

# Molecular mechanisms of MCM3AP-AS1 targeted the regulation of mir-708-5p on cell proliferation and apoptosis in gastric cancer cells

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**Abstract. – OBJECTIVE:** Gastric cancer (GC) is a common malignancy of the digestive tract. Accumulated studies proved that long non-coding RNA MCM3AP-AS1 (MCM3AP-AS1) may affect the mechanism of the progression of GC. However, the molecular mechanism is still unclear. Therefore, we aimed to explore the molecular mechanism of MCM3AP-AS1 targeting the regulation of microRNA-708-5p on cell proliferation and apoptosis in GC cells.

**MATERIALS AND METHODS:** The expression levels of MCM3AP-AS1 (MCM3AP antisense RNA 1) in gastric mucosal cells GES-1 and gastric cancer cell lines of MGc-803 and SGC-7901 cells were detected by qRT-PCR. The protein levels of Cyclin D1, P21 and Bcl-2 in MGc-803 and SGC-7901 cells after transfection were detected by Western blot. MTT assay was performed to detect cell proliferation and flow cytometry was carried out to determine GC cell apoptosis *in vitro*. In the end, the targeting relationship between MCM3AP-AS1 and microRNA-708-5p was detected by Dual-Luciferase reporter assay.

**RESULTS:** The level of MCM3AP-AS1 was significantly increased in GC cell lines. Knockdown of MCM3AP-AS1 curbed cell proliferation and enhanced apoptosis in MGc-803 and SGC-7901 cells. Furthermore, the effect of the downregulation of MCM3AP-AS1 on cell proliferation and apoptosis was reversed by knockdown of miR-708-5p, which was targeted by MCM3AP-AS1 *in vitro*.

**CONCLUSIONS:** MCM3AP-AS1 regulates the proliferation and apoptosis of gastric cancer cells by targeting the expression of microRNA-708-5p. The study may be useful to the therapy target of human GC.

**Keywords:**

MCM3AP-AS1, miR-708-5p, GC cell lines, Proliferation, Apoptosis

**Abbreviations**

GC = Gastric Cancer, MCM3AP-AS1 = MCM3AP antisense RNA 1, lnc RNA = long non-coding RNA.

## Introduction

Gastric cancer (GC) is a common malignant tumor of the digestive tract<sup>1</sup> and the second cancer for mortality rates worldwide<sup>2</sup>. At present, the main means of treating gastric cancer are surgery and chemotherapy, but the severe side effects and drug resistance caused by chemotherapy seriously affect the prognosis of patients<sup>3</sup>. Although researches revealed the partial mechanism of pathogenesis in human GC, further investigations to discover the potential regulatory mechanism in GC is needed.

Long non-coding RNA (lnc RNA) played an important role in cell proliferation, metastasis, invasion and apoptosis of various tumors<sup>4</sup>, including GC<sup>5</sup>. Such as lncRNA FOXD2-AS1, a tumor promoter in colorectal cancer<sup>6</sup> and involved in the poor prognosis of GC<sup>7</sup>. LncRNA H19 and miR-141 regulated cell proliferation and migration in GC cells<sup>8</sup>. What's more, MCM3AP-AS1 was also proved to enhance hepatocellular car-

cinoma growth<sup>9</sup>. Studies have applied bioinformatics methods to screen gastric cancer and its adjacent non-cancerous tissues and found that MCM3AP-AS1 was up-regulated in gastric cancer. All evidence suggested that MCM3AP-AS1 may participate in the pathogenesis in human GC. However, the effects of MCM3AP-AS1 on biological behaviors such as cell proliferation and apoptosis of GC have not been studied completely. The study aimed to discover the role of MCM3AP-AS1 on cell proliferation and apoptosis in GC cell lines *in vitro*.

Over the past decades, microRNA (miRNA) has been proved to be involved in tumorigenesis<sup>10,11</sup>. The expression of miR-181a-5p effected on cell proliferation in GC<sup>12</sup>. Meanwhile, miR-708-5p regulated prostate cancer cell apoptosis<sup>13</sup>; overexpression of miR-708-5p had a certain inhibitory effect on cell proliferation<sup>14</sup>. MiR-708-5p referred to the process of GC. The study demonstrated miR-708-5p expression was downregulated in GC cell lines. However, it has not been reported whether MCM3AP-AS1 targeted and regulated the biological behavior of miR-708-5p regulating GC cells *in vitro*.

In this study, bioinformatics methods and molecular biology techniques were used to investigate whether MCM3AP-AS1 targeted miR-708-5p to regulate the proliferation and apoptosis of gastric cancer cells, in order to find new therapeutic targets for clinical.

## Materials and Methods

### Cell Culture and Transient Transfection

Normal gastric mucosal cells GES-1 and gastric cancer cells MGC-803 and SGC-7901 were purchased from the Chinese Academy of Sciences Cell Bank (Shanghai, China). Of which GES-1, MGC-803 and SGC-7901 cells were cultured in Dulbecco's Modified Eagle's Medium (DMEM; HyClone, South Logan, UT, USA) medium containing 10% (volume fraction) fetal bovine serum (FBS, Gibco, Fisher Scientific, Waltham, MA, USA) and 100 U/ml antibiotics (penicillin 100 U/ml, streptomycin 100 µg/ml) (Gibco, Rockville, MD, USA) at a 5% CO<sub>2</sub> in cell incubator.

