miR-206 regulates 5-FU resistance by targeting Bcl-2 in colon cancer cells

Introduction: Previous studies have found that miRNAs play a key role in drug resistance. Multiple reports show that miRNAs act as regulators in colorectal cancer (CRC) cells, but the role of miR-206 in CRC is still not well understood. The current study aimed to explore the potential function of miR-206 in 5-FU resistance.

Methods: To indentify the role of miR-206 in 5-FU resistance, the expression of miR-206 was examined by real-time polymerase chain reaction (RT-PCR) in 5-FU-resistant (FR) CRC (HCT116/FR and RKO/FR) and their parental cell lines. miR-206 mimic was transfected to 5-FU-FR CRC, and the 5-FU sensitivity was detected by MTS and flow cytometry. Using miRNA target prediction software, we found that miR-206 could target the 3′ untranslated region (3′UTR) sequence of Bcl-2.

Results: miR-206 was found to be downregulated in 5-FU-FR CRC in comparison with their parental cell lines, suggesting its crucial relevance for colon cancer biology. Downregulation of miR-206 promoted drug resistance and decreased apoptosis of parental cells, while overexpression of miR-206 promoted drug cytotoxicity and apoptosis of HCT116/FR cells. We also identified miR-206 targeting Bcl-2 directly in CRC, which is required for miR-206 mediated-5-FU resistance.

Conclusion: Our results show that miR-206 targets Bcl-2 to mediate chemoresistance, proliferation, and apoptosis in CRC. This study provides a novel promising candidate for colon cancer therapy.

Keywords: miR-206, 5-FU, resistance, Bcl-2, colorectal cancer

Introduction

Colorectal cancer (CRC) is a leading cause of cancer incidence and death worldwide. The overall relative 5-year survival of patients with CRC is about 50%. Almost half of the patients with CRC are confronted with liver metastasis either at the time of diagnosis or later during the course of the disease. 5-Fluorouracil (5-FU)-based chemotherapy is the key therapy for advanced disease. The researchers used a hydrogen that simulated the location of a pyrimidine ring in the form of a carbon-5 uracil and fluorine, to meet the expected biochemical, pharmacological, and clinical anticancer drug activity.

miRNAs are endogenous 21–24-nucleotide single-strand noncoding RNAs that bind to the target mRNAs at 3′-UTR through imperfect complementary match to decrease protein expression by either translation repression or mRNA cleavage. A previous report indicated that miRNAs participate in various biological processes, such as tissues homeostasis, organ development, cellular proliferation, apoptosis, and human diseases. In cancer cells, miRNAs may function as tumor promoters by targeting tumor suppressor genes as well as tumor suppressors by inhibiting cellular oncogene expression.
Right now, miR-206 is one of the well-studied and excellently characterized miRNAs.\textsuperscript{14,15} miR-206 downregulation has been shown in many types of cancers.\textsuperscript{14} A previous study showed that miR-206 suppresses cancer cell invasion and migration via Cdc42.\textsuperscript{16} Moreover, miR-206 also inhibits VEGF expression and regulates the migration and apoptosis of laryngeal cancer cells.\textsuperscript{17} However, the role of miR-206 in 5-FU resistance regulation remains unclear.

In this study, the downregulation of miR-206 was found in the 5-FU-resistant CRC. Our results shown that miR-206 regulates 5-FU resistance by changing the protein levels of its target gene Bcl-2. Our findings indicated that miR-206/Bcl-2 as a novel regulator mediates 5-FU resistance in CRC.

**Materials and methods**

**Cell culture and reagents**
CRC cell lines including HCT116 and RKO were obtained from American Type Culture Collection (Rockville, MD, USA). The cells were cultured in Dulbecco’s Modified Eagle’s Medium (Gibco, Gaithersburg, MD, USA) supplemented with 10% fetal bovine serum (Gibco), and 1% penicillin–streptomycin at 37°C in a humidified environment with 5% CO\textsubscript{2}. miR-206 inhibitor, miR-206 mimic, or the appropriate negative controls (NCs) of miRNA inhibitor and miRNA mimic were obtained from GenePharma (Shanghai, People’s Republic of China).

5-FU-resistant cell lines, HCT116/FR and RKO/FR, were derived by incubation with stepwise increasing concentrations of 5-FU. Briefly, HCT116 and RKO cells were treated with increasing dose of 5-FU from 10 to 100 \( \mu \)g/mL for more than 12 months. During the treatment, after each round, when the surviving cells reached >70% confluence they were passaged by trypsinization, and the dose of 5-FU was increased.

