

TUMORIGENESIS AND NEOPLASTIC PROGRESSION

Inhibition of Hepatic Stellate Cell Activation Suppresses Tumorigenicity of Hepatocellular Carcinoma in Mice

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Transdifferentiation (or activation) of hepatic stellate cells (HSCs) to myofibroblasts is a key event in liver fibrosis. Activated HSCs in the tumor microenvironment reportedly promote tumor progression. This study analyzed the effect of an inhibitor of HSC activation, retinol-binding protein-albumin domain III fusion protein (R-III), on protumorigenic functions of HSCs. Although conditioned medium collected from activated HSCs enhanced the migration, invasion, and proliferation of the hepatocellular carcinoma cell line Hepa-1c1c7, this effect was not observed in Hepa-1c1c7 cells treated with conditioned medium from R-III—exposed HSCs. In a subcutaneous tumor model, larger tumors with increased vascular density were formed in mice transplanted with Hepa-1c1c7+HSC than in mice transplanted with Hepa-1c1c7 cells alone. Intriguingly, when Hepa-1c1c7+HSC-transplanted mice were injected intravenously with R-III, a reduction in vascular density and extended tumor necrosis were observed. In an orthotopic tumor model, co-transplantation of HSCs enhanced tumor growth, angiogenesis, and regional metastasis accompanied by increased peritumoral lymphatic vessel density, which was abolished by R-III. In vitro study showed that R-III treatment affected the synthesis of pro-angiogenic and anti-angiogenic factors in activated HSCs, which might be the potential mechanism underlying the R-III effect. These findings suggest that the inhibition of HSC activation abrogates HSC-induced tumor angiogenesis and growth, which represents an attractive therapeutic strategy. (Am J Pathol 2021, 191: 2219-2230; https://doi.org/10.1016/j.ajpath.2021.08.004)

Cancers develop in complex tissue environments, which is essential for sustained growth, invasion, and metastasis.¹ Tumor angiogenesis is an essential process for tumor progression, and lymphatic vessels provide an alternate route for tumor cell dissemination.² The tumor microenvironment (TME) comprises a mass of heterogeneous cell types, among which the two most prominent types of protumorigenic cells are cancer-associated fibroblasts (CAFs) and tumor-associated macrophages.³ CAFs, also called myofibroblasts, are reportedly associated with the progression of several types of cancers, and can originate as a result of the activation of resident fibroblasts, bone marrow-derived fibrocytes, epithelial cells, endothelial cells, or from certain specialized cells such as stellate cells (SCs) in the pancreas and liver.⁴ The presence of activated SCs has been shown in the stroma surrounding cancer cells, and bidirectional interactions between SCs and cancer cells, by which tumor-derived factors activate SCs, and, in turn, activated SCs promote metastatic growth.^{5,6} Conditioned media from activated SCs promotes the proliferation, migration, and invasion of tumor cells in vitro, and co-transplantation of

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these activated SCs and tumor cells into mice resulted in an enlarged tumor mass that correlated with enhanced angiogenesis.^{7,8}

Hepatic stellate cells (HSCs) are pericytes residing in the space of Disse between the sinusoidal endothelial cells and the parenchymal cells, and constitute 5% to 10% of the total number of cells in the liver.⁹ In normal liver, HSCs maintain a nonproliferative, quiescent phenotype and store approximately 80% of vitamin A (retinol) in the whole body as retinyl esters in lipid droplets in the cytoplasm. In response to fibrogenic stimuli, HSCs become activated, transdifferentiating from vitamin A-storing cells to myofibroblast-like cells.¹⁰ Upon activation, HSCs lose cytoplasmic vitamin A-containing lipid droplets, proliferate vigorously, and produce a large amount of extracellular matrix proteins. When cultured on plastic, HSCs undergo spontaneous activation in vitro. Activation of HSCs is a widely accepted key event in liver fibrosis, which is characterized by excessive accumulation of extracellular matrix proteins.¹¹ Cirrhosis is the advanced stage of liver fibrosis and is the leading risk factor for the development of hepatocellular carcinoma (HCC).¹² Cells resembling HSCs were isolated from the pancreas in the late 1990s,¹³ and these pancreatic stellate cells also play a central role in pancreatic fibrogenesis in a manner similar to HSCs.¹⁴

Albumin is the most abundant plasma protein with a mol. wt. of approximately 66 kDa produced in the liver.¹⁵ It is composed of three homologous domains (I to III) and performs a variety of functions. A previous study showed that albumin was expressed in quiescent SCs, but not in activated SCs, and that its forced expression in activated SCs induced the phenotypic reversion to fat-storing, early activated cells.¹⁶ Building on these results, a recombinant fusion protein (designated R-III; mol. wt., approximately 45 kDa) was developed as an antifibrotic agent, in which the domain III of albumin was fused to the C-terminus of retinol-binding protein.¹⁷ Retinol-binding protein was adopted for targeted delivery to SCs because the protein and its membrane receptor STRA6 coordinate the cellular uptake of retinol into HSCs.¹⁸ A follow-up study showed that R-III inhibited SC activation in vitro and reduced liver and kidney fibrosis in vivo.^{19–21} In this study, the effect of R-III on the protumorigenic functions of activated HSCs was examined, and R-III was found to suppress HSC-induced HCC angiogenesis and growth.

