RORα inhibits adipocyte-conditioned medium-induced colorectal cancer cell proliferation and migration and chick embryo chorioallantoic membrane angiopoiesis

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1Beijing Key Laboratory of Gene Resource and Molecular Development and College of Life Sciences, Beijing Normal University, Beijing, People’s Republic of China; 2Cardiovascular Research Center, School of Medicine, Xi’an Jiaotong University, Xi’an, China; 3The Second Artillery General Hospital of Chinese People’s Liberation Army, Beijing, China; and 4Institute of Aging Research, Hangzhou Normal University, Hangzhou, China

Submitted 25 March 2014; accepted in final form 5 December 2014

Xiao L, Wang J, Li J, Chen X, Xu P, Sun S, He D, Cong Y, Zhai Y. RORα inhibits adipocyte-conditioned medium-induced colorectal cancer cell proliferation and migration and chick embryo chorioallantoic membrane angiopoiesis. Am J Physiol Cell Physiol 308: C385–C396, 2015. First published December 10, 2014; doi:10.1152/ajpcell.00091.2014.—Lipid metabolic disturbances are related to many diseases, such as obesity, diabetes, and certain cancers. Notably, lipid metabolic disturbances have been reported to be a risk factor for colorectal cancer. Nuclear receptors act as ligand-dependent transcription regulators and play roles in signal integration, including the modulation of neurogenesis, homeostasis, and disease by regulating subsets of gene expression patterns both positively and negatively (13, 24). The retinoic acid-related orphan nuclear receptor α (RORα) is a member of the orphan nuclear receptor family and can functionally interact with potent oncogenic systems (34). Compared with other classes of nuclear receptors, the function and related signaling pathways in the development of colorectal cancer for RORα have not yet been studied extensively. Meanwhile, the effects of RORα on colorectal cancer cell proliferation, migration, and angiopoiesis are largely unclear.

In this study, we investigated the effect of the conditioned medium of 3T3-L1 on the proliferation and migration of colorectal cancer diagnosed each year (39). Numerous factors, including genetic abnormalities and environmental factors, have been implicated in the etiology of colorectal cancer (37). Recently, increasing evidence suggests that a lipid metabolic disturbance is a critical risk factor in the occurrence and development of colorectal cancer (5). Adipose tissue is the main organ in lipid metabolic homeostasis. Until recently, adipocytes were mainly considered as an energy storage depot, but there is now clear evidence suggesting that adipocytes act as endocrine cells, producing hormones, growth factors, cytokines, and other molecules named adipocytokines (36). Notably, increased adipose tissue mass provokes adipocytokine dysregulation, which results in metabolic complications, such as type 2 diabetes, and certain cancers, such as colorectal cancer (21). Taken together, these compelling observations suggest that excess adiposity may favor tumor invasiveness via a yet uncharacterized mechanism(s). However, the link between the local lipid metabolism of colorectal tissue and the development of colorectal cancer remains unclear.

3T3-L1 is an immortalized cell line that is committed to the adipocyte lineage (14–16). In the presence of a hormonal cocktail consisting of 3-isobutyl-1-methylxanthine (IBMX), dexamethasone (DEX), and insulin (INS), 3T3-L1 preadipocytes differentiate into adipocyte-like cells, expressing adipocyte-specific genes, accumulating triacylglycerol-rich lipid droplets, and secreting several adipocytokines (9). Herein, we used the conditioned medium of 3T3-L1 cells, which contain similar factors to the adipocytokines in the surrounding colorectal tumor milieu, to explore the link between the local lipid metabolism of colorectal tissue and the development of colorectal cancer (40, 42).

Members of the orphan nuclear receptor family play various roles in signal integration, including the modulation of neurogenesis, homeostasis, and disease by regulating subsets of gene expression patterns both positively and negatively (13, 24). The retinoic acid-related orphan nuclear receptor α (RORα) is a member of the orphan nuclear receptor family and can functionally interact with potent oncogenic systems (34). Compared with other classes of nuclear receptors, the function and related signaling pathways in the development of colorectal cancer for RORα have not yet been studied extensively. Meanwhile, the effects of RORα on colorectal cancer cell proliferation, migration, and angiopoiesis are largely unclear.

