

Flurbiprofen suppresses the inflammation, proliferation, invasion and migration of colorectal cancer cells via COX2

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Abstract. Colorectal cancer is an aggressive disease with a poor prognosis and low survival rate at the advanced stage, therefore new innovative targets are urgently required. Flurbiprofen has been reported to exhibit therapeutic effects in other types of cancer, such as esophageal cancer, breast cancer and colorectal cancer. Therefore, the present study aimed to investigate the function of flurbiprofen in colorectal cancer. SW620 colorectal cancer cells were treated with different concentrations of flurbiprofen to determine the optimum concentration. Subsequently, COX2 expression affected by flurbiprofen was tested using western blotting, reverse transcription-quantitative PCR and immunofluorescence. Enzyme-linked immunosorbent assay was used to determine the levels of tumor necrosis factor- α , interleukin (IL)-6 and IL-1 β . Cell Counting Kit-8, colony formation and flow cytometry assays were used to assess the proliferation and apoptosis of SW620 cells in various groups. Western blotting was performed to investigate the expression of proliferation-, apoptosis- and migration-related proteins after different treatments. Wound healing and Transwell assays were performed to measure the invasion and migration of colorectal cancer cells, respectively. The results demonstrated that flurbiprofen inhibited colorectal cancer cell proliferation. Furthermore, it was identified that flurbiprofen inhibited the expression of COX2. Notably, flurbiprofen suppressed the expression of inflammatory factors by inhibiting COX2. Moreover, flurbiprofen inhibited the proliferation, invasion and migration of colorectal cancer cells by inhibiting COX2. In conclusion, the present study revealed that flurbiprofen inhibited COX2 expression in colorectal cancer, and affected the proliferation, invasion, migration and apoptosis of colorectal cancer cells. These results expand the understanding of the

function of COX2 in colorectal cancer and the effect of flurbiprofen on COX2 expression.

Introduction

Colorectal cancer is a common gastrointestinal malignancy with a high mortality rate, originating from the mucosal epithelium and glands of the large intestine (1). Colorectal cancer is currently the third most common form of cancer and the second most common cause of mortality globally as reported by research conducted in 2019 (2). In the early stage of the disease, the tumor can be effectively treated through minimally invasive surgery and radical surgery, which avoids the inconvenience of radiotherapy and chemotherapy, or even by implantation of an artificial sphincter (3,4). However, due to the occurrence of distant metastasis, such as lymph node metastasis and liver metastasis, the treatment options for numerous patients are poor, resulting in high recurrence rates and low survival rates, which markedly affects the life quality of patients (5,6). The exact pathogenesis of colorectal cancer remains unclear; however, a large number of studies have reported that the pathogenesis of colorectal cancer involves multiple genes and is a multi-stage process (7,8). Therefore, it is necessary to search for treatment methods and drugs for colorectal cancer, which may help to improve the therapeutic effect of colorectal cancer treatments, prolong the survival period and improve the quality of life (7,9,10).

Flurbiprofen is a commonly used non-steroidal anti-inflammatory drug; it is a non-selective cyclooxygenase (COX) inhibitor and has a high affinity for immune organs (11,12). Flurbiprofen is commonly used to treat inflammatory diseases, including osteoarthritis and rheumatoid arthritis, as well as pain control during surgery (13). Flurbiprofen has been shown to significantly reduce postoperative pain scores and improve post-chemotherapy immunity in patients with cancer (14,15). Flurbiprofen therapy can promote the natural circulation of dendritic cells in patients with esophageal squamous cell carcinoma and inhibit the rise of IL-10 and TGF- β (16). In addition, flurbiprofen increases the postoperative IL-8 expression in patients with esophageal cancer, reduces the pain score and improves the oxidation level (17). Furthermore, flurbiprofen inhibits the activation and aggregation of leukocytes in patients with a tumor and reduces the formation of venous

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embolism (18). Additionally, flurbiprofen inhibits the increase of TNF- α and IL-1 β in patients with breast cancer (19,20). Toxicity studies on breast and lung cancer cells have demonstrated that anti-inflammation can damage the integrity and permeability of cell membrane structures, thereby inhibiting the growth and proliferation of cells (21,22). Flurbiprofen has been reported to reduce IL-6 in patients with colorectal cancer, increase IL-2 secretion and reduce surgical complications (23). However, to the best of our knowledge, the direct effect of flurbiprofen on colorectal cancer and its mechanism remain unknown. Therefore, the present study investigated the effect of flurbiprofen on the proliferation and migration of colorectal cancer cells by regulating COX2.

At present, there are three reported COX isozymes, COX1, COX2 and COX3, which are known to regulate fever and pain (24). COX2 is an inducible enzyme that is mainly distributed on the cell nuclear membrane (25). COX2 is predominantly expressed in the kidney and brain at general physiological conditions, and remains silent in other tissues (26). The expression of COX2 changes only after various stimulations (27). It has been reported that COX2 is highly expressed in colorectal cancer (28); however, to the best of our knowledge, the function of COX2 in colorectal cancer and the mechanism of its function remain unclear. Therefore, further studies are required to clarify the role of COX2 in colorectal cancer. Additionally, inhibitors of COX2 should be investigated to potentially provide a new therapeutic target for colorectal cancer.

