Suppression of scinderin modulates epithelial-mesenchymal transition markers in highly metastatic gastric cancer cell line SGC-7901

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Abstract. Scinderin is a Ca²⁺-dependent filamentous actin (F-actin) severing and capping protein, which has a key role in regulated secretion. However, little is known regarding the function and mechanism of scinderin in human carcinoma development and progression. In the present study, the biological function of scinderin was investigated using a cell proliferation assay, flow cytometric analysis and a Transwell assay in highly tumorigenic and the metastatic human gastric cancer cell line SGC-7901 transfected with scinderin-small hairpin RNA lentivirus. The changes in the expression of epithelial-mesenchymal transition (EMT) markers were also investigated. The results indicated that scinderin knockdown effectively suppressed proliferation, reduced migration and arrested the cell cycle of the SGC-7901 cells at G2/M phase. Furthermore, scinderin knockdown altered the expression of EMT markers; the expression of E-cadherin was significantly upregulated, along with an evident decrease in N-cadherin and β-catenin expression. In conclusion, the present study suggested that suppression of scinderin impaired proliferation and migration of gastric cancer SGC-7901 cells and attenuates its EMT process. Scinderin may therefore be a potential target for tumor EMT and therapy against gastric cancer.

Introduction

Scinderin (or adseverin), a member of the gelsolin superfamily, is a Ca²⁺-dependent filamentous actin (F-actin) severing and capping protein with three actin-, two PIP₂- and two Ca²⁺-binding sites (1‑4). Scinderin is expressed largely in endocrine tissues and secretory cells. During human development, scinderin expression is different from that in adults: It is highly expressed in the human fetal kidney, very weakly in the brain and intestine but nowhere else; in human adult tissues, scinderin exhibits strong expression in the kidney and low expression in the heart; however, no expression is observed in the adrenal gland (5).

It has been demonstrated that scinderin has important roles in exocytosis, megakaryopoiesis, autoimmune disorders and tumorigenesis. Scinderin is known to regulate translocation of secretory vesicles by controlling F-actin dynamics (disassembly ↔ assembly) during secretion (6‑8). Overexpression of scinderin on human airway epithelial cells may inhibit mucin secretion (9). Scinderin has also been found in platelets. Previous studies have suggested that the expression of scinderin in the megakaryoblastic cell line MEG-01 induces cell differentiation, polyploidization, maturation and apoptosis with release of plateletlike particles, while cell proliferation and tumorigenesis in nude mice are inhibited (10). It has been reported that single-nucleotide polymorphisms (SNPs) in scinderin have a prominent effect in multiple sclerosis (11). Overexpression of scinderin triggers human non-small-cell lung carcinoma cell line IGR-Heu resistance to cytolytic T lymphocytes (12). Furthermore, scinderin is capable of preventing mitochondria-mediated apoptosis by directly binding to voltage-dependent anion channels in ciplatin-resistant cells (13).

Gastric cancer is one of the most common primary gastrointestinal cancers in the world. Despite improvements in diagnostic modalities, outcomes for patients with gastric cancer remain extremely poor, with the majority of cancer patients dying from the occurrence of metastases rather than their primary tumors. Due to its complexity, the process of tumor metastasis remains poorly understood. Epithelial-mesenchymal transition (EMT) is a transdifferentiation process during which epithelial cells lose their epithelial polarity and cell-cell contacts, gain mesenchymal

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Abbreviations: F-actin, filamentous actin; EMT, epithelial-mesenchymal transition; RTCA, real-time cell analyzer; RT-qPCR, real-time quantitative polymerase chain reaction

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properties, and become motile and invasive (14,15). In this process, fundamental changes in gene expression are involved in the disruption of epithelial polarity (e.g., E-cadherin and cytokeratins) and establishment of a mesenchymal phenotype (e.g., N-cadherin and vimentin) (16). EMT is physiologically relevant during embryogenesis and in adult tissue for wound healing. However, EMT is also associated with tumor metastasis. Of note, the majority of solid carcinoma cells (including breast, prostate, pancreatic, colon and gastric cancer, etc.) have been reported to undergo either partial or complete EMT so as to become motile and invasive. Currently, EMT is regarded as an important mechanism of tumor metastasis, having crucial roles in the early process of therioma metastases (15). Therefore, it is essential to find molecules involved in both the positive and negative regulation of EMT. It has been demonstrated that signaling pathways, including TGF-β, Wnt/β-catenin, Notch, Hedgehog, interleukin-6/signal transducer and activator of transcription 3 and nuclear factor-κB trigger EMT by inducing Snaill, Snai1, Twist1, Twist2, zinc finger E-box binding homeobox (ZEB1) and ZEB2 expression (17-19). Furthermore, non-coding RNAs (including microRNAs and long non-coding RNAs) also have critical roles in the regulation of the EMT (20-22).

