

# Krüppel-like factor 4 negatively regulates $\beta$ -catenin expression and inhibits the proliferation, invasion and metastasis of gastric cancer

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**Abstract.** Krüppel-like factor 4 (KLF4) is a zinc-finger protein that plays an important role in the progression of gastric carcinoma. The abnormal activation of  $\beta$ -catenin frequently occurs in gastric cancer and has been associated with the promotion of tumor growth, invasion and metastasis. However, the potential interaction between KLF4 and  $\beta$ -catenin during gastric cancer development is unknown. In this study, a lentiviral KLF4 expression vector was constructed and utilized to transfect the human gastric cancer cell lines, SGC-7901, BGC-823, MKN-28 and MKN-45. KLF4 and  $\beta$ -catenin expression levels were measured by quantitative real-time RT-PCR and western blot analysis. Cell proliferation, colony formation and invasive potential were determined in the KLF4-transfected gastric cancer cells. The expression of E-cadherin and matrix metalloproteinase 2 (MMP2) was determined by western blot analysis. The overexpression of KLF4 significantly inhibited the expression of  $\beta$ -catenin in the MKN-45 gastric cancer cells. The restored expression of KLF4 suppressed proliferation, colony formation and inhibited the invasion and metastatic properties of MKN-45 gastric cancer cells. Furthermore, the forced expression of KLF4 in gastric tumor cell lines restored E-cadherin expression and inhibited MMP2 expression. Consistent with the *in vitro* findings, the enforced expression of the KLF4 gene in MKN-45 gastric carcinoma cells by lentiviral vector-mediated gene transfer effectively suppressed tumor growth *in vivo*. Our results show that KLF4 inhibits  $\beta$ -catenin expression and regulates the  $\beta$ -catenin-mediated biological behaviors of gastric cancer cells. The modulation of KLF4 expression may represent a novel therapeutic approach for  $\beta$ -catenin-driven malignancies.

## Introduction

Although the incidence and mortality of gastric cancer is declining, it remains the fourth most common epithelial malignancy and the second leading cause of cancer-related mortality following lung carcinoma throughout the world (1,2). Gastric cancer is characterized by high regional lymph node invasion and distant metastasis, which are important factors that lead to recurrence and negatively affect the prognosis of gastric carcinomas (3-5). Therefore, there is a critical need to understand the molecular mechanisms that regulate regional lymph node invasion and distant metastasis of gastric cancer in order to develop novel treatment options for this deadly disease.

Invasion and metastasis are often associated with a number of molecular abnormalities, including the inactivation of the tumor suppressor, Krüppel-like factor 4 (KLF4), and activation of the oncogene,  $\beta$ -catenin (6,7). KLF4 has been identified as a zinc-finger transcription factor that is highly expressed in the epithelial cells of several organs (8-10), especially in the epithelium of the gastrointestinal tract (11-13). Previous studies have confirmed that KLF4 is a negative regulator of the cell cycle, and its expression pattern correlates with terminal differentiation via the suppression of proliferative genes and the up-regulation of inhibitors of proliferation (14). KLF4 also plays a key role in the differentiation of various organs during embryonic development (15). KLF4-null mice die from dehydration immediately after birth due to severe defects in late-stage differentiation of epidermal cells (8). In addition, KLF4 functions to maintain self-renewal through the regulation of pluripotency gene expression (15). Several studies have discovered KLF4 expression is decreased in benign and malignant tumors in the digestive tract, including early intestinal adenomas, colonic adenomas and adenocarcinomas, as well as esophageal and gastric cancer (16-19). These findings suggest that KLF4 plays an important role as a suppressor of tumor development and that the lack of KLF4 expression may play a role in tumor progression.

As a multifunctional protein,  $\beta$ -catenin is a transcriptional co-activator of T cell factor (TCF)/lymphoid enhancer factor (LEF) transcription factors and acts as a crucial factor in the cadherin-catenin complex that mediates intercellular adhesion (20).  $\beta$ -catenin also plays an important role in many processes, such as cell growth, differentiation, embryogenesis,

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tumorigenesis, and the growth and maintenance of stem cells (21-23). In normal cells, the cytoplasmic levels of  $\beta$ -catenin are tightly regulated by interactions with adenomatous polyposis coli (APC), glycogen synthase kinase 3 $\beta$  (GSK-3 $\beta$ ) and axin proteins. The APC-axin-GSK-3 $\beta$  complex normally promotes the degradation of  $\beta$ -catenin via the phosphorylation of serine and threonine residues in the N-terminal region of  $\beta$ -catenin (21). When activated by Wnt-signaling, the degradation of  $\beta$ -catenin is inhibited and the cytoplasmic levels of  $\beta$ -catenin stabilize, resulting in the formation of the  $\beta$ -catenin-LEF/TCF complex in the nucleus. In the nucleus,  $\beta$ -catenin acts as a transcription factor and activates target genes, such as cyclin-D1 and c-Myc (24).  $\beta$ -catenin has been shown to function as an oncogene in a variety of tumor types (25,26). In several types of aggressive cancers including gastric, lung, and breast cancer, the aberrant expression and localization of  $\beta$ -catenin expression may trigger the weakening of cell-cell junctions and promote cell dedifferentiation, hyperproliferation, invasion and metastasis (24). Therefore, it is critical to understand the central role of the Wnt/ $\beta$ -catenin signaling pathway and interaction between its target genes in the invasion and metastasis of gastric cancer. Previous studies have shown that KLF4 interacts with  $\beta$ -catenin and represses  $\beta$ -catenin-mediated gene expression, which plays an important role in maintaining homeostasis of the normal intestine, and is implicated in the regulation of stem cells and cancer progression (27,28). However, a clear link between KLF4 and  $\beta$ -catenin in gastric cancer progression has not yet been demonstrated.

In the present study, we demonstrate a correlation between the expression levels of KLF4 and  $\beta$ -catenin. We also demonstrate that KLF4 expression plays a role in the inhibition proliferation, invasion and metastasis of gastric cancer in *in vivo* and *in vitro* models. Our results suggest that the anticancer properties of KLF4 are due to its ability to inhibit the  $\beta$ -catenin expression levels, which subsequently alters  $\beta$ -catenin signaling mechanisms.

