

Collagen triple helix repeat containing 1 promotes tumor angiogenesis in gastrointestinal stromal tumors

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Received November 15, 2015; Accepted November 25, 2016

DOI: 10.3892/ol.2017.7111

Abstract. Collagen triple helix repeat containing 1 (Cthrc1) is a secreted protein that has been observed to lead to poorer prognosis by inducing the invasion and metastasis in different tumors; however, it has not been demonstrated that Cthrc1 is involved in tumor angiogenesis. Immunohistochemical staining of Cthrc1 and CD31 in gastrointestinal stromal tumor tissue demonstrated that Cthrc1 is associated with microvascular density. Overexpression of Cthrc1 protein may alter the properties of human umbilical vein endothelial cells (HUVECs), including migration, invasion, tubule formation and aortic ring sprouting. Small interfering RNA-mediated knockdown of Cthrc1 was performed to verify the opposite effects. Migration and tubule formation induced by Cthrc1 overexpression in HUVECs was attenuated by inhibition of phosphorylation in extracellular-signal-regulated protein kinase (ERK) and c-Jun N-terminal kinase (JNK) signaling pathways. The pro-angiogenic effect of Cthcr1 is associated with increased phosphorylation of ERK and JNK in HUVECs. Silencing the expression of Cthrc1 protein may be a promising strategy to inhibit tumor angiogenesis.

Introduction

Angiogenesis is a complex process that includes adhesion, migration, invasion, proliferation and differentiation in endothelial cells (1). Neovascularization is necessary for tumor

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growth by providing oxygen anxd nutrients (2). The tumor and the surrounding microenvironment, including cancer cells, endothelial cells, fibroblasts and immune cells, are associated and interact constantly (3,4). Tumors may influence the microenvironment by releasing extracellular signals, including vascular endothelial growth factor, tumor necrosis factor α , fibroblast growth factor 2 and interleukin 6 into the extracellular matrix (ECM), and promoting tumor-associated angiogenesis (5,6).

Collagen triple helix repeat containing 1 (Cthrc1) was identified to encode a secreted protein that serves a role in the cellular response to arterial injury through vascular remodeling (7). It was demonstrated that Cthrc1 is related to vascular remodeling by inhibiting collagen production and fibrogenesis, and by promoting cell migration (8). The majority of Cthrc1 studies have focused on its effects and underlying molecular mechanism, in promoting tumor cell invasion and metastasis (9-14). In the present study, it was demonstrated that increased expression of Cthrc1 protein in gastrointestinal stromal tumor (GIST) is associated with increased microvascular density (MVD) in a tissue microarray; however, to the best of our knowledge, no previous study has demonstrated the effect of Cthrc1 in endothelial angiogenesis.

Materials and methods

Immunohistochemical (IHC) staining and evaluation. Two continuous tissue microarrays (no. HDgS-GIST060CS-01) including surgical tissues from 60 GISTs were purchased from Shanghai Outdo Biotech Co., Ltd. (Shanghai, China). The 60 patients included 33 females and 27 males, with a median age of 53.4±13.70 years. The location of the samples included 5 cases in the stomach, 21 cases in the small intestine, 7 cases in the colon, 3 cases in the peritoneum and 2 cases in the mesentery. Sections were dewaxed in dimethylbenzene, hydrated in ethanol, incubated with 3% oxydol to inactivate endogenous peroxidase and incubated with a citrate solution for 30 min at 95°C for antigen retrieval. Sections were blocked with goat serum for 30 min and incubated with Anti-Cthrc1 (dilution, 1:200; cat. no. AP8778a; Abgent, San Diego, CA, USA) or anti-CD31 (dilution, 1:200; 11265-1-AP; ProteinTech, Chicago, IL, USA) at 4°C overnight. Sections were then incubated with a horseradish peroxidase (HRP)-labeled

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Key words: collagen triple helix repeat containing 1, extracellular-signal-regulated kinases, c-Jun N-terminal kinase, human umbilical vein endothelial cells, angiogenesis

goat anti-mouse/rabbit antibody (dilution, ready to use: cat. no. D-3004; Shanghai Long Island Biotec. Co., Ltd., Shanghai, China) at 30°C for 30 min and DAB for 30 sec (Fuzhou Maixin Biotechnology Development Co., Ltd., Fuzhou, China) at room temperature, followed by hematoxylin staining and mounting. Images were captured using a microscope (CX31-LV320; Olympus Corporation, Tokyo, Japan). CD31 staining results were evaluated by the amount of vessels per x200 fields (Fig. 1A-C). Cthrc1 staining results were evaluated as follows: -, no tumor cells stained; +, <25% tumor cells stained (Fig. 1D); ++, 25-50% tumor cells stained (Fig. 1E); +++, >50% tumor cells stained (Fig. 1F).

