Knockdown of Serum- and Glucocorticoid-Regulated Kinase 1 Enhances Cisplatin Sensitivity of Gastric Cancer Through Suppressing the Nuclear Factor kappa-B Signaling Pathway

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Background: Previous studies have published the promoting effect of serum and glucocorticoid-regulated kinase 1 (SGK1) in various malignant tumors. However, whether SGK1 promotes gastric cancer remains a mystery.

Aims: To clarify the function of SGK1 in gastric cancer and its potential regulatory mechanism.

Study Design: Cell culture study.

Methods: The SGK1-silenced model was generated in two gastric cancer cell lines and further evaluated their malignant behavior and susceptibility to cisplatin. The interaction between miR-15a-5p and SGK1 was evaluated by the luciferase reporter assay. The knockdown efficiency of SGK1 was confirmed by RT-qPCR and Western blot assays. Cell proliferation rate was assessed with CCK-8 assay, and flow cytometry was used to determine cell cycle progression and apoptosis.

Results: Western blot data displayed an elevated level of SGK1 in gastric cancer cell lines. Functionally, SGK1 deficiency suppressed gastric cancer cell proliferation (P < .01) by acting on cell-cycle progression. Moreover, SGK1 deficiency suppressed cell invasion and migration of gastric cancer cells (P < .01). Further, the silencing of SGK1 obviously suppressed cell proliferation and induced apoptosis of the cells after cisplatin treatment (P < .01), indicating that SGK1 deficiency facilitated the chemosensitivity of these 2 gastric cancer cell lines to cisplatin. Mechanically, downregulation of SGK1 repressed the cytoplasm-to-nucleus translocation of NF-kB p65. Interestingly, we found that miR-15a-5p binds to the 3'UTR of SGK1, which was confirmed using luciferase activity assay (P < .05). Moreover, the data suggested that SGK1 reversed the suppression effect of miR-15a-5p on gastric cancer cell migration (P < .01).

Conclusion: Loss of SGK1 suppresses the malignant behavior of gastric cancer cells and increases cisplatin sensitivity by restraining the NF-kB signaling pathway. Moreover, SGK1 may exert an inhibitory effect in gastric cancer by being targeted by miR-15a-5p. Therefore, SGK1 may be a prospective target for future gastric cancer therapy.

INTRODUCTION

Gastric cancer has a high incidence and poor prognosis, and ranks third in contributing to cancer death in Asia.1 In recent years, the leading cause of failure for gastric cancer treatment is the presence of drug resistance in gastric cancer, imposing a huge economic burden on patients and the healthcare system.2-4 Therefore, it is urgent to search for potential molecular targets to improve the chemotherapeutic sensitivity of gastric cancer cells, gradually improving the therapeutic effect of gastric cancer treatment.

Previous researches have indicated that serum- and glucocorticoid-regulated kinase 1 (SGK1) plays a carcinogenic role in the development of a large number of cancers, such as colorectal cancer,5 prostate cancer,6 lung adenocarcinoma,5 etc. Additionally, recent research has found that SGK1 is implicated in the drug resistance of several cancer cells. For example, knockdown of SGK1 can reduce doxorubicin resistance in colorectal cancer cells.7 The SGK1-inhibitor SI113 inhibits cancer cell proliferation and offsets the blooming of paclitaxel resistance in ovarian cancer.
cells. Importantly, an existing study has shown that SGK1 is significantly elevated in gastric cancer tissues, which is involved in the differentiation of Th2 and Th17 in the tumor microenvironment of gastric cancer. Nevertheless, the function and mechanism of SGK1 on gastric cancer are still unclear.

NF-κB, which is highly expressed in various cancers, sparks tumor growth via stimulating the production of angiogenesis factors. Moreover, there are several types of research indicating that SGK1 downregulation can inhibit the activity of the NF-κB pathway. Meanwhile, the repression of NF-κB by curcumin can enhance the antitumor effects of chemotherapeutics in gastric cancer. Hence, NF-κB blocking is a helpful strategy for tumor suppression via promoting apoptosis and repressing cell growth.

