Targeting of cancer-associated fibroblasts enhances the efficacy of cancer chemotherapy by regulating the tumor microenvironment

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Abstract. Cancer-associated fibroblasts (CAFs), key components of the tumor stroma, can regulate tumorigenesis by altering the tumor microenvironment in various ways to promote angiogenesis, recruit inflammatory immune cells and remodel the extracellular matrix. Using a murine xenograft model of colon carcinoma, the present study observed that oxaliplatin increased the accumulation of CAFs and stimulated the production of cytokines associated with CAFs. When oxaliplatin was combined with the small-molecule dipeptidyl peptidase inhibitor PT-100, which inhibits CAFs by targeting fibroblast activation protein (FAP), the accumulation of CAFs was markedly reduced, xenograft tumor growth was significantly suppressed and the survival of the mice increased, compared to those of mice treated with oxaliplatin or PT-100 alone. Furthermore, the xenograft tumor tissues of mice treated with oxaliplatin and PT-100 contained lower numbers of tumor-associated macrophages and dendritic cells, expressed lower levels of cytokines associated with CAFs and had a lower density of CD31+ endothelial cells. The present study demonstrated that pharmacological inhibition of CAFs improved the response to chemotherapy, reduced the recruitment of immune tumor-promoting cells and inhibited angiogenesis. Combining chemotherapy with agents which target CAFs may represent a novel strategy for improving the efficacy of chemotherapy and reducing chemoresistance.

Introduction

Tumors are complex structures composed of malignant cancer cells surrounded by the tumor stroma. The cells and components of the tumor stroma have received increasing attention due to their roles in tumor development, invasion and metastasis (1), and in the response to cancer therapy (2,3). The tumor stroma contains various cell types, including myeloid cell sub-populations, inflammatory cells, immunocytes, endothelial cells, epithelial cells and fibroblasts (4,5), which communicate between themselves and also directly with the cancer cells through cell-cell contacts and indirectly through paracrine/exocrine signaling, release of proteases and modulation of the extracellular matrix (ECM). Together, the stromal cells constitute the complex tumor microenvironment, which has a key role in tumor development (1,6). On this basis, identification of stromal targets for cancer therapeutics is of great interest, and such strategies may complement therapies directed against cancer cells. Among the potential targets in the stroma, cancer-associated fibroblasts (CAFs) have received intensive interest.

It is thought that the fibroblasts in the tumor stroma acquire a modified phenotype, which can be utilized for distinguishing them from normal tissue fibroblasts. Such ‘activated’ fibroblasts have been termed peritumoral fibroblasts, reactive stromal fibroblasts, myofibroblasts or CAFs (7). The precise origin of CAFs remains to be elucidated; however, a previous study has identified two potential pathways: Resident fibroblasts may be converted into CAFs through cytokine-induced, including transforming growth factor-beta (TGF-β) and stromal cell-derived factor-1 (SDF-1) (8); furthermore epithelial or endothelial cells may transform into CAFs via epithelial-mesenchymal transition (EMT) and endothelial-to-mesenchymal transition (EndMT), which are also mediated by cytokines, including fibroblast growth factor (FGF), osteopontin (9), TGF-β and SDF-1 (10,11).

Numerous studies have provided evidence for the cancer-promoting role of CAFs. In contrast to resting fibroblasts, CAFs are characterized by an increased rate of proliferation and differential expression of ECM components and growth factors, including TGF-β, vascular endothelial growth factor (VEGF), platelet-derived growth factor (PDGF) and fibroblast growth factor 2 (FGF-2) (12,13). These cytokines and growth factors have all been shown to have important roles in synchronizing...
key events which continuously occur in the tumor microenvironment. For instance, TGF-β promotes the infiltration of inflammatory/immune cells and CAFs into the tumor microenvironment, directly leading to changes in tumor cells (14). CAFs orchestrate tumor-promoting inflammation in a nuclear factor (NF)-κB signaling-dependent manner (15). VEGF induces microvascular permeability, leading to the extravasation of plasma proteins such as fibrin, which subsequently attracts CAFs, inflammatory immune cells and endothelial cells, leading to tumor angiogenesis (16). While CAFs and inflammatory cells are the principal sources of host-derived VEGF (16), PDGF and FGF-2 also have significant roles in angiogenesis. Therefore, CAFs are key factors that can promote tumor growth by inducing angiogenesis, recruiting inflammatory/immune cells and remodeling the ECM.