Although it has been demonstrated that scinderin expression is low or absent in the human adult stomach, no studies, to the best of our knowledge, have reported the biological effect of scinderin on human gastric cancer. In the present study, the effects of scinderin knockdown on cell cycle, proliferation and migration of the highly tumorigenic and metastatic human gastric cancer cell line SGC-7901 (scinderin is highly expressed) were investigated. Furthermore, the changes in E-cadherin, N-cadherin and β-catenin expression following scinderin knockdown were examined and the role of scinderin in the tumor EMT process was analyzed.

Materials and methods

SGC-7901 cell line. The human gastric cancer cell line SGC-7901 was purchased from Chinese Academy of Sciences (Shanghai, China). The cells were cultured in RPMI-1640 medium (Invitrogen Life Technologies, Carlsbad, CA, USA) with 10% fetal bovine serum (FBS) in a humidified incubator containing 5% CO₂ at 37℃.

Construction of scinderin-small hairpin (sh)RNA lentiviral plasmid. The target shRNA against the scinderin gene (Gene Bank accession NM_033128) was designed as follows: 5’-CGA GAT GAG CTG ACA ACA T-3’ and 5’-GAG GCC AGA TCT TGG GTG-3’. Oligonucleotides encoding shRNA sequences were synthesized (Genechem Corporation, Shanghai, China) and annealed into double strands. Double-stranded DNAs were inserted into Hpal/XhoI restriction sites of lentiviral frame plasmid (Genechem Corporation), encoding green fluorescent protein (GFP). They were then transfected into Escherichia coli and positive recombinant lentiviral plasmids were selected by polymerase chain reaction (PCR), using the primers 5’-GCC CCG GTT AAT TTG CAT AT-3’ and 5’-GAG GCC AGA TCT TGG GTG-3’. The conditions for PCR were denaturation at 94℃ for 30 sec, then 94℃ for 30 sec, 55℃ for 30 sec and 72℃ for 30 sec, for 30 cycles and extension at 72℃ for 6 min. The products were then verified by electrophoretic analysis on a 1.5% agarose gel containing ethidium bromide and DNA sequencing. The lentiviral vectors expressing GFP alone were used as the control.

Packaging and titration of lentiviral vectors. The recombinant lentiviral plasmid was co-transfected into 293T cells with packaging plasmids (pHelper 1.0 including gag/pol and pHelper 2.0 including VSVG; Genechem Corporation) by Lipofectamine 2000 (Invitrogen Life Technologies) to produce target lentivirus. Following 48 h, the virus in the supernatant was collected and the virus titer was measured following the dilution method: i) a total of 4x10^6 cells/well of the 293T cells were seeded in a 96-well plate in Dulbecco's modified Eagle's medium (Gibco Life Technologies, Carlsbad, CA, USA) supplemented with 10% FBS and cultured overnight; ii) the serial diluted vectors were added to the cultured cells and continued to culture for 48 h; iii) the GFP-positive cells were counted. The titer was represented as transduction unit (TU) per milliliter concentrated vector (TU/ml): Titer = GFP-positive cell number/dilution x 10^7.

Establishment of the stable scinderin-silenced cell line SGC-7901. A preliminary experiment identified that the best multiplicity of infection (MOI) of SGC-7901 cells was 10 (MOIs of 1, 10 and 20 were tested). Under these best transfection conditions, scinderin-shRNA lentivirus was transfected into the SGC-7901 cells. The efficiency of transfection was observed under a fluorescence microscope (MicroPublisher 3.3 RTV; Olympus, Tokyo, Japan). The effects of gene silencing were confirmed by quantitative (q)PCR and western blot analyses. The cells were divided into three groups: The CON group (uninfected), the NC group (transfected with negative control lentivirus) and the KD group (transfected with scinderin-shRNA lentivirus).