The vectors of MCM3AP-AS1 overexpression (pcDNA3.1-MCM3AP-AS1), overexpression empty control (pcDNA3.1), MCM3AP-AS1 knockdown (si-MCM3AP-AS1), knockdown negative control (si-NC), miR-708-5p inhibitor (anti-miR-708-5p) and inhibitor negative control

(anti-miR-NC) were purchased from Sangon Biotech (Shanghai, China). Subsequently, 48 h after cell passage, the cells were in logarithmic growth; next, Lipofectamine 2000 (Thermo Fisher Scientific) was carried to transfect vectors into cells according to manufacturer's manual.

### Quantitative Real-Time Polymerase Chain Reaction (qRT-PCR)

The logarithmic growth phase and transfected 48 h cells were collected. The total RNA was extracted by Trizol (Thermo Fisher Scientific). SuperScript<sup>III</sup> First-Strand Synthesis System (Thermo Fisher Scientific) was used to reverse transcribe into cDNA for qRT-PCR, reacting solutions were mixed using iQ<sup>TM</sup> SYBR<sup>®</sup> Green Supermix (Bio-Rad, Hercules, CA, USA), U6 and GAPDH were performed as internal reference genes. The process was as follow: 95°C for 30 minutes, 40 cycles at 95°C for 15 s, 60°C for 30 s and 72°C for 20 s, and then signals were collected by QuantStudio 3 QS3 (Thermo Fisher Scientific). The special primers were as follows: miR-708-5p: (Forward: 5'-GGC GCG CAA GGA GCT TAC AAT C-3', Reverse: 5'-GTG CAG CAG CAG AGG TAT-3'); U6: (Forward: 5'-GAG CCA CAG CGG AAC G-3', Reverse: 5'-CTA CCA CAT AGT CCA GG-3'); MCM3AP-AS1: (Forward: 5'-GCT GCT AAT GGC AAC ACT GA-3', Reverse: 5'-AGG TGC TGT CTG GTG GAG AT-3'); GAPDH: (Forward 5'- GGT GAA GGT CGG AGT CAA CG-3', Reverse: 5'-CAA AGT TGT CAT GGA TGA ACC-3')

### Western Blot

The total protein was extracted from MGC-803 and SGC-7901 cells condition in the logarithmic growth phase after transfecting for 48 h. The total protein concentration was determined by BCA Protein Assay Kit (Pierce, Waltham, MA, USA). SDS-polyacrylamide gel was prepared, 50 µg denatured protein was added per well, and electrophoresis was run. The isolated proteins were transferred onto polyvinylidene difluoride (PVDF) membrane (Roche, Basel, Switzerland); the voltage was controlled at 100 mV for 70 minutes. Then skim milk was used (Sangon Biotech) to block PVDF for 60 min; the diluted primary antibody was incubated at 4°C overnight. Tris Buffered Saline Tween (TBST) (Sangon Biotech) was washed three times for 5 min each. The diluted secondary antibody was incubated at room temperature for 60 minutes in the shaker, TBST

was washed three times for 5 min each. In the end point, protein signals were identified by using SuperSignal Chemiluminescent Substrates (Thermo Fisher Scientific) through Image Lab software (Bio-Rad). The special antibodies used in the study are listed as follows: the primary antibodies of Cyclin D1 (ab16663, 1:100), P21 (ab109520, 1:5000), Bax (ab32503, 1:5000), Bcl-2 (ab182858, 1:2000) and GAPDH (ab9485, 1:2500) were purchased from Abcam (Cambridge, MA, USA).

### MTT Assay

2.5% trypsin (Gibco) was carried out to digest MGc-803 and SGC-7901 cells, the cell concentration was adjusted to  $1 \times 10^5$ /ml and 100  $\mu$ l per well were added in 96-well plates. Whereafter, they were mixed with 20  $\mu$ l MTT reagent (Beyotime, Shanghai, China) and incubated for 4 hours before detection. Finally, the absorbance was measured at 490 nm (OD) by SpectraMax<sup>®</sup>iD3 (Molecular Devices, San Jose, CA, USA).

### Flow Cytometry

After transfection, MGc-803 and SGC-7901 cells were cultured for 48 h, then the medium was discarded and washed twice with PBS (phosphate-buffered saline) (HyClone). Subsequently, 500  $\mu$ l  $1 \times$  Binding Buffer, 5  $\mu$ l Annexin V-APC (Biosea Biotechnology, Beijing, China) were added in solutions and incubated for 15 minutes in the dark place; 2.5  $\mu$ l PI (Propidium iodide) was added and incubated for 5 minutes, then flow cytometry was performed to detect cell apoptosis *in vitro*.