Compared to normal fibroblasts, extensive changes in the expression of genes that encode certain extracellular matrix proteins and proteases have been observed in CAFs in numerous types of carcinoma (17-19); among these, fibroblast activation protein (FAP), a type II membrane-bound serine protease, has recently gained attention. FAP has been shown to possess dipeptidyl peptidase-20 (20) and collagenase-like (21) activity in vitro, and has been implicated in ECM remodeling. However, the in vivo substrates of FAP remain to be identified. Due to its tightly regulated pattern of expression in the stroma of malignant solid tumors (22,23), FAP has been classified as a candidate protein for targeting CAFs. The present study hypothesized that FAP inhibition may be useful for cancer therapy. FAP belongs to the post-proline dipeptidyl aminopeptidase family and has the highest similarity to dipeptidyl peptidase IV (DPPIV/CD26) (24). The catalytic sites of CD26/DPPIV and FAP contain the characteristic catalytic triad of Ser(DPPIV), Asp708/702, His740/734 (the residues are numbered according to human CD26/DPPIV and FAP, respectively), and the active serine is situated in a nucleophilic elbow motif within the sequence Gly-Trp-Ser-Tyr-Gly (13-15) (25,26). The aminopeptidase family and a previous study has recently gained attention. FAP has been shown to possess dipeptidyl peptidase IV (DPPIV/CD26) (24). The catalytic sites of CD26/DPPIV and FAP contain the characteristic catalytic triad of Ser(DPPIV), Asp708/702, His740/734 (the residues are numbered according to human CD26/DPPIV and FAP, respectively), and the active serine is situated in a nucleophilic elbow motif within the sequence Gly-Trp-Ser-Tyr-Gly (13-15) (25,26). The aminopeptidase family and a previous study has recently gained attention. FAP has been shown to possess dipeptidyl peptidase IV (DPPIV/CD26) (24). The catalytic sites of CD26/DPPIV and FAP contain the characteristic catalytic triad of Ser(DPPIV), Asp708/702, His740/734 (the residues are numbered according to human CD26/DPPIV and FAP, respectively), and the active serine is situated in a nucleophilic elbow motif within the sequence Gly-Trp-Ser-Tyr-Gly (13-15) (25,26). The aminopeptidase family and a previous study has recently gained attention. FAP has been shown to possess dipeptidyl peptidase IV (DPPIV/CD26) (24). The catalytic sites of CD26/DPPIV and FAP contain the characteristic catalytic triad of Ser(DPPIV), Asp708/702, His740/734 (the residues are numbered according to human CD26/DPPIV and FAP, respectively), and the active serine is situated in a nucleophilic elbow motif within the sequence Gly-Trp-Ser-Tyr-Gly (13-15) (25,26). The aminopeptidase family and a previous study has recently gained attention. FAP has been shown to possess dipeptidyl peptidase IV (DPPIV/CD26) (24). The catalytic sites of CD26/DPPIV and FAP contain the characteristic catalytic triad of Ser(DPPIV), Asp708/702, His740/734 (the residues are numbered according to human CD26/DPPIV and FAP, respectively), and the active serine is situated in a nucleophilic elbow motif within the sequence Gly-Trp-Ser-Tyr-Gly (13-15) (25,26). The aminopeptidase family and a previous study has recently gained attention. FAP has been shown to possess dipeptidyl peptidase IV (DPPIV/CD26) (24). The catalytic sites of CD26/DPPIV and FAP contain the characteristic catalytic triad of Ser(DPPIV), Asp708/702, His740/734 (the residues are numbered according to human CD26/DPPIV and FAP, respectively), and the active serine is situated in a nucleophilic elbow motif within the sequence Gly-Trp-Ser-Tyr-Gly (13-15) (25,26). The aminopeptidase family and a previous study has recently gained attention. FAP has been shown to possess dipeptidyl peptidase IV (DPPIV/CD26) (24). The catalytic sites of CD26/DPPIV and FAP contain the characteristic catalytic triad of Ser(DPPIV), Asp708/702, His740/734 (the residues are numbered according to human CD26/DPPIV and FAP, respectively), and the active serine is situated in a nucleophilic elbow motif within the sequence Gly-Trp-Ser-Tyr-Gly (13-15) (25,26). The aminopeptidase family and a previous study has recently gained attention. FAP has been shown to possess dipeptidyl pep

Fig. 1. Chemical structure of PT-100.