In this study, an increased level of SGK1 was first detected in gastric cancer cells. Based on our results and on previous studies, we postulated a hypothesis that SGK1 silencing might suppress the progress of gastric cancer, likely via blocking the NF-κB pathway. To confirm this hypothesis, we conducted an SGK1 silencing model in 2 cell lines to investigate the effect of SGK1 in the progress of gastric carcinoma.

MATERIAL AND METHODS

Cell Culture and Transfection

The gastric carcinoma cell lines SNU-1 (Procell, Wuhan, China), NCI-N87 (Procell), MKN45 (FuHeng, Shanghai, China), and the normal gastric epithelial cells GES-1 (FengHui BioLogy, Hunan, China) were cultured in RPMI-1640 medium (31800, Solarbio, China) containing 10% fetal bovine serum (FBS, SH30084.03, HyClone, USA). AGS cell lines (ZQXZ, Shanghai) were grown in an F12K medium (ZQ-599, ZQXZ) containing 10% FBS. HGC-27 cell lines (Procell) were grown in an RPMI-1640 medium containing 20% FBS. The cells were cultured in a controlled environment (5% CO₂, 37 °C).

Vector Construction and Cell Transfection

The sequences of siRNA are listed in Supplementary Table 1. The HGC-27 and NCI-N87 cell lines were transfected with SGK1 siRNAs, or si-NC. SGK1-overexpressed vectors were constructed by inserting the CDS sequence into pcDNA3.1 vectors. To examine the function of miR-15a-5p in regulating SGK1, HGC-27 and NCI-N87 cell lines were also co-transfected with the miR-15a-5p mimic (JTS scientific, Wuhan) or/and SGK1-overexpressed vectors. Blank vectors were used as the negative control. Lipofectamine 3000 reagents (L3000015, Invitrogen, USA) were employed for cell transfection.

qRT-PCR

TRIPure (RP1001, BioTeke Corporation, China) was used for total RNA or miRNA extraction. Then, total RNA and miRNA were synthesized into cDNA by the SuperScript M-MLV reverse transcriptase (PR6502, BioTeke Corporation) and the miRNA cDNA Synthesis kit (#B532451, Sangon Biotech, Shanghai), respectively. Finally, the qRT-PCR was conducted according to the protocols. GAPDH served as a control. The relative level of the specific mRNAs was analyzed using the 2−ΔΔCT method. The sequences of the primers are listed in Supplementary Table 2.

Western Blot

The proteins were abstracted with RIPA (R0010, Solarbio), separated by SDS-polyacrylamide gel system, and electrically transferred to PVDF membranes (IPVH00010, Millipore, USA). After blocking with 5% fat-free milk (A600669, Sangon), the proteins were incubated at 4 °C for 18 hours with the following antibodies: SGK1 (1 : 400, A1025, AbClonal, China), cyclin D (1 : 1000, A19038, AbClonal, China), cyclin E (1 : 500, A14225, AbClonal), N-cadherin (1 : 1000, A19083, AbClonal), Vimentin (1 : 1000, AF7013, Affinity, China), caspase-3 (1 : 1000, CST, No. 14220), caspase-9 (1 : 1000, CST, No. 9508), MMP2 (1 : 500, 10373-2-AP, Proteintech), MMP9 (1 : 1000, 10375-2-AP, Proteintech), E-cadherin (1 : 500, A3044, AbClonal), p-IκB (1 : 500, AP0707, AbClonal), IkB (1 : 500, A1187, AbClonal), p65 (1 : 1000, A2547, AbClonal), and p-p65 (1 : 1000, AF2006, Affinity). The next day, the proteins were immuno-blotted with HRP-IgG at 37 °C for 1 h. The relative protein level was normalized to GAPDH (1 : 10000, sc-47778, Santa Cruz). Bands were visualized by a gel image system (WD-9413B, LIU YI, Beijing, China).

Cell Proliferation Assay

To study the function of SGK1 on cell proliferation, the cells at the density of 5×10⁴ cells/well were seeded in a 96-well plate of y13B (LIU YI, Beijing, China; AF2006, Affinity). Vector construction transfection or cells were treated with cisplatin (MB1055, meilunbio, Dalian) at 0, 0.25, 0.5, 1, 1.5, 2, 3, 4, and 6 μM after cell transfection. After further growth for different durations, 10 μL CCK-8 solution (96992, Sigma, USA) per well was added. After treatment at 37 °C for 1 h, the absorbance value at 450 nm was detected by a microplate reader (800Ts, BioTek, USA).