Materials and Methods

Male BALB/c mice and female BALB/c-nu/nu mice were purchased from Orient Bio, Inc. (Sungnam, Korea) and maintained under temperature-, humidity-, and lightcontrolled conditions. All animal experimental protocols were approved by the institutional review board and performed in accordance with the NIH *Guide for the Care and Use of Laboratory Animals.*²² Mouse hepatoma Hepa-1c1c7 cells were obtained from the ATCC (Rockville, MD) and cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum. The fusion protein R-III (Supplemental Figure S1) was synthesized using a previously described method.¹⁹

Isolation of Mouse HSCs

HSCs were isolated from male BALB/c mice (age, 14 weeks) as described previously²¹ and cultured in DMEM supplemented with 10% fetal bovine serum. The purity of the HSCs was assessed by microscopic observation. The HSCs harvested on day 1 after seeding were used as preactivated HSCs. Cells were passaged before reaching 70% confluence in the primary culture and used as activated HSCs. The activation status of the HSCs was assessed on the basis of their increased expression of α -smooth muscle actin (α -SMA) and collagen type I, as well as through their morphologic changes.

Collection of Conditioned Medium

HSCs after passage 1 (activated HSCs) were incubated for 24 hours in low-serum DMEM (containing 0.5% fetal bovine serum and 0.75% bovine serum albumin) in the presence or absence of R-III (0.25 μ mol/L). Conditioned medium from HSCs (HSC-CM) was centrifuged for 10 minutes at 750 \times g, passed through a sterile filter, and frozen at -80° C for future use. Low-serum DMEM without cells incubated for 24 hours in cell culture dishes was used as the negative control.

Monolayer Wound Healing Assay

HSCs were grown to confluency in 24-well plates and mechanically scratched using a sterile 200- μ L pipet tip. The media and the dislodged cells were aspirated and replaced by CM. After incubation at 37°C for 24 or 48 hours, the cells were photographed using a microscope in five random fields. The width of the wounded cell monolayers was measured from the images, and the rates of migration were calculated using the following formula: % migration = (initial scratch wound width – scratch wound width at time point X)/initial scratch wound width \times 100.

Cell Invasion Assay

Biocoat Matrigel-coated invasion chambers (BD Biosciences, Bedford, MA) were used to examine cell invasiveness. Briefly, Hepa-1c1c7 cells (2.5×10^4) in 500 µL serum-free medium were seeded into the upper chamber, and 700 µL CM was added to the lower chamber. The cells were allowed to invade the Matrigel for 20 hours at 37°C. The noninvading cells on the upper surface of the membrane were removed with a cotton swab. Cells were fixed with 4% paraformaldehyde, stained with 0.5% crystal violet, and air-dried. Invading cells were counted for each membrane in four adjacent microscopic fields.

Cell Proliferation

Hepa-1c1c7 cells (8×10^3 cells/well) were seeded into 96well plates, incubated at 37°C for 24 hours, and then treated with CM. After 24 hours of treatment, cell viability was measured using the Cell Counting Kit-8 (Dojindo Laboratories, Tokyo, Japan).

Mouse HCC Subcutaneous Implantation Model

Six-week-old female BALB/c-nu/nu mice were injected subcutaneously in the flank with 1×10^6 Hepa-1c1c7 cells alone (n = 6) or in combination with 2×10^5 activated HSCs (n = 12). Hepa-1c1c7+HSC-transplanted mice were divided randomly into 2 groups (n = 6 per group) and injected via the tail vein with saline or R-III (30 µg) once every 3 days, starting on day 6 after tumor cell transplantation. Mice were euthanized 22 days after injection. Tumor growth was monitored twice a week with electronic calipers, and the tumor volume was calculated using the following standard formula: tumor volume $(mm^3) = longest diameter \times (shortest diameter)^2 \times 0.5.$ Hematoxylin and eosin-stained slides of tumors were subjected to reading and determination of areas of necrosis, and necrotic area was calculated with ImageJ software version 1.53a (NIH, Bethesda, MD; http://imagej.nih.gov/ ij). All experiments were performed in duplicate.