Materials and methods

Cell culture. Human SW620 colorectal cancer cells were cultured in DMEM (Hyclone; GE Healthcare Life Sciences) containing 10% FBS (Gibco; Thermo Fisher Scientific Inc.) and 1% penicillin/streptomycin (Invitrogen; Thermo Fisher Scientific Inc.), in a cell culture chamber at 37°C with 5% CO₂. Following counting and inoculation in plates, the cells grew to the required cell density of 80-90% and were treated accordingly. For determining the optimal concentration of flurbiprofen for treating SW620 cells; 2, 4, 10 and 20 nM of flurbiprofen were used for 12, 24 or 48 h at 37°C. Cells treated with 0.1% DMSO were the control group.

Cell transfection. A total of 3x10⁵ SW620 cells (cell density, 5x10⁴ cells/well) were plated in a 6-well plate, and transfection with an overexpression plasmid was performed when the cells grew to 70-80% density. During transfection, the original cell culture medium was removed and the cells were cultured in medium without serum and antibiotics. The overexpression plasmid pcDNA-COX2 (20 nM) and negative control pcDNA (1 μ g) from Shanghai Genechem Co., Ltd., were diluted with 125 μ l Opti-MEM medium (Gibco; Thermo Fisher Scientific Inc.), and Lipofectamine 2000 (Invitrogen; Thermo Fisher Scientific, Inc.) was diluted with 125 μ l Opti-MEM medium at room temperature for 5 min. These were then mixed for 15 min before addition to the cell culture medium. The liquid was changed after 4-6 h, and the cells were collected for further experiments after 24-48 h.

RNA extraction and reverse transcription-quantitative PCR (RT-qPCR). RNA was extracted from the treated cells using

TRIzol (Invitrogen; Thermo Fisher Scientific Inc.). RT was performed with PrimeScript™ RT Reagent kit with gDNA Eraser (Takara Bio, Inc.) at 37°C for 15 min and at 98°C for 5 min. qPCR was performed using SYBR reagent (Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer's protocol. Thermocycling conditions were as follows: 95°C for 3 min, 40 cycles of 95°C for 5 sec followed by 60°C for 30 sec. GAPDH was used as the internal reference and quantification was performed using the 2^{- $\Delta\Delta$ C_q} method (29). The primer sequences used were: COX2 forward, 5'-ATCATAAGCGAG GGCCAGCT-3' and reverse, 5'-AAGGCGCAGTTTACGCTG TC-3'; GAPDH forward, 5'-GACAGTCAGCCGCATCTT CT-3' and reverse, 5'-GCGCCCAATACGACCAAATC-3'.

Cell Counting Kit-8 (CCK-8) assay. SW620 cells were seeded into a 96-well plate 24 h post-transfection at a density of 3,000 cells/well. CCK-8 was used to assess the cell viability at 12, 24 and 48 h. A total of 10 μ l CCK-8 solution (Dojindo Molecular Technologies) was added to each well and the cells were cultured for a further 4 h according to the manufacturer's protocol. An enzyme-linked immune detector was used to measure the absorbance value of each well at a wavelength of 450 nm.

Western blot analysis. The treated cells were lysed with RIPA lysis buffer (cat. no. 9806; Cell Signaling Technology, Inc.), placed in a 1.5-ml EP tube, and centrifuged at 4°C and 12,000 x g for 10 min. The supernatant was isolated and the protein concentration was measured using a BCA Protein assay kit (cat. no. 23227; Thermo Fisher Scientific, Inc.). Next, the protein was denatured at 100°C for 10 min with 10% SDS-polyacrylamide gel electrophoresis, and 25 μ g protein sample per lane was separated. Following membrane transfer, the PVDF membrane was incubated with non-fat milk for 30 min at room temperature, incubated with primary antibody for 4°C overnight, and washed with TBS containing 0.05% Tween-20 (TBST) 3 times for 10 min each time. Primary antibodies were all purchased from Abcam and were against COX2 (1:1,000; cat. no. ab179800), cyclin E (1:1,000; cat. no. ab33911), CDK2 (1:1,000; cat. no. ab32147), p21 (1:1,000; cat. no. ab109520), Bcl-2 (1:5,000; cat. no. ab196495), Bax (1:1,000; cat. no. ab199677), cleaved caspase-3 (1:500; cat. no. ab49822), caspase-3 (1:5,000; cat. no. ab32351), MMP2 (1:1,000; cat. no. ab181286), MMP9 (1:1,000; cat. no. ab137867) and GAPDH (1:3,000; cat. no. ab9485). The membrane was then incubated with horseradish peroxidase-conjugated secondary antibody goat anti-rabbit IgG (1:10,000; cat. no. ab205718; Abcam) at room temperature for 2 h and washed with 1X TBST three times for 10 min each time. Enhanced chemiluminescence reagents (Beyotime Institute of Biotechnology) was added for visualization and next quantified with Quantity One version 4.1 software (Bio-Rad Laboratories Inc.).

Immunofluorescence assay. In the immunofluorescence assay, changes in COX2 expression were examined. SW620 cells were subjected to fixation with 4% paraformaldehyde for 15 min at room temperature and permeabilization with 0.5% Triton X-100. Subsequently, 5% normal goat serum (PH0424; Scientific Phygene) was used to block cells for 20 min at 37°C, followed by incubation overnight at 4°C with anti-COX2 (1:1,000; cat. no. 4842; Cell Signaling Technology, Inc.). Subsequently,