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COLORECTAL CANCER is one of the most prevalent malignancies in the world (35), with more than 1,000,000 new cases of colorectal cancer diagnosed each year (39). Numerous factors, including genetic abnormalities and environmental factors, have been implicated in the etiology of colorectal cancer (37). Recently, increasing evidence suggests that a lipid metabolic disturbance is a critical risk factor in the occurrence and development of colorectal cancer (5). Adipose tissue is the main organ in lipid metabolic homeostasis. Until recently, adipocytes were mainly considered as an energy storage depot, but there is now clear evidence suggesting that adipocytes act as endocrine cells, producing hormones, growth factors, cytokines, and other molecules named adipocytokines (36). Notably, increased adipose tissue mass provokes adipocytokine dysregulation, which results in metabolic complications, such as type 2 diabetes, and certain cancers, such as colorectal cancer (21). Taken together, these compelling observations suggest that excess adiposity may favor tumor invasiveness via a yet uncharacterized mechanism(s). However, the link between the local lipid metabolism of colorectal tissue and the development of colorectal cancer remains unclear.

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colorectal cancer cells and revealed that the conditioned medium promoted the proliferation and motility of colorectal cancer cells and significantly enhanced the formation of blood vessels (BVs) in chicken embryonic chorioallantoic membranes (CAMs) in vitro. Next, we demonstrated that the conditioned medium led to a decrease in the expression of RORα. In addition, activating RORα with cholesterol sulfate (CS), putative ligand of RORα, inhibited the proliferation and motility of colorectal cancer cells, which was induced by conditioned medium via the regulation of cell cycle-regulatory molecules. Furthermore, CS inhibited the formation of BVs in chicken embryonic CAMs via the regulation of the vascular endothelial growth factor (VEGF) and hypoxia-inducible factor (HIF-1α) signaling pathway. The expression of lipogenic genes was elevated in colorectal tumor tissue compared with control colorectal tissue, and CS inhibited the expression of lipogenic genes in colorectal cancer cells.

MATERIALS AND METHODS

Reagents. 3-[4,5-Dimethylthiazol-2-yl]-2,5-diphenyl tetrazolium bromide (MTT), TO-901317, CS, IBMX, DEX, and INS were purchased from Sigma (St. Louis, Mo). Dimethyl sulfoxide (DMSO) was purchased from Merck (Darmstadt, Germany). TRizol, Dulbecco’s modified Eagle’s medium (DMEM), fetal bovine serum (FBS), and an antibiotic mixture (penicillin-streptomycin) were purchased from Gibco-BRL (Grand Island, NY). The BCA-100 protein quantitative analysis kit was purchased from Pierce (Rockford, IL). RORα rabbit polyclonal antibody, β-actin rabbit polyclonal antibody, and all secondary antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Protein molecular weight markers were obtained from Pharmacia (Saclay, France). The polyvinylidifluoride (PVDF) membrane for Western blot analysis was purchased from Millipore (Bedford, MA).

Cell culture and preparation of conditioned medium. The human colon cancer SW480 cells were generously provided by Dr. Junjie Zhang (Beijing Normal University, College of Life Science). The mouse colon carcinoma cell line C26 and the murine 3T3-L1 preadipocyte cell line was purchased from the Chinese Academy of Medical Sciences & Peking Union Medical College (Beijing, China). The human

Oil Red O staining. For the Oil Red O staining, the cells were washed three times with PBS and fixed with 3.7% formaldehyde for 1 h. Oil Red O (0.5% in isopropanol) was diluted with water (3:2), filtered two times with a 0.45-μm filter, and added to the fixed cells for 15 min at room temperature (RT). Cells were washed with 70% ethanol and water before visualization via light microscopy and photographed.