Mouse Hepatocellular Carcinoma Orthotopic Implantation Model

Six-week-old female BALB/c-nu/nu mice were used for implantation. Orthotopic implantation was performed as previously described.²³ Briefly, after anesthetizing the mice (n = 18), the median lobe of the liver was exposed and a piece of sterile Gelfoam (Technew, Rio de Janeiro, Brazil) was placed to maintain the surface moisture in the liver and to prevent leakage and bleeding. Small-volume Matrigel suspensions (20 μ L) containing 1 \times 10⁶ Hepa-1c1c7 cells alone (n = 6) or in combination with 2×10^5 HSCs (n = 12) were injected into the subcapsular region of the liver parenchyma in the median lobe. The mice injected with Hepa-1c1c7+HSC were divided randomly into 2 groups (n = 6 per group) and injected via tail vein with saline or R-III (30 µg) once every 3 days, starting on day 6 after tumor transplantation. Mice were euthanized 35 days after surgery and weighed. The livers were harvested, and the total liver tumor volume was calculated as the sum of the volumes of all tumors; tumor volume (mm³). To assess the metastatic spread of green fluorescent protein (GFP)-labeled Hepa-1c1c7 cells, lung and pancreas were examined ex vivo using the EVOS FL imaging system (Thermo Fisher Scientific, Bothell, WA).

Generation of a Stable GFP-Expression Hepa-1c1c7 Cell

Hepa-1c1c7 cells were transfected with pEGFP vector using Lipofectamine 2000 (Invitrogen, Carlsbad, CA) and subject to drug selection by the addition of G418 to the medium. The high-GFP—expressing cell among stably transfected cells was selected and used for the experiments.

Immunohistochemistry and Immunofluorescence

Sections (5-µm thick) of formalin-fixed, paraffin-embedded tissues were prepared and subjected to immunohistochemistry with the antibodies listed in Table 1. Seven randomly selected high-power microscopic fields were photographed, and immunohistochemistry staining intensities were quantified using a three-tiered system of none-to-weak, moderate, and strong immunoreactivity, as described previously.²⁴ For immunofluorescence, after antibody incubation, nuclei staining was performed with DAPI. The cells were observed under a Leica TCS SP8 microscope and the fluorescent intensity quantification was analyzed using LasX software (Leica, Mannheim, Germany).

Quantitative Real-Time PCR

Total RNA was prepared using TRIzol (Ambion, Austin, TX), and was used to synthesize the cDNA. Real-time PCR was performed on an ABI QuantStudio 3 Real-Time PCR System (Thermo Fisher Scientific, Waltham, WA). To control for variations in the reaction, the PCR products were normalized against the mRNA levels of glyceraldehyde 3-phosphate dehydrogenase. The primers used are listed in Table $2.^{25-28}$

 Table 1
 Primary Antibodies Used in This Study

	Primary	Company, catalog no.
Method	antibody	(location)
Immunohistochemistry	CD31	Cell Signaling, 77699 (Beverly, MA)
	LYVE-1	Novus Biological, NB100- 725 (Centennial, CO)
	Ki-67	Abcam, ab16667 (Cambridge, MA)
Immunofluorescence	α-SMA	Abcam, ab32575
	Cygb/STAP	Generous gift from Dr. Norifumi Kawada (Osak City University)
	PDGFRa	Abcam, ab93531
Western blot	Histidine ⁴⁹ -tag	AB Frontier, LF-MA-20209 (Seoul, Korea)
	α-tubulin	Cell Signaling, 2125

 $\alpha\text{-SMA},$ $\alpha\text{-smooth}$ muscle actin; Cygb/STAP, cytoglobin/stellate cell activation-associated protein (Cygb/STAP); LYVE-1, lymphatic vessel endothelial receptor-1; PDGFR α , platelet-derived growth factor receptor alpha.

Western Blot Analysis

Cell lysates were prepared for analysis by electrophoresis and immunoblotting as described previously.¹⁹ The primary antibodies used are listed in Table 1.

Proteome Profiler Angiogenesis Antibody Array

The proteome profiler mouse angiogenesis antibody array was performed following the manufacturer's instructions (R&D Systems, Minneapolis, MN).

Establishment of Primary Culture from Tumor Tissue and CAF Isolation

Primary culture was established as previously described.²⁹ Briefly, tumor tissues were dissected from the livers of Hepa-1c1c7+HSC-injected mice and minced with a curved scissor. After enzymatic digestion with collagenase, the tissue/media mixture was centrifuged at $450 \times g$ for 5 minutes at 4°C and resuspended in red blood cell lysis buffer. After centrifugation, cell pellets were resuspended in DMEM supplemented with 10% fetal bovine serum and seeded on culture plates to establish primary cultures. For CAF isolation, cell pellets were resuspended in 0.5 mL fluorescence-activated cell sorting buffer. After removing 0.05 mL for use as unstained control, the rest of the solution was incubated with anti-phycoerythrin-conjugated platelet-derived growth factor receptor alpha (PDGFR α) antibody (ab93531, 10 µg/mL; Abcam, Cambridge, MA) for 1 hour at 4°C. The cells were washed with fluorescence-activated cell sorting buffer and the PDGFRa-positive cells were sorted using BD FACS Aria III (BD Biosciences, San Jose, CA). These cell-sorting experiments were performed using the sort precision mode set to purity mode to exclude droplets containing unwanted cells.

