

Aspirin inhibits cholangiocarcinoma cell proliferation via cell cycle arrest *in vitro* and *in vivo*

TINGTING SHI¹, JIAN GONG², KOJI FUJITA¹, NORIKO NISHIYAMA¹, HISAKAZU IWAMA³, SHI LIU², MAI NAKAHARA¹, HIROHITO YONEYAMA¹, ASAHIRO MORISHITA¹, TAKAKO NONURA¹, HIDEKI KOBARA¹, KEIICHI OKANO⁴, YASUYUKI SUZUKI⁴ and TSUTOMU MASAKI¹

¹Department of Gastroenterology and Neurology, Faculty of Medicine, Kagawa University, Kida, Kagawa 761-0793, Japan; ²Department of Gastroenterology, The First Affiliated Hospital of Dalian Medical University, Dalian, Liaoning 116011, P.R. China; ³Life Science Research Center, Kagawa University; ⁴Department of Digestive Surgery, Faculty of Medicine, Kagawa University, Kida, Kagawa 761-0793, Japan

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Abstract. Cholangiocarcinoma is the most common biliary duct malignancy and the second most common primary liver cancer, accounting for 10-20% of hepatic malignancies. With high mortality and poor prognosis, the 5-year survival rate of cholangiocarcinoma is only 10%. A previous study demonstrated a significant association between aspirin use and a decreased risk of cholangiocarcinoma. However, the effect of aspirin on cholangiocarcinoma remains unknown. Therefore, the aim of the present study was to investigate the effects of aspirin on cholangiocarcinoma *in vitro* and *in vivo*. Three cholangiocarcinoma cell lines were used to analyze the effect of aspirin on cell proliferation, cell cycle progression, apoptosis, and the regulation of microRNAs. MicroRNAs are known to regulate the development and progression of various types of cancer. An HuCCT-1 xenograft model was used for the *in vivo* study. It was determined that aspirin inhibited the proliferation of human cholangiocarcinoma cells (except TKKK cells). Aspirin induced cell cycle arrest in the G₀/G₁ phase and regulated cell-cycle related proteins in cholangiocarcinoma cells

(HuCCT-1 cells) but did not induce apoptosis. The expression of miR-340-5p was significantly upregulated after treatment, and overexpression of miR-340-5p inhibited the proliferation of HuCCT-1 cells and decreased the levels of cyclin D1. TKKK cells had low miR-340-5p expression, which may explain why aspirin had no effect on their proliferation. *In vivo*, aspirin reduced the growth of xenografted tumors. In conclusion, the present study indicated that aspirin partially inhibited cholangiocarcinoma cell proliferation and tumor growth by inducing G₀/G₁ phase cell cycle arrest, potentially through the miR-340-5p/cyclin D1 axis.

Introduction

Cholangiocarcinoma (CCA) is the most common biliary duct malignancy and the second most common primary liver cancer, accounting for 10-20% of all primary hepatic malignancies (1). CCA arises from bile duct epithelial cells, and there are three subtypes based on anatomic location: Intrahepatic (iCCA), perihilar (pCCA), and distal (dCCA). The etiology of CCA includes lithiasis (2), primary sclerosing cholangitis (3), parasitic infection (4), congenital abnormalities (5), chronic liver disease (6), cirrhosis (6), metabolic abnormalities (7), and lifestyle (7). The incidence of iCCA appears to be increasing (8); the rates of CCA in North America, Japan, and Australia have been rising over the past two decades (9). Although surgery is the preferred treatment, the 5-year postoperative survival rate is markedly low. Chemotherapy can be used for inoperable cases; however, the highly desmoplastic nature, rich tumor microenvironment, and profound genetic heterogeneity all contribute to iCCA therapeutic resistance (9).

The nonsteroidal drug aspirin is an anti-inflammatory and anticoagulation agent used to prevent and reduce the risk of cardiovascular events. According to some clinical analyses, long-term use is also associated with a reduction in cancer risk, including colon (10), breast (11), and hepatocellular carcinoma (12). Aspirin use also has a significant inverse association with the development of all three CCA subtypes and an approximately 3-fold reduction in CCA risk (13). Numerous anticancer mechanisms for aspirin have been

Correspondence to: Dr Tsutomu Masaki, Department of Gastroenterology and Neurology, Faculty of Medicine, Kagawa University, 1750-1 Ikenobe, Miki, Kida, Kagawa 761-0793, Japan
E-mail: tmasaki@med.kagawa-u.ac.jp

Abbreviations: CCA, cholangiocarcinoma; NSAID, non-steroidal anti-inflammatory drug; FCM, flow cytometer; AMPK, AMP-activated protein kinase; miRNA, microRNA; NF-κB, nuclear factor κ-light-chain-enhancer of activated B cells; JNK, c-Jun-N-terminal kinase; GSK-3, glycogen synthase kinase-3; mTOR, mammalian target of rapamycin; STAT3, signal transducers and activators of transcription 3; p-Bcl-2, phosphorylated B-cell lymphoma-2; ELISA, enzyme-linked immunosorbent assay; Cdk, cyclin-dependent kinase

Key words: aspirin, cell cycle, cholangiocarcinoma, *in vivo*, microRNAs

identified, such as inhibition of cyclooxygenase (14), activating key molecular targets in the AMPK, mTOR, STAT3, and NF- κ B pathways (15), decreasing the levels of reactive oxygen species and glucose consumption (16), inducing autophagy via JNK/p-Bcl2/Beclin-1, AMPK/mTOR, and GSK-3 signaling (17), inducing apoptosis and mitochondrial dysfunction by increasing oxidative stress (18), and changing the tumor microenvironment by affecting platelets (19,20). However, the effect of aspirin on CCA remains unknown. Elucidation of the mechanism of aspirin in CCA could contribute to the development of new therapeutic agents in the future.

