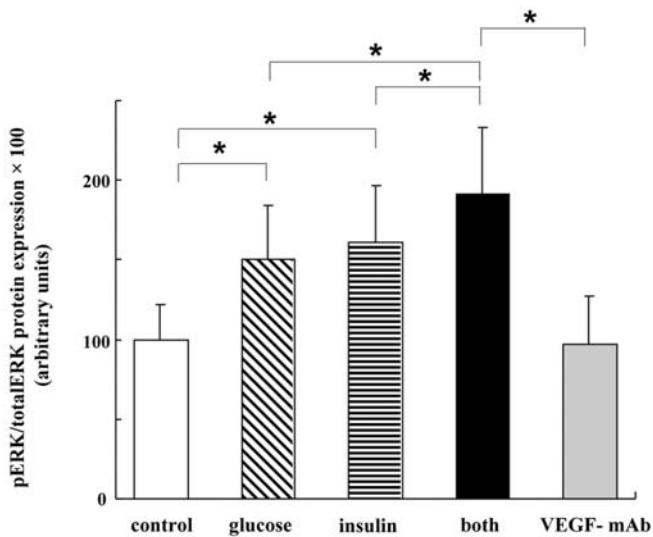



Figure 7. Glucose- and insulin-induced ERK1/2 phosphorylation in the EC. ERK1/2 phosphorylation was significantly increased in either glucose- or insulin-treated EC, and the combination treatment of glucose and insulin exerted more ERK1/2 phosphorylation as compared with either single treatment. Similar to tubular formation, this augmentation was almost attenuated by treatment with VEGF-neutralizing monoclonal antibody (VEGF-mAb). The results are expressed as the ratio of phosphorylated ERK1/2 (pERK) to the total ERK1/2 in HUVEC. The data represent the mean \pm SD (n=8). Statistically significant difference between the indicated groups (* p <0.01).

liver fibrosis development and hepatocarcinogenesis significantly progressed in the presence of IR. It is known that activated HSCs play a central role on the basis of their ability

to undergo activation during development of liver fibrosis, and that HSCs produce several important molecules such as TGF- β_1 (31,32). Previous investigations have shown that high glucose and insulin stimulated the proliferation of HSCs (12,13). In this study, we also observed that both glucose and insulin stimulated the proliferation of activated HSCs. Furthermore, we first noticed that the combination treatment with glucose and insulin significantly promoted the proliferation of activated HSCs as compared with either single agent. We did not detect these promoting effects of glucose and insulin on quiescent HSCs. The exact reason of this discrepancy is not clear at this time. It has been reported that insulin receptors were up-regulated during the activation of HSCs (13), and that high glucose stimulated the proliferation at least partly through the NADPH oxidase-mediated reactive oxygen species (ROS), which is known to be associated with the activation of HSCs (12). These phenomena may be involved. Recently, it has been reported that the IR status accelerated the development of steatohepatitis and fibrosis in the NASH experimental model (33). The study showed a close relationship between steatohepatitis and fibrosis through up-regulation of lipogenesis-related genes. In contrast, in the current study, we did not observe steatosis or steatohepatitis in the OLETF and LETO rats. The pig serum model induces liver fibrosis without severe inflammation (34). As we focused mainly on the effect of IR without any additional effect such as inflammation or steatohepatitis-induced cytokines, we employed this model in the present study.

We also observed the promoting effect of IR on the development of pre-neoplastic lesions along with augmentation of intrahepatic neovascularization and VEGF. We previously

 SPANDIDOS PUBLICATIONS hat hepatic neovascularization and VEGF exhibited increase during hepatocarcinogenesis (17). A recent

study suggested that alterations in the hepatic micro-circulation occur at a very early stage of liver carcinogenesis (35). As such, we found that neovascularization was augmented mainly in the GST-P-positive pre-neoplastic lesions. Notably, the size of the pre-neoplastic lesions significantly increased under IR whereas the number was not altered. It is possible that the IR status is involved in the so called 'angiogenic switch' rather than in cell transformation itself. It has been reported that both insulin and glucose can induce VEGF expression (36,37). In this study, we found an additive effect of high insulin and glucose on EC tubular formation, and this promoting effect was markedly cancelled by treatment with VEGF mAb. These results indicated that the IR status induced neovascularization at least partly via VEGF. It has been reported that high glucose and insulin stimulate tubular formation of ECs via the ERK1/2 pathway (18). In the current study, we also observed that high glucose and insulin stimulated the ERK1/2 phosphorylation of ECs, and these augmentations were almost attenuated by treatment with VEGF mAb, indicating that the glucose- and insulin-mediated ERK1/2 phosphorylation was, at least partly, mediated by VEGF.

Although previous studies conducted to determine the molecular processes associated with fibrosis and angiogenesis were performed independently, recent studies have revealed that both biological phenomena emerge synergistically (38). We and other groups reported that neovascularization significantly increased during liver fibrosis development (38-40). As in the ECs, VEGF has been shown to act on activated HSCs (39,40). Accordingly, it is possible that augmentation of VEGF-mediated neo-vascularization may also play some role in the development of IR-induced liver fibrosis development. Further studies to elucidate the exact mechanism are required in the future.

In conclusion, we demonstrated that the IR status directly accelerated liver fibrosis development and hepatocarcinogenesis via stimulation of activated HSCs and neo-vascularization, respectively. The augmentation of neo-vascularization was associated with VEGF up-regulation. Our *in vitro* study revealed that high glucose and insulin increased the proliferation of HSCs and EC tubular formation. The co-existence of both factors exerted more potent activities than either single status. These results suggested that an anti-IR therapy may become an alternative new therapeutic strategy against the progression of chronic liver diseases.

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