Cell culture. Human umbilical vein endothelial cells (HUVECs) were purchased from China Center for Type Culture Collection (Beijing, China). HUVECs were cultured in Dulbecco's modified Eagle's medium (DMEM; Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) supplemented with 10% fetal bovine serum (FBS; ScienCell Research Laboratories, Inc., Carlsbad, CA, USA) at 37°C in a humidified atmosphere of 5% CO₂ for 72 h prior to use. U0126 (Selleck Chemicals, Houston, TX, USA) and SP600125 (Selleck Chemicals) were used to inhibit extracellular-signal-regulated kinase 1/2 (ERK1/2) and c-Jun N-terminal kinase (JNK) phosphorylation, respectively. U0126 and SP600125 were dissolved in Dimethyl sulfoxide (DMSO; Sigma-Aldrich; Merck Millipore, Darmstadt, Germany), so DMSO was used as a phosphorylation control. The final concentration of U0126 and SP600125 was 10 μ M in the culture medium for the experiments.

Cthrc1-expressing adenoviral vector. An adenoviral vector expressing the human Cthrc1 coding sequence under the control of a mouse cytomegalovirus promoter was constructed as described previously (15). A control adenoviral vector expressing green fluorescent protein was prepared in parallel (15).

Small interfering RNA (siRNA) experiments. Cthrc1 was knocked down using siRNA (siCTHRC1) and the synthetic duplex oligomers 5'-CCCAUUGAAGCUAUAAUUU-3' and 5'-AAAUUAUAGCUUCAAUGGG-3', purchased from Shanghai GenePharma Co., Ltd. (Shanghai, China). HUVECs were transiently transfected with siRNAs using GenMute[™] reagent (SignaGen Laboratories, Rockville, MD, USA), according to the manufacturer's protocol.

Aortic ring assay. A total of 6 male Sprague-Dawley rats (8-weeks old) weighing 320-345 g were purchased from Shanghai SLAC Laboratory Animal Co., Ltd. The mice were sacrificed on the day of purchase. Chloral hydrate (10%) was injected in to the abdominal cavity and the mice were sacrificed via the cervical dislocation method under the influence of the anesthetic. All animal experiments were approved by the Institutional Animal Care and Use Committee of Renji Hospital, School of Medicine, Shanghai Jiao Tong University (Shanghai, China). Aortas were harvested and cut into aortic rings of ~1 mm thickness. A total of ½ were immersed in DMEM with Cthrc1-expressing adenovirus (2x10⁷ pfu/ml), and the other ½ were immersed in DMEM with the control adenovirus (2x10⁷ pfu/ml) at 37°C. Following a total of 4 h, the rings were placed in 96-well culture plates

and coated with liquid MatrigelTM (BD Biosciences, Franklin Lakes, NJ, USA). Following solidification of the Matrigel, 200 μ l Opti-MEM (Thermo Fisher Scientific, Inc.) with 2% FBS was added. The aortic rings were incubated at 37°C, and medium was changed every two days. Images of the aortic rings (3 fields per ring) were captured (original magnification, x200) on the eighth day with a microscope (Eclipse Ti-U; Nikon Corporation, Tokyo, Japan), the images were captured directly from the 96-well culture plates. Photoshop (version CS6; Adobe, San Jose, CA, USA) was used to sharpen the images and count the number of sprouting endothelia from the aortic ring for statistical analysis.

Cell proliferation assay. HUVECs were seeded in 96-well plates at a density of 2,000 cells/well and were incubated at 37° C overnight. Culture medium containing Cthrc1-expressing adenovirus or control adenovirus (1x10⁷ pfu/ml) was added into various wells, and the culture medium was changed after 2 h. The optical density value of each group (six wells/group) was determined using the Cell Counting Kit-8 (CCK-8; Dojindo Molecular Technologies, Inc., Kumamoto, Japan) on days 0, 1, 2 and 3, following the manufacturer's protocol.