Enzyme-linked immunosorbent assay. Enzyme-linked immunosorbent assay (ELISA) was performed using a commercial VEGF ELISA kit (R&D Systems). Assays were performed in duplicate, and readings were performed with standard curves obtained with standard proteins provided with the kits.

Transfection of small-interfering RNA duplexes. The small-interfering RNA (siRNA) duplexes targeting all isoforms of human RORα and mouse RORα as well as control nonspecific siRNA were synthesized and purified by Genepharm. SW480 and C26 cells were transfected with siRNAs using the Lipofectamine 2000 (Invitrogen) reagent according to the manufacturer’s protocol. The oligonucleotide sequences used in siRNA assays are listed in Table 2.

Overexpression transfection assays. In overexpression experiments, SW480 or C26 cells were transfected by p3xFLAG-CMV10-RORα (RORα) or control p3xFLAG-CMV-10 (p3X) using Lipofectamine 2000 transfection reagent (Invitrogen) according to the manufacturer’s instructions. After 72 h, they were assessed by quantitative real-time PCR and Western blot analysis.

Cell viability test (MTT assay). To determine cell viability, cells (5 × 10^3/well) were seeded in sterile 96-well tissue culture plates and treated as indicated. Cell viability was evaluated in each well via the addition of 20 μl of MTT (2.5 mg/ml in PBS) 4 h before the reading. For analysis, the cell-free supernatants were removed completely from each well, and 100 μl of DMSO were added. The optical densities of
the wells were measured using a spectrophotometric multiwell microplate reader (Multiskan MS; Thermo Electron, Waltham, MA) at a wavelength of 570 nm.

Cell count. Harvested cells were washed with 1 × PBS and mixed with trypan blue dye, and the number of viable cells was determined manually using a hemocytometer. Briefly, a suspension of ~1 × 10^6 cells/ml was prepared, and a 1:1 mixture of the cell suspension and the 0.4% trypan blue solution was made. The solution was gently mixed and allowed to stand for 5 min at RT. Cell suspension (15 μl) was applied to the edge of the chamber between the cover slip and the V-shaped groove in the chamber. The cell suspension was drawn into the chamber by capillary action and then allowed to sit for 1–2 min and counted.

Cell cycle analysis. At 48 h after induction, SW480 and C26 cells were harvested and fixed with 75% cold ethanol. The fixed cells were stained with propidium iodide (Calbiochem, San Diego, CA), and the cell cycle distribution was determined using a FACS Calibur flow cytometer (BD Biosciences, Heidelberg, Germany).

Transwell assay. A Transwell system (8 μm pore size; Corning, Corning, NY) was used to evaluate the cell migration. First, a 100-μl cell suspension (1 × 10^6 cells/ml) in FBS-free medium was added to the upper compartment while 500 μl of growth medium (GM),
adipocyte-conditioned medium (mCM), or mCM with CS was placed in the lower compartment. The total number of cells that migrated into the lower chamber was counted after 48 h of incubation at 37°C with 5% CO₂. The filters were fixed with methanol and stained with 0.1% Giemsa. After being gently rinsed with water, the cells on the upper surfaces of filters were removed by wiping the filters with a cotton swab. Cells remaining on the lower surface of the filters were photographed under a microscope. The number of migrated cells was counted with 10 randomly selected fields. Experiments were performed in triplicate.

Hematoxylin and eosin staining. Tumor and normal colon tissue samples were fixed in 10% buffered formalin and embedded in paraffin. Sections (4–5 μm) were prepared and stained with hematoxylin and eosin according to standard protocol.

Immunohistochemical staining. Human colon sample paraffin-embedded sections of 5 μm thickness were obtained from the Second Artillery General Hospital of Chinese PLA. For villin staining, paraffin-embedded tissues were first deparaffinized in xylene for 10 min, followed by 5 min each in serial dilutions of ethanol (100, 95, and 75%). After antigen retrieval, sections were incubated for 1 h at RT with antivillin antibody at then counterstained with hematoxylin, dehydrated, and mounted. Negative control studies were done by incubating samples without the primary antibody.