Cell migration assay. Cell migration was assessed using a Transwell assay (Corning Incorporated Life Sciences, Lowell, MA, USA). Transwell insert chambers were firstly rehydrated with 100 µl serum-free medium at 37℃ for 1 h. Following removing the medium from the chambers, 1x10^4 cells in 100 µl serum-free medium were added to the upper chamber, with 600 µl medium containing 30% FBS in the lower chamber, and then cultured for 24 h. Subsequently, the cells were removed from the top side of the membrane using a cotton-tipped swab, and the invaded cells attached to the bottom of the membrane were fixed with 10% glutaraldehyde, stained with 0.1% crystal violet (Shanghai Genebase Gene-Tech Co., Ltd, Shanghai, China) and photographed under an inverted microscope (MicroPublisher 3.3 RTV; Olympus). Next, the cells were destained with 10% acetic acid, and the optical density (OD) of the solution was measured at 570 nm. The OD value was directly proportional to the cell quantity and indirectly represented the cellular migration ability.

RT-qPCR. Total RNA was extracted from cells using TRIzol (Invitrogen Life Technologies), according to manufacturer's instructions. Quantitative RNA expression was measured by the SmartSpec™ Plus Spectrophotometer (Bio-Rad, Hercules, CA, USA). Reverse transcription in a 20 µl system was performed using the GoTaq® 2-step RT-qPCR System kit (Promega Corporation, Madison, WI, USA) following the manufacturer's
instructions. qPCR was performed using the MX3005P qPCR System (Stratagene, La Jolla, CA, USA). Primers for qPCR were scinderin, forward 5'-TCT GCG TTC CTG ACT GTT C-3' and reverse 5'-GAC CTC TTT TCT TTG ATG TTC C-3'; GAPDH, forward 5'-TGA CTT CAA CAG CGA CAC CCA-3' and reverse 5'-CAC CCT GTT GCT GTA GCC AAA-3'; E-cadherin, forward 5'-CCA TCG CTT ACA CCA TTC T-3' and reverse 5'-GCT GTT GCT GTT GTG TT-3'; N-cadherin, forward 5'-CTC CTA TGA GTG GAA CAG GAA CG-3' and reverse 5'-TTG GAT CAA TGT CAT AAT CAA GTG CTG TA-3'; β-actin, forward 5'-TGC TGC GTG GAA CGA GAA CG-3' and reverse 5'-AAG GAA GGC TGG AAG AGT-3'. The relative mRNA expression was calculated with the \(2^{-\Delta\Delta C_{T}}\) method. GAPDH or β-actin were used as normalizers for each sample. All of the experiments were repeated in biological duplicate.

**Western blot analysis.** The cells were lysed with radioimmuno precipitation assay (RIPA) lysis buffer (Biotek Corporation, Beijing, China), and the protein concentration was measured by bicinchoninic acid assay (Beyotime Corporation, Nantong, China). The total cell proteins were separated by 10% SDS-PAGE and transferred to nitrocellulose filter (NC) membranes (Millipore Corporation, Bedford, MA, USA). The NC membranes were firstly blocked by 5% skimmed milk for 1 h and then incubated overnight at 4°C with primary antibodies, including scinderin rabbit mAb (1:1,000, Santa Cruz Biotechnology, Dallas, TX, USA), GAPDH mouse mAb (1:1,000, Santa Cruz Biotechnology), E-cadherin rabbit mAb (1:1,000, Cell Signaling Technology, Danvers, MA, USA), N-cadherin rabbit pAb (1:1,000, Cell Signaling Technology), β-catenin rabbit mAb (1:1,000, Cell Signaling Technology) and β-actin rabbit mAb (1:1,000, Cell Signaling Technology). Thereafter, blots were stained with fluorescent secondary antibodies IRDye 800CW Goat anti-Rabbit IgG (H+L) (1:10,000, LI-COR Bioscience, Lincoln, NE, USA) or IRDye 800CW Goat anti-Mouse IgG (H+L) (1:10,000, LI-COR Bioscience) and protein bands were visualized using an Odyssey Imaging system (LI-COR Bioscie). GAPDH or β-actin was used as the internal control.

**Real-time cell proliferation assay (RTCA).** The real-time cell proliferation experiment was performed with a RTCA single plate (SP) instrument (Roche Applied Science, Mannheim, Germany), which was placed in a 37°C incubator with 5% CO₂. Based on the manufacturer's instructions, the background signal of the culture medium was first measured by adding 100 μl medium to 96-well plates (E-plate 96, Roche Applied Science) containing gold microelectrodes on its bottom. Next, the cells were seeded in the special 96-well plates at density of 8,000 cells/well. Following 30 min of incubation at room temperature, the cells were placed in the RTCA SP and monitored every 15 min for 160 h. Data analysis was performed using RTCA software 1.2 supplied with the instrument. The RTCA software comprised the xCELLigence system, which converts impedance values into cell index (CI) values corresponding to each well. The CI value is directly proportional to the quantity of cells and was used as an indirect measure for cellular proliferation capability.