### Dual-Luciferase Reporter Assay

Constructed vectors: wildtype 3'UT-MCM3AP-AS1 3'UT and mutant (MCM3AP-AS1-MCM3AP-AS1 3'UT) expression vectors of MCM3AP-AS1, renilla luciferase (pGL3) were purchased from Promega (Madison, WI, USA). Subsequently, they were transfected with one of them into MGc-803 and SGC-7901 cells by using Lipofectamine<sup>™</sup> 2000 and cultured for 48 hours. Dual-Luciferase assay kit (Promega) was added to lyse cells and stimulate luciferase signals were captured by SpectraMax<sup>®</sup>iD3 (Molecular Devices, San Jose, CA, USA).

### Statistical Analysis

SPSS 22.0 (SPSS Inc., IBM, Armonk, NY, USA) was used for data analysis. The Student's *t*-test was used for comparison between the two

groups. One-way analysis of variance with post-hoc test (ANOVA) was followed by Tukey's test for multiple comparisons of groups. GraphPad Prism (San Diego, CA, USA) was performed to plot images.  $p < 0.05$  indicates that the difference was statistically significant.

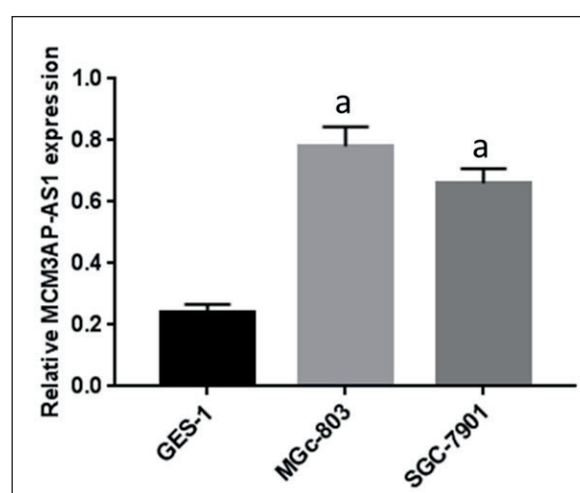
## Results

### The Level of MCM3AP-AS1 Was Strikingly Increased in GC Cell Lines

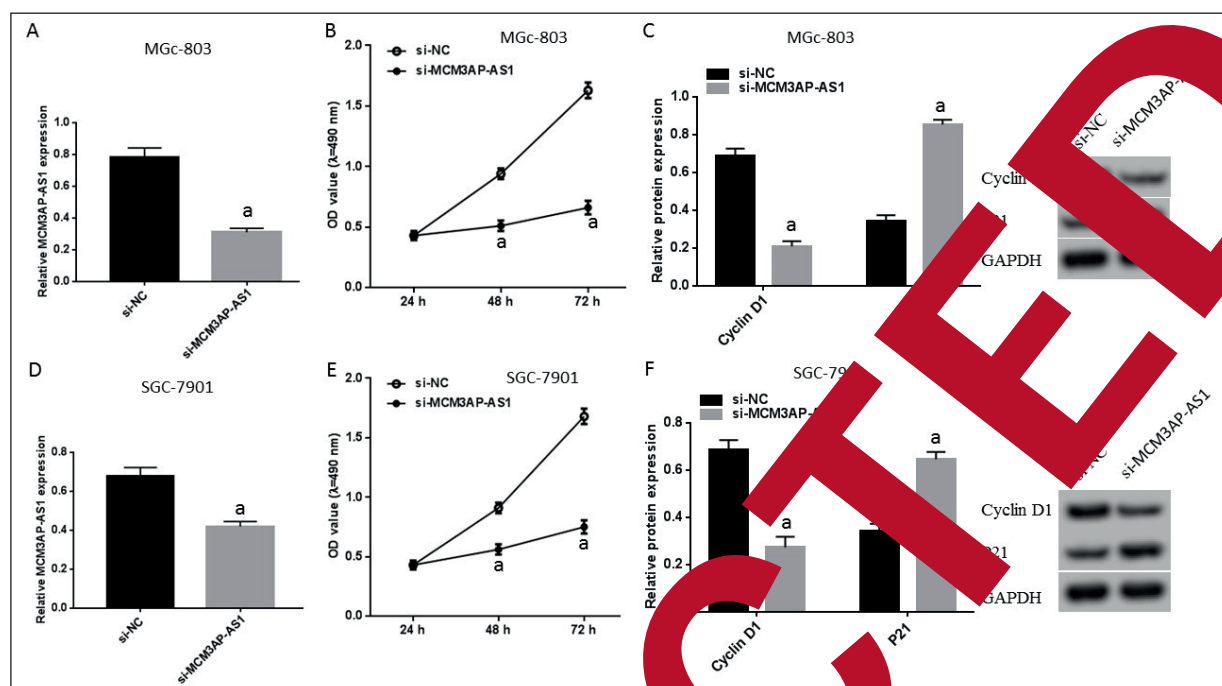
GC cell lines and GES-1 were cultured for 48 hours at 37°C, 5% CO<sub>2</sub>. Total RNA was extracted and the level of MCM3AP-AS1 was measured by qRT-PCR. The results showed that it was notably induced in MGc-803 and SGC-7901 cells (Figure 1).