Cell Cycle and Apoptosis Assay

Cells (4×10⁴ cells/well) were planted in a 6-well plate. The cell cycle and apoptosis were determined by using the commercial kit (C1052, Beyotime) according to the users’ manual.

Wound-healing Assay

Cells were incubated with 1 μg/mL mitomycin C (M0503, Sigma) for 60 min. Next, the cells were wounded using a 200-μL pipette tip and the migration distance was photographed and recorded at 0 h, 24 h, and 48 h.

Invasion Assay

The transwell invasion assay was conducted to detect the function of SGK1 on cell invasion. Briefly, the upper chamber (3422, Corning) was coated with Matrigel matrix (356234, Corning, USA) for 2 hours at 37 °C. After that, 200 μL of cell suspension was added to the upper chamber, and 800 μL of medium supplemented with 10% FBS was added to the bottom chamber. After that, the cells were cultured for 24 hours at 37 °C, fixed with 4% paraformaldehyde, and next incubated with crystal violet (0.4%, 0528, Amresco, Dalian, China) for 24 h, and 48 h.

USA) for 5 min. Finally, the cells were photographed and the invasion cell number was counted.

**Immunofluorescence Assay**
Firstly, the cells were fixed with 4% paraformaldehyde at 25 °C for 15 minutes. Next, the cells were incubated for 30 minutes with Triton X-100 (ST795, Beyotime, China), the nonspecific proteins were then blocked with Normal Goat Serum (SL038, Solarbio, China) for 15 min at 25 °C. After incubation with the primary antibody p65 (1 : 200, A2547, ABclonal) at 4 °C for 18 h, the fluorescent antibody was applied and incubated at 25 °C for 1 h. Finally, the cell nuclei were counterstained with DAPI (C1002, Beyotime), and the pictures were obtained by a fluorescent microscope (DP73, OLYMPUS, Japan).

** Luciferase Reporter Assay**
The interaction between miR-15a-5p and SGK1 was evaluated by the luciferase reporter assay. Briefly, the SGK1 sequence with the targeted site of miR-15a-5p was inserted into the pmirGLO vector (Promega, USA) and named the wt-SGK1-site. Then, the binding site was mutated and named the mut-SGK1-site. After cell transfection, a luciferase detection kit (KGAF040, KeyGen, China) was applied to determine the luciferase activity, according to the manufacturer’s instructions.

**Statistical Analysis**
The quantitative values were shown as the means ± SD. Statistical analysis was conducted using GraphPad Prism 8.0 (GraphPad, CA, USA). The qRT-PCR, cell apoptosis assay, and luciferase reporter assay were analyzed with one-way ANOVA. The CCK-8, invasion, and wound-healing data were analyzed by two-way ANOVA. A value of P < .05 was identified as statistically significant. Moreover, all data were in a normal distribution, which was tested by the Kolmogorov–Smirnov test (P > .1).

**RESULTS**

**SGK1 Silencing Inhibited Gastric Cancer Cell Proliferation**
To determine the effect of SGK1 in gastric cancer, we investigated its level in 5 gastric cancer cell lines. Our results revealed that when compared with normal gastric epithelial cells, SGK1 expression was elevated in gastric cancer cells (Figure 1A), indicating that SGK1 might act as an activator in gastric cancer. Hence, we constructed an SGK1-silenced model in 2 gastric cancer cell lines. The knockdown efficiency of SGK1 was confirmed by qRT-PCR and Western blot assays (Figure 1Band C, P < .01). To disclose the effects of SGK1 in gastric cell proliferation, the CCK-8 was first conducted to evaluate cell proliferation. In the SGK1-silenced NCI-N87 cells, cell viability was significantly suppressed at 24 h, 48 h, and 72 hours (P < .05). In the SGK1-silenced HGC-27 cells, cell proliferation was also prominently repressed (Figure 1D, P < .01). Moreover, the cell cycle assay exhibited that SGK1 silencing results in a larger numbers of cells in the G1 phase, and to the contrary, fewer cells in the S and G2 phases after SGK1 silencing, than that in the control cells (Figure 1E). Western blot data suggested that the levels of cyclin D and cyclin E were reduced on account of SGK1 depletion (Figure 1F). Overall, we concluded that SGK1 deficiency repressed gastric cancer cell proliferation via mediating cell cycle progression.