The purpose of the present study was to determine the antitumor effects of aspirin in CCA and identify the key molecular targets and microRNAs (miRNAs) associated with this effect.

Materials and methods

Chemicals. Aspirin was obtained from Wako Pure Chemical Industries, Ltd. The prepared solution was diluted in cell culture medium as per the requirement of cells and fresh pH 7.2 to 7.5, within the range suitable for cell growth was used.

Cell lines and cell culture. Human CCA cell lines (HuCCT-1, RBE, and TKKK) were obtained from the Japanese Research Resources Bank. HuCCT-1 and RBE cells were grown in RPMI-1640 media (Gibco-Invitrogen; Thermo Fisher Scientific, Inc.) supplemented with 10% fetal bovine serum (FBS) (product no. 557-30355; FUJIFILM Wako Pure Chemical Industries, Ltd.) and penicillin/streptomycin (100 mg/l; Invitrogen; Thermo Fisher Scientific, Inc.). TKKK cells were maintained in DMEM (Gibco-Invitrogen; Thermo Fisher Scientific, Inc.) supplemented with 10% FBS and penicillin/streptomycin. All cell lines were grown in a humidified incubator at 5% CO₂ and 37°C.

Cell viability assay. Cell viability assays were performed using the Cell Counting Kit-8 (Dojindo Molecular Technologies, Inc.) according to the manufacturer's instructions. Briefly, cells (5,000 cells in 100 μ l/well) were seeded into 96-well plates and allowed to adhere, followed by treatment with 0, 2.5, 5, or 10 mmol/l aspirin for 48 h. The medium was replaced with 100 μ l of fresh medium containing 10% of CCK-8 reagent, and cells were incubated at 37°C for 3 h. The absorbance was measured at 450 nm using a multi-grating microplate reader SH-9000Lab (CORONA Electric Co., Ltd.). Experiments were carried out three times. Compared with RBE, HuCCT-1 cells were more sensitive to aspirin treatment in the cell viability assay. Moreover, considering the xenograft model, HuCCT-1 cells can be easily transplanted. Thus, HuCCT-1 cells were selected for further study *in vitro* and *in vivo*.

Flow cytometric analysis of the cell cycle. Flow cytometric analyses were performed using the Cycle Phase Determination kit (Cayman Chemical Company). HuCCT-1 and TKKK cells (1.0x10⁶ cells/100-mm dish) were treated with 2.5 mmol/l aspirin or without for 24 to 48 h. Cells were trypsinized and resuspended in phosphate-buffered saline (PBS) at a density of 1x10⁶ cells/ml. Approximately 1x10⁶ cells were stained

in 100 μ l of PBS with 10 μ l RNase A (250 μ g/ml) and 10 μ l propidium iodide (PI) stain (100 μ g/ml) and incubated at room temperature in the dark for 30 min. Flow cytometry (FCM) was performed to compare the proportion of aspirin-treated and control cells in each phase of the cell cycle. FCM was performed using a Cytomics FC 500 flow cytometer (Beckman Coulter, Inc.) with an argon laser (488 nm), and the percentages of cells were analyzed using the Kaluza software version v2.1 (Beckman Coulter, Inc.). The experiments were repeated thrice.

Apoptosis analysis. Aspirin-mediated apoptosis was analyzed using FCM and Annexin V-FITC Early Apoptosis Detection kit (Cell Signaling Technology, Inc.). HuCCT-1 cells (1.0x10⁶ cells/100-mm dish) were treated with 2.5 mmol/l aspirin or without for 48 h. Apoptotic and necrotic cells were analyzed by double staining with FITC-conjugated Annexin V and PI per the manufacturer's instructions. FCM was conducted using a Cytomics FC 500 flow cytometer with an argon laser (488 nm) to compare the proportion of apoptotic cells in the aspirin-treated and control groups, and data were analyzed using the Kaluza software version v2.1. The experiments were repeated thrice.

Apoptosis analysis by ELISA. ELISA was performed to analyze the levels of caspase-cleaved cytokeratin 18 (cCK18) using the M30 Apoptosense ELISA kit (cat. no. 10011; Peviva; Diapharma) according to the manufacturer's instructions. HuCCT-1 cells (5,000 cells/well) were seeded into 96-well plates and treated with 2.5 mmol/l aspirin for 48 h. Subsequently, the cells were lysed in polyoxyethylene octyl phenyl ether (Pure Chemical Industries, Ltd.) and further analyzed according to the manufacturer's instructions. The experiments were repeated thrice.