Statistical Analysis

Results are expressed as means \pm SD. A two-way analysis of variance with the Dunnett multiple comparison test was used for *in vitro* experiments. Data from *in vivo* studies were analyzed by either the one-way ANOVA, Wilcoxon rank-sum test, or the Kruskal-Wallis test, followed by the Dwass-Steel-Critchlow-Fligner multiple comparison. A *P* value < 0.05 was considered statistically significant.

Results

Effect of R-III on the Protumorigenic Functions of Activated HSCs with Respect to the HCC Cell Line Hepa-1c1c7 *in Vitro*

Because activated HSCs in the TME promote tumor progression,⁶ the study investigated whether the recombinant

Table 2	Primers	Used	in	This	Study
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Gene name	Sequence	Reference
ACTA2	F: 5'-CCAGCACCATGAAGATCAAG-3'	
	R: 5'-tggaaggtagacagcgaagc-3'	
COL1A1	F: 5'-CGACCTCAAGATGTGCCACT-3'	
	R: 5'-CTTGGTTAGGGTCGATCCAG-3'	
GFAP	F: 5'-CACGAACGAGTCCCTAGAGC-3'	
	R: 5'-gcaagtgcctcctggtaact-3'	
P21	F: 5'-CAGATCCACAGCGATATCCA-3'	
	R: 5'-AGACAACGGCACACTTTCGT-3'	
TIMP1	F: 5'-GCATCTGGCATCCTCTTGTT-3'	
	R: 5'-TGGGGAACCCATGAATTTAG-3'	
TIMP2	F: 5'-TTCTTGACATCGAGGACCCG-3'	
	R: 5'-AGCGTGAACCCACTTGGATG-3'	
VEGFR2	F: 5'-TGGGCAGTCAAGTCCGAATC-3'	
	R: 5'-TTGGACTCAATGGGCCTTCC-3'	
PDGFD	F: 5'-GATGACTACTTTGTGGCAAAACC-3'	
	R: 5'-GTGACTGATTCCCAGTTGGTC-3'	
PDGFC	F: 5'-GCCCGAAGTTTCCTCATACA-3'	25
	R: 5'-ACACTTCCATCACTGGGCTC-3'	
VEGFA	F: 5'-AGTCCCATGAAGTGATCAAGTTCA-3'	25
	R: 5'-ATCCGCATGATCTGCATGG-3'	
VEGFD	F: 5'-GAACACAAGCACCTCCTACAT-3'	26
	R: 5'-CACAGGCTGGCTTTCTACTT-3'	
CXCL12	F: 5'-GAGCCAACGTCAAGCATCTG-3'	27
	R: 5'-CGGGTCAATGCACACTTGTC-3'	
ANGPT1	F: 5'-gggggaggttggacagtaa-3'	28
	R: 5'-CATCAGCTCAATCCTCAGC-3'	

F, forward; R, reverse.

fusion protein R-III suppresses the protumorigenic functions of HSCs. HSCs were isolated from the liver of BALB/c mice by enzyme perfusion/density gradient centrifugation and culture-activated. In agreement with previous studies,¹⁷ real-time PCR showed that the mRNA expression of α -SMA and collagen type I, well-known markers of HSC activation,^{30,31} was greatly increased in HSCs after passage 1 (activated HSCs) compared with that in HSCs on day 1 after seeding (pre-activated HSCs). This was decreased in HSCs after R-III treatment (Figure 1A). By contrast, the mRNA expression of glial fibrillary acidic protein, a marker of quiescent HSCs,³² was decreased in activated HSCs. CM was collected from activated HSCs with or without R-III treatment for 24 hours, and Hepa-1c17 cells (HCC cell line) were treated with CM. CM from activated HSCs enhanced the migration, invasion, and proliferation of Hepa-1c1c7 cells (Figure 1, B-D), similar to that observed in different HCC cell lines.^{7,33,34} In contrast, CM from R-III-treated HSCs did not exert significant stimulatory effects on tumor cell migration, invasion, or proliferation. The expression of cellular senescence markers,³⁵ such as p21, tissue inhibitor of metalloproteinase-1 (TIMP-1), and TIMP-2, was not changed significantly, thereby indicating that this is perhaps not a result of tumor cell senescence by CM from R-III-treated HSCs (Figure 1E). The change in growth