Statistical analysis. The data were presented as means ± SE. The significance of paired data was determined by Student’s t-test. Data with more than two groups were analyzed by ANOVA followed by the post hoc Dunnett’s multiple-comparison test. Differences were considered significant at P < 0.05 and 0.01.

RESULTS

The distribution of adipocytes in human colorectal cancer tissue is elevated compared with normal colorectal tissue. Many studies in lipid metabolism and cancer have shown that an obvious correlation between lipid metabolism disorders and cancer exists. However, whether the local lipid metabolic disorder induces colorectal cancer has not been reported. To explore the relationship of local colorectal lipid metabolism disorders and colorectal cancer, we analyzed the fat cell distribution in the colorectal tissue, since the fat tissue is an essential organ in lipid metabolic homeostasis. Histological analysis confirmed the presence of adipocytes (Bodipy staining, green) surrounding the colorectal cancer in a human patient sample (Fig. 1, A and B). Figure 1C showed the villin staining of colorectal cancer tumor cells surrounding the adipocytes in the human samples. In addition, detailed patient information, including body weight, body mass index, cholesterol, low density lipoprotein, high density lipoprotein, and triglyceride level, is included in Table 3. These observations suggest that adipocytes are clinically relevant cells that may provide a physiological basis for the occurrence of colorectal cancer. Here, we investigated the possible association of adipocytes with colorectal cancer.

Colorectal cancer cells cultured with mCM exhibit an enhanced proliferative and motile phenotype. The results in Fig. 1 suggest that the occurrence of colorectal cancer may be linked to fat cells. Because adipose tissue is a functional part of the endocrine system, and because of the increased number and size of adipocytes, colorectal cancer tissue will have a higher level of secreted adipocytokines. Previous studies reported the secretion profiles during adipocyte differentiation involving a complex and dynamic process, mostly adipokines, such as adiponectin and other inflammation-related molecules, exhibit a temporally ordered pattern during differentiation (41, 42). To better define how adipocytokines in local colorectal tissue may induce colorectal cancer, we collected the entire replacement medium during 8-day differentiation (days 2, 4, 6, and 8 postdifferentiation) of 3T3-L1 and mixed them together, termed mCM (as mentioned in MATERIALS AND METHODS). Next, we cultured colorectal cancer cells (SW480 and C26) with mCM for 24 or 48 h. Upon incubation of SW480 and C26 cells with mCM, the proliferation of the cells was greatly increased (Fig. 2A) compared with GM. In addition, cell migration is an additional critical step in tumor metastasis (6). Thus, we determined the effect of mCM on cell migration. When cell motility was examined using a standard Transwell assay (28), microscopy (Fig. 2B) and quantification (Fig. 2C) of migrated cell numbers showed that more numbers of mCM-treated SW480 and C26 cells than GM-treated cells migrated toward the bottom chamber in a time-dependent manner. Tumor progression is closely linked with microvessel formation (2). Overgrowth of BVs may lead to the development and progression of tumor growth (3, 20, 22, 27). The avian egg offers an efficient model to assess new compounds and scaffolds, particularly on the avian CAM, a well-known site of experimentation. Therefore, we examined whether mCM could affect the progress of angiogenesis using a chick CAM assay. As demonstrated in Fig. 2D, relative to GM, the mCM-treated CAM significantly enhanced the formation of BVs in vitro, and the tube area increased three times as shown in Fig. 2E. Both obesity and a high-fat diet have been proposed to be critical risk factors for the occurrence and development of colorectal cancer. Consistent with these reports, the results of our study, as demonstrated in Figs. 1 and 2, suggest that the adipocytokines that are secreted by adipocytes in the local colorectal tissue have an important function in the proliferation and migration of colorectal cancer cells and the angiogenesis observed in tumor formation and growth. mCM decreases the expression of RORα in colorectal cancer cells. RORα, a member of the orphan nuclear receptor family, functions in lipid metabolism and certain cancers. RORα is expressed in many tissues in mammals, including cerebellar Purkinje cells, liver, thymus, skeletal muscle, skin, lung, and kidney (17, 38). We examined the expression of RORα in human tissue samples. Western blot and real-time PCR analyses revealed that the expression levels of RORα protein and mRNA were markedly lower in colorectal cancer compared with control colorectal tissue in humans (Fig. 3, A–C). Based on the different expression patterns of RORα between the human colorectal tumor tissue and control colorectal tissue, we concluded that RORα might function as a warning sign for local colorectal cancer. Subsequently, we determined the expression of the RORα target genes, ApoC3 and Bmal1, in colorectal tissues. Real-time PCR analysis revealed that the expression patterns of these genes were similar to those observed for RORα in human colorectal cancer tissue as well as normal tissue distant from the tumor (Fig. 3C).