Western blot analysis. HuCCT-1 and TKKK cells were seeded (1.0x10⁶ cells/100-mm dish) and treated with 2.5 mmol/l aspirin for 24 or 48 h. The cells were lysed with PRO-PREP complete protease inhibitor mixture (iNtRON Biotechnology, Korea). Supernatants were collected, and the protein concentrations were measured using a NanoDrop 2000 spectrofluorometer (Thermo Fisher Scientific, Inc.). Protein aliquots (10 μ g) were resuspended in sample buffer and separated on 12% Tris-glycine gradient gels via sodium dodecyl sulfate-polyacrylamide gel electrophoresis. The resolved proteins were transferred to a nitrocellulose membrane and blocked with a blocking buffer containing 2% skimmed milk (GE Healthcare) in TBST with 0.1% Tween 20 (cat. no. T9142; Takara Bio, Inc.) for 30 min at room temperature. Subsequently, the membranes were incubated overnight at 4°C with primary antibodies in 5% serum (cat. no. 9048-46-8; FUJIFILM Wako Pure Chemical Industries, Ltd.) followed by incubation with horseradish peroxidase (HRP)-conjugated secondary antibodies in 2% skimmed milk. The following primary antibodies were used: Cyclin D1 (SP4) (1:2,000 dilution; cat. no. MA5-14512), retinoblastoma (Rb) (1:1,000 dilution; cat. no. MA1-34070), and cyclin E (HE-12) (1:1,000 dilution; MS-870-P1) were obtained from Thermo Fisher Scientific, Inc.; phosphorylated Rb (pS780) (1:1,000 dilution; cat. no. 558554) was obtained from BD Biosciences; Cdk2

(1: 5,000 dilution; cat. no. sc-163) was obtained from Santa Cruz Biotechnology, Inc.; and anti- β -actin (1:5,000 dilution; product no. A5441) was purchased from Sigma-Aldrich; Merck KGaA. The membranes were washed again with TBST and incubated with HRP-conjugated anti-mouse (dilution 1:2,000; product no. 7076) and anti-rabbit (dilution 1:2,000; product no. 7074) IgG secondary antibodies obtained from Cell Signaling Technology, Inc. for 1 h at room temperature. Finally, the signals were visualized using a typically enhanced chemiluminescent (ECL) kit (cat. no. 45-000-999; Cytiva), and blots were imaged using ImageQuant LAS 4010 (GE Healthcare).

miRNA microarray. Cells were treated with 2.5 mmol/l aspirin for 48 h, and total RNA was extracted using the miRNeasy Mini kit (Qiagen GmbH) according to the manufacturer's instructions. After confirming the purity and quantity of each sample using an Agilent 2100 Bioanalyzer and an RNA 6000 Nano kit (both from Agilent Technologies), respectively, the samples were labeled using a miRCURY Hy3 Power Labeling kit (Exiqon A/S) and hybridized to a human miRNA Oligo Chip (v.21; Toray Industries, Inc.). Chips were scanned using the 3D-Gene Scanner 3000 (Toray Industries, Inc.). The 3D-Gene extraction software version 1.2 (Toray Industries, Inc.) was used to calculate the raw signal intensity of the images. The raw data were analyzed using the GeneSpring GX 10.0 software (Agilent Technologies, Inc.) to assess the differences in miRNA expression between the aspirin-treated and control samples. Global normalization was performed on raw data obtained above the background level. Differentially expressed miRNAs were determined using Welch's t-test.

Reverse transcription-quantitative polymerase chain reaction (RT-qPCR) analysis of miRNAs. qPCR was used to validate the expression levels obtained from the miRNA assay. Total RNA was extracted as previously described above and diluted to 2.0 ng/ μ l. TaqMan microRNA assays (Applied Biosystems; Thermo Fisher Scientific, Inc.) were adopted to determine the expression of miRNAs using U6 small nuclear RNA (*RNU6B*) as an internal control. miRNAs were reverse transcribed using the TaqMan microRNA Reverse Transcription kit (Applied Biosystems; Thermo Fisher Scientific, Inc.). Reverse transcription was performed in 20- μ l reaction volumes consisting of 5 μ l of RNA, 3 μ l of 5X RT primer, and 12 μ l of reverse transcription Master Mix. qPCR was performed in the MicroAmp Fast Optical 96-Well Reaction Plate (Applied Biosystems; Thermo Fisher Scientific, Inc.). Each well contained a 20- μ l reaction consisting of 2 μ l of cDNA, 1 μ l of 20X qPCR assay, 7 μ l of nuclease-free water, and 10 μ l of TaqMan Fast Advanced Master Mix (Applied Biosystems; Thermo Fisher Scientific, Inc.). PCR was performed using the ViiA7 real-time PCR system (Applied Biosystems; Thermo Fisher Scientific, Inc.) with the following reaction steps: Hold at 50°C for 2 min, denaturation at 95°C for 20 sec followed by 40 cycles of 1 sec at 95°C and 20 sec at 60°C. The relative expression of miR-340-5p was calculated using the comparative Cq method according to the following formula: $2^{-\Delta\Delta Cq}$ ($\Delta Cq = \text{miRCq} - \text{U6Cq}$, $\Delta\Delta Cq = \Delta Cq - \text{average control } \Delta Cq$) (21). The primer sequences are as follows: miR-340-5p forward, 5'-GCGTTATAAAGCAATGAGA-3' and reverse, 5'-GTG

CGTGTCGTGGAGTCG-3'; U6 forward, 5'-GCTTCGGCA GCACATATACTAAAAT-3' and reverse, 5'-CGCTTCACG AATTTGCGTGCAT-3'.