Figure 1 R-III reduces the protumorigenic effects of activated hepatic stellate cells (HSCs) *in vitro*. **A** and **F**: Total RNA was isolated from HSCs on day 1 after seeding (pre-activated), and from HSCs after passage 1 (activated) with or without R-III treatment (0.25 μ mol/L) for 24 hours. Real-time PCR was used to analyze mRNA levels of collagen type I, α -smooth muscle actin (α -SMA), and glial fibrillary acidic protein (GFAP) (**A**), and platelet-derived growth factor (PDGF)-c, PDGF-d, and CXCL12 (**F**). **B**–**D**: Conditioned medium (CM) was collected from activated HSCs with or without R-III treatment, and was used to stimulate the Hepa-1c1c7 cells. The migratory and invasive abilities of the cells were analyzed using a monolayer wound healing assay (**B**) and a Matrigel invasion assay (**C**), respectively. **B: Black lines** indicate the margins of migrating cells. **D:** The Cell Counting Kit-8 was used to analyze cell proliferation. **E:** Hepa-1c1c7 cells were incubated with CM from activated HSCs that were treated with or without R-III for 24 hours, and the mRNA levels of p21, tissue inhibitor of metalloproteinase-1 (TIMP-1), and TIMP-2 were analyzed by real-time PCR. Data are expressed as means \pm SD for three independent experiments. **P* < 0.05, ***P* < 0.01, and ****P* < 0.001 by two-way analysis of variance (ANOVA) with the Dunnett multiple comparison test. Scale bars: 400 μ m (**B**); 200 μ m (**C**). Ctrl, control.

factor expression was assessed because of previous reports indicating that activated HSCs stimulate the migration, invasion, and proliferation of cancer cells by producing an increased number of growth factors and cytokines.^{6,8,36} The mRNA expression of PDGF-c, PDGF-d, and CXCL12 (also known as stromal cell—derived factor 1) was found to be up-regulated during HSC activation, but was down-regulated in response to R-III (Figure 1F). It should be noted that R-III did not enter Hepa-1c1c7 cells or affect their growth (Supplemental Figure S2), which is consistent with the previous report that R-III does not enter carcinoma cells.¹⁷ These findings show that R-III—mediated inhibition of HSC activation is accompanied by reduced expression of PDGF-c, PDGF-d, and CXCL12, which correlates with a decrease in the protumorigenic functions of HSCs.

Effect of R-III on the Protumorigenic Functions of HSCs in the Subcutaneous Tumor Model

To explore the antitumor activity of R-III *in vivo*, a subcutaneous xenograft model was used. Hepa-1c1c7 cells $(1 \times 10^6 \text{ cells})$ were transplanted either alone or in combination with activated HSCs ($2 \times 10^5 \text{ cells}$) into the flanks of nude mice (ratio of Hepa-1c1c7:HSCs, 5:1), and the Hepa-1c1c7+HSC-transplanted mice were injected intravenously with saline or R-III once every 3 days, starting on day 6 after transplantation. Palpable tumors were not detected in mice injected with HSCs alone. Although the Hepa-1c1c7 cells alone formed tumors, the tumors were larger when these cells were transplanted along with HSCs (Figure 2, A–C). Importantly, intravenous administration of R-III induced large areas of hemorrhagic necrosis within the tumors, which was accompanied with reduced tumor weight (Figure 2, C-E). It is well known that tumor necrosis results from tissue hypoxia that occurs during tumor progression owing to rapid tumor growth and an inadequate blood supply. Because R-III did not induce apoptotic cell death (terminal deoxynucleotidyl transferase-mediated dUTP nick-end labeling assay, data not shown), the study tested whether R-III could affect tumor angiogenesis. Consistent with previous reports,^{33,34} co-transplantation with HSCs significantly promoted tumor angiogenesis and proliferation, as shown by immunohistochemistry for CD31 and Ki-67, a cellular proliferation marker (Figure 2, F and G). However, this effect of activated HSCs was abrogated by R-III administration. Real-time PCR analysis showed that the expression level of vascular endothelial growth factor (VEGF) receptor-2 in tumors roughly paralleled the intensity of CD31 immunostaining (Figure 2H).



Figure 2 R-III induces necrosis in subcutaneous tumors. **A:** Representative images of subcutaneous tumors in mice transplanted with Hepa-1c1c7 alone or with Hepa-1c1c7+hepatic stellate cells (HSCs) in the presence or absence of R-III treatment. **B:** Tumor volumes were measured at the indicated time points. **C:** Tumor weights were determined after being harvested from the mice 22 days after tumor transplantation. **D** and **E:** Tumor sections were stained with hematoxylin and eosin, and the necrotic areas are indicated by **black dashed lines** and measured. R-III induces large areas of hemorrhagic necrosis within tumors. **F** and **G:** Tumor sections were subjected to immunohistochemistry with CD31 and Ki-67 antibodies. **G:** Semiquantitative analysis of each immunostaining. **H:** Total RNA was extracted from the tumors, and the expression level of vascular endothelial growth factor receptor (VEGFR)-2 was analyzed by real-time PCR. Data are expressed as means \pm SD by one-way analysis of variance (**C**) and by the Kruskal-Wallis test, followed by Dwass-Steel-Critchlow-Fligner multiple comparison (**H**). **P* < 0.05, ***P* < 0.01, and ****P* < 0.001. Scale bars: 5 mm (**A**); 50 µm (**F**).