**MTS assay**
Indicated cells were transfected with miR-206 mimic/inhibitor. After 24 hours, cells were cultured in 96-well plates at a density of 4x10\textsuperscript{3} cells/well. Then the cells were treated with increasing concentration of 5-FU for 72 hours. Subsequently, MTS assay was performed using the MTS assay kit (Promega, Fitchburg, WI, USA) according to the manufacturer’s instructions. Luminescence was measured with a Wallac Victor 1420 Multilabel Counter (Perkin Elmer, Bridgeville, PA, USA). Each assay was conducted in triplicate and repeated three times.

**siRNA transfection**
Bcl-2 siRNA and control siRNA were purchased from Santa Cruz Biotechnology (Dallas, TX, USA). HCT116 cells were seeded in 12-well plates for 24 hours to reach 30%–40% confluence and transiently transfected cells with Bcl-2 siRNA or control siRNA with Lipofectamine 2000 (Invitrogen, Carsbad, CA, USA) according to the manufacturer’s instructions.

**RNA extraction and real-time reverse transcription-polymerase chain reaction (PCR)**
Total RNA was extracted using the TRIzol RNA Kit (Invitrogen) according to the manufacturer’s protocol. Briefly, total RNA was used to generate cDNA using SuperScript II reverse transcriptase (Invitrogen). Polymerase chain reaction (PCR) was performed in triplicate using SsoFasrTM Probes Supermix (Bio-Rad, Hercules, CA, USA) in a final reaction volume of 20 \( \mu \)L with gene-specific primer/probe sets and a standard thermal cycling procedure (35 cycles) on a Bio-Rad CFX96™ Real-time PCR System. Relative gene mRNA levels were assessed using TaqMan Gene Expression Real-Time PCR assays. Result was expressed as the threshold cycle (C\textsubscript{t}). The relative quantification of the target transcripts was determined by the comparative C\textsubscript{t} method (\( \Delta \Delta C\text{t} \)) according to the manufacturer’s protocol. The 2\textsuperscript{-}\( \Delta \Delta C\text{t} \) method was used to analyze the relative changes in gene expression. Control experiments were conducted without reverse transcription to confirm that the total RNA was not contaminated with genomic DNA. \( \beta \)-Actin expression served as the internal reference for Bcl-2. U6 RNA served as an internal control for miRNA. The primers used were as follows: Bcl-2: forward: 5\textsuperscript{′}-GACCTGACACACTACCTCAT-3\textsuperscript{'}, reverse: 5\textsuperscript{′}-CTGCACCTGACCCCTTCACC-3\textsuperscript{'}, \( \beta \)-actin: forward: 5\textsuperscript{′}-GACCTGACACACTACCTCAT-3\textsuperscript{'}, reverse: 5\textsuperscript{′}-AGA CAGCACTGTGTTGGCATA-3\textsuperscript{′}.

**Western blotting**
Western blotting was performed as previously described,\textsuperscript{18,19} with antibodies for Bcl-2 (ab692; Abcam, Cambridge, MA, USA) and \( \beta \)-actin (#4970; Cell signaling, Danvers, MA, USA).

**Apoptosis assays**
Annexin V/propidium iodide staining was performed using annexin–Alexa 488 (Invitrogen) and PI.\textsuperscript{20}

**Luciferase reporter assay**
To evaluate the function of miR-206, the 3\textsuperscript{′}-UTR of Bcl-2 was amplified and inserted downstream of the luciferase reporter gene in the luciferase reporter pmIR-REPORT luciferase reporter vector. The mutant 3\textsuperscript{′}-UTR of Bcl-2 was amplified using wild-type (WT) Bcl-2 3\textsuperscript{′}-UTR as the template, and the mutant plasmid was created using a Site-Directed Mutagenesis
Kit. For luciferase reporter assays, the cells were cotransfected with miR-206 mimics and WT or mutant Bcl-2 3’-UTR, along with Renilla luciferase pRL-TK vector (Promega) as an internal control. Following transfection for 24 hours, the cells were collected and lysed using RIPA buffer. Luciferase activity was then measured by using Dual Luciferase Assay System (Promega) according to the manufacturer’s instructions.