### Knockdown of MCM3AP-AS1 Strikingly Suppressed Cell Proliferation in GC Cell Lines

Cell interfering RNA (si-RNA) of MCM3AP-AS1 (si-MCM3AP-AS1) and negative control (si-NC) were constructed and transfected into GC cell lines, respectively. The level of MCM3AP-AS1 was strikingly decreased in MGc-803 and SGC-7901 cells (Figure 2A and 2D). What's more, MTT assay was performed to detect GC cell proliferation *in vitro*. Evidence revealed that knockdown of MCM3AP-AS1 inhibited cell proliferation in GC cell lines (Figure 2B and 2E). Furthermore, the expression of



**Figure 1.** The level of MCM3AP-AS1 was strikingly increased in GC cell lines. The level of MCM3AP-AS1 in MGc-803 and SGC-7901 cells was measured by qRT-PCR. <sup>a</sup> $p < 0.05$ .



**Figure 2.** Knockdown of MCM3AP-AS1 strikingly suppressed cell proliferation in GC cell lines. Constructed knockdown vector of MCM3AP-AS1 (si-MCM3AP-AS1) and negative control (si-NC) were transfected into MGc-803 and SGC-7901 cells (A-F), respectively. The level of MCM3AP-AS1 was measured in MGc-803 (A) and SGC-7901 (D) cells via qRT-PCR. What's more, MTT assay was carried out to evaluate the cell proliferation of MGc-803 (B) and SGC-7901 (E) cells. Finally, the expression of Cyclin D1 and P21 was detected by Western blot *in vitro* (C and F). <sup>a</sup> $p < 0.05$ .

Cyclin D1 and P21, which was relative to cell cycle, was measured by Western blot and the effect of MCM3AP-AS1 downregulation on cell proliferation *in vitro* (Figure 2B and 2F) was verified.

### Downregulation of MCM3AP-AS1 Significantly Promoted Cell Apoptosis in GC Cell Lines

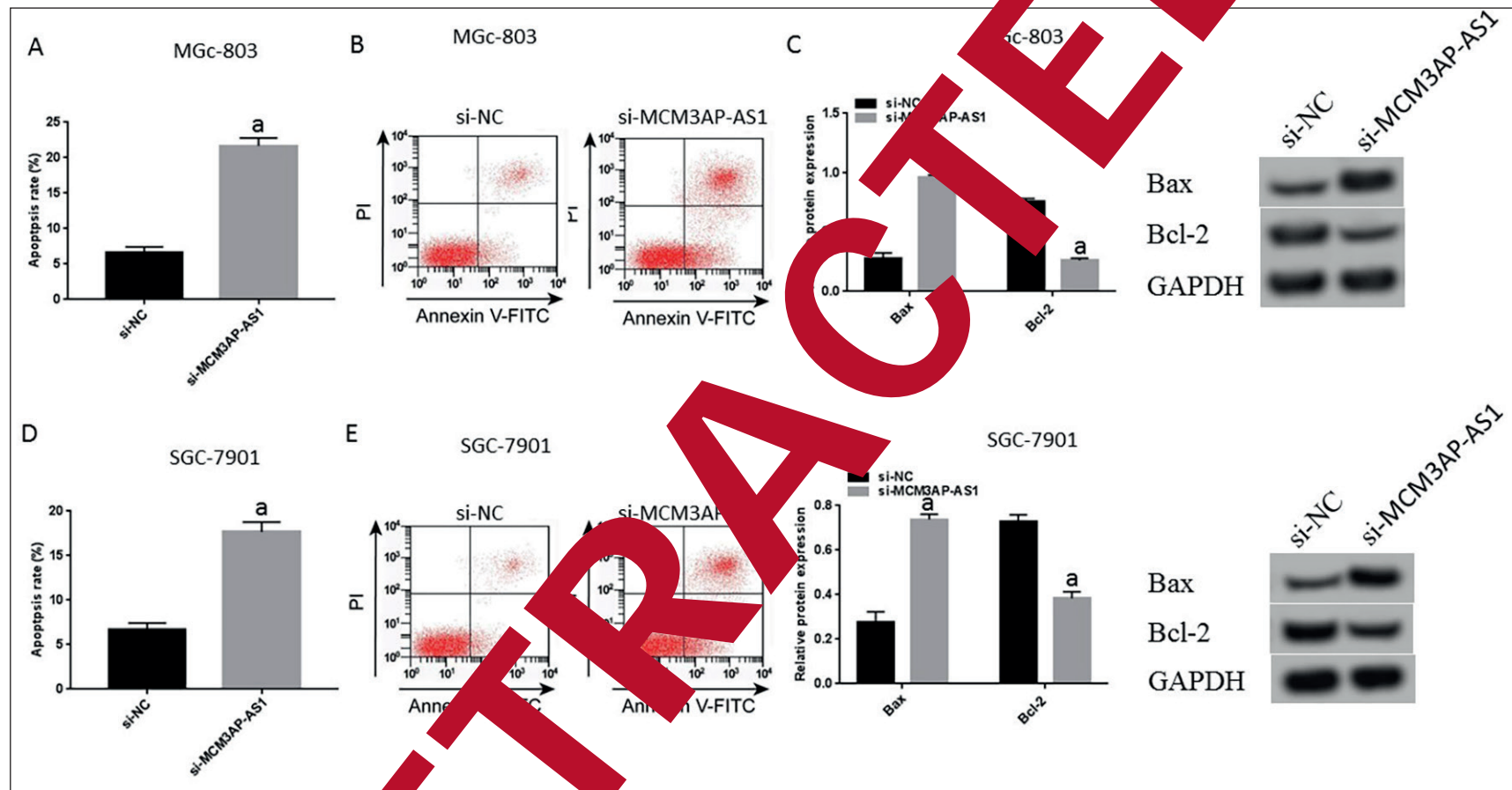
After transfecting with si-NC or si-MCM3AP-AS1 into MGc-803 and SGC-7901 cells, cell apoptosis was analyzed by flow cytometry. The results suggested that knockdown of MCM3AP-AS1 enhanced cell apoptosis in GC cell lines (Figure 3A, 3B, and 3C). Meanwhile, the expression of Bax and Bcl-2 was measured, which was referred to apoptosis by Western blot. It proved the result that knockdown of MCM3AP-AS1 on cell apoptosis *in vitro* (Figure 3D and 3F).