Effect of R-III on the Protumorigenic Functions of HSCs in the Orthotopic Tumor Model

To assess the growth and metastasis of HCC cells, an orthotopic HCC model in nude/athymic mice was established using GFP-expressing HCC reporter cell lines. Hepa-1c1c7 cells, either alone or in combination with activated HSCs, were transplanted directly into the median liver lobe of nude mice. When co-transplanted with both Hepa-1c1c7 and HSCs, the orthotopic tumors grew larger and bloody ascites developed in all mice 3 to 5 weeks after transplantation (Figure 3A). When evaluating the metastatic spread, GFP-positive cells were detected in the enlarged lymph nodes-near the pancreas-only in the cotransplanted mice, which indicates occurrence of regional metastasis (Figure 3). Lung metastasis was not detected. On the other hand, mice treated with R-III had smaller tumors, developed lesser ascites, and had fewer GFP-positive lymph nodes, indicating that R-III inhibited HSC-induced HCC tumorigenicity. There were no significant differences in the body weights of treatment groups (Supplemental Figure S3).

Inhibition of Tumor Angiogenesis and Lymphangiogenesis by R-III

Because lymphangiogenesis is another mode of vascularization in tumors and lymphatic vessels represent an alternate route for dissemination of cancer cells,³⁷ sections of orthotopic tumors were immunostained with antibodies against lymphatic vessel endothelial receptor-1 (LYVE-1), a lymphatic endothelial cell marker, as well as CD31. Intratumoral and peritumoral blood vessel density, and peritumoral lymphatic vessel density, were increased significantly in tumors from Hepa-1c1c7+HSC-transplanted mice compared with that in tumors from mice that received Hepa-1c1c7 alone (Figure 4). The occurrence of intratumoral lymphatic vessels was rare. However, R-III administration abrogated the effect of HSCs on vascular density.

Regulation of Pro-Angiogenic and Anti-Angiogenic Factors in HSCs by R-III

The *in vivo* results from both subcutaneous and orthotopic tumor models showed that co-transplantation with HSCs significantly promoted tumor angiogenesis, which was abolished by R-III administration. Because activated HSCs reportedly release multiple pro-angiogenic factors into the TME,^{34,36,38,39} the expression of these factors was assessed by real-time PCR. It was found that, similar to the expression of PDGF-d and CXCL12 (Figure 1E), the expression of VEGF-d and angiopoietin-1 was upregulated during HSC activation, but down-regulated in response to R-III (Figure 5A). The expression of VEGF-a was increased slightly during HSC activation, but was not altered significantly with R-III. To determine the secretion of angiogenesis-related factors, CM from activated HSCs treated with or without R-III was analyzed by a proteome profiler angiogenesis antibody array. Intriguingly, R-III treatment enhanced the secretion of 12 proteins, such as ADAMTS1, angiogenin, Cyr61, endostatin, IGFBP-3, CXCL1, CXCL4, CXCL10, MMP-9, PDGF-AA,



Figure 3 R-III inhibits tumor progression in the orthotopic model. **A:** Representative images show the entire mouse and dissected organs such as gross liver and pancreas with surrounding lymph nodes. Tumor volumes were determined at 35 days after orthotopic transplantation of Hepa-1c1c7 (1×10^6) \pm hepatic stellate cells (HSCs) (2×10^5) (**right**). The **black** and **yellow arrows** indicate ascites and swollen lymph nodes, respectively. **B:** Fluorescence microscopy images show green fluorescence protein (GFP)-expressing Hepa-1c1c7 cells in lymph nodes near the pancreas (**left**, **yellow arrows**), and quantification of GFP-positive nodules is shown (**right**). Images were acquired using the EVOS FL Imaging System. **A** and **B:** Data are expressed as means \pm SD. **P* < 0.05 by Kruskal-Wallis test, followed by the Dwass-Steel-Critchlow-Fligner multiple comparison. [†]*P* < 0.05 by Wilcoxon rank-sum test. Scale bars: 5 mm (**A**); 2 mm (**B**).

thrombospondin-2, and TIMP-4, among which 7 proteins (ADAMTS-1, endostatin, IGFBP-3, CXCL4, CXCL10, thrombospondin-2, and TIMP-4) reportedly are angiogenesis inhibitors⁴⁰⁻⁴² (Figure 5, B and C). In contrast, the secretion of monocyte chemoattractant protein-1 (MCP-1), an angiogenic chemokine,⁴³ was decreased by R-III. Changes in the expression and secretion of proangiogenic and anti-angiogenic factors by R-III correlated with the inhibition of tumor angiogenesis and growth.