## Materials and methods

**Cell culture and lentiviral infection.** Human gastric cancer cell lines (MKN-28, BCG-823, SGC-7901 and MKN-45) were purchased from the cell bank of Chinese Academy of Sciences and cultured in RPMI-1640 medium (Gibco Biomedical, Carlsbad, CA, USA) supplemented with 10% fetal bovine serum (FBS) at 37°C in the presence of 5% CO<sub>2</sub>. All gastric cancer cell line cultures were infected with the concentrated viral stocks of pLv-UbC-IRES2-KLF4-enhanced green fluorescent protein (EGFP) or pLv-UbC-IRES2-EGFP in RPMI-1640 for 8 h at 37°C. The cell lines were renamed 28-KLF4, 28-EGFP, 823-KLF4, 823-EGFP, 7901-KLF4, 7901-EGFP, 45-KLF4, 45-EGFP after stable transduction with KLF4 or the control EGFP using a concentration of 3x10<sup>5</sup> transducing units/well for 24-well plates at a multiplicity of infection (MOI) of 50. Over 95% cells infected with pLv-UbC-IRES2-KLF4-EGFP or pLv-UbC-IRES2-EGFP were green fluorescent protein (GFP)-positive after 96 h of infection.

**Quantitative real-time PCR.** Total RNA isolation and real-time PCR were performed as previously described (29). Briefly, total RNA from the gastric cancer cell line was isolated according

to the manufacturer's instructions for TRIzol<sup>®</sup> reagent (Takara, Dalian, China). RNA (1  $\mu$ g) was purified with DNase I (Takara) and then reverse-transcribed using SuperScript<sup>™</sup> II Reverse Transcriptase (Takara), according to the manufacturer's instructions. Duplicates of the 4-fold serial dilution of cDNA samples were used in a 25- $\mu$ l total volume containing 12.5 l SYBR<sup>®</sup>-Green PCR master mix (Applied Biosystems, Foster City, CA, USA) and 300 nM primers for each RT-PCR reaction. Amplification was performed in the 7300 Real-Time PCR System (Applied Biosystems). Cycle parameters were: 50°C (2 min) and 95°C (10 min), followed by 38 cycles of 95°C (15 sec) and 60°C (1 min), and a dissociation stage of 95°C (15 sec), 60°C (30 sec) and 95°C (15 sec). Fold inductions were calculated using the formula  $2^{-(\Delta\Delta C_t)}$ , where  $\Delta\Delta C_t$  was the  $\Delta C_t(\text{stimulus}) - \Delta C_t(\text{control})$ ,  $\Delta C_t$  was  $C_t(\text{KLF4, } \beta\text{-catenin}) - C_t(\text{GADPH})$  and  $C_t$  was the cycle at which the threshold was crossed. The gene-specific primer pairs are shown in Table I. PCR product quality was monitored by post-PCR melt curve analysis.

**Flow cytometry.** The effect of KLF4 overexpression on cell viability was assessed using flow cytometry after staining with propidium iodide and Annexin V-FITC (Roche) according to the manufacturer's instructions. Briefly, following KLF4 overexpression, cells were trypsinized and washed twice in ice-cold phosphate-buffered saline (PBS). A total of 1x10<sup>6</sup> cells were resuspended in 100  $\mu$ l binding buffer (10 mM HEPES/NaOH pH 7.4, 140 mM NaCl, 2.5 mM CaCl<sub>2</sub>) to which was added 5  $\mu$ l 2 mg/ml Annexin V-FITC and 10  $\mu$ l 50 mg/ml propidium iodide. Following a 15-min incubation in the dark, flow cytometry was performed using a CyAn ADP analyzer (Dako, Dublin, Ireland) using a 515/545-nm filter set for FITC detection and a 620/640-nm set for propidium iodide. The analysis was repeated in triplicates.

**Western blot analysis.** The following antibodies were used for western blot analysis: anti-KLF4 (rabbit polyclonal antibody, Abcam Biotechnology, Cambridge, UK), anti- $\beta$ -catenin (mouse monoclonal antibody; Millipore Biotechnology, Billerica, MA, USA); anti-matrix metalloproteinase 2 (MMP2) (rabbit polyclonal antibody; Santa Cruz Biotechnology, Santa Cruz, CA, USA), anti-E-cadherin (rabbit polyclonal antibody; Santa Cruz Biotechnology), and anti- $\beta$ -actin (mouse monoclonal; Santa Cruz Biotechnology). Cell lysates were minced and washed 3 times with ice-cold PBS. They were gently centrifuged and soaked in 500  $\mu$ l of hypotonic buffer (1 mM NaHCO<sub>3</sub>) containing 2 mM phenylmethanesulfonyl fluoride (PMSF) for 30 min and centrifuged at 15,000 x g for 15 min. The supernatant was mixed with half its volume of 33 loading buffer [30% glycerol, 6% sodium-dodecylsulfate (SDS), 62.5 mM Tris-HCl, and pH 6.8]. The protein concentrations were measured with a 5000002 Protein Assay Kit II (Bio-Rad, Hercules, CA, USA). Before centrifugation, the protein was mixed with 750  $\mu$ l of loading buffer (10% glycerol, 2% SDS, 62.5 mM Tris-HCl, pH 6.8) and boiled for 5 min in the presence of 2-mercaptoethanol. A total of 50  $\mu$ g of each sample was electrophoresed on 10% SDS-polyacrylamide gels and transferred onto PVDF membranes (Millipore Biotechnology). The membranes were first blocked with 5% skim milk, then incubated with primary antibodies, followed by secondary antibodies coupled with horseradish peroxidase (HRP) (Amersham, Arlington Heights,

Table I. Primer sequences of KLF4 and  $\beta$ -catenin and their qRT-PCR products.

Gene		Primer sequence	Product size
KLF4	Forward	5'-TCGGACCACCTCGCCTTACA-3'	133 bp
	Reverse	5'-CTGGGCTCCTTCCCTCATCG-3'	
$\beta$ -catenin	Forward	5'-CCAAGTGGGTGGTATAGAGG-3'	156 bp
	Reverse	5'-AGTCCATAGTGAAGGCCAAC-3'	
GADPH	Forward	5'-ATCAAGAAGGTGGTGAAGCAGG-3'	113 bp
	Reverse	5'-AAGGTGGAGGAGTGGGTGTCG-3'	

KLF4, Krüppel-like factor 4.

IL, USA), and finally visualized with an enhanced chemiluminescence (ECL) reagent.