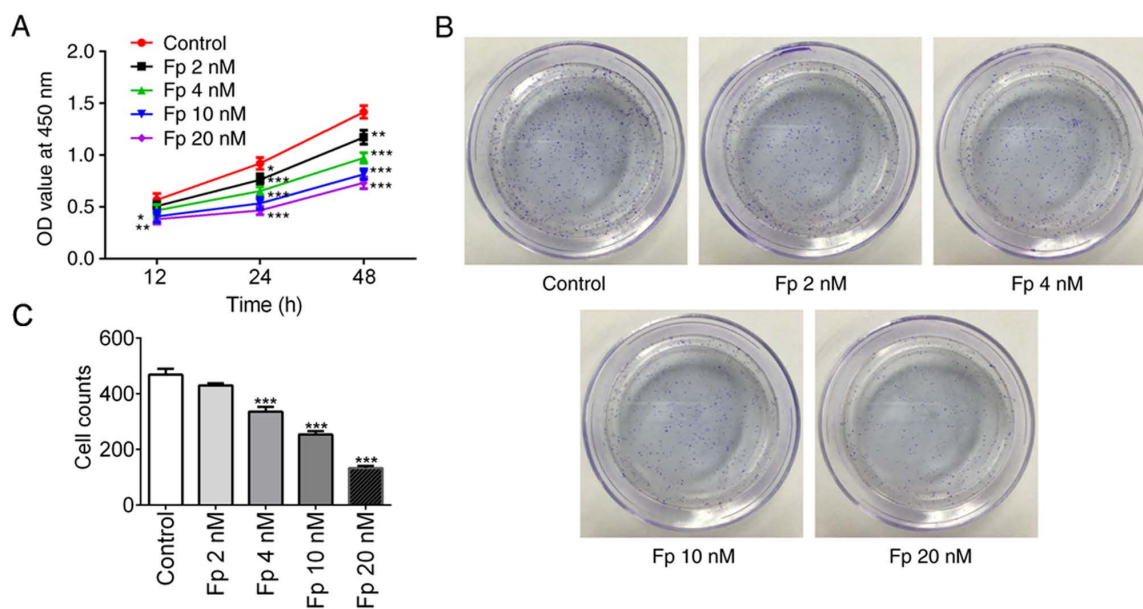


Figure 1. Fp inhibits colorectal cancer cell proliferation. (A) Cell viability of SW620 cells treated with Fp were evaluated by Cell Counting Kit-8 assay. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ vs. control. (B and C) Colony formation ability was explored by colony formation assay. *** $P < 0.001$ vs. control. Fp, flurbiprofen; OD, optical density.

the cells were probed with an Alexa Fluor 488 labeled-goat anti-rabbit IgG (1:500; cat. no. A0423; Beyotime Institute of Biotechnology) for nearly 2 h at room temperature. Nuclei were counterstained using DAPI (Invitrogen; Thermo Fisher Scientific Inc.) at room temperature for 5 min, and images were captured using a fluorescence microscope (magnification, $\times 20$).

ELISA. Human IL-6 Quantikine (cat. no. D6050), human IL-1 β Quantikine (cat. no. DLB50) and human TNF- α Quantikine ELISA kits (cat. no. DTA00D) (all R&D Systems, Inc.) were used to measure the levels of IL-6, IL-1 β and TNF- α in cell supernatants of SW620 according to the manufacturer's protocols.

Colony formation assay. A total of 200 SW620 cells/well were plated in 12-well plates and cultured in DMEM with 10% FBS for 14 days. PBS was used to wash the cells twice, and then the cells were fixed with methanol for ~ 10 min at room temperature. Following fixation, the cells were washed twice with PBS, stained with crystal violet for 30 min at room temperature, and then washed with double-distilled water. The number of the colonies ≥ 50 was counted using the naked eye.

Flow cytometry. For flow cytometric analysis, FITC Annexin V Apoptosis Detection kit (Guangzhou RiboBio Co., Ltd.) was used. After 30 min of culture in the dark, SW620 cells (1×10^5 cells/well) in 200 μ l of binding buffer were mixed with 5 μ l of propidium iodide (PI) for 15 min at 4°C and apoptosis was detected using a BD Accuri C6 Plus personal flow cytometer (Becton, Dickinson and Company) and analyzed using FlowJo version 10 software (Tree Star, Inc.).

Wound healing assay. Cell migratory ability was analyzed using a wound healing assay. Cells in control, Fp 10 nM, Fp 10 nM + pcDNA and Fp 10 nM + pcDNA-COX2 groups were cultured and then a 5-mm horizontal scratch was made

using a pipette tip. Subsequently, cells were incubated for 1 day at 37°C in a serum-free medium, and the gap width was assessed and recorded at 0 and 24 h using a light microscope (Leica Microsystems) with magnification, $\times 100$. Relative migration rate was calculated as the percentage of wound healing.

Transwell invasion assay. The cell invasion assay was performed using 24-well Transwell inserts (Corning, Inc.) pre-coated with Matrigel (BD Biosciences at room temperature for 15-30 min. Following cell suspension in 200 μ l serum-free DMEM, cells (1×10^5) were added to the upper chamber and medium supplemented with 10% FBS was placed in the bottom chamber. Subsequently, cells on the upper surface were removed, and cells on the bottom surface were fixed and stained with crystal violet at room temperature for 20 min. Cell numbers in five random fields were calculated using a light microscope (Nikon Corporation) with magnification, $\times 100$.