Inactivation of HSC-Derived CAFs by R-III

Immunostaining of Hepa-1c1c7+HSC orthotopically transplanted mice with antibodies against cytoglobin/stellate cell activation-associated protein (Cygb/STAP), an



Figure 4 R-III suppresses tumor angiogenesis and lymphangiogenesis. **A:** Tumor sections were stained with hematoxylin and eosin (H&E) and subjected to immunohistochemistry for CD31 and lymphatic vessel endothelial receptor-1 (LYVE-1). Selected areas (**dotted boxes**) of CD31 staining (intratumoral or peritumoral areas) and of LYVE-1 staining (peritumoral area) are shown above or below the original tumor sections at higher magnification. **B:** Semi-quantitative analysis of CD31 (intratumoral area) (**upper**) and LYVE-1 staining (peritumoral area) (**lower**) is shown. Data are expressed as means \pm SD. **P* < 0.05 by Kruskal-Wallis test, followed by the Dwass-Steel-Critchlow-Fligner multiple comparison. Scale bar = 200 µm. HSC, hepatic stellate cell; LV, liver; T, tumor.

HSC marker,⁴⁴ and PDGFR- α , a marker for CAFs,^{45,46} led to an extensive overlap in staining, indicating that HSCs are the major contributor to the myofibroblast pool (Figure 6A). Cygb/STAP staining showed that HSCs had infiltrated the HCC stroma and peritumor tissues. Next, a series of experiments were performed to confirm the effect of R-III on HSC-derived CAFs. First, liver sections from Hepa-1c1c7+HSC orthotopically transplanted mice were subject to immunohistochemistry for α -SMA. R-III administration reduced the intensity of α -SMA staining in the tumor and peritumoral areas (Supplemental Figure S4). Second, primary cultures were established

from orthotopic tumors (Hepa-1c1c7+HSC) and treated with R-III. As seen in Supplemental Figure S4, the staining intensity for α -SMA was decreased significantly after R-III treatment (Figure 6B). Third, cells were isolated from orthotopic tumors (Hepa-1c1c7+HSC) by fluorescence-activated cell sorting using PDGFR- α antibody. R-III treatment induced the morphologic changes and the reappearance of cytoplasmic lipid droplets, a characteristic feature of pre-activated or early activated HSCs, in approximately 30% of isolated cells (Figure 6C). Thus, all of these findings suggest that R-III inactivates activated/myofibroblastic HSCs, affecting the



Figure 5 R-III regulates angiogenesis-related factors in hepatic stellate cells (HSCs). **A:** Total RNA was isolated from the HSCs on day 1 after seeding (preactivated), and from HSCs after passage 1 (activated) with or without R-III treatment for 24 hours. Real-time PCR was used to analyze the mRNA levels of vascular endothelial growth factor (VEGF)-a, VEGF-d, and angiopoietin 1. **B** and **C:** Conditioned medium (CM) from activated HSCs treated with or without R-III was analyzed using a proteome profiler mouse angiogenesis array kit. Thirteen of 53 angiogenesis-related proteins with significant change are marked by **square boxes. A** and **B:** Data are expressed as means \pm SD. **C:** The quantitative densitometric data are presented as the mean fold change \pm SD of two independent experiments. n = 3. *P < 0.05, **P < 0.001 by two-way analysis of variance with the Dunnett multiple comparison test.

expression of pro-angiogenic and anti-angiogenic growth factors (Figure 6D). In summary, consistent with the *in vitro* results, the data obtained from the experiments

using nude mice indicate that the inhibition of HSC activation abrogates the protumorigenic functions of HSC-derived CAFs.



Figure 6 R-III inactivates hepatic stellate cell (HSC)-derived cancer-associated fibroblasts (CAFs). **A:** Liver tissues from Hepa-1c1c7+HSC—transplanted mice were analyzed using double-immunofluorescence staining for platelet-derived growth factor receptor α (PDGFR α) (red) and cytoglobin/stellate cell activation-associated protein (Cygb/STAP) (green). The **white dotted line** demarcates the stromal—tumor border. Semiquantitative scoring of CYGB⁺ PDGFR α^+ double-positive cells as a percentage of PDGFR α -stained cells is shown on the **right. B:** Primary cultures from orthotopic tumors were treated with R-III and analyzed by immunofluorescence. **C:** Cells were isolated from orthotopic tumors (Hepa-1c1c7+HSC) by fluorescence-activated cell sorting using the PDGFR α^- antibody and treated with R-III. R-III induces phenotypic conversion from myofibroblast to fat-storing cells (**arrows**). **D:** Proposed model for the anticancer effect of R-III. R-III inactivates HSC-derived CAFs, thus inhibiting the secretion of angiogenic and growth factors. **B:** Data are expressed as means \pm SD. **P* < 0.01 by Wilcoxon rank-sum test. Scale bar = 200 µm. HCC, hepatocellular carcinoma; PS, peritumoral stroma; T, tumor.