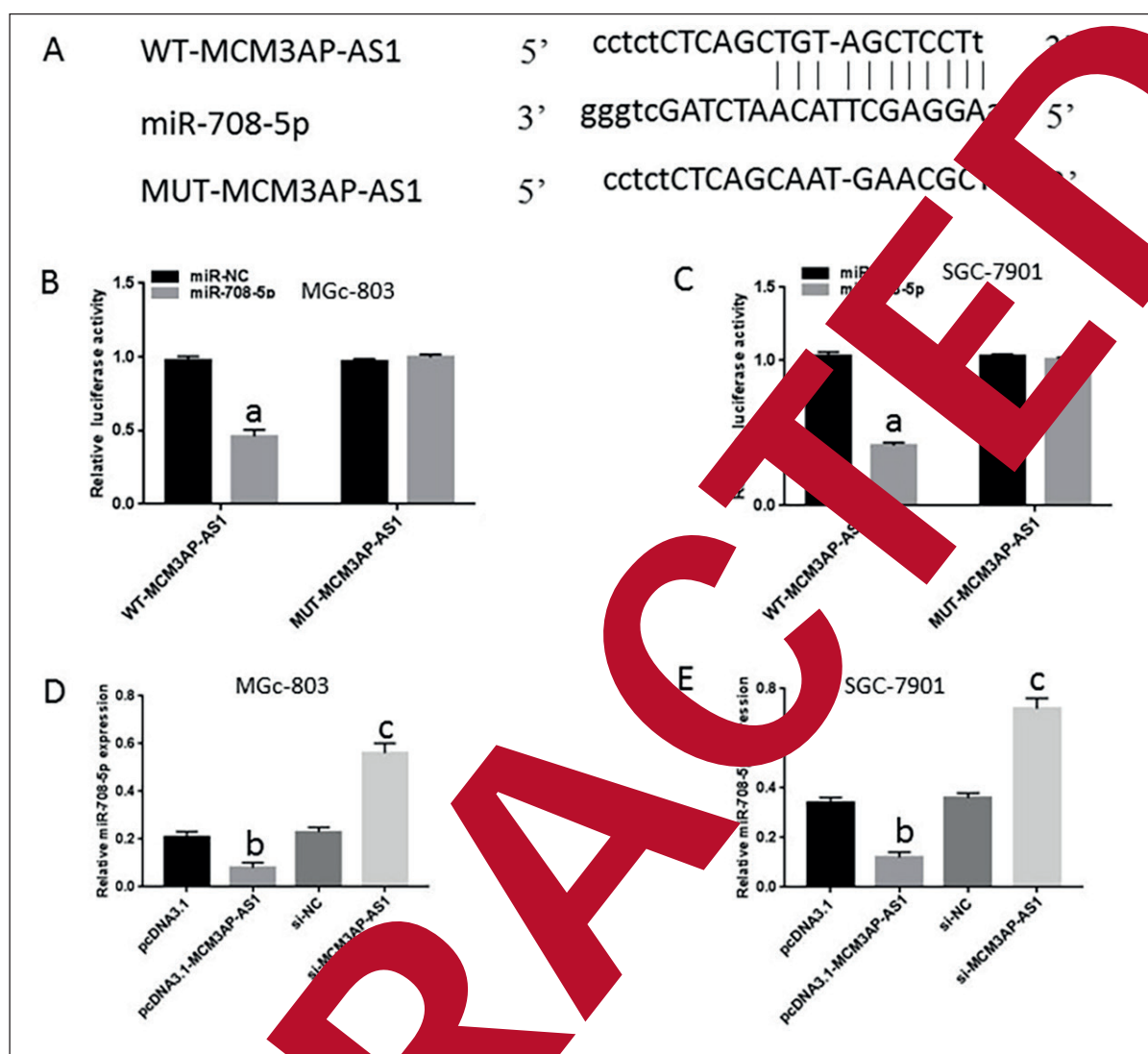
### miR-708-5p Was a Target Gene of MCM3AP-AS1

bioinformatics software was used to predict the relationship between MCM3AP-AS1 and miR-

### Knockdown of miR-708-5p Rescued the Effect of MCM3AP-AS1 on MGc-803 and SGC-7901 Cell Proliferation

After transfection with si-NC, si-MCM3AP-AS1, si-MCM3AP-AS1 + anti-miR-NC or si-MCM3AP-AS1 + anti-miR-708-5p into GC cell lines, the level of miR-708-5p was promoted by si-MCM3AP-AS1 while rescued *via* knockdown of miR-708-5p *in vitro* (Figure 5A and 5D). In addition, the downregulation of miR-708-5p reversed the effect of si-MCM3AP-AS1 on cell proliferation in GC cell lines (Figure 5B and 5E). Cell cycles relative proteins of Cyclin D1 and P21 also verified the conclusion of GC cell proliferation *in vitro*.