**Xenograft study**

All procedures and experiments involving animals in this study were approved by the Committee on the Ethics of Animal Experiments of Northeast Electric Power University and were performed in compliance with the institutional ethical guidelines for animal experimentation according to the guide of the Committee on the Ethics of Animal Experiments of Northeast Electric Power University. HCT116 and HCT116/FR cells were suspended in 100 μL of phosphate-buffered saline at a concentration of 4×10⁶ cells/mL and injected into either flank of the same NOD/SCID mice at 5–6 weeks of age (n=6). 5-FU (20 mg/kg), polyplex containing 500 nM miR-206 mimic in 50 μL of phosphate-buffered saline, or their combination was administrated by intraperitoneal injection twice a week. Tumor growth was monitored using calipers, and tumor volumes were calculated according to the formula: 1/2 x length x width². Mice were euthanized when tumors reached ~1.0 cm³ in size. Tumors were dissected and fixed in 10% formalin and embedded in paraffin. TUNEL immunostaining was performed on 5 μM paraffin-embedded tumor sections by using an AlexaFluor 488-conjugated secondary antibody (Invitrogen) for signal detection.

**Statistical analysis**

All data are represented as the mean of at least triplicate samples ± standard deviation. Statistical analysis included a one-way analysis of variance or Student’s t-test using GraphPad Prism V software (GraphPad Software, La Jolla, CA, USA). *P*<0.05 was considered statistically significant.

**Results**

**Generation of 5-FU-resistant CRC**

To investigate the mechanism of 5-FU resistance in CRC, we generated HCT116 and RKO 5-FU-resistant cell lines (HCT116/FR and RKO/FR cells). To test the resistance properties of HCT116/FR and RKO/FR cells and the parental cells, we exposed these cells to 5-FU. As expected, 5-FU treatment could remarkably inhibit the proliferation and induce apoptosis of parental cells but not resistant cells (Figure 1A and B). Moreover, significant changes were observed in the distribution of cell cycle phases in HCT116/FR and RKO/FR after 5-FU treatment. As shown in Figure 1C, 5-FU treatment could induce G1 phase arrest strikingly in HCT116 and RKO cells, but not in HCT116/FR and RKO/FR. These data demonstrate that 5-FU-resistant cells (HCT116/FR and RKO/FR) were generated successfully.

**miR-206 was downregulated in HCT116/FR and RKO/FR cells**

To determine the involvement of miR-206 in 5-FU-resistant HCT116/FR and RKO/FR cells, we used qRT-PCR to examine miR-206 expression in these cells. Compared to parental cells, the expression level of miR-206 was significantly decreased in 5-FU-resistant cells (Figure 2A and B). These results indicate that miR-206 downregulation may contribute to 5-FU resistance in CRC.

**5-FU resistance can be overcome by miR-206 overexpression in CRC**

Next, we investigated the role of miR-206 in regulating resistance to 5-FU in CRC. miR-206 was overexpressed in HCT116/FR and RKO/FR cell lines (Figure 3A). Compared to miR-mimic-NC cells, MTS of the miR-206-overexpression (miR-206 mimic) HCT116/FR and RKO/FR cell lines was inhibited to a greater extent by 5-FU (Figure 3B and C). Furthermore, miR-206 mimic recovered apoptosis induced by 5-FU in 5-FU-resistant cells (Figure 3D). Thus, the results demonstrate that miR-206 mimic overcame 5-FU resistance in 5-FU-resistant cells.

**Inhibition of miR-206 promotes resistance to 5-FU in CRC**

Then, to determine the association of 5-FU resistance and miR-206 in HCT116 and RKO cells, we examined the role of inhibition of miR-206 in the 5-FU-sensitive cells. As shown in Figure 4A, qRT-PCR results indicated that miR-206 inhibitor decreased levels of miR-206, suggesting that miR-206 is efficiently the downregulated in HCT116 and RKO cells. HCT116 and RKO cells transfected with miR-206 inhibitor demonstrated significant resistance to 5-FU compared with the NC group (Figure 4B). Moreover, HCT116 and RKO cells treated with miR-206 inhibitor had a lower apoptosis rate than the NC group in response to 5-FU treatment (Figure 4C). Collectively, these data indicate that miR-206 downregulation contributes to the resistance to 5-FU in CRC.