Western blotting. Proteins were extracted from tissue samples using total protein extraction kits (cat. no. C510003; Sangon Biotech Co., Ltd., Shanghai, China). The protein concentration was determined using a BCA Protein Assay kit (cat. no. 23252; Thermo Fisher Scientific, Inc.). Proteins (40 μ g) were loaded and electrophoresed on a 10% SDS-PAGE gel. Proteins were transferred to a nitrocellulose membrane (EMD Millipore, Billerica, MA, USA), which was blocked using 5% milk for 2 h at room temperature, and then incubated with the primary antibodies at 4°C overnight. An HRP-labeled secondary antibody (dilution, 1:5,000; cat. no. 7074; Cell Signaling Technology, Inc.) was added and incubated at room temperature for 1 h. The immunoreactive signals were detected using Super Signal West Femto Maximum Sensitivity Substrate (Thermo Fisher Scientific, Inc.). The primary antibodies were anti-Cthrc1 (dilution, 1:1,000; cat. no. AP8778a; Abgent), anti-ERK1/2 (dilution, 1:1,000; cat. no. 9102; Cell Signaling Technology, Inc.), anti-phospho (P)-ERK1/2 (dilution, 1:1,000; cat. no. 4370; Cell Signaling Technology, Inc.), anti-JNK (dilution, 1:1,000; cat. no. 9252; Cell Signaling Technology, Inc.), anti-P-JNK (dilution, 1:1,000; cat. no. 4668; Cell Signaling Technology, Inc.) and anti-β-actin (dilution, 1:4,000; cat. no. 8457; Cell Signaling Technology, Inc.). Western blots were analyzed densitometrically using ImageJ software version 1.45S (National Institutes of Health, Bethesda, MD, USA).

Tubule formation assay. Matrigel was pipetted into 48-well plates and allowed to polymerize for 30 min at 37°C. HUVECs ($6x10^4$) were seeded on Matrigel in 200 μ l culture medium. The plates were incubated at 37°C for 6-8 h, prior to the capture of images using an Eclipse Ti-U microscope (magnification, x100). ImageJ was used to measure the tube length in μ m. The sum of tube length was determined for statistical analysis.

Scratch wound assay. HUVECs were cultured to full confluence in 6-well culture plates. The cell layer was scratched using a sterile pipette tip and the scratched area was imaged





Figure 1. (A-C) IHC staining of different MVDs in gastrointestinal stromal tumor tissue arrays. Representative images of (D) +, (E) ++ and (F) +++ Cthrcl IHC staining. (G) Expression of phosphorylated and total ERK1/2 and JNK in HUVECs treated with Con or Cthrc1-expressing adenovirus determined using western blotting. β -Actin was used as a loading control. (H) Quantitative analysis of the P-JNK/total JNK and P-ERK1/2/total ERK ratios determined from western blots. (I) Cell proliferation of HUVECs transfected with Con or Cthrc1-expressing adenovirus was determined using the Cell Counting Kit-8 assay. (J) Images of aortic rings treated with Con or Cthrc1-expressing adenovirus after 8 days. (K) Quantitative analysis of aortic ring-sprouting endothelial cells transfected with control or Cthrc1-expressing adenovirus after 8 days. ^{**}P<0.001. IHC, immunohistochemical; MVD, microvascular density; Cthrc1, collagen triple helix repeat containing 1; ERK, extracellular-signal-regulated kinase; JNK, c-Jun N-terminal kinase; HUVEC, human umbilical vein endothelial cell; P-, phospho.

with an Eclipse Ti-U microscope (original magnification, x200). Following incubation at 37°C for 24 h, the scratched area was imaged again. ImageJ was used to measure the scratched area in μ m². The healing area was evaluated for statistical analysis.

Cell invasion assay. A Transwell chamber (24-well insert; pore size 8 μ m; EMD Millipore) was coated with Matrigel for 6 h prior to the invasion assay. A total of 2x10⁵ HUVECs were plated in the Transwell chamber with 200 μ l FBS-free DMEM. DMEM (600 μ l) containing 20% FBS was added to the lower

chambers of the 24-well plates. After 36 h of incubation at 37°C, the cells that had invaded through the pores were fixed using 4% paraformaldehyde (25 min) and stained using 0.5% crystal violet (30 min) at room temperature, and the number of cells was counted under a x200 Eclipse Ti-U light microscope.

Statistical analysis. Each individual experiment was conducted with three or five replicates. All data are presented as the mean \pm standard error, and were analyzed using the Student's t-test or Mann-Whitney U test with SPSS (version 18.0; SPSS, Inc., Chicago, IL, USA). P<0.05 was considered to indicate a statistically significant difference.