Statistical analysis. Data obtained from triplicated independent experiments were presented as mean \pm standard deviation (SD). Two-sided unpaired Student's t-test was used to assess the differences between two groups. For multiple groups, one-way ANOVA followed by Tukey's post hoc test was used. GraphPad Prism 5.0 software (GraphPad Software, Inc.) and SPSS 13.0 software (SPSS, Inc.) were employed to analyze data. $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Flurbiprofen inhibits colorectal cancer cell proliferation. To evaluate the impact of flurbiprofen on colorectal cancer, SW620 cells were treated with different concentrations of flurbiprofen (2, 4, 10 and 20 nM) and the cell proliferation was assessed by a CCK-8 assay at 12, 24 and 48 h, respectively. As expected, flurbiprofen significantly decreased SW620 cell

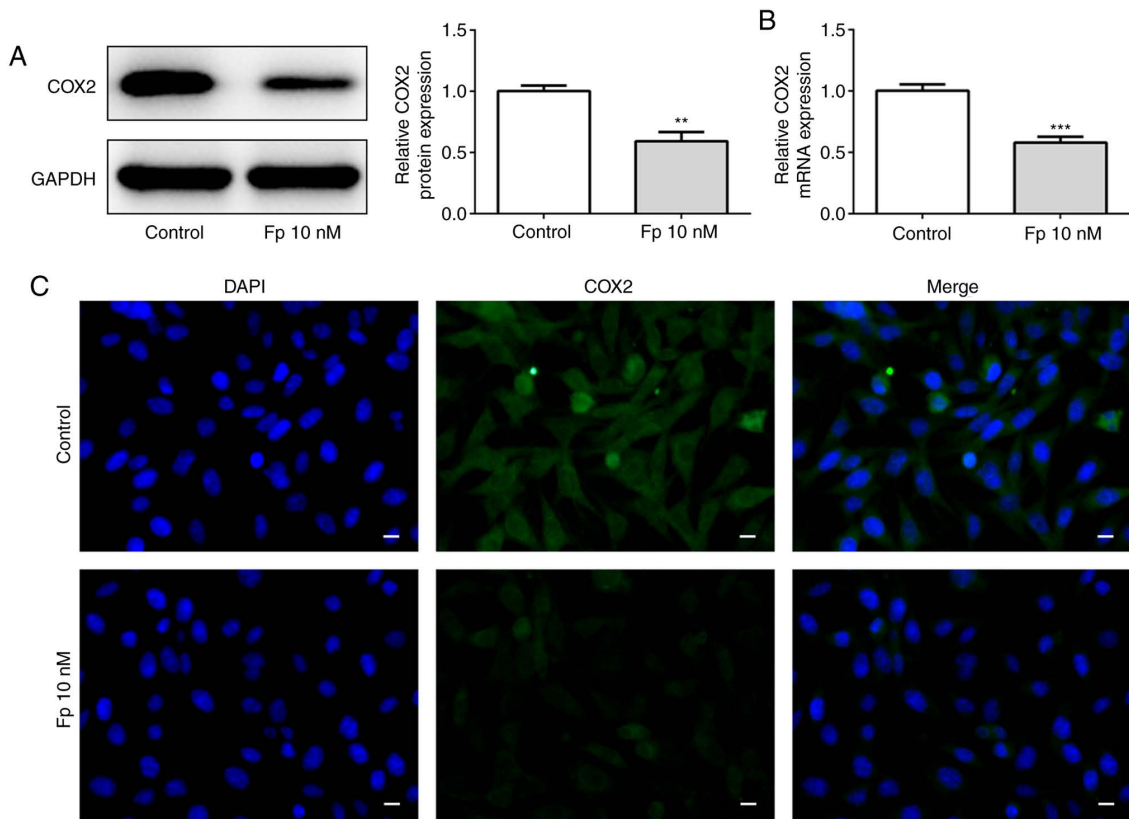


Figure 2. Fp inhibits COX2 expression. COX2 (A) protein and (B) mRNA expression in control and Fp-treated SW620 cells was analyzed via western blot analysis and reverse transcription-quantitative PCR. (C) An immunofluorescence assay detected the changes of COX2 expression in SW620 cells in response to Fp. Scale bar, 50 μ m. ** $P < 0.01$, *** $P < 0.001$ vs. control. Fp, flurbiprofen; COX2, cyclooxygenase 2.

proliferation in a concentration- and time-dependent manner (Fig. 1A). In the colony formation assay, it was observed that with an increasing concentration of flurbiprofen, the number of colonies gradually decreased (Fig. 1B-C). When the cells were treated with 10 nM flurbiprofen, the cell viability (at 24 h) and colony formation ability were reduced to ~50% of the levels observed in the control group, therefore 10 nM flurbiprofen was selected for the following experiments. These data suggest that flurbiprofen inhibited colorectal cancer cell proliferation.

Flurbiprofen decreases COX2 expression. It has been reported that flurbiprofen is a non-selective COX inhibitor (30). Therefore, the present study aimed to evaluate whether flurbiprofen has an effect on COX2 in SW620 colorectal cancer cells. Western blotting was used to measure the protein level of COX2 after cells were treated with 10 nM flurbiprofen (Fig. 2A). In addition, RT-qPCR was used to assess the mRNA level of COX2 following treatment with 10 nM flurbiprofen (Fig. 2B). As expected, the protein and mRNA levels of COX2 were significantly decreased. To confirm this result, an immunofluorescence assay with SW620 cells treated with 10 nM flurbiprofen or DMSO was performed (Fig. 2C). The fluorescence intensity of COX2 was markedly reduced when treated with 10 nM flurbiprofen. The data confirmed that flurbiprofen decreased COX2 expression.