**MTT assay.** For cell proliferation assay,  $2 \times 10^3$  cells/well were plated in triplicates in 24-well plates and cell proliferation was evaluated on days 1, 2, 3, 4, 5, 6 and 7. The proliferation of the KLF4 overexpressing cell line, 45-KLF4, was represented as a percentage of the cell count compared to the control cell line, MKN-45, set at 100% proliferation. The mean values from 3 different experiments were calculated to represent the proliferation difference.

**Colony formation assays.** To examine the effects of up-regulated KLF4 expression on the proliferation of gastric cancer cells, 45-KLF4 cells were used for colony formation assay as described previously (30,31) with MKN-45 cells as the control group. The cells were seeded onto 24-well plates (2,000 cells/well) and cultured for 2 weeks in RPMI-1640 medium containing 10% FBS. These cultures were washed twice with PBS, fixed with 4% paraformaldehyde for 15 min, dried at room temperature, and stained with Giemsa (Sigma-Aldrich, St. Louis, MO, USA). Clones larger than 2 mm were counted manually and the number of clones per well was averaged from 3 wells for each experiment, with 3 independent experiments performed.

**Transwell migration and invasion assays.** Cell migration and invasion assays were performed as described previously (32,33), using 8.0  $\mu$ m pore size polycarbonate membrane transwell inserts in a 24-well plate. MKN-45 (control), and 45-KLF4 cells ( $2 \times 10^5$ ) were plated on the upper well of a Matrigel invasion chamber in a 24-well chamber that contained RPMI-1640 medium with 5% FBS. The lower side of the separating filter was filled with RPMI-1640 medium with 20% FBS. The chamber was incubated in a tissue culture incubator, and after 24 h, cells on the upper surface (non-migrated cells) were removed by scrubbing with a cotton swab, and the cells remaining on the filter were fixed with 4% paraformaldehyde for 15 min, dried at room temperature, stained with Giemsa (Sigma-Aldrich, St. Louis, MO, USA) and manually counted. The cells that successfully migrated through the filter were photographed. At least 5 different x400 magnification fields were counted for each experiment, and the results were averaged over 3 independent experiments. Invasion assays were

performed as described above using BD Matrigel™ Invasion Chambers (BD Biosciences, Franklin Lakes, NJ, USA) as per the manufacturer's instructions.

**Tumorigenicity assay in nude mice.** The inhibition of tumor growth and tumorigenicity assay due to KLF4 forced expression was performed as previously described (30,34). BALB/c nude mice (4-6 weeks old) were obtained from the Department of Laboratory Animal Science, Chongqing University Health Science Center. All procedures were carried out according to the animal protocol approved by the Institutional Animal Care and Use Committee of the Chongqing Medical University. 45-KLF4 and MKN-45 (control) cells were trypsinized, washed in PBS, resuspended in saline solution, and  $5 \times 10^6$  cells per 0.2 ml were injected subcutaneously into each nude mouse (at least 6 mice in each experimental group). The tumor size was measured every third day. The tumor volume was calculated according to the formula  $V = (a \times b^2)/2$ , where a = the largest superficial diameter and b = the smallest superficial diameter. After 3 weeks, the mice were sacrificed and the tumors were collected and photographed.

**Statistical analysis.** All data are presented as the means  $\pm$  SEM. The significance of differences was determined using the Student's t-test, and analysis of variance (ANOVA) and appropriate post hoc tests were used when comparing multiple parameters.  $P < 0.05$  was considered to indicate statistically significant differences.

## Results

**KLF4 and  $\beta$ -catenin expression in parental gastric cancer cell lines.** In order to identify a suitable cell line for further analysis of KLF4 function and the association between KLF4 and  $\beta$ -catenin, KLF4 and  $\beta$ -catenin mRNA and protein expression levels were examined in parental gastric cancer cell lines. A high variability of KLF4 expression was observed for these cell lines (Fig. 1). Specifically, levels of KLF4 mRNA (Fig. 1B-1) and protein (Fig. 1C-1) expression were lowest in the MKN-45 cells. Simultaneously, we examined the level of  $\beta$ -catenin expression in the same cell line. By contrast, the expression levels of  $\beta$ -catenin mRNA (Fig. 1B-2) and protein (Fig. 1C-2) were highest in the MKN-45 cells. Therefore, the MKN-45 cell line was selected for further analysis.