Materials and methods

Animals and cell lines. A total of 40 female BALB/c mice (6-8 weeks old and weighing 20 g) were purchased from Beijing HFK Bioscience Co. Ltd., (Beijing, China) and maintained in a under a specific pathogen-free environment at the State Key Laboratory of Biotherapy (Chengdu, China) with controlled temperature (20-26°C), humidity (40-70%) and a 12-h light/dark cycle.

All animal experiments were approved by the Institutional Animal Care and Use Committee of Sichuan University (Chengdu, China). CT26, a BALB/c-derived murine colon carcinoma cell line, was purchased from the American Type Culture Collection (Manassas, VA, USA). CT26 cells were cultured in RPMI-1640 supplemented with 10% fetal bovine serum (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) in a humidified atmosphere containing 5% CO2 at 37°C. PT-100. PT-100 was purchased from Shanghai Speed Chemical Co. Ltd., (Shanghai, China). The purity was >97%. The molecular weight of PT-100 is 246.1 g/mol and its structure is shown in Fig. 1.

Marine tumor xenograft model. The flanks of the mice were shaved and subcutaneously inoculated with 1x106 CT26 cells. The tumor-inoculated mice were divided into four groups: The saline vehicle group; the oxaliplatin group, which was treated with 5 mg/kg oxaliplatin (Jiangsu Hengrui Medicine Co., Ltd., Lianyungang, China) three times a week; the PT-100 group, which was treated with 20 µg PT-100 per mouse daily; and a combined oxaliplatin and PT-100 group. All drugs were dissolved in saline and administered by intraperitoneal injection. The day of tumor inoculation was defined as day 0, the treatment began on day 8 and animals were treated for 14 days. Tumor growth was monitored every 3 days by measurement of the length (L) and the width (W) of the xenograft tumors using Vernier calipers (Shanghai Taihai Measuring Tools Co., Ltd., Shanghai, China). The tumor volume was calculated using the following formula: Volume (cm3) = W2x0.5L. The xenograft experiment was performed twice with 5 mice per group in each experiment.

Tissue preparation, histology and immunostaining. At the appropriate time-points, for paraffin embedding, the tumors were excised and post-fixed in 4% paraformaldehyde overnight. For cryopreservation, the tumors were excised and stored in optimum cutting temperature compound (OCT; Leica Microsystems GmbH, Wetzlar, Germany) at -20°C. FAP expression was assessed by immunohistochemistry in the CT26 colon cancer cell-derived xenograft tumors from female BALB/c mice. Tissues were cut into 5-µm sections, incubated with 3% H2O2 at 4°C for 20 min, washed using phosphate-buffered saline (PBS) containing 3.74 g 12H3O,Na2HPO4,
0.44 g NaH₂PO₄·2H₂O and 7.2 g NaCl in 11 distilled water (pH 7.4) and incubated with serum at 37°C for 20 min. Subsequently, the sections were incubated with the following primary antibodies at 4°C overnight: Polyclonal anti-FAP (1:500; Ab28244) (anti-F), monoclonal anti-vimentin (1:500; ab92547) (anti-F) and monoclonal anti-CD31 (1:50; ab7388; all Abcam, Cambridge, MA, USA) (anti-F). Following washing with PBS, the sections were incubated with a Bio-Tin-streptavidin horseradish peroxidase (HRP) detection system (SP-9000; OriGene Technologies, Inc., Beijing, China) at 37°C for 1 h, washed with PBS and subsequently incubated with 3,3-diaminobenzidine (DAB) for 2 min and washed using PBS. The HRP-conjugated secondary antibody and the DAB were included in the detection system (SP-9000; ZSGB-Bio Orgene Co., Ltd., Beijing, China), which was not diluted and was used according to the manufacturer's instructions. To assess the levels of apoptosis in CT26 tumors, in situ terminal deoxynucleotidyl transferase-mediated deoxy-UTP nick end labeling (TUNEL) staining was performed on the tumor sections using the the DeadEnd™ Fluorometric TUNEL system (G3250; Promega, Madison, WI, USA) according to the manufacturer's instructions.

For histological analysis, the paraffin-embedded tumor tissues were de-paraffinized in xylene, immersed in 100, 95 and then 75% ethanol, immersed in PBS (pH 7.4) and stained with hematoxylin (H3136-25G) and eosin (E4009-5G; both Sigma-Aldrich, St. Louis, MO, USA), according to the manufacturer's instructions. A Leica DM 2500 microscope was used for visualization (Leica Microsystems GmbH).