**Cell cycle analysis.** SGC-7901 cells stably transfected with scinderin-shRNA or empty lentiviral vector were firstly cultured in serum-free medium for 24 h for synchronization in G0/G1 phase, and then cultured in RPMI-1640 medium with 10% FBS for another 24 h. The collected cells were washed twice with phosphate-buffered saline (PBS) and fixed in ice-cold 70% ethanol overnight at -20°C. Next, the cell pellets were washed twice with ice-cold PBS and stained with propidium iodide solution (BD Biosciences, San Jose, CA, USA) for 20 min in the dark. Finally, the cells were analyzed using the BD FACSCalibur Flow Cytometer (BD Biosciences) for their DNA content. The percentages of cells in the different phases of the cell cycle were determined with Cell Quest Pro software 3.1 (BD Biosciences).

**Statistical analysis.** Data are presented as the mean ± standard deviation. The Student's t-test was used in quantitative data analysis, P<0.05 was considered to indicate a statistically significant difference. All statistical analysis was performed with SPSS 20.0 software (SPSS, Inc., Chicago, IL, USA).

**Results**

**Construction of scinderin-shRNA lentiviral plasmid.** The double-stranded DNAs encoding shRNA sequences were inserted into HpaI/XhoI restriction sites of the lentiviral frame plasmids. The electrophoresed PCR products of the recombinant lentiviral plasmids were as follows: The length of positive clones containing shRNA was 343 bp and the length of blank clones was 299 bp (Fig. 1). Furthermore, DNA sequencing demonstrated that the RNA interference sequence targeting scinderin was coincident with the anticipative results (data not shown). This suggested that the shRNA sequence targeting scinderin was successfully inserted into the lentiviral plasmid.

**Packaging and titration of lentivirus.** Recombinant lentivirus were produced in 293T cells co-transfected with the recombinant lentiviral plasmid and packaging plasmids. The viral titer indicated that the recombinant lentivirus was packaged successfully and the viral titer was \(6\times10^8\) TU/ml (Fig. 2).

**Transfection efficiency of lentivirus in gastric cancer cell line SGC-7901.** The preliminary experiment found that the optimal

![Figure 1. Electrophoresed polymerase chain reaction products of recombinant lentiviral plasmids. Lane 1, ddH₂O; lane 2, non-shRNA lentiviral plasmids; lane 3, marker; lane 4, scinderin-shRNA recombinant lentiviral plasmids. shRNA, small hairpin RNA.](image-url)
MOI of SGC-7901 cells was 10. Following transfection under these optimal conditions for 96 h, the transfection efficiency was determined to be >90% by counting GFP-positive cells under the fluorescent microscope. The result demonstrated that scinderin-shRNA lentivirus was transfected into SGC-7901 cells successfully.

Interference efficiency of scinderin in SGC-7901 cells. The silencing of scinderin was confirmed by qPCR and western blot analyses. The results demonstrated that following transfection into the SGC-7901 cells, both the mRNA and protein levels of scinderin were significantly decreased in the KD group compared with the CON and NC groups (Fig. 3). This suggested that stable scinderin-silenced cancer cells were established and appropriate for use in subsequent experiments.

Scinderin knockdown suppresses migration of SGC-7901 cells. The migration assay demonstrated that the number of cells penetrating the membrane of chambers was significantly lower in the KD group than that in the NC group (P<0.05, Fig. 4). This result indicated that scinderin knockdown may attenuate migration of SGC-7901 cells.

Scinderin mediates EMT marker expression in SGC-7901 cells. Considering scinderin knockdown inhibits the migration of the metastatic human gastric cancer cell line SGC-7901, it was investigated whether scinderin mediates EMT, a critical event in tumor metastases. qPCR and western blot analyses revealed that concomitant with an evident decrease in N-cadherin expression, E-cadherin expression in the KD group was significantly upregulated compared with that in the NC group (Fig. 5). Furthermore, it was observed that scinderin knockdown decreased the expression of β-catenin protein (Fig. 5B), an important regulatory molecule in EMT. Therefore, these results illustrated that scinderin knockdown may inhibit the EMT process in SGC-7901 cells.