Bioinformatics analysis used for prediction of the target genes of miR-340-5p and functional and network analyses. For bioinformatics analysis, miRDB database (www.mirdb.org) was used to predict the target genes of miR-340-5p and the Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway analyses was used for upregulated miRNAs depending on the Database for Annotation, Visualization, and Integrated Discovery (DAVID database) (22). miRNA regulatory network used mirPath v.3 (<http://snf-515788.vm.okeanos.grnet.gr>) (23); miRGator v3.0 (<http://mirgator.kobic.re.kr>) (24) and miTarBase (<http://mirtarbase.cuhk.edu.cn/php/index.php>) (25) online software to reveal the expression of miR-340-5p in related cancer and normal tissue and the relationship with the underlying target gene. The cut-off criterion was set as $P < 0.05$.

Cell transfection. For transfection, 5×10^5 HuCCT-1 cells were seeded in 6-well plates with antibiotic-free medium 1 day before transfection to reach a confluence of 90% at the time of transfection. Cells were transfected with miR-340-5p mimic (50 nM) (sense, 5'-UUAUAAAGCAAUGAGACUGAUU-3' and antisense, 5'-UCAGUCUCAUUGCUUUAUAATT-3'), inhibitor (50 nM) (5'-AAUCAGUCUCAUUGCUUUAUA A-3'), or negative control miRNA (50 nM) (mimic negative control 5'-UUGUACUACACAAAAGUACUG-3'; inhibitor negative control 5'-CAGUACUUUUGUGUAGUACAA-3') (Life Technologies; Thermo Fisher Scientific, Inc.) using Lipofectamine RNAiMAX Reagent (Invitrogen; Thermo Fisher Scientific, Inc.). The transfection complex was prepared according to the manufacturer's instructions and added to the cells, and the plates were incubated in a humidified atmosphere with 5% CO₂ at 37°C. Cells cultured with transfection reagent as untransfected control. The medium was replaced 6 h post-transfection. Cell samples were collected at 0 or 48 h after transfection for further analysis.

Colony formation assay. Cells were trypsinized for 5 min and resuspended at a density of 1×10^3 /ml. Five hundred microliters were seeded into 6-well plates, and 1.5 ml RPMI-1640 medium containing 10% FBS was added to each well. The plates were incubated at 37°C at 5% CO₂, and the medium was changed every 3 days until conspicuous colonies were observed. Colonies were stained with 0.1% crystal violet for 5 min at room temperature and positive colony formation (>50 cells/colony) was evaluated by counting the number of colonies.

Xenograft model analysis. The animal study was conducted in accordance with the guidelines set by the Committee on Experimental Animals of the Kagawa University. All experimental protocols were approved (approval no. 18674) by the Institutional Review Board of the Department of Laboratory Animal Science of Kagawa University (Kawaga, Japan). Thirty-five female athymic mice (BALB/c-nu/nu; 6 weeks old; 19-21 g) were purchased from Japan SLC (Shizuoka, Japan). The mice were maintained at 20-25°C with 30-60% humidity under a 12:12 h light/dark cycle, using a laminar airflow rack

and had continuous free access to sterilized (γ -irradiated) food (CL-2; CLEA Japan, Inc.) and autoclaved water. Mice were subcutaneously inoculated with 1.5×10^6 HuCCT-1 cells in the right flank. When the xenografts were palpable with an approximate diameter of 3 mm, we randomly assigned the animals to three groups of 10 animals each. These groups were treated with 60 mg/kg aspirin, 100 mg/kg aspirin, or vehicle (10% ethanol in PBS) by intraperitoneal injection every day. The tumor volume (mm^3) was calculated as tumor length (mm) \times tumor width (mm^2)/2. For humane endpoints, if difficulty in feeding and/or intake of water; apparent poor physical condition; a rapid and non-recoverable weight loss (over 20% body weight); and/or a significant increase in tumor size was observed (tumor size more than 10% of body weight and/or tumor diameter more than 20 mm), the experiments were discontinued. Animals were euthanized through CO_2 euthanasia with 20% displacement of cage volume/min. Sacrifice was confirmed by observation of unconsciousness, absence of heartbeat and absence of breathing. All animals were sacrificed on day 28 of treatment. The experiments were performed from February 20, 2019 to March 28, 2019.

Immunohistochemistry. Immunohistochemistry was performed on tumor tissues obtained from the xenografted mice. We prepared 5- μm -thick sections from formalin-fixed (10% formalin at room temperature for 24 h), paraffin-embedded tissue blocks which were deparaffinized, rehydrated, and subjected to immunohistochemistry studies. Following a blocking step at room temperature for 30 min using a Vectastain Elite ABC kit (Vector Laboratories, Inc.), tissue sections were incubated with primary antibodies [Cyclin D1 (EP12); 1:200 dilution; cat. no. 241R-4, Sigma-Aldrich; Merck KGaA]. Sections were washed and incubated with anti-rabbit IgG Antibody (1:50 dilution; cat. no. BP-9100-50; Vector Laboratories) at room temperature for 1 h and then with a streptavidin-peroxidase solution. Color reactions involved the use of 3,3'-diaminobenzidine (DAB) with Mayer's hematoxylin counterstaining at room temperature for 10 sec. The specificity of immunostaining was evaluated using non-immune mouse IgG (1:50 dilution; cat. no. BP-9200-50; Vector Laboratories) at room temperature for 1 h as a negative control for the primary antibody. Sections were examined microscopically for specific staining, and nuclei with a brown color regardless of staining intensity were regarded as positive. Cyclin D1 positivity was calculated at a magnification of $\times 40$ by dividing the number of positive cells by the total number of cells counted in five random fields and expressed as a percentage. Images were captured using an Olympus BX51 microscope and Olympus DP72 camera (magnification, $\times 40$; Olympus Corporation).