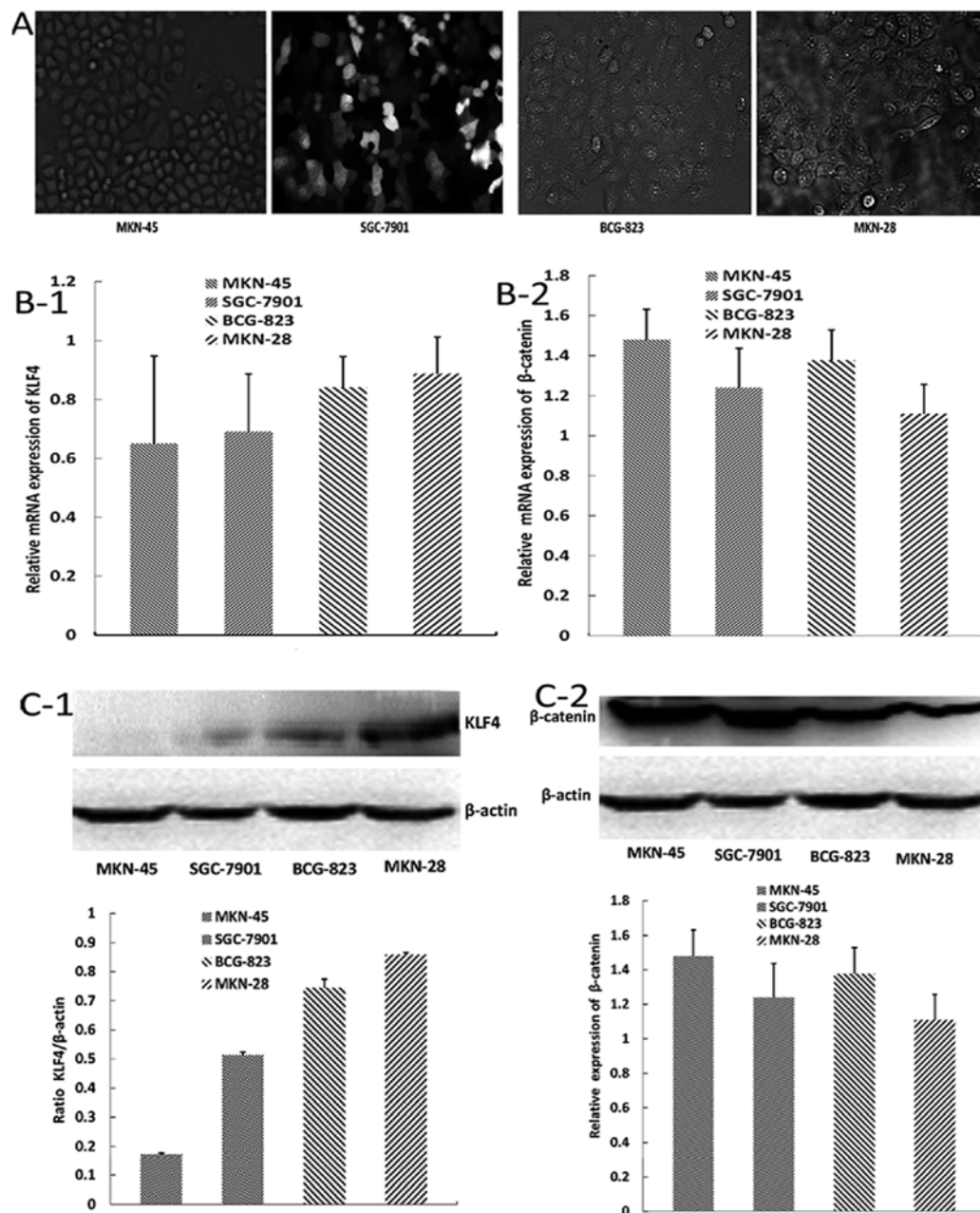


Figure 1. The mRNA and protein expression levels of KLF4 and  $\beta$ -catenin in parental gastric cancer cell lines. (A) Representative images of parental gastric cancer cells (x200). (B) Abnormal mRNA expression of KLF4 (B-1) and  $\beta$ -catenin (B-2). (C) Decreased expression of the KLF4 protein (C-1), and high levels of the  $\beta$ -catenin protein (C-2). The protein levels were consistent with the relative mRNA expressions. KLF4, Krüppel-like factor 4.

**Evaluation of KLF4 and  $\beta$ -catenin expression in MKN-45 and 45-EFGP cells.** The expression levels of KLF4 and  $\beta$ -catenin were evaluated in the parental gastric cancer cells that were treated with the empty lentiviral vector, MKN-45 (control untransduced) and 45-EFGP (transduced) cells. As shown in Fig. 2, there was no statistical difference in the mRNA and protein levels of KLF4 and  $\beta$ -catenin in the 45-EFGP cells compared to the parental MKN-45 cell line. Therefore, cells treated with the empty lentiviral vector were omitted in the subsequent experiments.

**Examination of the association between KLF4 and  $\beta$ -catenin expression in KLF4-overexpressing gastric cancer cell lines.** To clarify the association between KLF4 and  $\beta$ -catenin expres-

sion levels after successfully restoring the expression of KLF4, the mRNA and protein expression of KLF4 and  $\beta$ -catenin in the parental gastric cancer cell lines and the transduced cell lines, 28-KLF4, 823-KLF4, 7901-KLF4 and 45-KLF4, were examined. A high variability in KLF4 expression among these cell lines after lentiviral infection was observed (Fig. 3B-1 and C-1). Specifically, the levels of KLF4 mRNA and protein expression were highest in the 45-KLF4 cells. However, mRNA and protein levels of  $\beta$ -catenin decreased in the KLF4-expressing cells (Fig. 3B-1 and C-1).

**KLF4 overexpression inhibits proliferation, colony formation, invasion and migration.** To investigate the effects of KLF4 overexpression in gastric cancer cells, cell cycle, proliferation,