Table 2. Oligonucleotide sequences used in siRORα assays

<table>
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<tr>
<th>Species</th>
<th>Sequence (5′-3′)</th>
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<tbody>
<tr>
<td>Human</td>
<td>5′-GCCUUCCACACUGCUUATT-3′</td>
</tr>
<tr>
<td></td>
<td>5′-UAACUCACGUGUUGCAGUUTT-3′</td>
</tr>
<tr>
<td>Mouse</td>
<td>5′-GAGCUUGACGAGUAUGCUTT-3′</td>
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<td>5′-ACGUUAUGCGGAGCUCTT-3′</td>
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To further examine the role of RORα in the development of colorectal cancer, we determined whether mCM modulated the expression of RORα and its target genes. Real-time PCR analysis revealed that the expression levels of RORα, ApoC3, and Bmal1 were decreased in both SW480 and C26 cells upon culture in mCM compared with culture in control medium (GM) (Fig. 3D). These results provide a basis for the study of the role of RORα in the secretion of the adipocytokines linked to colorectal cancer. Importantly, nuclear receptors are major targets for drug discovery and have key roles in cell development and homeostasis, as well as in many diseases such as obesity, diabetes, and cancer. Therefore, we chose RORα as a potential target to test whether it has a role in antitumorigenesis.

CS (a putative RORα ligand) inhibits proliferation and motility in colorectal cancer cells. Based on the results in Fig. 3, we hypothesized that decreasing levels of RORα are associated with the development of colorectal cancer. To investi-
gate the roles of RORα in the development of colorectal cancer, we examined whether pharmacological activation of RORα affects the growth of colorectal cancer cells. Thus, we treated SW480 and C26 cells with mCM alone or mCM in the presence of different concentrations of CS (a putative RORα ligand, 5 and 10 μM) and examined the cell growth. As demonstrated in Fig. 4A, when SW480 and C26 cells were incubated with mCM and CS for 48 h, the proliferation of SW480 cells was greatly decreased in a dose-dependent manner. Treatment with CS at concentrations of 1–100 μM had little impact on the SW480 and C26 viability, indicating that CS inhibited cell proliferation at these concentrations without affecting cell viability (Fig. 4B). Meanwhile, when SW480 and C26 cells were grown in the presence of mCM, a significant stimulation of tumor motile abilities was observed compared with cells grown in GM. When SW480 and C26 cells were grown in the presence of mCM with different concentration of CS, fewer cells migrated toward the bottom chamber compared with cells treated with mCM alone (Fig. 4, C and D). The observation that CS perturbed the colorectal cancer cell proliferation led us to examine whether CS might modulate cell cycle progression in colorectal cancer cells. SW480 and C26 cells were incubated with mCM and different concentrations of CS, and the cell cycle states were analyzed using flow cytometry. At 48 h after induction, 45.68% of mCM-treated SW480 cells entered the S phase, whereas only 39.99% of 5 μM and 34.44% of 10 μM CS-treated SW480 cells entered the S phase. Meanwhile, at 48 h after induction, 39.28% of mCM-treated C26 cells entered the S phase, whereas only 36.08% of 5 μM and 33.83% of 10 μM CS-treated C26 cells entered the S phase. Therefore, compared with cells treated with mCM alone, SW480 and C26 cells treated with mCM in addition to CS exhibited a larger portion of the population in the G1 phase and a smaller fraction of the population in the S phase in a dose-dependent manner (Fig. 4, E and F). Thus, RORα induced cell cycle arrest at the G1/S boundary in colorectal cancer cells. Because CS arrested cell cycle progression at the G1 phase in colorectal cancer cells, we asked the question...
whether CS might affect the expression of cyclin-dependent kinase inhibitors, thus resulting in GI/S arrest. As shown in Fig. 4G, CS substantially promoted the expression of p21 and decreased cyclin D1 mRNA in SW480 and C26 cells. These data suggest that p21 and cyclin D1 are likely targets for RORα in the inhibition of cell proliferation and cell cycle arrest since they induce quiescent cells to progress through the cell cycle.