Statistical analysis. GraphPad Prism software version 6.0 (GraphPad Software, Inc.) was used for all analyses. A two-tailed unpaired Student's t-test was used to determine statistical significance between different groups. Two-way analysis of variance (ANOVA) or mixed ANOVA was performed to test the comparisons and corrected by Tukey's post hoc test. One-way analysis of variance before the Tukey's post hoc test was performed to test the comparisons. A P-value of <0.05 was considered to indicate a statistically significant difference.

Results

Aspirin inhibits the proliferation of most human CCA cells. The anti-proliferative effects of aspirin on human CCA cells were determined using the HuCCT-1, RBE, and TTKK CCA cell lines. Cells were treated with 2.5, 5, or 10 mmol/l aspirin for 48 h, and the anti-proliferative effect of aspirin was assessed using the cell viability assay. Untreated cells were used as controls. Aspirin inhibited cell proliferation in CCA cells in a dose and time-dependent manner, except in TTKK cells (Fig. 1).

Aspirin induces cell cycle arrest in the G_0/G_1 phase and regulates cell-cycle related proteins in HuCCT-1 cells. To determine whether aspirin affected the cell cycle in CCA cells, HuCCT-1 cells were treated with aspirin and FCM was performed to examine cell cycle progression. Western blotting was also performed to evaluate the expression of cell-cycle related proteins. Cells were treated with 2.5 mmol/l aspirin for 24 or 48 h, and untreated cells were used as the control. Following aspirin treatment, the population in the G_0/G_1 phase significantly increased, whereas cells in the S phase decreased (Fig. 2A and B). Western blot results indicated that treatment with aspirin for 48 h significantly modulated cyclin D1, a key protein expressed in the early G_1 phase. Cyclin D1 is a key regulator of the cell cycle and is involved in the transition from the G_1 phase to the S phase (26). The levels of phosphorylated Rb decreased with aspirin treatment, suggesting that the treated cells were in G_1 arrest (Fig. 2C).

Aspirin does not induce cell apoptosis in HuCCT-1 cells. To determine whether aspirin induced apoptosis in HuCCT-1 cells, FCM was used to detect apoptotic cells after aspirin treatment. The different quadrants represent living cells (lower left quadrant), early apoptotic cells (lower right quadrant), and late apoptotic cells (upper right quadrant). The proportion of early apoptotic cells with aspirin treatment and without treatment were similar after 48 h (Fig. 3A). Additionally, there was no obvious difference in cCK-18 levels between treated and untreated cells after 24- or 48-h treatments (Fig. 3B).

Aspirin affects miRNA expression in HuCCT-1 cells. A customized microarray platform was used to analyze the expression of 2,555 miRNAs in aspirin-treated and control HuCCT-1 cells. Treatment with 2.5 mmol/l aspirin for 48 h induced the overexpression of eight miRNAs, whereas the expression of six miRNAs was decreased (Table I; Fig. 4). Unsupervised hierarchical clustering analysis was conducted by calculating Pearson's centered correlation coefficient, and the results indicated that differentially expressed miRNAs in aspirin-treated HuCCT-1 cells clustered together. Previous studies have indicated that miR-340-5p has tumor-suppressive properties (27,28), and miR-340-5p exhibited functions in different parts in the KEGG pathway analyses of upregulated miRNAs, which included 'prion diseases', 'microRNA in cancer', 'proteoglycans in cancer', and 'signaling pathways regulating pluripotency of stem cells' as well as others (Fig. 5A). Although the expression of miR-340-5p was not clearly revealed in CCA, it was decreased in most types of cancer (Fig. 5B). The miRNA regulatory network and miRDB database predicted

