Original Article

Upregulation of CD147 promotes cell invasion, epithelial-to-mesenchymal transition and activates MAPK/ERK signaling pathway in colorectal cancer

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Abstract: Colorectal cancer (CRC) is one of the most common cancers in the world. CD147, a transmembrane protein, has been reported to be correlated with various cancers. In this study, we aimed to investigate the mechanism of CD147 in regulating drug resistance, cell invasion and epithelial-to-mesenchymal transition (EMT) in CRC cells. qRT-PCR and western blotting were used to evaluate the expression of CD147 in 40 CRC cases and 4 cell lines. Increased expression of CD147 at both mRNA and protein levels was found in CRC samples, and the level of CD147 was correlated with lymph node metastasis. CD147 overexpression increased the 5-Fluorouracil (5-FU) resistance, enhanced the invasion and EMT of CRC cells by regulating EMT markers and MMPs. Adverse results were obtained in CD147 knockdown CRC cell line. Further investigation revealed that CD147 activated MAPK/ERK pathway, ERK inhibitor U0126 suppressed the CD147-induced cell invasion, migration and MMP-2, MMP-9 expression. Taken together, our study indicates that CD147 promotes the 5-FU resistance, and MAPK/ERK signaling pathway is involved in CD147-promoted invasion and EMT of CRC cells.

Keywords: CD147, 5-FU resistance, invasion, EMT, MAPK/ERK pathway, colorectal cancer

Introduction

Colorectal cancer (CRC) is the third most common cancer in men (746,000 cases, 10.0% of the total) and the second in women (614,000 cases, 9.2% of the total) worldwide [1]. Every year nearly 0.7 million people die of CRC, the high mortality is mainly due to the chemoresistance and tumor metastasis. It is reported that approximate 50% of the CRC patients would develop distant metastasis eventually [2]. Chemotherapy is an essential treatment to metastatic cancer patients, however, drug resistance leads to treatments failure and cancer progression [2]. So better understanding of the molecular mechanism underlying CRC chemoresistance and metastasis is critical to the treatments.

CD147 (also named EMMPRIN, basigin, M6 and tumor cell-derived collagenase stimulatory factor), is a glycosylated cell surface transmembrane protein of the immunoglobulin superfamily (IgSF) [3]. The CD147 protein has 269 amino acids, exists in two forms: glycosylated form (HG, ~40-60 kDa) and core-glycosylated form (LG, ~32 kDa), the ratio is variable in different cell types, while HG-CD147 is considered to be the functional form [4]. CD147 has a broad tissue distribution, implicating in many physiological processes [5]. Aberrant CD147 expression has been observed to correlate with several diseases, especially cancers [6]. During tumorigenesis, CD147 contributes to cell proliferation, metastasis, drug resistance and angiogenesis [7-9], moreover, its level may also predict tumor relapse and patient outcome [10, 11].

CD147 has been shown to interact with hyaluronan, multidrug transporters of the ABC family and monocarboxylate transporters to mediate drug resistance [12]. In addition, CD147 is capable of inducing the expression of several matrix metalloproteinases (MMPs), including MT1-MMP, MMP-1, MMP-2 and MMP-9 [13].
MMPs are major proteases in degrading the extracellular matrix (ECM) and basement membrane, which are vital to tumor initiation, progression and invasion [14]. It was reported that CD147 was upregulated in CRC [15], nevertheless, whether the aberrant expression of CD147 correlates with drug resistance and metastasis of CRC are still unclear.

In the present study, we aimed to evaluate the effects of CD147 on drug resistance and metastasis of CRC cells. We assessed the expression pattern of CD147 and analyzed its correlation with clinicopathological factors of CRC patients. Next the effects of CD147 on drug resistance, cell invasion and EMT were elucidated. Further we investigated the involvement of MAPK/ERK signaling pathway in CD147-induced cell invasion and migration.