Scinderin knockdown suppresses proliferation and arrests cell cycle of SGC-7901 cells. Furthermore, the effect of scinderin knockdown on cell proliferation was also investigated. The results of RTCA proliferation (Fig. 6A) demonstrated that there were no significant differences in the CI values between the NC and KD groups at the initial stage (P>0.05). Following 24 h, however, the proliferation of SGC-7901 cells in the KD group was significantly lower than that in the NC group (P<0.01 or P<0.001) and entered the stationary stage in advance. These data clearly indicated that scinderin knockdown effectively suppressed the proliferation of SGC-7901 gastric cancer cells.

The cell cycle distribution based on DNA content was examined by flow cytometric analysis. It was identified that there was a 1.03% increase in the number cells in G2/M-phase in the KD group compared with the NC group (P<0.01, Fig. 6B). These results suggested that scinderin knockdown in SGC-7901 cells may arrest the cell cycle at G2/M phase.

Discussion

A study by Haifu Wu demonstrated that scinderin is the most differentially expressed gene in colon cancer patients with and without liver metastasis by gene-chip technology and qPCR assay (23). The study also identified, consistent with the results of the present study, that scinderin knockdown in colorectal cancer cell lines SW480 and DLD-1 evidently inhibited cell proliferation and migration. These results suggested that scinderin has important roles in the development and progression of cancer.

The present analyses of scinderin knockdown in highly metastatic SGC-7901 cells suggested that scinderin is associated with cell migration. It is well-established that the actin cytoskeleton contributes to maintaining the distinctive structures and functions of epithelial cells by mediating interactions with the cellular basement membranes and cell-cell contacts (24). Aberrance of the actin cytoskeleton is considered as one fundamental characteristic of the majority of malignant and metastatic cells. Studies performed in epithelial cells have demonstrated that gelsolin and villin, two proteins associated with the scinderin family, regulate cell migration by altering actin filament dynamics (25,26). Scinderin is an important actin-capping, -severing and -nucleating protein. Since it can depolymerize actin filaments and bind actin monomers, it is hypothesized that the loss of scinderin in epithelial cells may increase the cellular levels of F-actin, thus modifying actin filament dynamics. Another role of scinderin may serve to regulate the proper distribution of F-actin filaments in cells.

Furthermore, scinderin may also regulate the EMT process to affect cell migration. The present study indicated that scinderin knockdown in SGC-7901 cells resulted in a high increase of E-cadherin and reduced the expression of N-cadherin. E-cadherin as an important Ca²⁺-dependent adhesion molecule has key roles in the cell adhesion of solid tissues. It is anchored to the actin cytoskeleton via α-γ- and β-catenin, providing the physical structure for both cell-cell attachment and the recruitment of signaling complexes (27). Loss or inactivation of E-cadherin is generally regarded as the main trigger for the disruption of tight epithelial cell-cell contacts, causing migratory or invasive states of malignant cells. In several carcinoma types, a deficiency of E-cadherin expression is most commonly accompanied by the gain of N-cadherin, which has been established as the cadherin switch in the EMT process. Although the functional implication of this cadherin switch for tumor progression has remained unknown, the overexpression of N-cadherin may be equally necessary and sufficient to overcome E-cadherin-mediated cell-cell adhesion and to promote therioma development. Of note, gelsolin has also been found to regulate EMT in human mammary epithelial cells. However, this is inconsistent with the present results that gelsolin knockdown by siRNA in MCF10A cells induces EMT, controlling E-cadherin and N-cadherin conversion via Snail (28).