Angiogenesis is a key step in tumor growth, invasion, and metastasis. Thus, the antiangiogenic therapy was postulated to be an attractive approach for antitumor treatment (11). To examine the ability of CS to inhibit angiogenesis, we used a CAM assay. Three days after hatching, the BVs on the slides were counted. As shown in Fig. 4H, relative to the mCM-treated group, CS significantly inhibited the formation of BVs.

Genetic RORα inhibits proliferation and motility in colorectal cancer cells. To investigate whether genetic activation of RORα also affects the growth of colorectal cancer cells, we overexpressed RORα and knocked down RORα in SW480 and C26 cells, respectively.

Figure 5A suggested we have succeeded in knocking down and overexpressing ROR. As shown in Fig. 5B, when SW480 cells knocked down RORα, 41.09% of SW480 cells entered the S phase, whereas only 36.69% of negative control SW480 cells entered the S phase. When SW480 cells overexpressed RORα, 26.66% of SW480 cells entered the S phase, whereas 31.37% of vehicle control SW480 cells entered the S phase. RORα in SW480 cells were knocked down, the mRNA expression of p21 was decreased, and the mRNA expression of cyclin D1 was increased (Fig. 5C). In contrast, overexpression of RORα increased the expression of p21 mRNA and decreased the mRNA expression of cyclin D1 (Fig. 5D).

CS decreases the expression level of key transcriptional regulators of angiogenesis. Several lipogenic genes, including sterol-regulatory element-binding protein-1c (SREBP-1c) and fatty acid synthase (FAS), are upregulated in certain obesity-linked cancers, such as breast, prostate, endometrial, and colorectal (1, 8, 10, 23, 43). Consistent with these reports, the levels of lipogenic genes, such as liver X receptor α, SREBP-1c, peroxisome proliferator-activated receptor γ, fatty acid-binding protein-4 (aP2/FABP4), stearoyl-CoA desaturase-1 (SCD-1), acetyl-CoA carboxylase (ACC), CAAT/enhancer-binding protein α, and adipose differentiation-related protein were elevated in human colorectal cancer tissues compared with normal tissues from the same colon cancer patients (Fig. 6A). To explore the molecular mechanism through which CS inhibits angiogenesis in chicken embryonic CAM, we induced 3T3-L1 to differentiate via incubation in differentiation medium (DM) in the presence or absence of CS. 3T3-L1 cells