Western blot analysis. On day 22, the tumors were excised and proteins were extracted using radioimmunoprecipitation assay buffer (Beyotime Institute of Biotechnology, Inc., Haimen, China) supplemented with protease inhibitor cocktail (100:1; Sigma-Aldrich). Tumor tissues were homogenized prior to western blot and PCR analysis in liquid nitrogen using a mortar and pestle. Total protein concentrations in the supernatant were determined according to the manufacturer's instructions. Transcript expression was determined relative to GAPDH. The primers used were purchased from BGI-Shenzhen (Shenzhen, China) and were as follows: GAPDH, 5'-ACCCAGAAGCTTGATGATG-3' (forward) and 5'-TCT AGACGGCAAGTCAGTCT-3' (reverse); TGF-β, 5'-AAGTGG GTCCATGAACTCTAA-3' (forward) and 5'-GCTACATTATACA AGACG6CAT-3' (reverse); FGF-2, 5'-GGCTGCTGCTGGCTCTA AGTGTA-3' (forward) and 5'-CCGTTTGTGACCATGACATT-3' (reverse); and osteopontin, 5'-TGC ACC CAG AT CCTA TAGCC-3' (forward) and 5'-CTCCATGCTCATCATTAGC-3' (reverse). The 20 µl PCR system contained 2 µl genomic DNA, 2 µl dNTPs, 10 µl buffer, 0.5 µl RoX reference dye and 5 pmol of each primer. Thermal cycling was performed as follows: Denaturing at 95°C for 3 min, then 30 cycles of 95°C for 30 sec, 55°C for 30 sec and 72°C for 30 sec, with elongation at 72°C for 10 min using a CFX96 Real-Time C1000 thermocycler (Bio-Rad Laboratories, Inc.). PCR products were electrophoresed on 1% agarose gel (800669; Schwarz/Mann Biotech, Cleveland, OH, USA) at 200 mA until separation was achieved.

DNA fragments were visualized using a long wave UV light box and images were captured using a Universal Hood II Gel Doc™ XR camera (1708170) and were analyzed using Image Lab software (both Bio-Rad Laboratories, Ltd.).

Statistical analysis. SPSS version 11.5 (SPSS, Inc., Chicago, IL, USA) was used for statistical analyses. Values are expressed as the mean ± standard error of the mean. One-way analysis of variance was used to assess statistical significance. Survival curves were compared using the log-rank test. P<0.05 was considered to indicate a statistically significant difference.

Results

Oxaliplatin induces accumulation of CAFs in the tumor micro-environment. Chemotherapy is widely used for the treatment of colorectal cancer; however, drug resistance has become problematic due to its increasing frequency. Several mechanisms of drug resistance of cancers have been proposed (31,32); however, a consensus has not yet been reached and further elucidation is required. The present study hypothesized that CAFs may have an important role in drug resistance of cancers.
To test this hypothesis, Balb/C mice (n=5 per group) were inoculated with CT26 colon carcinoma cells (day 0) and then treated with the saline vehicle or oxaliplatin for 14 days (from days 8-22). Subsequently, the mice were sacrificed via cervical dislocation and the tumor xenografts were excised, sectioned and stained using an anti-FAP and anti-vimentin antibodies. As shown in Fig. 2, treatment with oxaliplatin markedly increased the amount of FAP and vimentin, which are specific markers of CAFs, expressed in the stroma of the tumors, indicating that oxaliplatin increased the accumulation of CAFs in the xenograft tumors.

Combined treatment with oxaliplatin and a pharmacological inhibitor of CAFs reduces tumor growth in vivo. In order to evaluate whether the ability of oxaliplatin to increase the accumulation of CAFs in the xenograft tumors was clinically relevant, the present study combined pharmacological inhibition of CAFs by PT-100 with the chemotherapeutic drug oxaliplatin, to which CT26 cells are partially sensitive, in a xenograft model. Mice (n=5 per group) were subcutaneously injected with CT26 cells on day 0 and then treated with saline vehicle, PT-100 alone, oxaliplatin alone, or PT-100 combined with oxaliplatin from days 8-32. As shown in Fig. 3A, PT-100 combined with oxaliplatin suppressed tumor growth more significantly than treatment with either oxaliplatin or PT-100 alone (P<0.05). PT-100 combined with oxaliplatin also significantly increased the survival of the mice compared to that in the other three treatment groups, particularly the mice treated with saline vehicle, which died after 45-60 days (Fig. 3B).