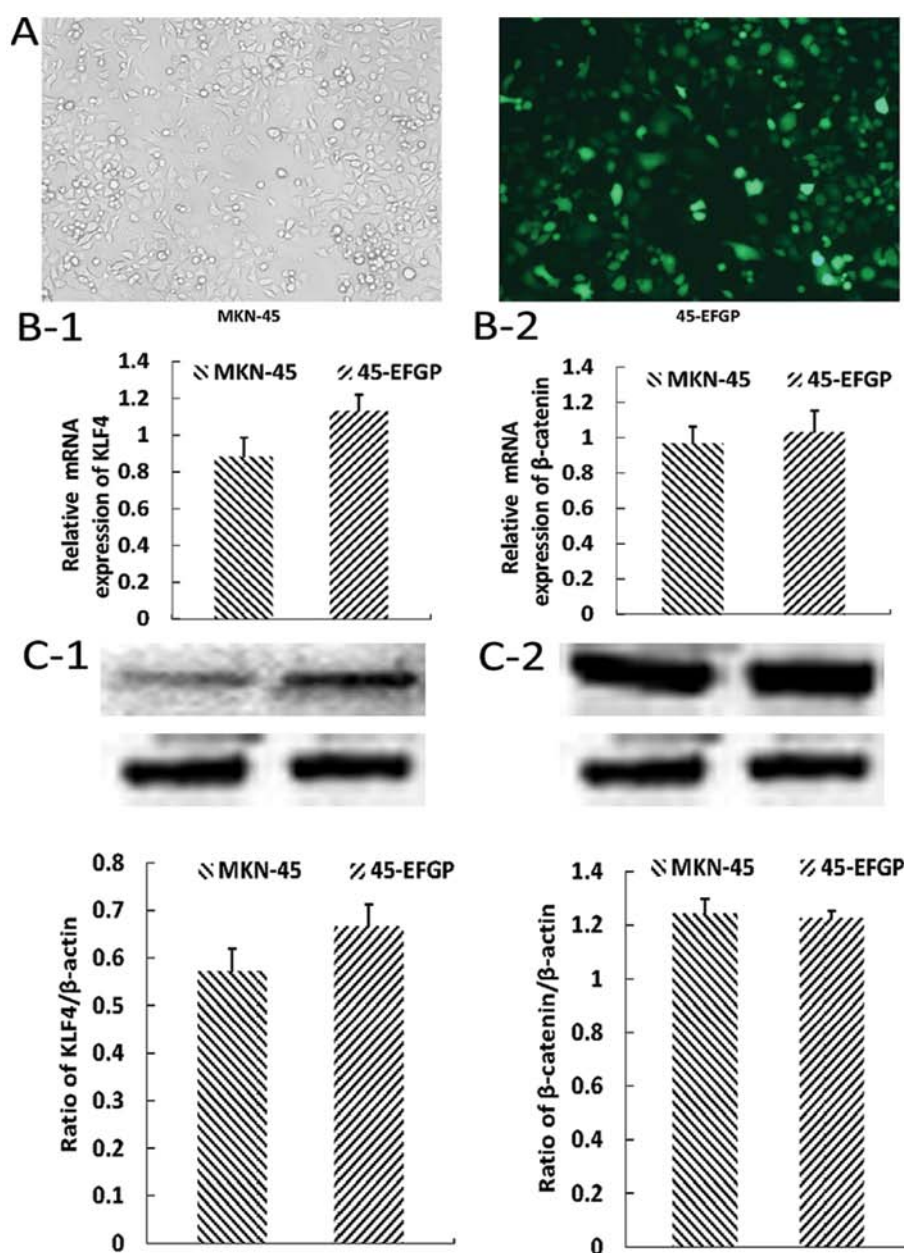


Figure 2. Expression of mRNA and protein of KLF4 and  $\beta$ -catenin in MKN-45 and 45-EFGP cells. (A) Representative images of MKN-45 cells (left) and 45-EFGP cells (right), (x200). (B) mRNA expression of KLF4 (B-1) and  $\beta$ -catenin (B-2). (C) Protein expression of KLF4 (C-1) and  $\beta$ -catenin (C-2). KLF4, Krüppel-like factor 4.

colony formation, invasion and migration *in vitro* were evaluated in the MKN-45 and 45-KLF4 cell lines. Compared with the MKN-45 cells, 45-KLF4 cells showed a 7-fold G1-block in the cell cycle (Fig. 4A). A significant difference in the percentage between MKN-45 and 45-KLF4 cells in the subG1 population was observed. 45-KLF4 cells displayed a  $61.8 \pm 0.0068\%$  inhibition of proliferation compared to MKN-45 cells, as determined by MTT assay (Fig. 4B). Consistent with reduced proliferation, the colony formation capability of 45-KLF4 cells was significantly weaker in comparison with the parental MKN-45 cells. As shown in Fig. 3C, 45-KLF4 cells showed an almost 12-fold inhibition of colony formation ( $P < 0.05$ ). The majority of clones from 45-KLF4 cells were significantly smaller than those of the parental MKN-45 cells. To deter-

mine whether the ability of invasion and metastases decreased in the 45-KLF4 cells, a transwell experiment was performed to assess the invasion and metastatic ability of the 45-KLF4 cell line compared to the MKN-45 cell line (Fig. 4D). To compare the migratory capacity of the MKN-45 cells, we found that 45-KLF4 cell lines exhibited significantly decreased migration and invasion, the 45-KLF4 cells being 4-fold less invasive ( $P < 0.05$ ) (Fig. 4D). Similarly, the overexpression of KLF4 in 45-KLF4 cells inhibited cell invasion compared to the MKN-45 control cells ( $P < 0.05$ ). Cell migration showed a similar profile (data not shown). Taken together, these results suggest that enhanced KLF4 expression plays a role in the suppression of the cell cycle, proliferation, colony formation, invasion and migration in gastric cancer cells *in vitro*.