Fig. 3. mCM decrease the expression of retinoic acid receptor-related orphan receptor α (RORα) and the target genes ApoC3 and Bmal1 in colorectal cancer cells. A: Western blot analysis of RORα expression in human colorectal adenocarcinoma tissues (C) and adjacent normal (N) tissues. B: RORα levels were normalized to the amount of β-actin. C: real-time PCR results evaluating mRNA levels of RORα, ApoC3, and Bmal1 expression in human colorectal adenocarcinoma tissues and adjacent normal tissues. D: real-time PCR analysis to measure relative mRNA expression of RORα, ApoC3, and Bmal1 in mCM-treated SW480 and C26 cells. The real-time PCR data are representative of 3 independent experiments. The blot shown is representative of 4 independent experiments. Data are means ± SE. The results are from 3 independent experiments. *P < 0.05 and **P < 0.01 vs. normal control (GM).
were maintained in DMEM containing 10% FBS. To induce differentiation, 2-day postconfluent preadipocytes were cultured in DMEM containing 10% FBS, 10 mg/ml INS, 0.5 mM DEX, and 0.5 mM IBMX (termed DM) for 2 days. The medium was then replaced with DMEM containing INS, and the cells continued to differentiate until day 8 (still termed DM). On day 8 after induction, the cells were stained with the lipophilic dye Oil Red O to demonstrate triacylglycerol accumulation. DM-treated cells resembled differentiated adipocytes and accumulated triacylglycerol as expected. In contrast, CS-treated cells had a reduced level of Oil Red O staining, indicating that CS reduced lipid accumulation in the cells (Fig. 6B). Thus, adipogenesis is tightly associated with angiogenesis, as shown by the findings that adipose tissue explants trigger BV formation, whereas adipose tissue endothelial cells promote preadipocyte differentiation (7). As shown above, angiogenesis can be enhanced by mCM. The expression levels of VEGF and HIF-1α were examined during adipocyte differentiation in the presence and absence of CS because of their known proangiogenic functions. Notably, CS treatment reduced the mRNA expression level of VEGF and HIF-1α in 3T3-L1 cells (**P < 0.05 and **P < 0.01 vs. mCM). J: SW480 and C26 cells were incubated with GM, mCM, or mCM with CS for 48 h. Cell supernatants were prepared and subjected to ELISA to detect VEGF. All data are presented as means ± SE; n = 3. **P < 0.05 and **P < 0.01 compared with mCM using 1-way ANOVA.
tissues. We examined whether CS was able to suppress the expression of lipogenic genes in colorectal cancer cells. In SW480 and C26 cells, CS decreased the mRNA expression of ACC, SCD-1, and FABP4 (Fig. 6E). Meanwhile, SW480 cells also displayed a reduction of VEGF and HIF-1α mRNA level in overexpressed RORα and significant upregulations of the expression of VEGF and HIF-1α mRNA level when RORα was knocked down (Fig. 6, F and G).

DISCUSSION

The dysregulation of adipocytokines in adipose tissue is involved in several pathological conditions, including diabetes and certain cancers. We examined the presence of adipocytes in human colorectal tissue and demonstrated a significant increase in both the number and size of the adipocytes in colorectal cancer tissue compared with normal colorectal tissue. In the present study, using the 3T3-L1 cells as a model of adipocytes, we created conditioned medium (mCM) to imitate the secretion of adipocytokines from the adipocytes to mimic the natural condition surrounding colorectal tissue. Considering the complexity of adipocyte tissue (different stage adipocytes or preadipocytes), we used the entire replacement medium during 8-day differentiation (days 2, 4, 6, and 8 postdifferentiation) of 3T3-L1 in all of our experiments. We observed that mCM promoted the proliferation and motility ability of colorectal cancer cells and enhanced the formation of BVs in chicken embryonic allantoic membranes in vitro. In addition, we checked the effects of different postdifferentiation day conditioned medium (days 0, 2, 4, and 8) on the growth and migration of C26 cells (data not shown). The results showed different stages of adipocyte media enhanced the growth and migration of colonic cells, particularly day 4 and day 8 medium. Thus, we concluded that adipocytokines that are secreted by the adipocytes in the local colorectal tissue contributed to the progression of colorectal cancer.