Consistent with the results shown in Fig. 2, FAP, a specific marker for CAFs, was most highly expressed in the stroma of the tumors of the mice treated with oxaliplatin alone, indicating that oxaliplatin increased the accumulation of CAFs in the xenograft tumors. However, PT-100 combined with oxaliplatin markedly reduced the expression of FAP and therefore prevented the accumulation of CAFs in the stroma of the tumor tissues. In addition, TUNEL staining was performed on the xenograft tumor sections, which revealed that PT-100 combined with oxaliplatin increased the number of apoptotic tumor cells compared with that in the three other treatment groups (Fig. 3C-F).

Next, to investigate how oxaliplatin induced the accumulation of CAFs, RT-qPCR analysis was performed to assess the expression of a number of cytokines which are associated with the accumulation of CAFs in the xenograft tumor tissues. CAFs are hypothesized to accumulate in the tumor microenvironment by the following major mechanisms: Resident fibroblasts transforming into CAFs, the EMT and the EndMT (8,10,11); these processes are mediated by several common cytokines, including TGF-β3, FGF-2 and osteopontin (9,12). Thus, the present study quantitatively measured the abundance these cytokines in the xenograft tumor tissues. As shown in Fig. 3G, oxaliplatin increased the expression of TGF-β3 and FGF-2 (basic FGF) by 1.79- and 2.63-fold, respectively, compared to those in the tumors of the saline vehicle-treated animals. Furthermore, the expression of TGF-β3 and FGF-2 in the tumors of animals treated with PT-100 alone or with PT-100 combined with oxaliplatin was lower than that in the saline vehicle-treated animals. Although oxaliplatin treatment did not significantly increase the expression of osteopontin mRNA, its expression was significantly decreased in the tumors of the animals treated with PT-100 only or with PT-100 combined with oxaliplatin (0.28- and 0.16-fold, respectively, of that in the saline vehicle-treated group). These results indicated that pharmacological inhibition of CAFs using PT-100 reduced the tumor expression levels of cytokines known to promote the accumulation of CAFs.

Combined treatment with oxaliplatin and a pharmacological inhibitor of CAFs reduces the recruitment of tumor-associated macrophages (TAMs) and dendritic cells. The tumor microenvironment can affect the malignant potential of the tumor; tumor-associated macrophages and dendritic cells have an important role in this process (33,34). Macrophages constitute an extremely heterogeneous population, including M2 (or alternatively activated) macrophages, which are now generally accepted to be TAMs. TAMs mostly exert pro-tumor functions by promoting tumor-cell survival, proliferation and dissemination. Tumor-associated dendritic cells (DCs) are another immune-regulatory cell population. A variety of sub-populations of tumor-associated DCs are known, among which CD11c+ DCs are able to promote tumorigenesis (35). In order to test whether CAFs interact with M2 macrophages or immature dendritic cells (imDCs) and tumor-infiltrating DCs (TIDCs), tumor cells from mice at day 22 were analyzed by flow cytometry. As shown in Fig. 4, the number of M2 macrophages (F4/80+) was highest in the saline vehicle-treated group and lowest in the xenograft tumors of animals treated with PT-100 combined with oxaliplatin; furthermore, tumors of animals

Figure 2. Effects of oxaliplatin on the accumulation of CAFs in CT26 colon cancer cell-derived murine xenograft tumors. Immunohistochemical analysis of tumors harvested on day 22 for FAP and an vimentin, which are specifically expressed in CAFs (magnification, x100 for FAP and x200 for vimentin). CAF, cancer-associated fibroblast; FAP, fibroblast activation protein.
treated with PT-100 combined with oxaliplatin contained significantly lower amounts of M2 macrophages compared with those in the groups treated with saline vehicle, oxaliplatin alone and PT-100 alone (P<0.01). In addition, the number of imDCs and TIDCs was highest in the saline vehicle-treated tumors, while the tumors of animals treated with PT-100