Results

MVD is associated with the tumor site and Cthrc1 protein expression level. Quantification of the IHC staining demonstrated that there was a significant decrease in MVD in tumors located in the stomach compared with non-stomach-located tumors (8.19 ± 6.97 vs. 16.48 ± 17.04 , respectively; P=0.003; Table I). GIST with increased expression of Cthrc1 protein exhibited increased MVD compared with GIST with decreased expression of Cthrc1 (14.44 ± 15.35 vs. 7.67 ± 7.05 , respectively; P=0.021; Table I). Statistical analysis indicated that MVD was not significantly associated with sex, mitotic index, tumor size or National Institutes of Health risk grade (Table I).

Effect of Cthrc1 expression on aortic ring sprouting. To investigate the effect of Cthrc1 on angiogenesis, an aortic ring assay was performed. The aortic ring assay demonstrated that Cthrc1-overexpressing aortic rings developed a significantly increased number of sprouting endothelial cells compared with the control (P<0.001), indicating that Cthrc1 serves an important role in angiogenesis (Fig. 1G and H).

Effect of Cthrc1 overexpression on HUVEC proliferation, tubule formation, migration and invasion. Densitometric analysis of western blots (Fig. 1I) demonstrated significantly increased ratios of P-JNK/total JNK (P=0.004) and P-ERK1/2/total ERK1/2 (P=0.005) in HUVECs transfected with Cthrc1-expressing adenovirus compared with HUVECs transfected with control adenovirus (Fig. 1J). CCK-8 results demonstrated that HUVEC proliferation was not significantly affected by Cthrc1-expressing adenovirus (Fig. 1K), indicating that Cthrc1 promotes aortic ring sprouting without promoting HUVEC proliferation. The scratch wound assay and the tubule formation assay revealed that Cthrc1 protein overexpression was able to increase the migratory function of HUVECs (Fig. 2A and B), and increase HUVEC tubule formation (Fig. 3A and B). The cell invasion assay demonstrated that Cthrc1 protein overexpression was able to increase HUVEC invasion (Fig. 3C and D). Decreased HUVEC migration and tubule formation were observed following treatment with the kinase inhibitors U0126 and SP600125 (Figs. 2A and B, 3A and B), suggesting that ERK1/2 and JNK phosphorylation may be the underlying molecular mechanism of promotion of angiogenesis by Cthrc1.

Effect of Cthrc1 knockdown on HUVEC tubule formation, migration and invasion. Following Cthrc1 knockdown using Table I. Characteristics of patients with gastrointestinal stromal tumors.

Characteristic	MVD (number of vessels/field)	P-value
Sex		0.899
Male (n=33)	12.45±13.70	
Female (n=27)	13.11±14.62	
Mitosis per 50 HPFs		0.947
<5 (n=32)	12.91±13.68	
≥5 (n=28)	12.57±14.62	
Tumor size, cm		0.169
<10 (n=44)	11.57±12.13	
≥10 (n=16)	16.00 ± 18.28	
NIH risk grade		0.111
Low and intermediate (n=41)	11.73±12.50	
High (n=19)	14.95±16.99	
Tumor site		0.003ª
Stomach (n=27)	8.19±6.97	
Non-stomach (n=33)	16.48±17.04	
Cthrc1 protein expression		0.021ª
-, + (n=15)	7.67±7.05	
++, +++ (n=45)	14.44±15.35	

^aP<0.05. MVD, microvascular density; HPF, high-power field; NIH, National Institutes of Health; Cthrc1, collagen triple helix repeat containing 1; -, no tumor cells stained; +, <25% tumor cells stained; ++, 25-50% tumor cells stained; +++, >50% tumor cells stained.

siRNA in HUVECs, the ability of HUVECs to form tubules, migrate and invade was significantly decreased (Fig. 4), indicating that inhibition of Cthrc1 protein expression may be a promising method of inhibiting GIST angiogenesis.

Discussion

Previous studies have demonstrated that patients with increased expression of Cthrc1 protein have a poorer prognosis compared with those with decreased expression, a phenomenon that has been observed in numerous types of tumor (9-12,16). Tumor MVD is negatively associated with tumor prognosis. The present study demonstrated that Cthrc1 protein expression is associated with GIST MVD. The aortic ring assay demonstrated that Cthrc1 was able to promote aortic ring sprouting. The wound healing assay, cell invasion assay and tubule formation assay demonstrated that Cthrc1 was able to promote HUVEC migration, invasion and tubule formation. The results of the present study indicated that Cthrc1 serves an important role in GIST angiogenesis and may explain why patients with increased expression of Cthrc1 protein have a poorer prognosis.