**Bcl-2 regulates miR-206-mediated 5-FU resistance**

Next, we used TargetScanHuman 6.2 (http://www.targetscan.org) to predict the targeting gene and found that Bcl-2 is a target gene of miR-206 (Figure 5A). To confirm whether
Bcl-2 is targeted by miR-206, a luciferase reporter vector with the target site or the mutant target site downstream of the luciferase gene (Bcl-2-3′-UTR-WT and Bcl-2-3′-UTR-mut) was generated. Compared with the Bcl-2-3′-UTR-mut and miR-206 mimic cotransfected group, a significant decrease was observed in Bcl-2-3′-UTR-WT and miR-206 mimic cotransfected group in HCT116 cells (Figure 5B). These data indicate that Bcl-2 is a target gene of miR-206.

Next, we analyzed whether miR-206 regulates Bcl-2 protein level in HCT116 and RKO cells. HCT116 and RKO cells transfected with miR-206 mimic downregulated mRNA and protein level of Bcl-2 in the cells (Figure 5C). Moreover, transfected miR-206 inhibitor upregulated mRNA and protein levels of Bcl-2 in HCT116 and RKO cells (Figure 5D). These results indicate a strong inverse and consistent correlation between the miR-206 levels and Bcl-2.

To investigate if Bcl-2 plays an important role in miR-206-mediated 5-FU resistance, we transfected the Bcl-2 or control siRNA into the miR-206 inhibitor-treated 5-FU-resistant A

B. ** **

C. ** **

Figure 1 Generation of 5-FU-resistant HCT116 and RKO cells. Notes: (A) The indicated cell lines were treated with increasing concentrations of 5-FU for 72 hours. Cell proliferation was determined by MTS assay. (B) The indicated cell lines were treated with 40 μg/mL 5-FU for 48 hours. Apoptosis was analyzed by Annexin V/PI staining followed by flow cytometry. (C) The indicated cell lines were treated with 40 μg/mL 5-FU for 48 hours. Cell cycle distribution was detected by flow cytometry. Results in (A–C) are expressed as means ± SD of three independent experiments. **P<0.01.

Abbreviations: 5-FU, 5-fluorouracil; PI, propidium iodide; SD, standard deviation.
Figure 2 miR-206 expression in 5-FU-resistant cells.

Notes: (A) Relative miR-206 level in HCT116 parental cells and 5-FU-resistant cells was detected by real-time PCR. (B) Relative miR-206 level in RKO parental cells and 5-FU-resistant cells was detected by real-time PCR. Results are expressed as means ± SD of three independent experiments. **P < 0.01.

Abbreviations: 5-FU, 5-fluorouracil; PCR, polymerase chain reaction; SD, standard deviation.

Figure 3 Effect of miR-206 in 5-FU-resistant cells.

Notes: (A) HCT116/Fr and RKO/Fr cells were transfected with miR-206 mimic or mimic-nc for 24 hours. Relative miR-206 level was detected by real-time PCR. (B) HCT116/Fr cells were transfected with miR-206 mimic or mimic-NC for 24 hours, and then treated with increasing concentrations of 5-FU for 72 hours. Cell proliferation was determined by MTS assay. (C) RKO/Fr cells were transfected with miR-206 mimic or mimic-NC for 24 hours, and then treated with increasing concentrations of 5-FU for 72 hours. Cell proliferation was determined by MTS assay. (D) HCT116/Fr and RKO/Fr cells were transfected with miR-206 mimic or mimic-NC for 24 hours, and then treated with 40 μg/mL 5-FU for 48 hours. Apoptosis was analyzed by Annexin V/PI staining followed by flow cytometry. Results are expressed as means ± SD of three independent experiments. *P < 0.05; **P < 0.01.

Abbreviations: 5-FU, 5-fluorouracil; NC, negative control; PCR, polymerase chain reaction; PI, propidium iodide; SD, standard deviation.
HCT116 and RKO cells, and cell survival rate was assessed. The results confirmed that Bcl-2 level was reduced in HCT116 and RKO cells, and Bcl-2 siRNA significantly decreased the survival rate of the 5-FU-resistant HCT116 and RKO cells compared with the control group (Figure 5E), suggesting that miR-206 downregulation modulates 5-FU resistance in HCT116 cells by upregulating Bcl-2.