Figure 3. Effects of oxaliplatin and PT-100 combination therapy on the accumulation of CAFs in CT26 colon cancer cell-derived murine xenograft tumors. (A) Tumor growth curves. Values are expressed as the mean ± standard error of the mean. *P<0.05 vs. the three treatment groups. (B) Survival curves. (C) Hematoxylin and eosin staining of tumors harvested on day 22 (magnification, x100). (D) Immunostaining for FAP, which is specifically expressed in CAFs, in tumors harvested on day 22 (magnification, x200). (E) Detection of apoptotic cells in tumors excised on day 22 using the DeadEnd™ fluorometric terminal deoxynucleotidyl transferase dUTP nick end labeling system. (F) Western blot of FAP in the lysates of tumors excised on day 22. (G) Effects of oxaliplatin and PT-100 combination therapy on the expression of cytokines associated with CAFs in CT26 colon cancer cell-derived murine xenograft tumors. Tumors were resected at day 22 and the cytokine expression levels were assessed using reverse-transcription quantitative polymerase chain reaction analysis. *P<0.05 vs. the saline vehicle control. Values are expressed as the mean ± standard error of the mean. CAF, cancer-associated fibroblast; FAP, fibroblast activation protein.
combined with oxaliplatin contained a significantly lower number of imDCs and TIDCs, as compared with the groups treated with saline vehicle, oxaliplatin only and PT-100 only (P<0.05). These results indicated that pharmacological inhibition of CAFs using PT-100 in combination with oxaliplatin significantly reduced the recruitment of M2 macrophages and CD11c+ DCs to the xenograft tumors.

Combined treatment of oxaliplatin with a pharmacological inhibitor of CAFs reduces tumor angiogenesis. Angiogenesis, a key event required for tumor progression, is dependent on ECM remodeling (1). As an important source of ECM components, CAFs are able to promote angiogenesis during tumor growth and metastasis (1,5). In order to test whether pharmacological inhibition of CAFs had any effect on angiogenesis in the tumor xenografts, immunohistochemical analysis was used to determine the density of CD31+ endothelial cells in the xenograft tumors at day 22. The tumors treated with PT-100 combined with oxaliplatin had a markedly lower density of CD31+ cells compared to that in the

Figure 4. Effects of oxaliplatin and PT-100 combination therapy on the number of pro-tumorigenic immune cells in CT26 colon cancer cell-derived murine xenograft tumors. (A-C) Analysis of CD11b+ CD11c+ tumor-associated dendritic cells. (D and E) Analysis of CD11b+ F4/80+ tumor-associated macrophages. Cells were gated according to the isotype control. Upper circle: M2 macrophage, CD11b+ and F4/80+; lower circle: M1 macrophage, CD11b+ and F4/80−. *P<0.05 vs. the saline vehicle control. Values are expressed as the mean ± standard error of the mean. imDC, immature dendritic cells; TIDC, tumor-infiltrating dendritic cells.

Figure 5. Effects of oxaliplatin and PT-100 combination therapy on tumor angiogenesis in CT26 colon cancer cell-derived murine xenograft tumors. The tumor tissues were resected at day 22 and the density of blood vessels was indicated by immunostaining with an anti-CD31 antibody (magnification, x400).
groups treated with saline vehicle, oxaliplatin alone and PT-100 alone (Fig. 5). These results indicated that pharmacological inhibition of CAFs using PT-100 in combination with oxaliplatin significantly reduced tumor angiogenesis.

Discussion

Chemotherapy has been a basic, vital and widespread means of treating colorectal cancer for numerous years; however, chemoresistance has become a significant obstacle. A number of studies have investigated the mechanisms of drug resistance to chemotherapeutic agents; it is mainly attributed to gene mutations, gene amplification, epigenetic changes that influence the uptake, metabolism or export of drugs from single cells (36,37), or alterations to the signaling pathways which protect tumor stem cells (31). The tumor microenvironment has become a research hot spot, as it may be involved in the resistance of solid tumors to chemotherapy (38).