Cthrc1 has been demonstrated to promote the phosphorylation of ERK and JNK in colon cancer (12) and GIST (9), respectively. Previous studies have revealed that the increase in phosphorylation of ERK and JNK may promote HUVEC





Figure 2. Effect of Cthrc1 overexpression on HUVEC migration. (A) Scratch wound assay of HUVECs treated with DMSO, U0126 and SP600125 following transfection with Con adenovirus and Cthrc1-expressing adenovirus. (B) Quantitative analysis of HUVEC wound healing. **P<0.001. HUVEC, human umbilical vein endothelial cell; DMSO, dimethyl sulfoxide; Cthrc1, collagen triple helix repeat containing 1.

migration and tubule formation (17-19). In the present study, it was demonstrated that Cthrc1 was able to promote ERK and JNK phosphorylation in HUVECs, and inhibition of ERK and JNK phosphorylation may decrease HUVEC migration and tubule formation. These results indicate that the pro-angiogenic effect of Cthrc1 is associated with the phosphorylation of ERK and JNK.

The planar cell polarity (PCP) signaling pathway is a highly conserved signaling cascade that coordinates epithelial and axonal morphogenic movements during organ development by regulating angiogenesis (20). Inhibition of the PCP signaling pathway disrupts endothelial cell growth, polarity and migration (20-22). Cthrcl is a Wnt co-factor protein that selectively activates the Wnt/PCP signaling pathway by stabilizing ligand-receptor interactions (9,23,24). Cthrcl may activate the PCP signaling pathway; however, it was demonstrated to suppress the canonical Wnt signaling pathway in human embryonic kidney-293T and GIST cells (9,23). Therefore, Cthrc1 may activate the PCP signaling pathway in HUVECs and promote tumor angiogenesis.

Collagen matrix deposition in ECM is an important process in the inhibition of tumor invasion and angiogenesis (25,26). Cthrc1 may reduce collagen type I mRNA and protein levels in fibroblasts, resulting in decreased collagen synthesis and contributing to vascular remodeling (27). Matrix metalloproteinase 9 (MMP9) is an important factor in collagen degradation (28). Cthrc1 has been demonstrated to promote MMP9 secretion in colon cancer (12). Therefore, inhibiting Cthrc1 may increase collagen matrix deposition and decrease tumor invasion and angiogenesis.



Figure 3. Effect of Cthrc1 overexpression on HUVEC proliferation, tubule formation, migration and invasion. (A) Tubule formation assay of HUVECs treated with DMSO, U0126 and SP600125 in control and Cthrc1-expressing adenovirus groups. (B) Representative images of HUVECs transfected with control or Cthrc1-expressing adenovirus that had invaded through Matrigel to the bottom of the Transwell filter. (C) Quantitative analysis of tubule formation assay results. **P<0.001. HUVEC, human umbilical vein endothelial cell; DMSO, dimethyl sulfoxide; Cthrc1, collagen triple helix repeat containing 1.



Figure 4. Cthrcl knockdown decreases HUVEC tubule formation, migration and invasion. (A) Scratch wound assay of HUVECs treated with control siRNA or siCTHRC1. (B) Quantitative analysis of scratch wound assay results. (C) Tubule formation assay of HUVECs transfected with control siRNA or siCTHRC1. (D) Quantitative analysis of tubule formation assay results. (E) Cell invasion assay images of HUVECs transfected with control siRNA or siCTHRC1. (F) Quantitative analysis of invasion assay results. *P<0.001. Cthrc1, collagen triple helix repeat containing 1; HUVEC, human umbilical vein endothelial cell; siRNA, small interfering RNA; siCTHRC1, siRNA targeting Cthrc1.



Although the present study has revealed certain underlying molecular mechanisms regarding the pro-angiogenic effect of Cthrc1, further studies *in vivo* and *in vitro* are required to validate these experimental findings.

Acknowledgements

The present study was supported by the National Natural Science Foundation of China (grant no. 81270474). The authors of the present study thank Professor Xiong Ma (Shanghai Jiao-Tong University, Shanghai, China) for providing the Cthrc1-expressing adenovirus and the control adenovirus.

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