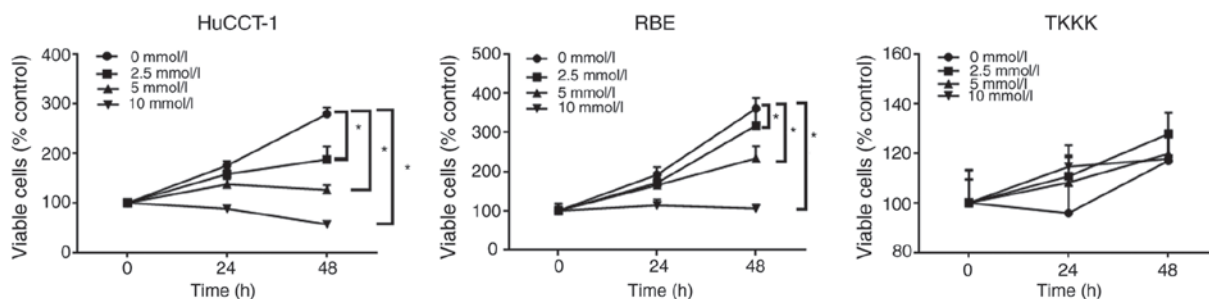


Figure 1. Aspirin inhibits the proliferation of some CCA cells. All cell lines were treated with 0, 2.5, 5, or 10 mmol/l aspirin for 24 or 48 h. The data points represent the mean cell number from three independent cultures, and the error bars represent standard deviations. The antiproliferative effect was significantly higher in cells treated with aspirin treatment for 48 h than in control cells (0 mmol/l) as determined by two-way analysis of variance, except for TKKK cells. *P<0.01, vs. the control. CCA, cholangiocarcinoma.

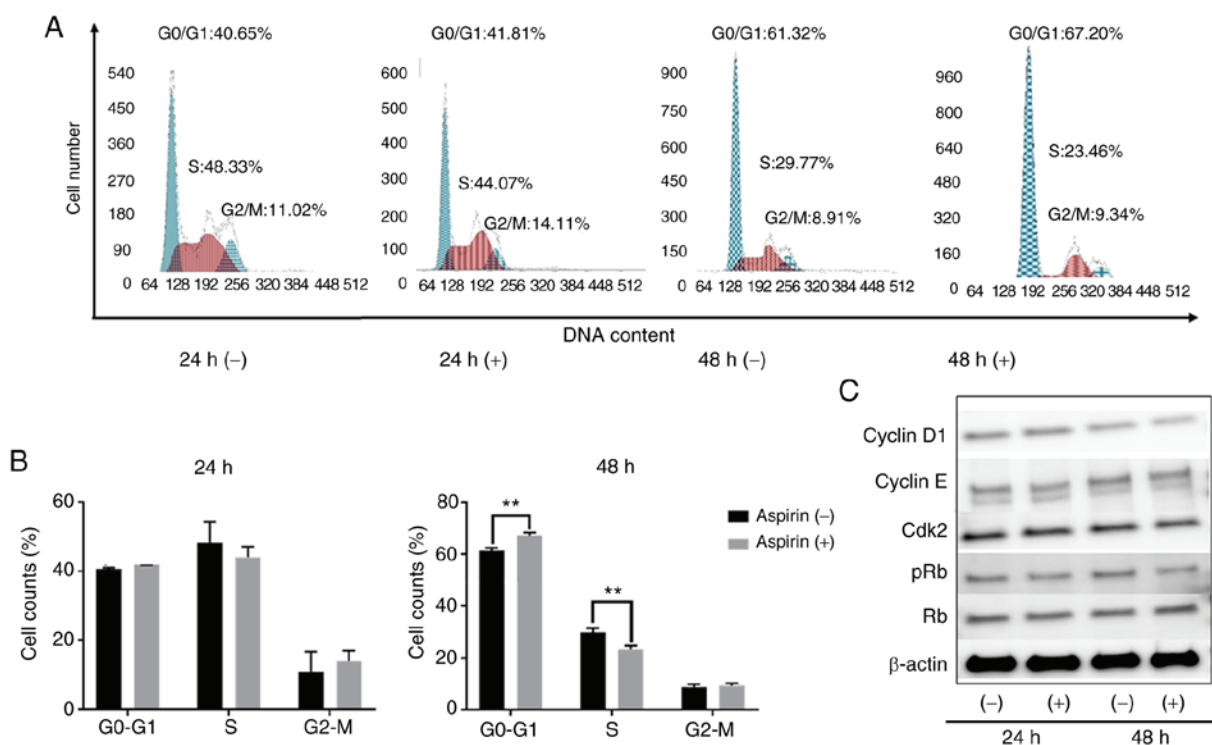


Figure 2. Aspirin causes cell cycle arrest in HuCCCT-1 cells. (A) HuCCCT-1 cells treated with or without 2.5 mmol/l aspirin were analyzed by flow cytometry to estimate the proportion of cells in each phase of the cell cycle. (B) Graphical representation of the proportion of cells in each phase of the cell cycle. Aspirin blocked the cell cycle at the G₀/G₁ phase. **P<0.01, vs. the control. (C) Expression of cyclin D1, cyclin E, Cdk2, Rb, and pRb in HuCCCT-1 cells 24 and 48 h after aspirin treatment. Rb, retinoblastoma; pRb, phosphorylated Rb.