**SGK1 Deficiency Suppressed Cell Migration and Invasion**
To determine whether SGK1 enhanced the capacity of gastric cancer cells to migrate and invade, we further conducted wound healing and transwell invasion assays. Our findings demonstrated that SGK1 deficiency led to the significant inhibition of cell migration at 48 hours, and invasion (Figure 2A-D, P < .01). Moreover, the decreased production of MMP9 and MMP2 were observed in SGK1-silenced NCI-N87 and HGC-27 cells (Figure 2E). Likewise, knockdown of SGK1 enhanced E-cadherin expression whereas it decreased N-cadherin expression (Figure 2F). Collectively, we revealed that SGK1 downregulation abrogated the cell migration and invasion activities of gastric carcinoma cells.

**SGK1 Deficiency Enhanced Cisplatin Sensitivity of Gastric Carcinoma Cells**
To assess the effect of SGK1 on cisplatin sensitivity, the SGK1 down-regulated cells were treated with cisplatin. Our CCK-8 assay showed that SGK1-silenced cells exhibited lessened cell viability relative to that in control cells after cisplatin treatment at different concentrations (0, 0.25, 0.5, 1, 1.5, 2, 3, 4, and 6 μM) (Figure 3A, P < .01). The IC50 in the control group was calculated based on the CCK-8 results in Figure 3A. To investigate whether knockdown of SGK1 could promote the cisplatin sensitivity, a low-dose cisplatin (1.5 μM, a quarter of the IC50 of the control cells, Figure S1) was chosen for the subsequent experiments. Compared with the negative control, SGK1 deficiency significantly promoted cell apoptosis in SGK1-silenced cells after cisplatin treatment (Figure 3B). Furthermore, the elevated production of active caspase-3 and caspase-9 were observed after cisplatin or si-SGK1 treatment. When cisplatin and si-SGK1 were used in combination, the production of active caspase-3 and caspase-9 was found to be further upregulated (Figure 3C). Collectively, our results showed that SGK1 deficiency enhanced cisplatin sensitivity of gastric cancer cell lines.

**SGK1 Silencing Repressed the NF-κB Signaling Pathway**
Mechanically, we detected the levels of p65, p-p65, IκBα, and p-IκBα. The immunofluorescence staining results showed that fewer p65-positive proteins were stained in the nucleus after SGK1 silencing, indicating that SGK1 loss of function repressed the nucleus translocation of p65 (Figure 4A). Consistent with the findings of immunofluorescence staining, Western blot results exhibited that SGK1 deficiency decreased the production of p-p65 and p-IκBα, suggesting an inhibitory effect of SGK1 silencing on the NF-κB pathway (Figure 4B). Taken together, these results affirmed that SGK1 accelerated gastric cancer deterioration, likely via regulating the NF-κB signaling pathway.
Interestingly, we also found that 3'UTR of SGK1 was regulated by miR-15a-5p, which was identified by the luciferase activity assay (Figure 5A and B, *P < .05). The qRT-PCR results implied that transfection with miR-15a-5p mimic declined the miR-15a-5p level (Figure 5C, **P < .01). Additionally, up-regulation of miR-15a-5p inhibited the migration activity whereas SGK1 overexpression reversed the changes induced by the miR-15a-5p mimic in NCI-N87 (Figure 5D and E, *P < .05) and HGC-27 cells (Figure 5D and E, **P < .01). The Western blot results displayed decreased expression of SGK1, cyclin E, and MMP9 in miR-15a-5p overexpressed cells, which was reversed by SGK1 up-regulation (Figure 5F). Thus, our results demonstrated that SGK1 might suppress the progression of gastric cancer via being targeted by miR-15a-5p.
DISCUSSION

As treatment strategies are constantly evolving, chemotherapy, such as platinum-based chemotherapy containing cisplatin and oxaliplatin, has become an effective method for the treatment of advanced gastric cancer. However, among patients undergoing treatment, there are obviously individual differences in response rates and survival rates, and drug resistance limits the productiveness of cancer chemotherapy and constitutes a major obstacle in clinical treatment. Therefore, enhancing the chemotherapy sensitivity of gastric cancer cells is essential for gastric cancer therapy.