**Materials and methods**

**Tissue specimens and cell culture**

All tissue samples were obtained from Shandong provincial hospital, consent forms were obtained from all patients. The study was approved by the Ethical Committee of Shandong Provincial Hospital. Cell lines were purchased from the Cell Bank of Chinese Academy of Sciences. Cells were cultured in RPMI 1640 and DMEM medium, respectively, with 10% fetal bovine serum, at 37°C in a humidified incubator with 5% CO₂.

**Quantitative real-time polymerase chain reaction (qRT-PCR)**

Total RNA was extracted with the TRizol reagent (Invitrogen, USA) according to the manufacturer’s protocol. Complementary DNA was reverse-transcribed using a reverse transcription kit (Takara, Japan). qRT-PCR was performed using SYBR Premix Ex Taq Kit (Takara, Japan) on ABI 7300. β-actin was used as endogenous control for the normalization of gene expression. The forward and reverse primers for CD147 were 5’-CAGCGGTTGGAGGTTGT-3’ and 5’-TTTGAGGTGGAGGTGG-3’, for β-actin were 5’-AAAGACTGTAGCCCAACAC-3’ and 5’-GTGATACTCTGCGTGTGAT-3’. The PCR cycle conditions consisted of an initial denaturation step at 95°C for 30 sec, followed by 40 cycles at 95°C for 30 sec, 60°C for 30 sec, and 72°C for 30 sec. Relative mRNA expression levels were determined by the 2⁻ΔΔCt method in comparison with control cells.

**Western blotting analysis**

Cells were harvested, washed by cold phosphate-buffered saline (PBS), and lysed in RIPA lysis buffer (Beyotime, China). Total proteins extracted were quantified with BCA Protein Assay Kit (Beyotime, China). Proteins were separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), then transferred to polyvinylidene difluoride (PVDF) membranes and blocked in 5% non-fat milk in TBST buffer for 1 h. Then the membranes were incubated overnight at 4°C with primary antibodies: CD147 (1:500), E-cadherin (1:1000), vimentin (1:500), MMP-2 (1:500), MMP-9 (1:500), ERK (1:500), p-ERK (1:500), β-actin (1:1000), followed by incubation of secondary antibodies (1:5000). Protein bands were detected using ECL detection system (Beyotime, China).

**Cell transfection and construction of CD147 downexpression and overexpression CRC cell lines**

Plasmids containing CD147 siRNA, CD147 negative control and CD147 cDNA, CD147 negative control were purchased from Genepharma (Genepharma, China). 2 × 10⁵ cells were seeded into 6-well plate. When the cells were 70-90% confluent 24 h later, 2 μg plasmids were transfected into cells using Lipofectamine 2000 (Invitrogen, USA). 48 h after transfection, cells were selected by 400 μg/ml G418 (Sigma, USA) for 1 month to form stable clones. Caco-2 cells transfected with CD147 expression vector and negative control were named Caco-OvCD147 and Caco-cCD147. SW480 cells transfected with CD147 siRNA vector and negative control were labeled as SW480-SiCD147 and SW480-cCD147.

**CCK-8 assay**

5-FU resistance was measured by Cell Counting Kit-8 (CCK-8). Cells were seeded in 96-well plates at a density of 1 × 10⁵ cells/100 μl/well. After 24 h incubation, cells were treated with different concentration of 5-FU (0, 2, 4, 6, 8, 10 μM). After 48 h, 10 μl CCK-8 (Beyotime, China) was added into the medium, then the supernatants were removed and measured at 450 nm
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Transwell assay

Cell invasion was measured by using transwell chambers (Costar, USA). $1 \times 10^5$ cells were suspended in serum-free medium and planted into the upper chamber, then 500 μl complete medium with serum was added to the lower well. Cells were incubated at 37°C for 24 h and the invading cells to the lower surface were stained with 0.1% crystal violet (Beyotime, China) for 30 min. The cells were photographed under a microscope.