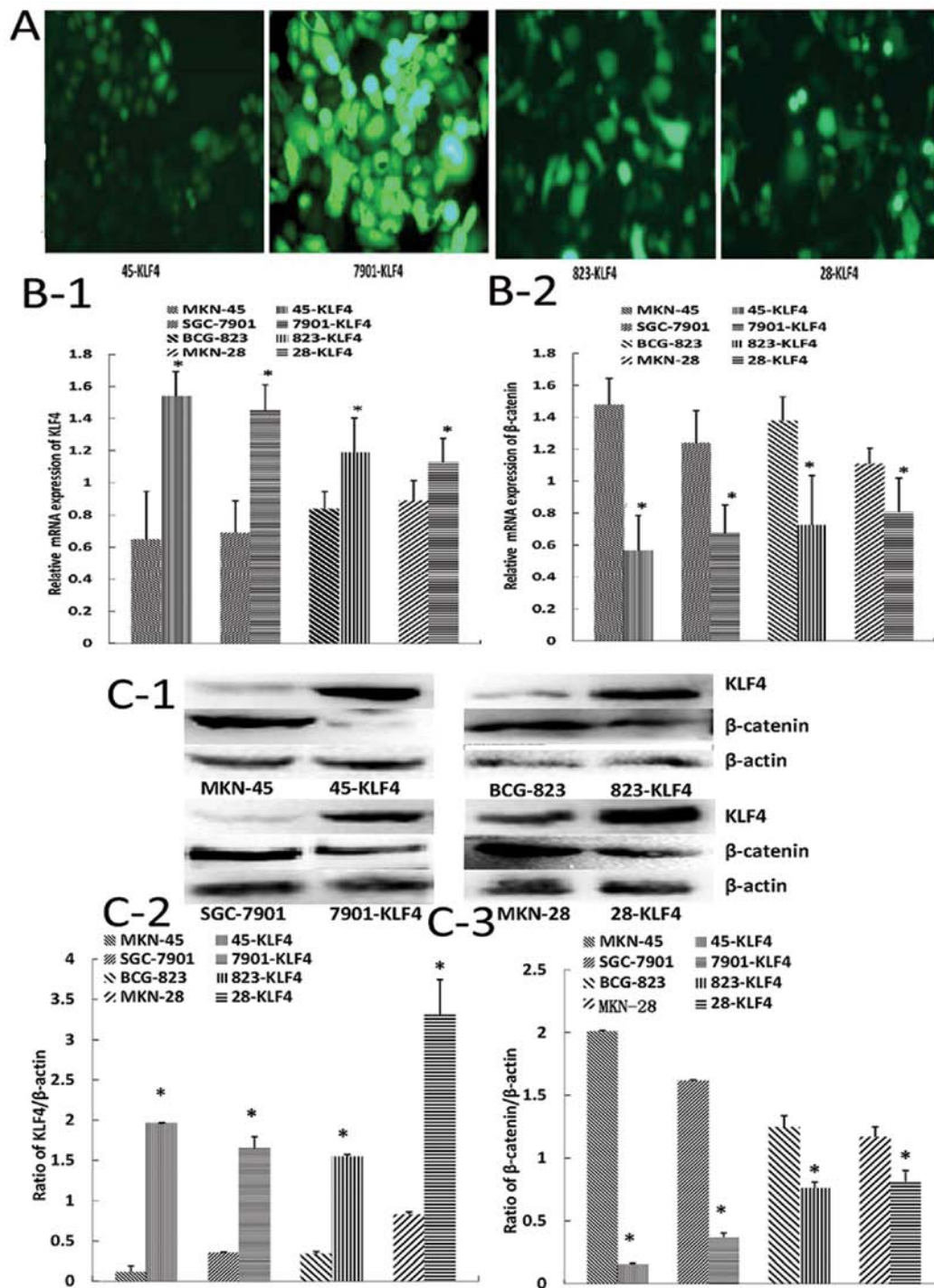


Figure 3. (A) Representative images of KLF4 overexpression in gastric cancer cells, (x200). (B) The expression of KLF4 mRNA (B-1) significantly increased, but levels of  $\beta$ -catenin mRNA (B-2) decreased. (C) KLF4 protein (C-1 and C-2) levels notably increased in stably transfected cells, but the protein level (C-1, C-3) of  $\beta$ -catenin decreased; these results were consistent with the mRNA levels. \* $P < 0.05$ , compared with parental cells. KLF4, Krüppel-like factor 4.

*Enhanced expression of KLF4 restores E-cadherin expression and inhibits MMP2 expression.* A previous study has indicated that KLF4 prevents the metastasis of breast cancer through the up-regulation of E-cadherin expression (33). In a similar fashion, the enhanced expression of KLF4 in the 45-KLF4 cells resulted in the up-regulation of E-cadherin expression and decreased MMP2 expression (Fig. 5). In the parental MKN-45 cells which have low KLF4 and higher  $\beta$ -catenin expression levels, we observed the opposite pattern with decreased

E-cadherin and elevated MMP2 expression levels (Fig. 5). These results support the concept that KLF4 positively regulates E-cadherin expression and inhibits MMP2 expression in gastric cancer cells.

*KLF4-mediated inhibitory effects on gastric cancer in vivo.* The inhibitory effect of KLF4 overexpression on tumor growth was analyzed in 45-KLF4 cells *in vivo*. There were at least 6 nude mice in the MKN-45 and 45-KLF4 treatment groups.

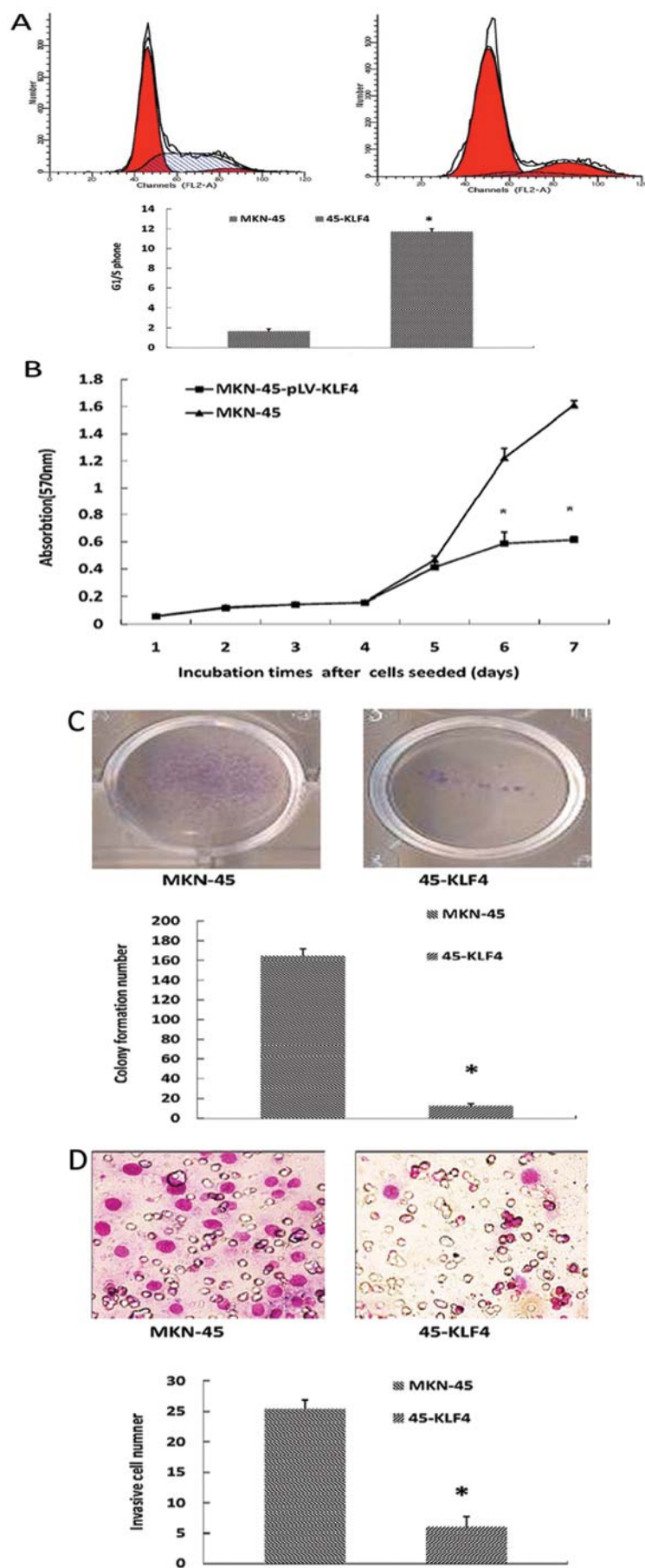


Figure 4. Inhibitory effects of KLF4 overexpression inhibited growth of gastric cancer cells *in vitro*. (A) Forced expression of KLF4 induced G1/S arrest. \* $P < 0.05$  compared with MKN-45 cells. (B) Growth curves of cells plotted by the MTT assay and the inhibition rate of cell proliferation by KLF4; \* $P < 0.05$  compared with MKN-45 cells. (C) Analysis of colony formation numbers, \* $P < 0.05$  versus the control group. (D) Representatives photomicrographs (x400) of cell invasion. The mean number of cells was summarized. \* $P < 0.05$  versus the parental cells. KLF4, Krüppel-like factor 4.