Discussion

Previous studies have shown that stromal HSCs promote tumor progression. The present study further supports the effect of activated/myofibroblastic HSCs on HCC progression in vitro and in vivo. The rapid growth of malignant HCC cells requires adequate nutritional supply and active angiogenesis.⁴⁷ The blood vessel density in the tumors derived from mice co-transplanted with HSCs was higher compared with that in the tumors originated from Hepa-1c1c6 cells alone. Activated HSCs reportedly promote tumor angiogenesis by secreting a number of angiogenic factors, thereby creating a protumorigenic microenvironment. The results of this study showed that angiogenic factors, such as VEGF-d, PDGF-d, angiopoietin-1, and CXCL12, were significantly up-regulated during HSC activation (Figures 1E and 5A). CXCL12 is a chemokine that recruits circulating endothelial precursor cells, and is involved in tumor growth, metastasis, angiogenesis, and cancer cell-TME interactions.³⁶

On the other hand, treatment of activated HSCs with R-III decreased expression of these angiogenic factors, decreased secretion of MCP-1, and increased secretion of antiangiogenic factors, which correlated with a reduction in vascular density and tumor growth delay. Because R-III did not affect Hepa-1c1c7 cells directly, the *in vivo* antitumor activity of R-III likely was attributed to its action on activated/myofibroblastic HSCs.

Quiescent SCs store retinoids in the form of retinyl esters in cytoplasmic lipid droplets. When the lipid droplets are lost rapidly during SC activation, a portion of their retinoid contents are likely released and metabolized into retinaldehyde and retinoic acid. The role of retinoids in HSC activation has been proposed, but previous reports regarding the effects of exogenous retinoids on HSCs and liver fibrosis were controversial.⁴⁸ However, Lee et al¹⁹ reported that endogenous retinoic acid was involved in the activation of HSCs and that R-III inhibited HSC activation by sequestering retinoic acid. Retinoic acid sequestration in turn caused a dramatic increase in IL-1 β expression, resulting in the inhibition of Smad3 nuclear translocation in HSCs.²¹ This shows that R-III can influence gene expression in HSCs. However, further study is needed to address how R-III affects the expression of pro-angiogenic and anti-angiogenic factors.

Extensive necrosis was found in the subcutaneous tumors harvested from Hepa-1c1c7+HSC+RIII groups compared with the other two groups namely, the Hepa-1c1c7–alone group and the Hepa-1c1c7+HSC group. Necrotic tissue is characterized by cell death with or without loss of structural integrity (coagulative necrosis versus liquefactive necrosis). Thus, tumors undergoing necrosis can undergo a reduction in tumor weight, caused by cell death, membrane disruption, and loss of nuclear and cytoplasmic content, without loss of tumor volume.⁴⁹ It is reasonable to conclude that the difference in tumor weights, without a concomitant change in

tumor size, was caused by the extensive necrosis in the Hepa-1c1c7+HSC+R-III tumors. This necrosis represents tumor cell death, and eventual resorption of the tumor debris is expected to occur at a later stage in tumor evolution, at a time point beyond the scope of the current experimental protocol.

The expression of VEGF-d, which enhances lymphangiogenesis and lymphatic metastasis,^{50,51} was up-regulated significantly during HSC activation and was down-regulated by R-III. PDGF-d is also reportedly involved in tumor lymphangiogenesis.⁵² The results of this study showed that the expression profile of VEGF-d and PDGF-d correlated with peritumoral lymphatic density and lymph node metastasis. As previously described,¹⁹ mice showed no apparent side effects after receiving R-III.

Most therapeutic strategies against cancer have focused on targeting tumor cells directly, but, recently, strategies also have been developed to target the TME. Because stromal cells are indispensable for tumor angiogenesis, antiangiogenic therapies must be aimed at targeting stromal cells as well as cancer cells. This study supports that stromal HSCs contribute significantly to HCC angiogenesis, and therefore suggests that inhibition of HSC-induced tumor angiogenesis may be useful for controlling tumors. The current study supports the presence of a protumor crosstalk between cancer cells and activated HSCs, which represents an attractive therapeutic target for anticancer therapy. Moreover, because SCs are also present in extrahepatic organs such as pancreas, kidneys, intestines, and lungs,⁵³ their potential contribution to cancer progression should be investigated.

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Author Contributions

J.O. conceived and supervised the study; M.-J.K. and J.O. designed the experiments and wrote the manuscript; M.-J.K., S.L., U.J., C.M., H.P., and J.W.M. performed the experiments and collected data; and W.G.S.-S., Y.-S.K., S.-H.P., and J.O. analyzed and interpreted data.

Supplemental Data

Supplemental material for this article can be found at *http://doi.org/10.1016/j.ajpath.2021.08.004*.

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