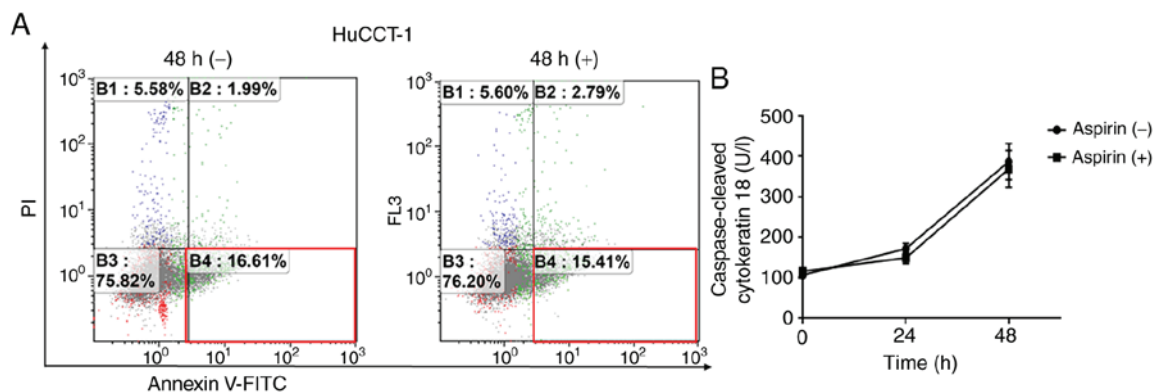


Figure 3. Aspirin does not induce apoptosis in HuCCCT-1 cells. (A) Aspirin treatment (2.5 mmol/l for 48 h) did not alter the proportion of early apoptotic HuCCCT-1 cells as determined by flow cytometry. (B) The expression of caspase-cleaved cyokeratin 18 as determined using ELISA after 24 or 48 h of 2.5 mmol/l aspirin treatment.

Table I. Statistical results and chromosomal locations of microRNAs in HuCCT-1 cells.

miRNA	Fold change (Treated/Untreated)	P-value	Chromosomal location
A, Upregulated			
hsa-miR-6504-3p	3.092	0.021	16
hsa-miR-4266	2.701	0.005	2q13
hsa-miR-4708-3p	2.158	0.028	14
hsa-miR-532-5p	2.055	0.039	X
hsa-miR-340-5p	2.029	0.039	5
hsa-miR-203a-3p	1.552	0.018	14
hsa-miR-1180-3p	1.547	0.009	17
hsa-miR-146a-5p	1.522	0.022	5
B, Downregulated			
hsa-miR-6744-3p	0.654	0.003	11
hsa-miR-6752-3p	0.652	0.000	11
hsa-miR-6859-5p	0.640	0.041	
hsa-miR-4669	0.636	0.030	9q34.2
hsa-miR-3607-5p	0.526	0.045	
hsa-miR-604	0.445	0.011	10p11.23

Cells treated with aspirin that exhibited a fold change (FC)>1.5, FC<0.67, or a P-value <0.05 compared with untreated cells are presented.

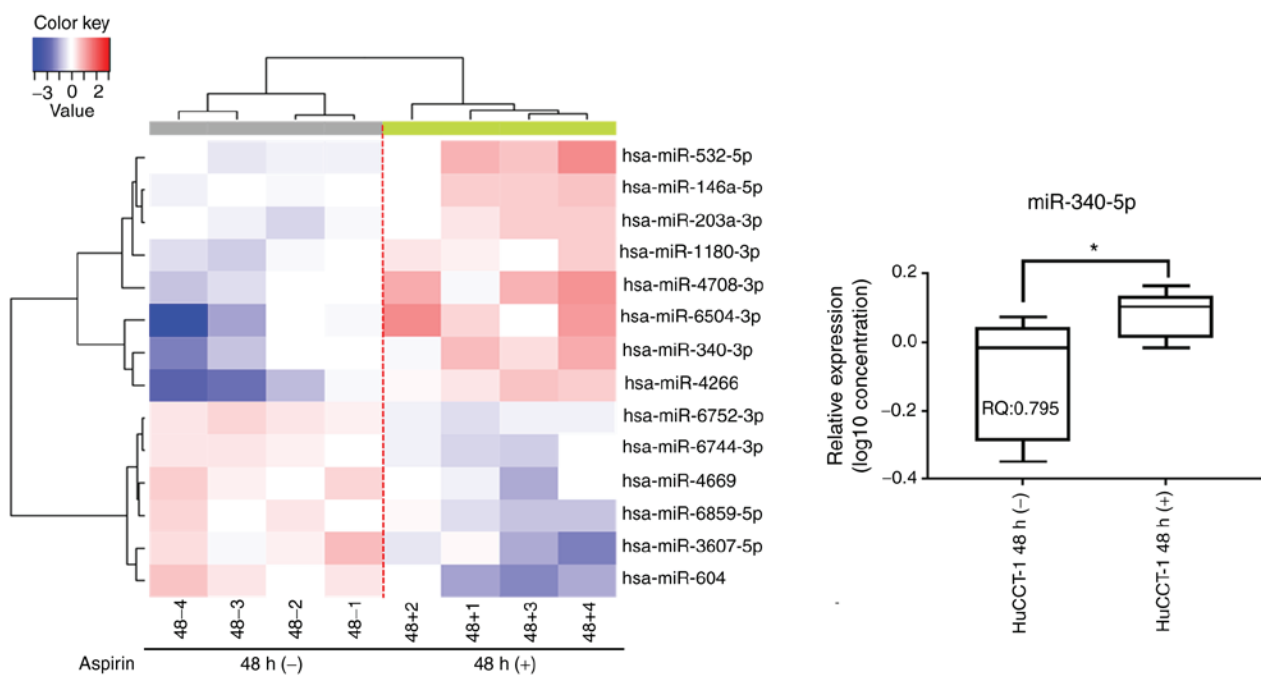


Figure 4. Aspirin affects microRNA expression in HuCCT-1 cells. Hierarchical clustering of the expression profiles of numerous differentially expressed miRNAs from HuCCT-1 cells cultured with or without aspirin. Relative quantification of miRNA following aspirin treatment. miR-340-5p expression was significantly upregulated after aspirin treatment. *P<0.05, vs. the control. The $\log_{10} 2^{-\Delta\Delta Cq}$ values for microRNAs were used to create the image, and the lines represent averages with interquartile ranges.

that CCND1 may be a target of miR-340-5p (Fig. 5C and D). Thus, it was selected for further study, and RT-qPCR indicated that miR-340-5p levels were significantly upregulated in the

aspirin-treated cells compared to the untreated cells (Fig. 4). Numerous other studies have revealed the effects of miRNAs on CCA in recent years (Table II) (29-47).