Flurbiprofen inhibits the expression of inflammatory factors by inhibiting COX2. COX2 was overexpressed in SW620 cells, and western blotting and RT-qPCR were used to assess

transfection efficiency (Fig. 3A and B). It has been reported that COX2 is a highly expressed inflammatory factor (31). Thus, the present study investigated whether flurbiprofen inhibits the expression of inflammatory factors by inhibiting COX2. ELISA was used to measure the levels of TNF- α , IL-1 β and IL-6. As presented in the Fig. 3C, the levels of TNF- α , IL-1 β and IL-6 were significantly downregulated in cells treated with flurbiprofen compared with control cells. However, when cells were co-treated with flurbiprofen and COX2 overexpression plasmid, the levels of TNF- α , IL-1 β and IL-6 were significantly higher compared with in cells treated with flurbiprofen alone. These results suggested that flurbiprofen inhibited the expression of inflammatory factors via repression of COX2.

Flurbiprofen inhibits the proliferation of colorectal cancer cells by inhibiting COX2. To determine whether flurbiprofen has an effect on colorectal cancer cell proliferation, a colony formation assay was performed, and it was identified that transfection of COX2 overexpression plasmid reversed the suppressed cell colony formation ability by flurbiprofen alone (Fig. 4A-B). Subsequently, western blotting was used to evaluate the changes of proliferation-associated proteins, including cyclin E, cyclin-dependent kinase 2 (CDK2) and p21. The results demonstrated that in cells treated with flurbiprofen, the expression levels of cyclin E and CDK2 were significantly decreased, and the expression of p21 was significantly increased. However, in cells co-treated with flurbiprofen and COX2 overexpression plasmid, the flurbiprofen-induced expression level changes of cyclin E, CDK2 and p21 were significantly reversed (Fig. 4C).

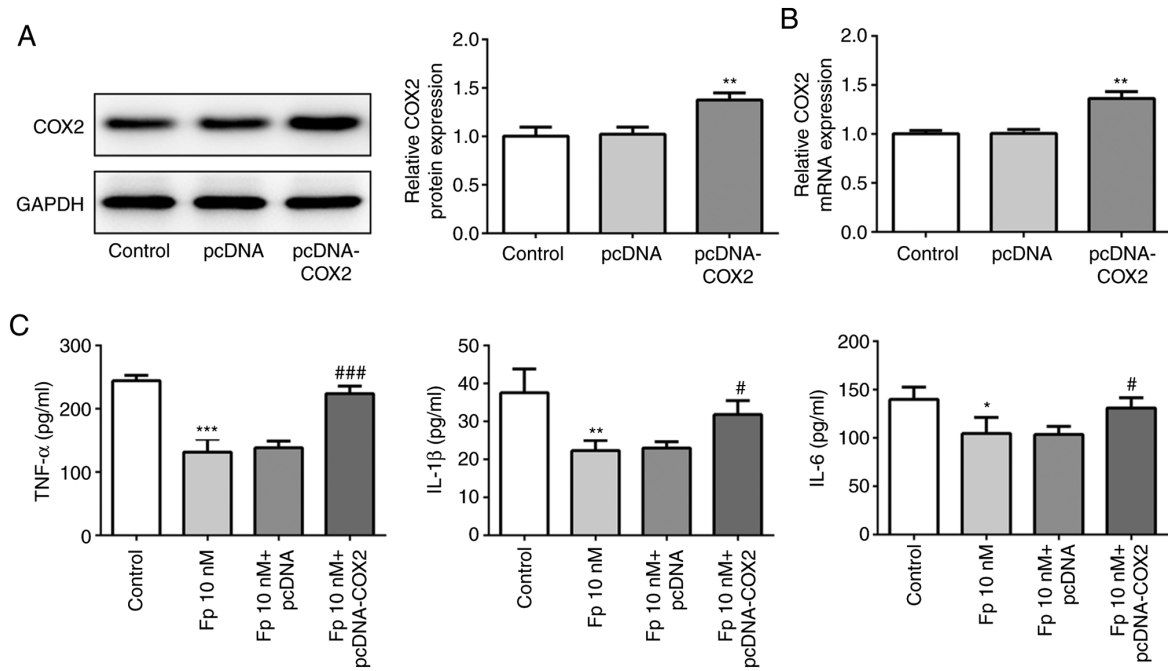


Figure 3. Fp restrains inflammation. (A) Western blot analysis and (B) reverse transcription-quantitative PCR were used to measure COX2 protein and mRNA expression levels. **P<0.01 vs. pcDNA. (C) ELISA was used to estimate the levels of the inflammatory factors TNF- α , IL-1 β and IL-6 following COX2 overexpression and/or treatment with Fp. *P<0.05, **P<0.01, ***P<0.001 vs. control. #P<0.05, ###P<0.001 vs. Fp 10 nM + pcDNA. Fp, flurbiprofen; COX2, cyclooxygenase 2.