**Figure 4.** miR-708-5p is a target gene of MCM3AP-AS1. The binding sites between MCM3AP-AS1 and miR-708-5p were predicted by bioinformatics software of miRanda (A). Dual-luciferase reporter assay was used to verify the relationship of MCM3AP-AS1 and miR-708-5p *in vitro* (B). Furthermore, the level of miR-708-5p was measured after transfecting with up or down regulation factors of MCM3AP-AS1 (C). <sup>a</sup> $p < 0.05$ , <sup>b</sup> $p < 0.01$ , <sup>c</sup> $p < 0.001$ .

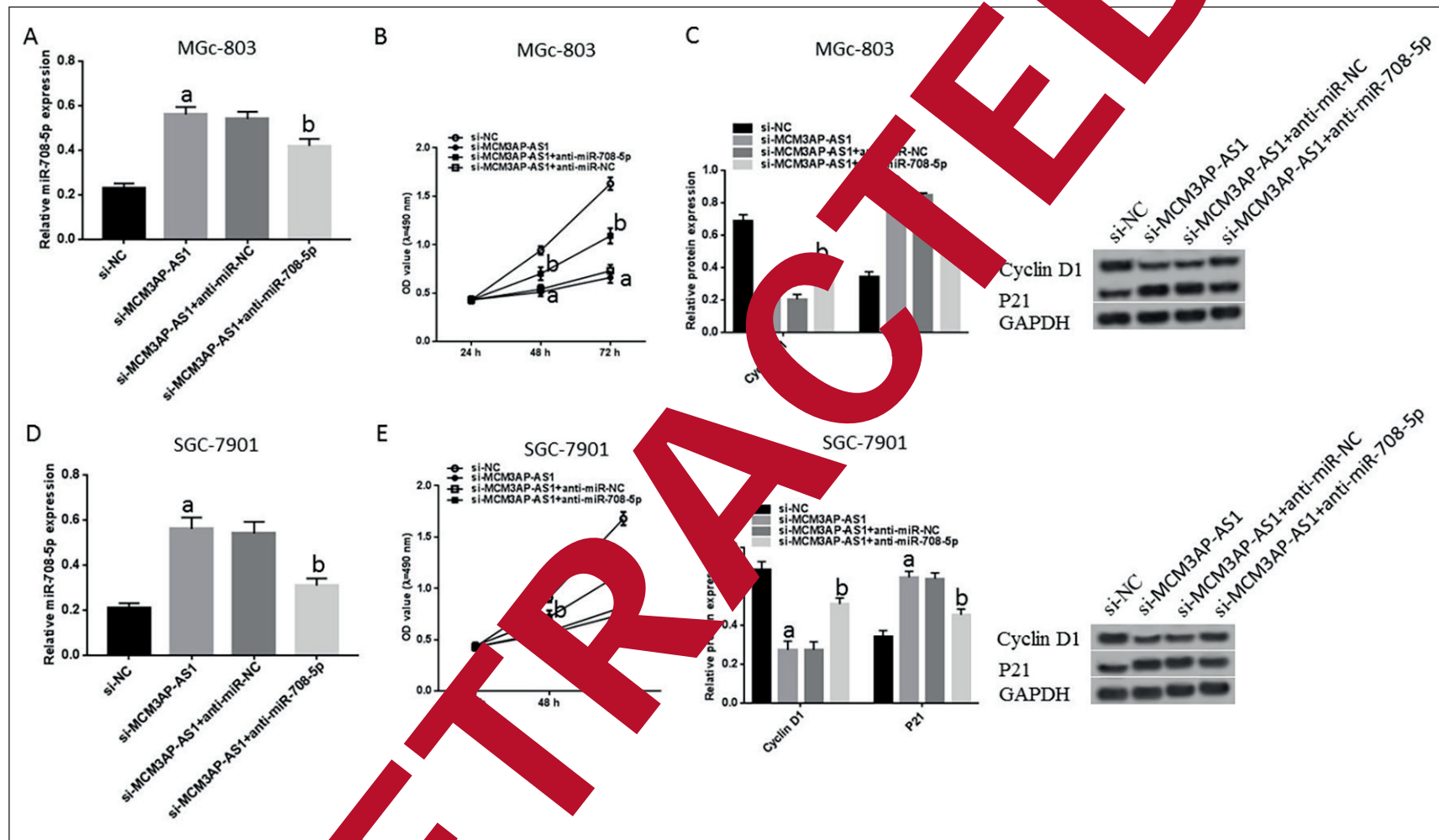
#### Downregulation of miR-708-5p Restored the Effect of MCM3AP-AS1 on Cell Apoptosis in GC Cell Lines