In our study, we have revealed that SGK1 deficiency improved the sensitivity of gastric cancer cells to cisplatin, alleviating the deterioration of gastric carcinoma.

Recent researches have shown that SGK1 serves as an activator in tumorigenesis. For example, SGK1 enhances the survival, invasiveness, and adhesiveness of colorectal carcinoma cells. SGK1 presumably regulates the survival of kidney cancer cells and cholangiocarcinoma cells. In our study, we found that SGK1 deficiency suppressed the proliferation, invasion, and migration of gastric cancer cells. Consistently, Lang et al have
expounded that SGK1 promotes the adhesion of MDCK cells to fibronectin and inhibits chemotherapy-induced apoptosis, which exhibits an ability to confer cell survival.\textsuperscript{18,20} Zhang et al have declared that SGK1 exerts the antiapoptotic effect in breast cancer cells, in part via regulating NF-κB signaling.\textsuperscript{21} Additionally, SGK1 is implicated in chemoresistance.\textsuperscript{22} For instance, tumors with SGK1 gain of function are resistant to paclitaxel-dependent cell death.\textsuperscript{23} Our study also showed that SGK1 deficiency inhibited gastric cancer cell proliferation and promoted apoptosis after cisplatin treatment, indicating that downregulation of SGK1 improved chemotherapy sensitivity of gastric cancer cells to cisplatin.
Furthermore, our study stated that SGK1 loss of function obviously decreased the phosphorylation of p65 and IκB, indicating the downregulation of NF-κB. In line with our results, it is reported in hematological and epithelial malignancies that suppression of NF-κB leads to enhanced apoptosis. In addition, NF-κB activation in tumor cells can promote the production of new vascular and metastatic factors, causing drug resistance and reduced effectiveness of therapy. It is noteworthy that stimulating upstream genes of NF-κB exhibits a promoting effect in cancer resistance. A previous study has shown that SGK1 can act as an upstream activator of the NF-κB pathway, indicating that SGK1 may promote cancer resistance via upregulating the NF-κB pathway. Interestingly, we also showed that SGK1 might exert the inhibitory effect in gastric cancer via being targeted by miR-15a-5p. Existing research has indicated that miR-15a-5p level is reduced in cisplatin-resistant gastric cancer cells, which participates in the prognosis of patients with adjuvant chemotherapy. Another research has implicated that miR-15a-5p inhibition reduces the cisplatin sensitivity of carcinoma cells. However, the regulatory effect of miR-15a-5p by targeting SGK1 in gastric cancer needs further exploration.

In a word, these data indicated that downregulation of SGK1 promoted the chemosensitivity of gastric cancer to cisplatin and alleviated the deterioration of gastric carcinoma by suppressing the NF-κB pathway (Figure 6). Thus, SGK1 might become a prospective target for future gastric cancer therapy.
FIG. 5. (A-F). SGK1 targeted by miR-15a-5p promoted gastric cancer cell migration. (A) The binding site of miR-15a-5p on the 3’UTR of SGK1. (B) The luciferase activities in HGC-27 cells. *P < .05, compared with SGK1 3’UTR wt + NC mimic. (C) The miR-15a-5p production in 2 gastric carcinoma cells lines **P < .01, compared with NC mimic. (D, E) The cell migration of gastric carcinoma cells (with or without miR-15a-5p mimic or/and SGK1 overexpression transfected). Scale bar: 200 μm. One-way ANOVA, **P < .01, compared with NC mimic + vector; #P < .05, ##P < .01, compared with miR-15a-5p + vector. (F) The protein levels of SGK1, cyclin E, and MMP9 (with or without miR-15a-5p mimic or/and SGK1 overexpression transfected). GAPDH served as a loading control. Scale bar in (D): 200 μm.
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**References**


SUPPLEMENTARY TABLE 1. The Sequences of siRNA

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SUPPLEMENTARY TABLE 2. The Sequences of the Primers

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