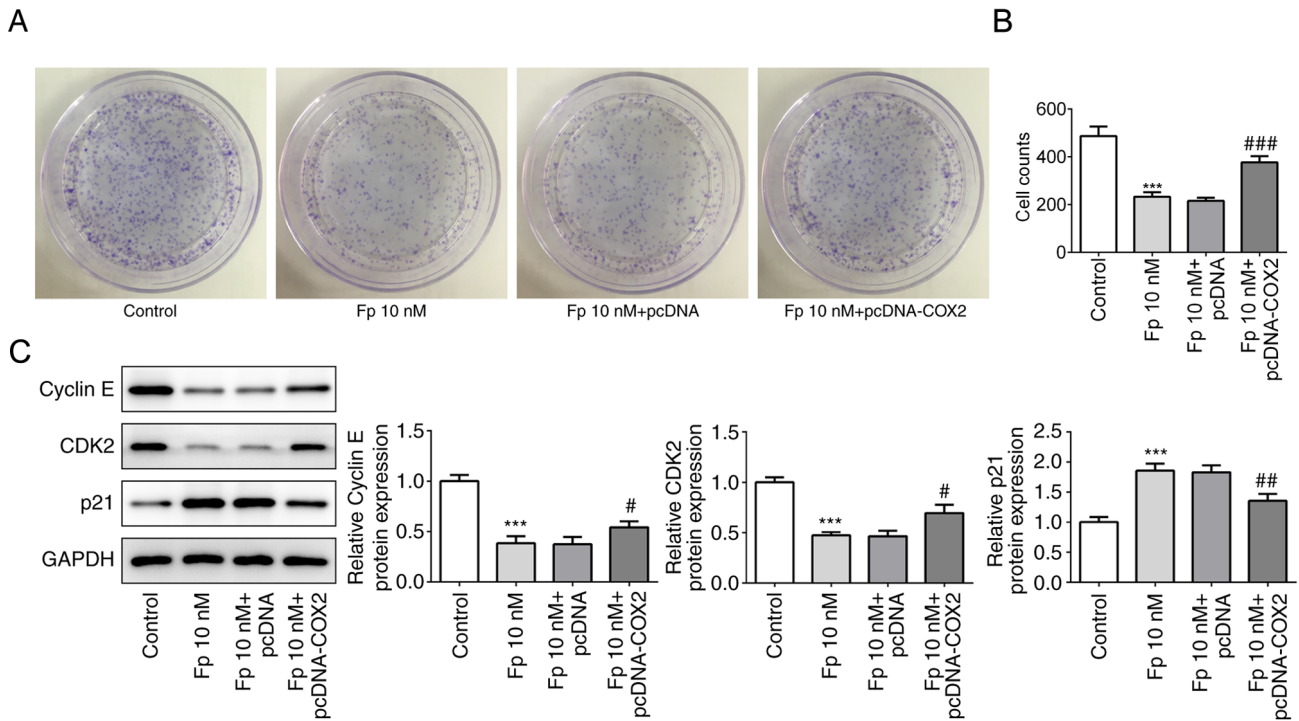


Figure 4. Fp suppresses cell proliferation. (A and B) Colony formation assay assessed the colony formation of SW620 cells in the four treatment groups. ***P<0.001 vs. control; ###P<0.001 vs. Fp 10 nM + pcDNA. (C) Apoptosis-related proteins in the different groups were examined via western blotting. ***P<0.001 vs. control. #P<0.05, ##P<0.01 vs. Fp 10 nM + pcDNA. Fp, flurbiprofen; CDK2, cyclin-dependent kinase 2; COX2, cyclooxygenase 2.

These results suggested that flurbiprofen could inhibit the proliferation of colorectal cancer cells by inhibiting COX2.

Flurbiprofen promotes the apoptosis of colorectal cancer cells by inhibiting COX2. It has been reported that COX2 can inhibit

apoptosis (32), therefore the present study investigated whether flurbiprofen has an effect on colorectal cancer cell apoptosis. The results indicated that the apoptosis of cells treated with flurbiprofen was significantly increased. However, in cells co-treated with flurbiprofen and COX2 overexpression vector,