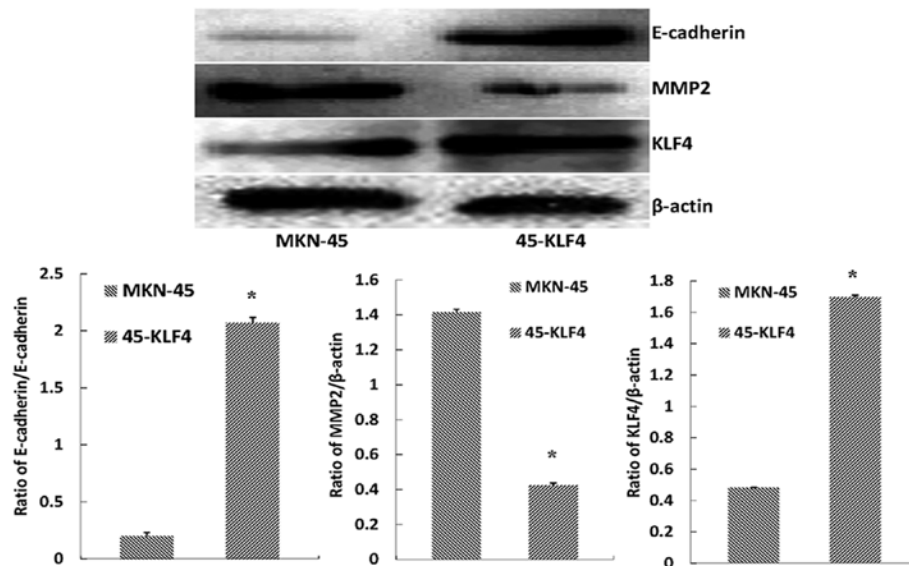


Figure 5. KLF4 is required to sustain E-cadherin expression and inhibit MMP2 expression in gastric cancer cells. (A) Western blot analysis of KLF4, E-cadherin and MMP2. (B) Values are the means  $\pm$  SD, \*different from MKN-45 cells; (B-1) increased E-cadherin expression in 45-KLF4 cells, \* $P < 0.05$  versus the control group; (B-2) restored KLF4 expression sharply decreased MMP2 expression, \* $P < 0.05$  versus the control group; (B-3) KLF4 expression was significantly increased in 45-KLF4 cells compared with MKN-45 cells ( $P < 0.05$ ). KLF4, Krüppel-like factor 4.

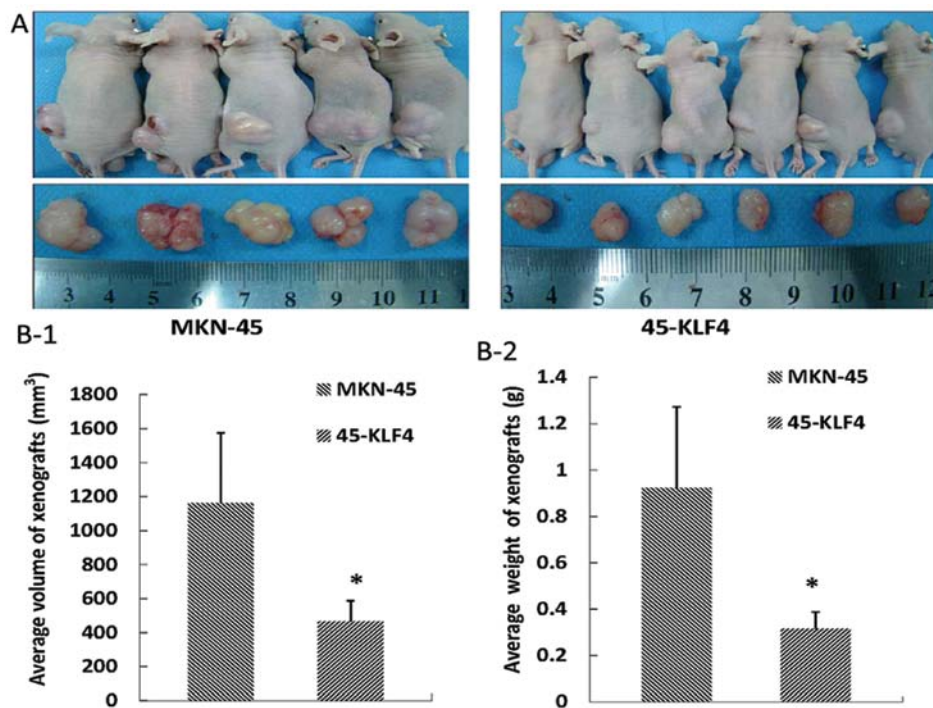


Figure 6. Suppression of tumor growth by KLF4 *in vivo*. (A) Images of tumor growth in MKN-45 and 45-KLF4 cells. (B) Compared with the control group, the average volume (B-1) and weight (B-2) of the xenografts were significantly smaller (\* $P < 0.05$ ), consistent with the findings *in vitro*, that restoring KLF4 gene expression suppresses tumor growth *in vivo*. KLF4, Krüppel-like factor 4.

One mouse died prematurely in the control group. MKN-45 and 45-KLF4 cells ( $5 \times 10^6$ ) were inoculated subcutaneously into the rear flanks of nude mice. Observation of tumor growth was performed at 3-day intervals. As described in Fig. 6, the tumor growth of 45-KLF4 cells was dramatically retarded compared with the parental MKN-45 cells. At the termination of the experiment, the mean size of KLF4-expressing tumors was

significantly smaller than that of the parental MKN-45 cells ( $P < 0.05$ ) (Fig. 6).