RORα is a member of the orphan nuclear receptor family and has been implicated in the regulation of several genes involved in lipid metabolism and tumorigenesis. Herein, we reveal that RORα is expressed in human colorectal tissues and that the adipocyte conditioned medium led to a decrease in the
expression of RORα. Meanwhile, expression levels of RORα and its target genes are lower in human colorectal tumor tissue compared with control colorectal tissue. Because of the difference in expression patterns in colorectal tumor and normal tissues, we concluded that RORα might have a potential role in the progression of colorectal cancer. Subsequently, we observed that CS, a putative ligand of RORα, inhibited SW480 and C26 cell proliferation and induced G1 arrest in colorectal cancer cells in a dose-dependent manner via the modulation of the expression of cell cycle-regulatory genes. Using a Transwell assay, we demonstrated that CS inhibited cell motility in colorectal cancer cells. Furthermore, angiogenesis is a key step in tumor growth and cancer development. VEGF and HIF-1α are the two important angiogenic cytokines that have not only angiogenic properties but also an autocrine ability to regulate the synthesis of multiple angiogenic proteins in multiple cell types (12, 25, 30, 44). Thus, the level of VEGF and HIF-1α could directly and indirectly indicate the ability of a tumor to cause angiogenesis. In fact, we observed that CS inhibited angiogenesis in chicken embryonic allantoic membranes by regulating the expression of both VEGF and HIF-1α. In addition, we examined the RORα agonist CS under regular growth condition on colorectal cancer cell proliferation and migration using flow cytometric and Transwell migration assay. The results showed there were no significant effects of the RORα agonist CS under regular growth conditions on the proliferation and migration of C26 cells (data not shown).

Many cancer cells exhibit an increased lipogenic activity related to rapid cell growth and division. Because the cell growth and division rates are higher in cancer cells than in normal cells, lipid metabolites such as phospholipids are crucial for maintaining membrane integrity in cancer cells (19).
There is a renewed interest in the ultimate role of FAS, a key lipogenic enzyme catalyzing the terminal steps in the de novo biosynthesis of fatty acids in cancer pathogenesis. Increased expression of the tumor-associated FAS gene, by conferring growth and survival advantages rather than functioning as an anabolic energy-storage pathway, appears to accompany the natural progression of most human cancers (26). Moreover, the development and progression of cancer is accompanied by marked changes in the expression and activity of enzymes involved in the cellular homeostasis of fatty acids. One class of enzymes that play a particularly important role in this process is ACC. In addition, evidence suggests that there is a critical implication of SCD-1 in the modulation of multiple biological mechanisms, specifically in lipid biosynthesis and the proliferation and survival signaling pathways that contribute to the development and progression of cancer (18). Recently, some studies have demonstrated that deficiencies in FABP4 substantially impaired metastatic tumor growth in mice, indicating that FABP4 has a key role in ovarian cancer metastasis (29). In this aspect, inhibitors of lipogenic genes have been proposed as anticancer drugs. For instance, FAS inhibitors, such as cerulenin and C75, exhibit potent inhibition of human cancer cell growth (31–33). Indeed, we observed that CS inhibited the expression of lipogenic genes, including FAS, ACC, SCD-1, and FABP4, and that this inhibition is potentially associated with retardation of colorectal cancer cell growth. When we repressed the expression of RORα via siRNA, the expression of lipogenic genes was significantly increased.

Therefore, this work suggests a more detailed mechanism of action of adipocytokines in the development of colorectal cancer. The accumulation of adipocytes may secrete elevated levels of adipocytokines, which provide the potential of a local action of adipocytokines in the development of colorectal cancer. The nuclear receptor peroxisome proliferator-activated receptor gamma (PPARγ) is a key regulator of adipogenesis, and its overexpression is associated with the development and progression of cancer (99). Moreover, the conditioned medium led to a decrease in the expression of RORα, which may be associated with the inhibition of tumor growth. When we repressed the expression of RORα via siRNA, the expression of lipogenic genes was significantly increased.

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REFERENCES