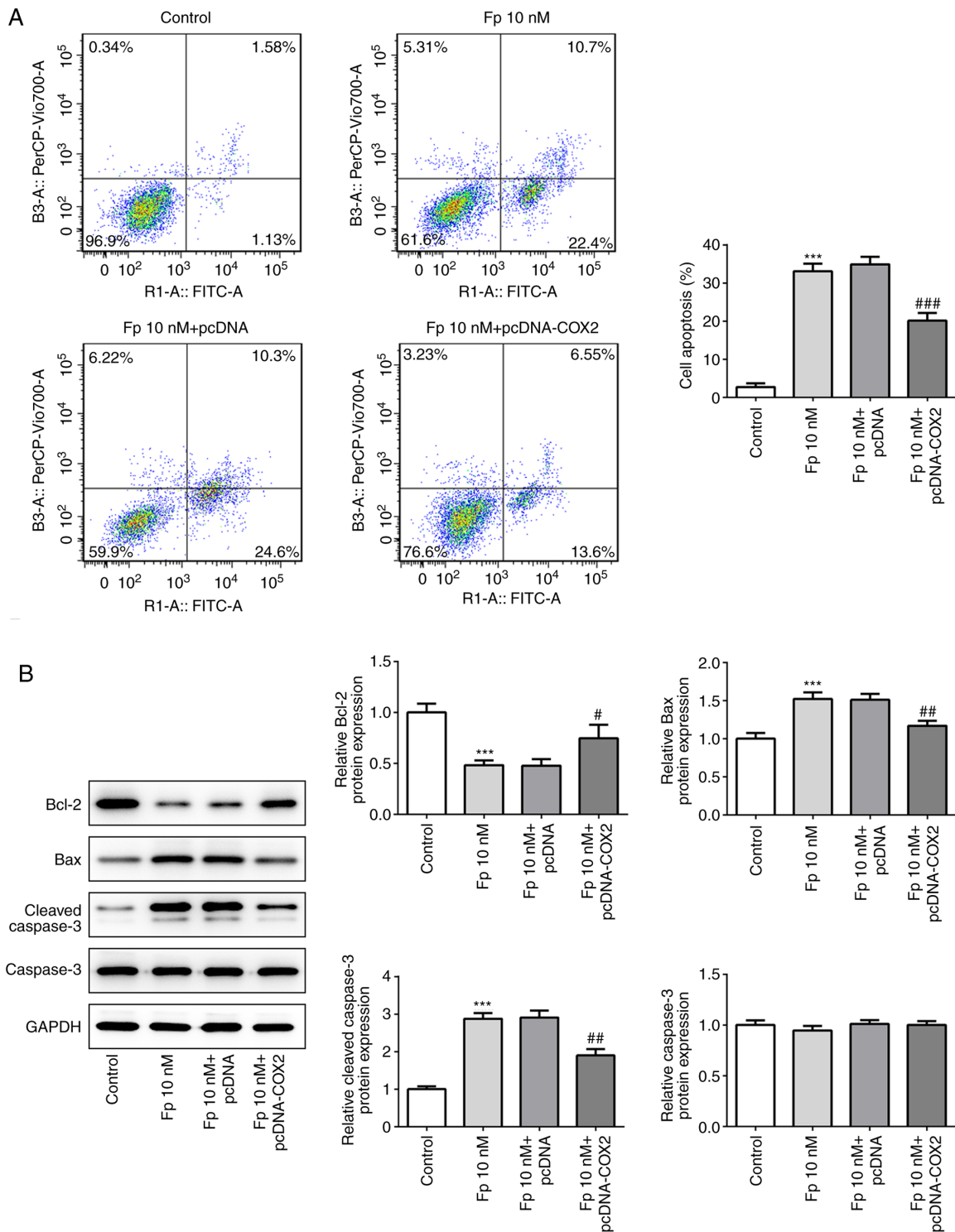


Figure 5. Fp induces cell apoptosis. (A) Cell apoptosis of SW620 cells was examined by flow cytometry analysis following COX2 overexpression and/or treatment with Fp. (B) Apoptosis-associated proteins were measured via western blot analysis in SW620 cells. *** $P < 0.001$ vs. control. # $P < 0.05$, ## $P < 0.01$, ### $P < 0.001$ vs. Fp 10 nM + pcDNA. Fp, flurbiprofen; COX2, cyclooxygenase 2.

the level of apoptosis was significantly reduced compared with in cells treated with flurbiprofen alone (Fig. 5A). Furthermore, the expression levels of apoptosis-associated proteins, including Bcl2, Bax, cleaved-caspase-3 and total caspase-3, were evaluated. The results demonstrated that flurbiprofen significantly decreased the expression of Bcl2 and significantly increased the expression of Bax and cleaved-caspase3, with no effect

on total caspase-3 (Fig. 5B). However, following co-treatment with flurbiprofen and COX2 overexpression vector, the flurbiprofen-induced changes of Bcl2, Bax and cleaved-caspase-3 expression levels were significantly reversed.

Flurbiprofen inhibits invasion and migration of colorectal cancer cells by inhibiting COX2. The aforementioned data