Wound healing detection

$5 \times 10^5$ cells were planted into 6-well plate, incubated overnight until the grown to confluent monolayer, a 200 μl eppendorf tip was used to scratch cell wounds, the wounded monolayers were washed with PBS three times to remove non-adherent cells, and incubated in serum-free medium for 48 h. Cell migration were observed and photographed under a microscope.

Statistical analysis

Data were presented as mean ± SD from three independent experiments. Two-sided Student’s t test was used to analyze the different expression of CD147. All statistical analyses were conducted using the SPSS version 17.0 (SPSS Inc. Chicago). $P < 0.05$ was considered statistically significant.

Results

Increased CD147 expression in CRC tissues and cell lines

CD147 expression level was evaluated in 40 CRC tissues and 4 CRC cell lines by qRT-PCR and western blotting. Results showed that CD147 mRNA expression was upregulated in 55% (22/40) tissues, and protein levels in...
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Table 1. Relationship between CD147 expression and clinicopathologic features of CRC patients

<table>
<thead>
<tr>
<th>Feature</th>
<th>Relative CD147 expression</th>
<th>P value</th>
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<tbody>
<tr>
<td>Gender</td>
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<td></td>
</tr>
<tr>
<td>Male</td>
<td>9</td>
<td>14</td>
</tr>
<tr>
<td>Female</td>
<td>6</td>
<td>11</td>
</tr>
<tr>
<td>Age</td>
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<tr>
<td>&lt; 60</td>
<td>6</td>
<td>9</td>
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<tr>
<td>≥ 60</td>
<td>9</td>
<td>16</td>
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<tr>
<td>Differentiation</td>
<td></td>
<td></td>
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<tr>
<td>Poorly</td>
<td>5</td>
<td>10</td>
</tr>
<tr>
<td>Moderately/Well</td>
<td>10</td>
<td>15</td>
</tr>
<tr>
<td>local invasion</td>
<td></td>
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<tr>
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<td>15</td>
</tr>
<tr>
<td>No</td>
<td>11</td>
<td>10</td>
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<tr>
<td>Lymph node metastasis</td>
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<tr>
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<td>5</td>
<td>17</td>
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<tr>
<td>No</td>
<td>10</td>
<td>8</td>
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<td>Yes</td>
<td>6</td>
<td>14</td>
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<tr>
<td>No</td>
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*P < 0.05.

62.5% (25/40) tissues (Figure 1A and 1B). In CRC cell lines, we found CD147 was expressed in all the 4 cell lines, SW480 cells exhibited highest expression level of CD147, while Caco-2 cells showed lowest CD147 level (Figure 1C and 1D). These results suggested upregulation of CD147 may play an important role in CRC progression.

Increased expression of CD147 is associated with lymph node metastasis of CRC

To further explore the relationship between the expression pattern of CD147 and clinicopathological features of patients with CRC, 40 patients were categorized into two groups according to the CD147 expression level difference (≥ 1.5-fold). The correlation was summarized in Table 1. Statistical analyses revealed that overexpression of CD147 is correlated with lymph node metastasis (P = 0.033) of CRC. The results indicated CD147 contributed to CRC metastasis.

Expression of CD147 in the stable transfected CRC cells

To further study the function of CD147, Caco-2 and SW480 cell lines were chosen to enhance and deplete CD147, respectively. RT-qPCR and western blotting were used to evaluate the relative mRNA and protein expression levels. Results showed that CD147 increased its expression by 2.3 fold in Caco-OvCD147 cells than control (Figure 2A). While CD147 reduced its expression by 0.68 fold in SW480-SiCD147 cells compared to SW480-cCD147 cells (Figure 2B).

CD147 enhanced the 5-FU resistance of CRC cells

CCK-8 kit was employed to detect the effects of CD147 on 5-FU resistance. The data revealed that increased expression of CD147 enhanced the 5-FU resistance of Caco-2 cells (Figure 3A), while reduction of CD147 decreased the SW480 cells sensitivity to 5-FU (Figure 3B). The results indicated CD147 regulates the 5-FU resistance of CRC cells.