Inhibition of miR-206 overcomes 5-FU resistance in vivo

Based on our findings that 5-FU combined with miR-206 mimic synergistically inhibited the growth of 5-FU-resistant cells in vitro, we confirmed the combination effect in vivo in mice. HCT116 and HCT116/FR tumors were treated with miR-206 mimic, 5-FU, or their combination.
Figure 5 Bcl-2 as a direct target gene of mir-206 in colon cancer.

Notes: (A) The predicted binding site of mir-206 in the 3′-UTR of Bcl-2. (B) The relative luciferase activity of HCT116 cells was detected after Bcl-2-3′-UTR-WT or mut was cotransfected with mir-206 mimic and mimic-NC. (C) HCT116 and RKO cells were transfected with mir-206 mimic or mimic-NC for 24 hours. Bcl-2 mRNA was detected by real-time PCR (left), and Bcl-2 protein level was analyzed by Western blotting. (D) HCT116 and RKO cells were transfected with mir-206 inhibitor or inhibitor-NC for 24 hours, mRNA of Bcl-2 was detected by real-time PCR (left), and Bcl-2 protein level was analyzed by Western blotting. (E) HCT116 and RKO cells transfected with mir-206 inhibitor with or without Bcl-2 sirna, and Bcl-2 expression were analyzed by Western blotting. Then, the cells were treated with increasing concentrations of 5-FU for 72 hours. Cell proliferation was determined by MTS assay. Results in (B–D) and (E) are expressed as means ± SD of three independent experiments. **P < 0.01.

Abbreviations: 5-FU, 5-fluorouracil; mut, mutation; NC, negative control; PCR, polymerase chain reaction; SD, standard deviation; WT, wild type.

The combination of 5-FU with miR-206 mimic inhibited the growth of tumors more efficiently than the 5-FU single treatment (Figure 6A), which indicates that miR-206 mimic might be a potential method for therapy in combination 5-FU in CRC. The apoptosis of the tumor cell also showed a similar trend (Figure 6B). These results further support the finding that miR-206 contributed to overcome 5-FU resistance in vivo.
Discussion

Usually, drug resistance is caused by the sequential addition of genetic mutations that reroute key signaling pathways promoting cancer cell infiltration and invasive phenotype.\(^{21}\)

Although the cause of drug resistance has been investigated for many years, there are still no remedies to improve drug resistance in clinical outcomes.\(^{21}\) In a previous study, miRNAs have been stratified into mediators that regulate the response of cancer cells to therapeutic cells.\(^{22}\)

It has been shown that the response of patients to chemotherapy is closely related to the functional status of miRNAs.\(^{23,24}\) Although the mechanism of miRNA regulation of drug resistance is unclear, current evidence suggests several roles of miRNAs, including changes in drug targets that affect treatment-induced cell death, regulation of multiple drug resistance (MDR)-related proteins, the use of drug concentrations, and the promotion angiogenesis.\(^{23,25}\) In the current study, we found that miR-206 was downregulated in HCT116/FR and RKO/FR cells compared with the parental cells.

In addition, our results also demonstrated that downregulation of miR-206 promotes resistance to 5-FU in CRC, but miR-206 overexpression reverses 5-FU resistance. Target prediction tools identified miR-206 targeting Bcl-2. It has been studied that Bcl-2 protein plays a key role in various cell processes, including apoptosis, invasion, cell proliferation, and metastases in many types of cancers.\(^{26}\) Thus, the function of miR-206 in targeting Bcl-2 is in agreement with the known
biological effects of miR-206. Actually, other miRNAs are able to confer resistance by targeting Bcl-2 in many other types of cancer. Previous studies have shown that Bcl-2 dysfunction has prognostic significance in several malignancies, such as colorectal cancer.27,28 These reports demonstrated a mechanism in which reduced expression of miR-206 should be associated with a reduced survival of colon cancer patients and also supports the development of miR-206 mimic as a potential target for reversing drug resistance. The overexpression of miR-206 can successfully induce cancer cell sensitivity to 5-FU. In the future, therapeutic strategies can be developed based on the predicted level of miR-206. In addition, miR-206 may potentially bind to 5-FU to prolong the drug susceptibility in CRC.

Conclusion

Our results uncovered a unique role of miR-206 in 5-FU resistance in CRC. The drug resistance is a major challenge for the 5-FU treatment, but the strategy of upregulating the miR-206 may be a potential way to increase the sensitivity of CRC to 5-FU.

Disclosure

The authors report no conflicts of interest in this work.

References