The present study demonstrated that the chemotherapeutic drug oxaliplatin increased the accumulation of CAFs in colon cancer xenograft tumor tissues, which may, to a certain extent, be responsible for drug resistance. The mechanisms by which oxaliplatin increased the accumulation of CAFs may be due to the upregulation of cytokines associated with the accumulation of CAFs. Studies have demonstrated that chemotherapy can increase hypoxia in the tumor microenvironment, and that such hypoxic conditions can induce a molecular response that drives the activation of a key transcription factor, namely the hypoxia-inducible factor (HIF). HIF regulates a variety of genes encoding cytokines that influence the growth, progression and metastasis of tumors (39,40). In the present study oxaliplatin may possibly have induced hypoxia in the tumors, which may explain for the upregulation of TGF-β3, FGF-2 and osteopontin observed in the xenograft tumors. TGF-β is a multifunctional cytokine that regulates tissue morphogenesis and differentiation by influencing cell proliferation, differentiation, apoptosis and ECM production (8,11). FGF-2, a prototypical pro-angiogenic factor, regulates a variety of important intracellular signal-transduction pathways, which mediate a series of cellular and molecular changes associated with the EMT (9). Osteopontin is a soluble ECM protein with known effector functions in tumor growth, angiogenesis and metastasis (12). All of these cytokines have been associated with the accumulation of CAFs (8,9). As a result of the increased accumulation of CAFs, the secretion of these cytokines is elevated and eventually accelerates tumor growth (10-12).

Although the importance of CAFs in tumor progression has been recognized (41), the understanding of the mechanisms by which CAFs influence tumorigenesis remains limited. One generally accepted hypothesis is that an increase in the accumulation of CAFs, a vital component of the structure of the tumor microenvironment, may increase fibrosis in the tumor microenvironment, which may reduce drug uptake by the tumor cells (30). However, the present study suggested that CAFs induce the recruitment of inflammatory immune cells, which may facilitate tumor growth and progression. Tumor-associated macrophages and dendritic cells have been reported to produce matrix remodeling enzymes, reactive oxygen species and other bioactive molecules that influence cancer-cell proliferation, angiogenesis, invasion and metastasis (35,42). In the present study, pharmacological inhibition of CAFs markedly suppressed the recruitment of pro-tumorigenic immune cells in vivo, which indicated that CAFs are able to influence the vital signaling pathways associated with the recruitment of pro-tumorigenic immune cells. It has been reported that CAFs orchestrate signaling processes associated with tumor-promoting inflammation in incipient neoplasia via a NF-κB signaling-dependent mechanism (15), while CAFs also modulated the tumor immune microenvironment in a 4T1 murine breast cancer model (43). In addition, as mentioned above, hypoxic tissues, in which the accumulation of CAFs is enhanced, secrete cytokines that undermine normal anti-tumor surveillance by macrophages, turning the macrophages into accomplices and facilitators of invasion and angiogenesis (44). However, in contrast to previous studies on tumor-associated immune cells (30,43), no significant differences between the number of CD4+/CD8+ T cells were identified among the four experimental groups of the present study. Therefore, the role of CAFs as tumor immune modulators in tumorigenesis requires further study.

Apart from recruiting inflammatory immune cells, the pro-tumor function of CAFs may also be associated with the induction of angiogenesis within the tumor stroma. Angiogenesis, the formation of new blood vessels, is required for malignant tumor growth and metastasis (45). The findings of the present study demonstrated that pharmacological inhibition of CAFs simultaneously suppressed angiogenesis in the tumor xenografts. CAFs may affect angiogenesis at several levels. CAFs secret growth factors, including VEGF and FGF-2, which are known to be important for endothelial cell migration (13). Moreover, CAFs are capable of remodeling connective tissue, and interact with epithelial cells and other cell types in connective tissues, which may promote angiogenesis (46). In addition, inflammatory immune cells may also affect tumor angiogenesis (15,42). The present study demonstrated that pharmacological inhibition of CAFs reduced the recruitment of inflammatory immune cells, which indicated that CAFs may promote tumor angiogenesis by recruiting inflammatory immune cells. It has recently been reported that vascular endothelial cells and CAFs express similar genes (47). Thus, it is necessary to study the similarities and differences between CAFs and tumor endothelial cells in order to accurately guide the clinical application of anti-angiogenic agents.

In conclusion, the present study demonstrated that chemotherapy with oxaliplatin increased the accumulation of CAFs in the stroma of colon tumors in vivo, whereas pharmacological inhibition of CAFs alongside oxaliplatin treatment reduced the accumulation of CAFs, enhanced the response to oxaliplatin chemotherapy as well as reduced the expression of cytokines associated with the accumulation of CAFs, the recruitment of pro-tumorigenic immune cells and angiogenesis. Chemotherapy combined with treatments which inhibit the recruitment, formation or activity of CAFs may represent a novel method of improving the tumor response to chemotherapy and help to overcome resistance to chemotherapeutic agents.

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