## Discussion

KLF4 (also known as Gut-enriched Krüppel-like factor) is an important member of the zinc finger-containing transcription



factor family (35). The Krüppel-like factor family, including KLF4, is characterized by 3 zinc-finger domains in its C-terminus (36). Depending on the molecular environment, KLF4 acts as either a tumor suppressor gene or an oncogene (37). High expression levels of KLF4 are present in normal gastrointestinal epithelial cells, which are associated with growth arrest, cell proliferation and late-stage cell differentiation (14,38). However, KLF4 expression levels have been shown to be significantly reduced in a number of cancer types (16,19,39-42) and tumor size has been shown to negatively correlate with KLF4 expression levels (43,44). Alterations in KLF4 expression levels lead to aberrant proliferation and differentiation in the gastric epithelium (45). Accumulating evidence has suggested that KLF4 is a putative tumor suppressor in the digestive tract. KLF4 knockout mice display defects in gastric cell differentiation and precancerous changes in the stomach (19,45). In addition, the forced expression of KLF4 in colon cancer cell lines induces colon cancer cell growth arrest (14), and represses  $\beta$ -catenin transcription (46). These results suggest that KLF4 is a tumor suppressor participating in the progression and development of gastric tumors. In our current study, we discovered that KLF4 mRNA and protein expression was low in parental gastric cancer cell lines by qRT-PCR and western blot analysis (Fig. 1). However, KLF4 mRNA and protein levels increased after stable overexpression of KLF4, when compared with parental cells (Fig. 2;  $P < 0.05$ ).

Over the years,  $\beta$ -catenin has been recognized not only for mediating intercellular adhesion (20), but also for its role in the regulation of cell growth, differentiation, invasion and metastasis (24).  $\beta$ -catenin is a key downstream molecule in the Wnt signaling pathway and binds to TCF/LEF. The  $\beta$ -catenin-TCF/LEF complex regulates and activates its downstream target transcription genes which are involved in development, tissue self-renewal and cancer (47), and plays a pivotal role in intracellular signaling and represents a key element in one of the most important pathways of epithelial carcinogenesis (22,48).

$\beta$ -catenin has been recognized as an oncogene in a variety of tumors (25,49). In many types of aggressive and lethal cancers, aberrant  $\beta$ -catenin expression may weaken the cell-cell junctions and promote carcinoma cell dedifferentiation, hyperproliferation, invasion and metastasis. This feature is commonly found in gastric cancer tumors (50). Previous studies have confirmed that high  $\beta$ -catenin expression is an independent indicator of poor prognosis for these carcinomas and closely correlates with enhanced tumor progression (49,51). In this study, we found that the expression of  $\beta$ -catenin was increased notably in both mRNA and protein gastric cancer cell lines with different gastric cancer cell differentiation stages (Fig. 1). However, the mRNA (Fig. 3B-2) and protein (Fig. 3C-1 and C-3) expression levels of  $\beta$ -catenin decreased significantly in the cell lines with enhanced KLF4 expression, when compared with the parental cells (Fig. 3).

In this study, we investigated whether the zinc-finger transcription factor, KLF4, plays a role in the inhibition of invasion and metastasis of gastric cancer by identifying its upstream regulators and downstream target genes, and characterizing its effects on gastric cancer cells. It is possible that the functional activation or inhibition of KLF4 may be determined not only by the activation status of its upstream genes, but also through a complex interplay of both positive and negative downstream

target genes. Our results strongly suggest that KLF4 is an important negative regulator of  $\beta$ -catenin expression in gastric cancer cells, for the following reasons: i) KLF4 and  $\beta$ -catenin expression were mutually exclusive in parental gastric cancer cell lines (for example, we observed a markedly reduced expression of KLF4 but the expression levels of  $\beta$ -catenin were abnormally increased); ii) the enhanced expression of KLF4 in the gastric cancer cell line, MKN-45, resulted in a significant increase in KLF4 at both the mRNA and protein levels, while the mRNA and protein expression of  $\beta$ -catenin were sharply decreased in the same cell lines; iii) in the KLF4-overexpressing cells (with low activity of  $\beta$ -catenin), many biological characteristics of gastric cancer were inhibited. These findings show that the reduced expression of KLF4 contributes to the inhibition of  $\beta$ -catenin expression in human gastric cancer cell lines. To our knowledge, this is the first report demonstrating that KLF4 negatively inhibits the activity of  $\beta$ -catenin in gastric cancer cell lines and regulates the malignant transformation of gastric cancer.

It is intriguing that the zinc-finger transcription factor, KLF4, regulates the invasion and metastasis-related genes, such as E-cadherin and MMP2. E-cadherin is a major component of the Wnt/ $\beta$ -catenin signal pathway and is an important effector of the E-cadherin/ $\beta$ -catenin complex that mediates intercellular adhesion (20). Thus, it is considered a hallmark of epithelial cell phenotype. Recent evidence supports the hypothesis that E-cadherin is a bona fide KLF4-regulated gene (33). In gastric cancer patients, decreased E-cadherin expression in tumors is associated with advanced stage, lymph node-positive metastasis and aggressiveness (52). We found that E-cadherin protein expression levels were down-regulated, but the level of MMP2 protein was high in gastric cancer parental cell lines. When KLF4 was overexpressed, the expression pattern of E-cadherin and MMP2 was reversed. A potential mechanism for the anti-invasive and metastatic effects was that KLF4 enhanced expression caused the up-regulation of E-cadherin expression while down-regulating MMP2 expression, two important invasion and metastasis-related genes (Fig. 5). Our results suggest that the metastasis-associated genes, E-cadherin and MMP2, are highly dependent upon KLF4 overexpression, and that KLF4 can affect metastasis in gastric cancer by regulating metastasis-mediated genes.

The results of MTT and flow cytometry assay, colony formation and transwell assay indicate that the up-regulation of KLF4 inhibits gastric cancer cell proliferation, colony formation and invasiveness (Fig. 4). Importantly, the anti-invasive effects of KLF4 observed *in vitro* (Fig. 6) were highly consistent with its *in vivo* expression pattern in primary human tumors. Moreover, KLF4 expression could also be reactivated in the Wnt/ $\beta$ -catenin-hyperactive gastric cancer cell lines MKN-45 by modulation of  $\beta$ -catenin expression levels, indicating that KLF4 expression is determined by complex interactions between signaling pathways.

Taken collectively, the results confirm that Wnt/ $\beta$ -catenin signaling is active and plays a key role in the progression of cell proliferation, dedifferentiation, invasion and metastasis. In summary, the enhanced expression of KLF4 inhibits invasion and metastasis of gastric cancer by negatively regulating the expression of  $\beta$ -catenin and subsequently  $\beta$ -catenin signaling mechanisms.

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