Notably, the present study identified that the loss of scinderin reduced the expression of β-catenin protein, a key nuclear effector of canonical Wnt signaling in the nucleus. The Wnt/β-catenin pathway is one of the fundamental signaling pathways controlling EMT, which may facilitate the expression of key transcriptional repressors that target E-cadherin, including Snail2, ZEB1 and Twist (29-31). Based on present knowledge, it is suggested that overexpression of scinderin may enhance free cytoplasmic levels of β-catenin by altering the actin cytoskeleton or other mechanisms. β-catenin, escaping cytoplasmic degradation, translocates into the cell nucleus and triggers transcription of Wnt-specific genes (32).
As a result, the expression of E-cadherin, a key hallmark of EMT, is highly aberrant in epithelial cells. Cadherins appear to directly affect the function of each other. N-cadherin, one of the mesenchymal cadherins, is highly expressed at the same time and facilitates the acquisition of a migratory phenotype. Whether this conjecture is valid remains to be investigated in the future. The physiological function of scinderin is remains limited to actin filaments. Thus far, it has been confirmed that changes of megakaryoblastic cells brought about by scinderin expression are mediated through the activation of scinderin.
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of rho family of guanosine triphosphatase/p21 activated kinase/mitogen-activated protein kinase kinase kinase. Mitogen-activated protein kinase kinase 4/c-Jun N-terminal kinase/c-Jun, c-fos and Raf/mitogen-activated protein kinase kinase/extracellular signal-regulated kinase pathways, the only molecular mechanisms reported that scinderin participates in (10). However, the present study indicated that scinderin may act as a potential positive regulator in the EMT process of gastric cancer cells.

In addition, the present study identified that the cell cycle was arrested in G2/M phase (based on DNA content), which may lead to impairment of proliferation of SGC-7901 cells with low scinderin expression. However, a recent study showed that scinderin silencing reduced the proliferation and colony formation of prostate cancer cell line PC3 by arresting the cell cycle at the G0/G1 phase (33). The mRNA level of cell cycle-related molecules was also assessed by RT-qPCR analysis, indicating that scinderin silencing induces upregulation of p21wa1/cip1 and p16 and downregulation of cyclin A2 in PC3 cells. In view of the selectivity of gene expression in different tissues, these two studies perhaps do not appear to be consistent: SGC-7901 cells are derived from lymph node metastasis of gastric cancer, and PC3 cells are from bone metastasis in prostate cancer, both of which are in different microenvironment for tumor growth and survival. Flow cytometry is unable to distinguish between G2 and M phase of the cell cycle, which may explain why the present study demonstrated that scinderin knockdown significantly suppressed the proliferation of SGC-7901 cells.

Figure 5. Expression of important EMT markers including E-cadherin, N-cadherin and β-catenin. (A) Quantitative reverse-transcription polymerase chain reaction analysis of E-cadherin and N-cadherin expression in SGC-7901 cells treated with scinderin knockdown. (B) Western blot analysis of E-cadherin, N-cadherin and β-catenin expression. β-actin was used as a loading control. Values are presented as the mean ± standard deviation of triplicate assays (*P<0.01, **P<0.001 as compared with NC group). EMT, epithelial-mesenchymal transition; NC, empty vector group; KD, experimental (scinderin-shRNA) group.

Figure 6. Effect of scinderin knockdown on cell proliferation and cell cycle distribution. (Ai) RTCA proliferation results and (ii) data analyses after 0, 24, 48, 72 and 96 h; scinderin knockdown significantly inhibited cell proliferation. (B) Stable transfected SGC-7901 cells were stained with propidium iodide and analyzed for cell-cycle distribution. All results represent the average of three independent experiments. *P<0.05, **P<0.01 and ***P<0.001 compared with NC group. RTCA, real-time cellular analysis; NC, empty vector group; KD, experimental (scinderin-shRNA) group.
but the changes in cell-cycle distribution were not evident. As is well-established, microfilaments have a number of important roles in the telophase of cell division. The function of scinderin in regulating actin dynamics has already been clearly demonstrated. It was therefore suggested that scinderin may effect cell proliferation by regulating F-actin; however, further studies are required to confirm this hypothesis.

The metastases of carcinomas are formed following a complex succession of cell-biological events including local invasion and metastatic colonization (34). As mentioned previously, scinderin expression is low or absent in the human adult stomach. The results of the present study suggested that: i) scinderin knockdown is able to reverse EMT process and effectively prevent migration of highly metastatic SGC-7901 cells, which also can be understood to weaken the local invasion in primary gastric cancer; ii) scinderin knockdown is able to inhibit the proliferation of SGC-7901 cells, greatly impairing the proliferation of cancer cells at metastatic sites and the formation of metastatic colonization. If human gastric cancer tissues provide similar results to those of SGC-7901 cells highly expressing scinderin, which is closely associated with carcinoma metastasis, scinderin may be a useful prognostic biomarker to distinguish the progression of gastric cancer and guide personalized therapy for patients.

In conclusion, the present study demonstrated that suppression of scinderin impaired the proliferation and migration of gastric cancer SGC-7901 cells and attenuated their EMT process. Therefore, scinderin may be a potential target for tumor EMT and therapy against gastric cancer.

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References