CD147 promoted the invasion and migration of CRC cells

Since CD147 expression was positively correlated with lymph node metastasis of CRC patients, we further verified the biological function of CD147 on CRC cells. The transwell assay indicated CD147 overexpression increased cell invasion of Caco-2 cells. Adverse results were found in CD147 knockdown cells (Figure 4A and 4B). In wound healing assay, the migratory activity of Caco-OvCD147 cells was induced compared to the Caco-cCD147 cells. By contrast, silencing of CD147 inhibited SW480-SiCD147 cells migration as compared with SW480-cCD147 cells (Figure 4C and 4D). Taken together, our data revealed that CD147 could promote the invasion and migration of CRC cells.

Expression of EMT markers, MMPs and pERK mediated by CD147

To identify the mechanism of CD147 in the metastasis of CRC cells, we further examined the expression of EMT markers and MMPs. Results demonstrated that overexpression of CD147 in Caco-2 cells led to decreased expression of E-cadherin and increased level of vimentin, and the level of MMP-2, MMP-9 were increased (Figure 5A). Knockdown of CD147 in SW480 cells induced the expression of E-cadherin and inhibited the level of vimentin,
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Figure 2. The expression of CD147 in CRC cell lines after stable transfection. A. The expression of CD147 was upregulated in mRNA and protein level in Caco-2 cells. B. The level of CD147 was downregulated in mRNA and protein level in SW480 cells.

while the expression of MMP-2 and MMP-9 were suppressed (Figure 5B). Previous study had reported that the MAPK/ERK pathway is involved in modulating the cancer metastasis and drug resistance [16], so we studied the effects of CD147 on ERK expression in CRC cell lines. The data indicated that upregulation of CD147 induced the phosphorylation of ERK (pERK), whereas downregulation of CD147 reduced the expression of pERK, but there were no obvious changes of the total-ERK (tERK) in the 2 cell lines (Figure 5C and 5D). These data suggested CD147 promoted cell EMT, stimulated the expression of MMP-2, MMP-9 and activated the MAPK/ERK pathway.

MAPK/ERK pathway is involved in regulating the CD147-induced cell invasion, EMT and MMPs expression

To validate whether the CD147-mediated cell invasion, migration and MMPs expression in CRC cells were regulated by MAPK/ERK signaling pathway, CD147 overexpressed Caco-2 cell lines were treated with ERK inhibitor (U0126; 10 µM). We found inhibition of ERK suppressed the invasive and migratory ability of Caco-OvCD147 cells, as compared with the Caco-cCD147 (Figure 6A and 6B). Western blot showed U-0126 treatment attenuated the increased expression of MMP-2, MMP-9 in
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Figure 3. Effects of CD147 on 5-FU resistance in CRC cells. A. CD147 overexpression enhanced the 5-FU resistance of Caco-2 cells. B. CD147 knockdown suppressed the 5-FU resistance of SW480 cells.

Figure 4. Effects of CD147 on invasion and EMT of CRC cells. A. CD147 overexpression increased the invasive ability of Caco-OvCD147, as compared with Caco-cCD147 cells (200×). B. CD147 knockdown inhibited the invasive ability of SW480-SiCD147, as compared with SW480-cCD147 cells (200×). C. Upregulation of CD147 promoted the migration of Caco-2 cells (40×). D. Downregulation of CD147 inhibited the migration of SW480 cells (40×).

Caco-2 cells (Figure 6C). The data indicated that U0126 inhibits CD147-induced invasion, migration, and MMP-2, MMP-9 expression of CRC cells. Taken together, MAPK/ERK signaling pathway may involve in CD147-stimulated cell invasion, EMT and MMPs expression of CRC cells.