To further investigate the role of MCM3AP-AS1 and miR-708-5p on GC progression, a rescue experiment was carried out with MCM3AP-AS1 and miR-708-5p vectors, including si-NC, si-MCM3AP-AS1, si-MCM3AP-AS1 + anti-miR-708-5p in MGc-803 and SGC-7901 cells. The expression of miR-708-5p regained the effect of si-MCM3AP-AS1 on cell apoptosis *in vitro* (Figure 6A and 6C). Meanwhile, the

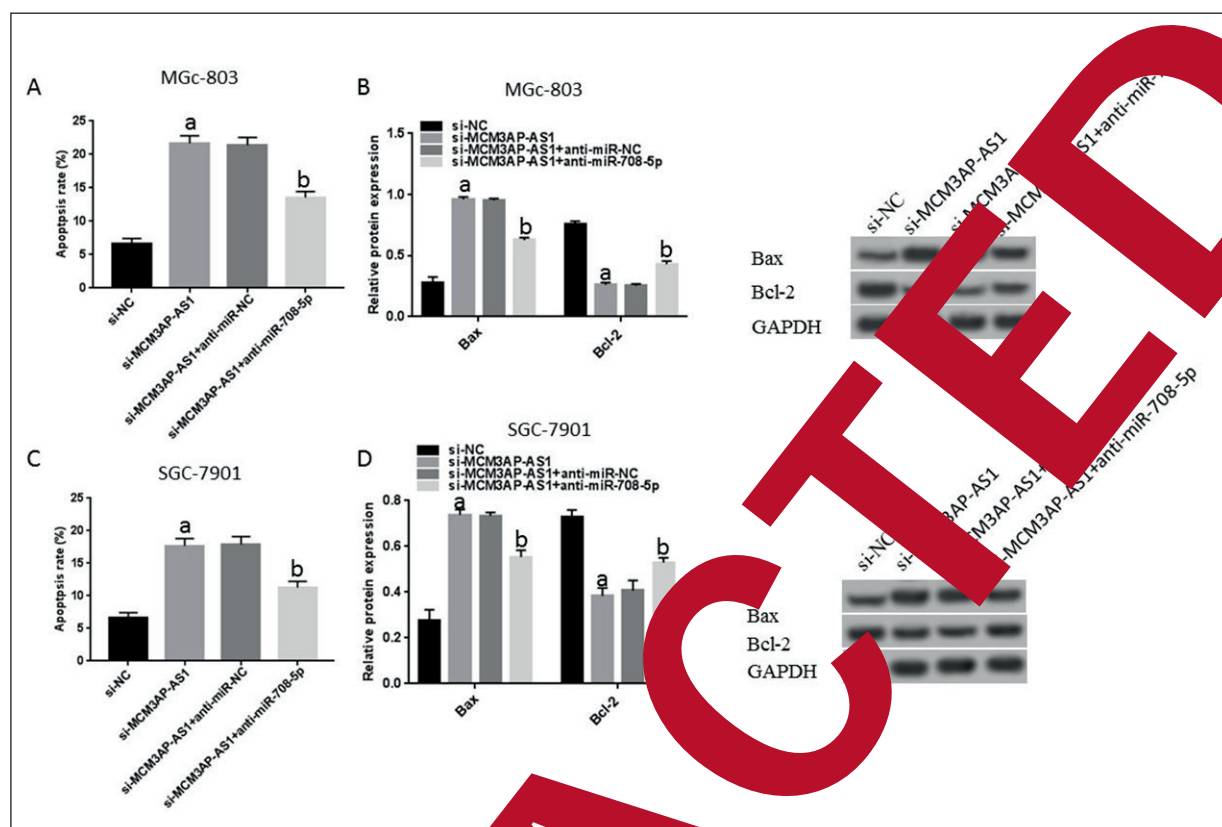
conclusion was also verified by measuring the expression of Bax and Bcl-2 in GC cell lines (Figure 6B and 6D).

#### Discussion

Gastric cancer is a malignant tumor that occurs in the gastric mucosa. Its incidence rate ranks first among all malignant tumors in China. Due to the different living habits and dietary structure in the north and the south, the incidence of gastric cancer in coastal areas and northwestern China



**Figure 5.** Knockdown of miR-708-5p reversed the effect of MCM3AP-AS1 on MGc-803 and SGC-7901 cell proliferation. Transfected vectors of si-NC, si-MCM3AP-AS1, si-MCM3AP-AS1 + anti-miR-NC or si-MCM3AP-AS1 + anti-miR-708-5p into GC cell lines (A-F). The level of miR-708-5p was measured by qRT-PCR (A and D). Moreover, MTT was used to detect cell proliferation of GC cell lines (B and E). Finally, the expression of apoptosis-related proteins of Cyclin D1 and P21 was examined by Western blot (C and F). <sup>a</sup> $p < 0.05$ , <sup>b</sup> $p < 0.01$ .