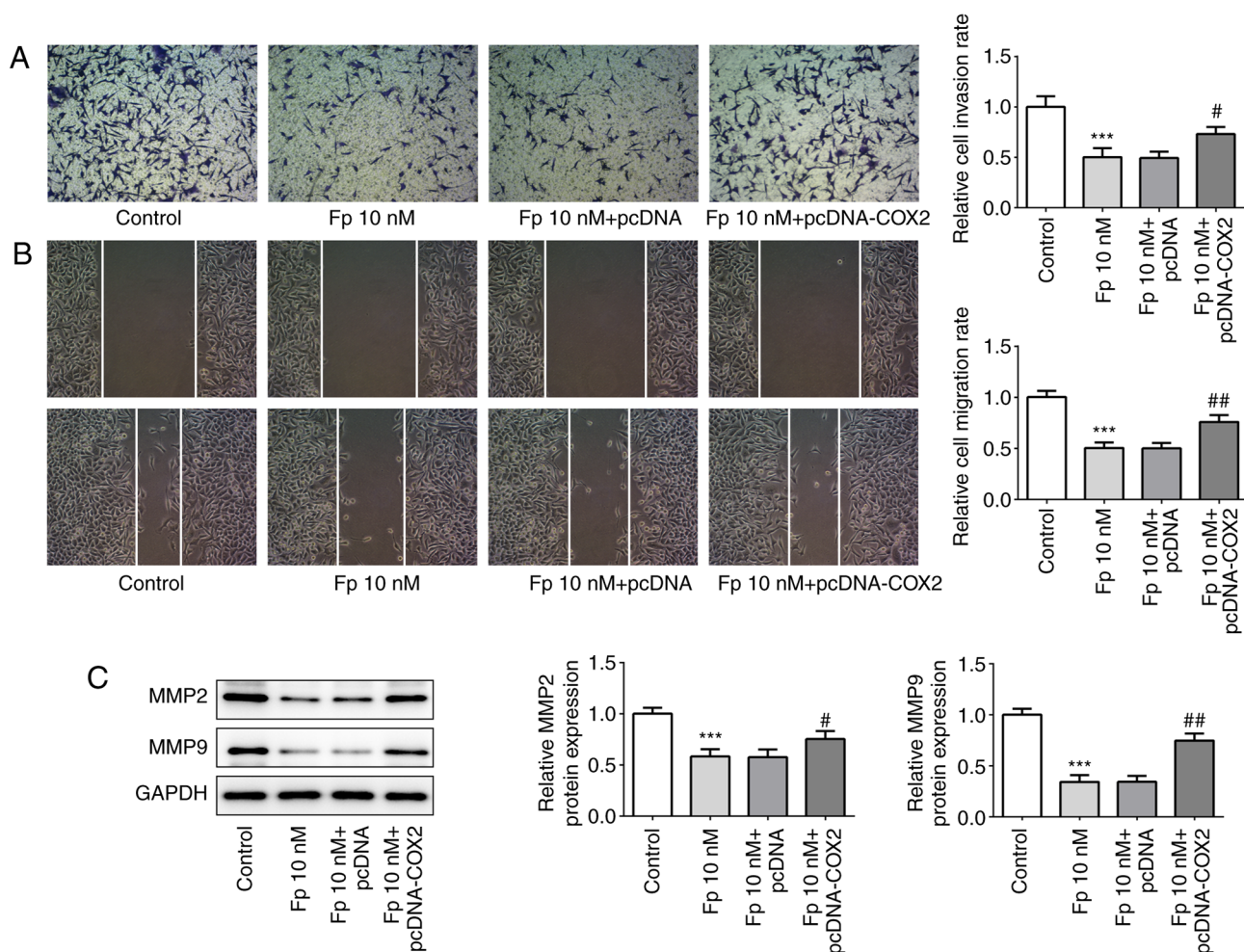


Figure 6. Fp suppresses cell invasion and migration. (A) Transwell (magnification, x100) and (B) wound healing (magnification, x100) assays were performed to assess cell invasion and migration following COX2 overexpression and/or treatment with Fp. (C) MMP2 and MMP9 protein expression levels were detected via western blotting. *** $P < 0.001$ vs. control. # $P < 0.05$, ## $P < 0.01$ vs. Fp 10 nM + pcDNA. Fp, flurbiprofen; COX2, cyclooxygenase 2; MMP, matrix metalloproteinase.

demonstrated the effects of flurbiprofen on cell proliferation and apoptosis in colorectal cancer, therefore the effects of flurbiprofen on colorectal cancer cell invasion and migration were then examined. Transwell and wound healing assays with SW620 cells were performed (Fig. 6A and B). When the cells were treated with flurbiprofen, the invasion and migration abilities were significantly reduced compared with the control cells. However, when cells were co-treated with flurbiprofen and COX2 overexpression plasmid, the invasion and migration abilities of cells were significantly reversed compared with the effect of flurbiprofen alone. Since matrix metalloproteinase (MMP)2 and MMP9 are key proteins associated with invasion and migration (33), whether flurbiprofen effected MMP2 and MMP9 expression was then investigated. The western blotting results demonstrated that the expression levels of MMP2 and MMP9 were significantly downregulated by flurbiprofen. However, when cells were co-treated with flurbiprofen and COX2 overexpression plasmid, the expression levels of MMP2 and MMP9 were significantly increased compared with in cells treated with flurbiprofen alone (Fig. 6C). In summary, flurbiprofen inhibited the invasion and migration of colorectal cancer cells by inhibiting COX2.

Discussion

Colorectal cancer is one of the most common malignancies of the digestive system (10). The etiology of colorectal cancer includes high nutrition, low fiber diet, genetic factors, some chronic intestinal diseases with intestinal mucosal hyperplasia, and a molecular biological basis of gradual canceration of colorectal mucosal epithelium (34). In recent years, the incidence of colorectal cancer has rapidly increased due to people's high protein diet and a lack of physical exercise, as well as exposure to various pathogenic factors (8). The onset process of colorectal cancer is relatively unidentified and generally limited of specific clinical manifestations (35). The majority of patients at the early stage do not have obvious clinical symptoms, and patients at end-stage or with distant metastasis lose the opportunity of surgery (36). Certain patients with colorectal cancer can receive chemotherapy, although chemotherapy is not always effective and is prone to resistance (37). All in all, the recurrence rate is high and the majority of patients cannot tolerate. Therefore, finding new treatments is important.

The present data suggested that flurbiprofen inhibited colorectal cancer cell proliferation. Flurbiprofen has several

advantages, including specific targeting, a quick onset, long duration and few side effects; therefore, it is widely used in the treatment of advanced analgesia, postoperative analgesia and cancer pain (12). Flurbiprofen is a non-steroidal intravenous drug that can inhibit COX1 and COX2 non-selectively (30). It has been reported that flurbiprofen can inhibit the growth of cervical cancer, and COX2 is also involved in this process (38). The current study demonstrated that flurbiprofen could inhibit the proliferation of colorectal cancer cells, and COX2 was involved in this process. As a non-selective inhibitor of COX2, the anticancer mechanism of flurbiprofen may be associated with the following aspects (39). First, specific targeting is a characteristic of flurbiprofen; it can inhibit tumor cell growth, but has little effect on normal cells (40). Secondly, flurbiprofen inhibits the expression of COX2, thus decreasing prostaglandin E2 and thromboxane A2, and even reducing vascular endothelial growth factor and fibroblast growth factor to inhibit tumor angiogenesis (27). Thirdly, it promotes the apoptosis of tumor cells (41); and finally, it enhances the effect of tumor immunotherapy (42).

In conclusion, the present study demonstrated that flurbiprofen decreased COX2 expression in colorectal cancer, and that flurbiprofen also inhibited the inflammation response, proliferation, invasion and migration, and promoted the apoptosis of colorectal cancer cells. Therefore, flurbiprofen may be a useful therapeutic method for colorectal cancer. However, the absence of the pcDNA-COX2 only group (i.e. without 10 nM flurbiprofen) in several assays is a limitation of the current study, resulting in a lack of comparative explanation for the function of flurbiprofen in colorectal cancer.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

XW wrote the majority of the manuscript. XY and YZ performed the experiments. XW and FJ analyzed and interpreted the data. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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