Discussion

Drug resistance and tumor metastasis are two important causes of treatment failure and mortality in cancer patients [16]. Tumor metastasis is a complex process includes multiple steps and molecules. Cell invasion and EMT are two important processes, in which cells migrate and invade to nearby tissues, eventually form new metastatic sites [8]. Many molecules, including CD147, MMPs, hyaluronan, are involved in these processes.

CD147 was reported to implicate in numerous physiological and pathological processes [5], it correlated with the metastasis process of vari-
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Drug resistance in tumor leads to chemotherapy failure, CD147 was reported to implicate in chemoresistance [20]. In this study, we selected 5-FU as an experimental drug. 5-FU is a classic chemotherapy drugs which could restrain the DNA synthesis processing of tumor cell [21]. Data revealed that CD147 decreased the sensitivity of CRC cells to 5-FU. It was reported that CD147 induced drug resistance by stimulating the hyaluronan production [22], further studies are needed to explore the potential mechanism in CD147-mediated 5-FU resistance in CRC cells. Our finding may assist in choosing applicable chemotherapeutics.

Given that CD147 related to the lymph node metastasis, we next examined the impacts of CD147 on CRC cells. Results revealed that CD147 promoted the invasion and migration of CRC cells. Loss of E-cadherin is a common feature of EMT [23]. We found overexpression of CD147 downregulated epithelial marker E-cadherin and upregulated mesenchymal marker vimentin, suggesting CD147 participate in regulating the EMT of CRC cells. One of the most important role of CD147 is the stimulation of MMPs [24]. MMPs are positive regulators of tumor invasion and metastasis [25]. MMP-2 and MMP-9 belong to the gelatinase subclass which can degrade the gelatin, the degradation products then induce multiple cellular signals, thus promote cancer metastasis [26]. Increased expressions of MMP-2 and MMP-9 have been reported to contribute to invasion or metastasis of various cancers [26, 27]. In this study, we found CD147 induced the expression of MMP-2 and MMP-9 which suggests that CD147 may activate MMP-2 and

Figure 5. The expression of EMT markers, MMPs and ERK. A. CD147 overexpression decreased the level of E-cadherin and increased vimentin, enhanced the expression MMP-2 and MMP-9 in Caco-2 cells. B. CD147 knockdown upregulated E-cadherin and downregulated vimentin, suppressed the expression MMP-2 and MMP-9 in SW480 cells. C. CD147 overexpression induced the level of p-ERK in Caco-2 cells. D. CD147 knockdown reduced the level of p-ERK in SW480 cells.
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Figure 6. Effects of ERK inhibitor on CD147 overexpressed cells. U0126 inhibited the migratory (40×) (A) and invasive (200×) (B) ability of Caco-OvCD147 cells. (C) U0126 suppressed the expression of MMP-2 and MMP-9 in Caco-OvCD147 cells.
MMP-9 to enhance the invasion and EMT in CRC cells. MAPK/ERK signaling pathway is reported to contribute to cell migration, angiogenesis and chemoresistance [28].

In hepatocellular carcinoma, CD147 enhances the secretion of MMP-2 via the activation of ERK, FAK, and PI3K/Akt signaling pathway [24]. In macrophages, CD147 induces the expression of MMP-9 through ERK and NF-κB [29]. Our studies indicate that CD147 promoted the expression of pERK in CRC cells; further research demonstrated that ERK inhibitor abrogated CD147-induced cell invasion, migration and MMPs expression. So we presumed that MAPK/ERK pathway is involved in CD147-induced cell invasion and EMT, but whether MAPK/ERK pathway regulates the cell invasion and EMT by direct mediation MMP-2, MMP-9 still needed to investigate.

In conclusion, we demonstrated that CD147 is upregulated in CRC tissues and associated with the lymph node metastasis. CD147 promotes the 5-FU resistance, and enhance cell invasion and EMT through activating MAPK/ERK pathway.

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Disclosure of conflict of interest

None.

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