**Figure 6.** Downregulation of miR-708-5p restores the effect of MCM3AP-AS1 on cell apoptosis in GC cell lines. GC cell lines transfected with si-NC, si-MCM3AP-AS1, si-MCM3AP-AS1+anti-miR-NC or si-MCM3AP-AS1+anti-miR-708-5p into MGC-803 and SGC-7901 cells (A-D), respectively. Cell apoptosis was measured by flow cytometry *in vitro* (A and C). Meanwhile, Bax and Bcl-2 expression was also identified by western blot (B and D). <sup>a</sup> $p < 0.05$ , <sup>b</sup> $p < 0.01$ .

is significantly higher than that in the youth. The recurrence and metastasis of tumor are the main reason for the difficulty of treatment. It is reported that tumor metastasis and invasion are affected by various lncRNAs in the past decades. Therefore, the role of lncRNA in gastric cancer is of great significance for the targeted treatment of gastric cancer.

lncRNA is a non-coding RNA that occupies 98% of human gene transcripts and is greater than 200 nt in length<sup>15</sup>. In recent years, it has been found that lncRNA can regulate the activity of various proteins at multiple levels, affecting various biological behaviors such as cell proliferation, differentiation and apoptosis of tumor cells<sup>16-18</sup>. Studies have shown that lncRNA MCM3AP-AS1 expression in primary liver cancer is significantly higher than adjacent tissues and normal liver tissues<sup>19</sup>. Further, MCM3AP-AS1 is significantly up-regulated in glioblastoma, and inhibition of MCM3AP-AS1 expression inhibits cell viability, migration, and angiogenesis

in glioblastoma<sup>20</sup>. The qRT-PCR assay showed that the level of MCM3AP-AS1 in two GC cells was higher than that in normal GC cells, which was consistent with the above results, suggesting that MCM3AP-AS1 was a key factor in the development and progression of gastric cancer. Meanwhile, Wang et al<sup>9</sup> reported that MCM3AP-AS1 was elevated in hepatocellular carcinoma, which promoted the growth of hepatocellular carcinoma by targeting inhibited expression of mir-194-5p. To further investigate the role of MCM3AP-AS1 in GC, this study repressed the expression of MCM3AP-AS1 in GC cell lines and detected cell proliferation and apoptosis *in vitro*. The results showed that after inhibiting the expression of MCM3AP-AS1, GC cell proliferation was significantly decreased, and the cell apoptosis rate was enhanced. The levels of Cyclin D1 and Bcl-2 were significantly decreased, and the levels of P21 and Bax were notably increased, suggesting that MCM3AP-AS1 may promote cell proliferation and inhibit apoptosis. In addition, the expression



of miR-708-5p was significantly increased after inhibiting the expression of MCM3AP-AS1, suggesting that MCM3AP-AS1 may regulate the expression of miR-708-5p.

MiR-708-5p is a newly discovered microRNA that plays an important role in cell proliferation, migration, invasion, and apoptosis in tumor<sup>13,21</sup>. The miRcod bioinformatics software predicted that MCM3AP-AS1 and miR-708-5p had common fragment sequences, speculating that miR-708-5p may be the target gene of MCM3AP-AS1. Subsequently, Dual-Luciferase report assay and western blot were carried out to further verify MCM3AP-AS1 negatively regulated the expression of miR-708-5p by directly binding to the miR-708-5p 3'UTR region.

Studies have reported that miR-708-5p expression is decreased in human mesenchymal stem cells, and overexpression of miR-708-5p promoted cell proliferation and migration. The expression of miR-708-5p was down-regulated in renal cell carcinoma, and it could inhibit both cell apoptosis of renal cancer and the occurrence of renal cancer by inhibiting the oncogene zinc finger E-box binding to homologous box protein 2<sup>22</sup>. Furthermore, miR-708-5p inhibited migration of breast cancer by inhibiting endoplasmic reticulum membrane protein and reducing the activation of extracellular signal-regulated kinases<sup>23</sup>. Afterward, the effect of MCM3AP-AS1 targeting miR-708-5p in GC cell lines was verified, anti-miR-708-5p was transfected into GC cells, which stable knockdown of MCM3AP-AS1. The results showed that miR-708-5p inhibitor reversed the effect of downregulation of MCM3AP-AS1 on cell proliferation and apoptosis *in vitro*. Furthermore, the expression of Cyclin D1, P21 and Bcl-2 further verified the above conclusion.

In summary, MCM3AP-AS1 targeted miR-708-5p to regulate cell proliferation and apoptosis of GC cells. This study clarified the mechanism of MCM3AP-AS1 targeting miR-708-5p and the target therapy for GC, providing potential therapeutic strategies. However, further investigations in clinical treatment of human GC are needed.

### Conclusions

The level of MCM3AP-AS1 was notably enhanced in GC cell lines. Further, the downregulation of MCM3AP-AS1 decreased cell proliferation and boosted apoptosis in MGC-803 and

SGC-7901 cells. miR-708-5p was a target gene of MCM3AP-AS1; its knockdown reversed the effect of the downregulation of MCM3AP-AS1 on cell proliferation and apoptosis in GC cell lines.

### Conflict of Interest

The Authors declare that they have no conflict of interest.

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