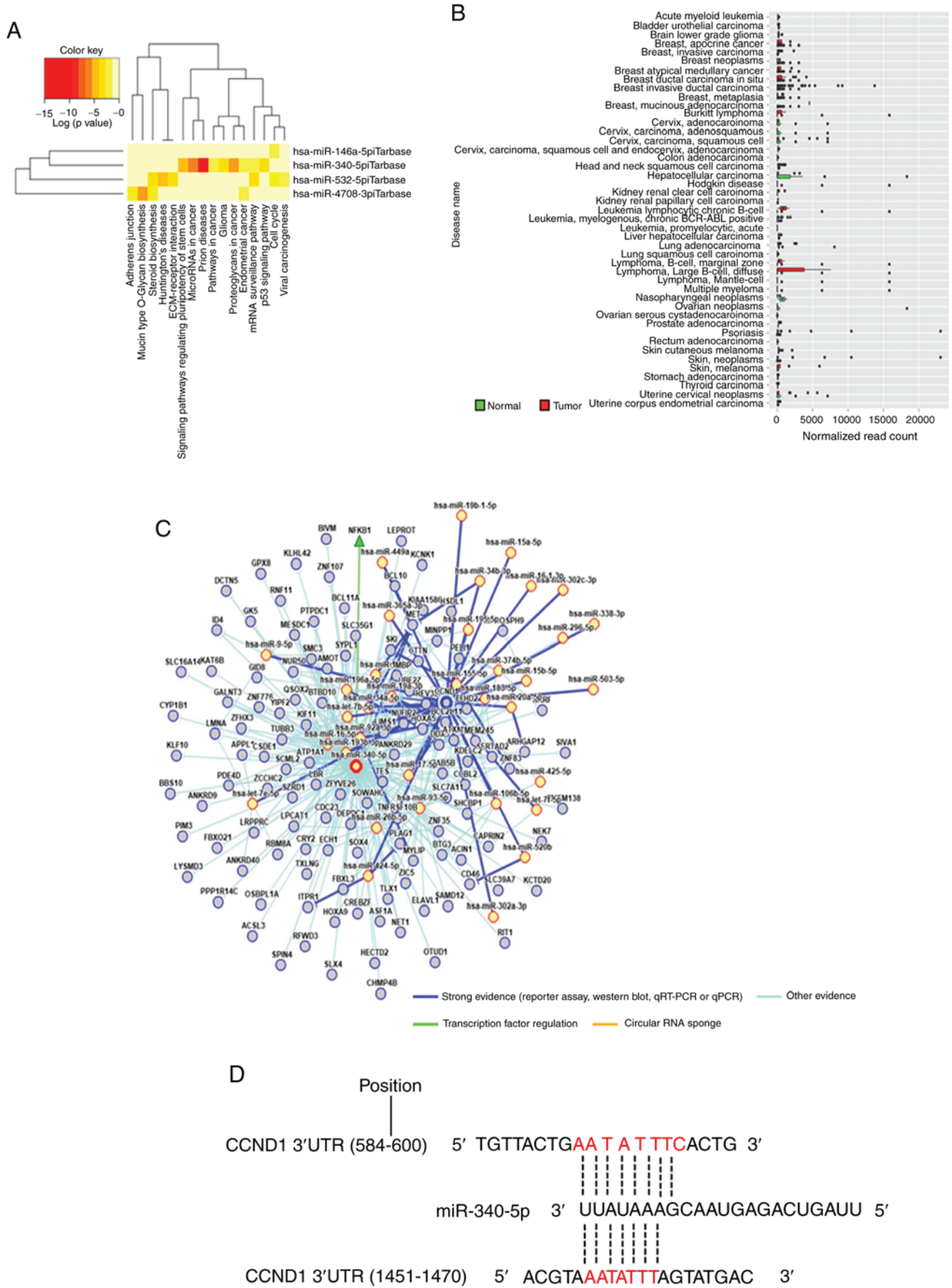


Figure 5. Bioinformatics analysis and the predicted target gene of miR-340-5p. (A) The Kyoto Encyclopedia of Genes and Genomes pathway analyses of upregulated miRNAs. (B) Expression levels of miR-340-5p in related cancer and normal tissue. Compared with normal tissue, the expression of miR-340-5p was revealed to be decreased in most types of cancer. (C) miRNA regulatory network of miR-340-5p. The intensity of evidence between miR-340-5p with predicted target genes and relationships with associated miRNAs are presented in this network. (D) The seed sequence of human miR-340-5p was complementary to the 3'UTR of CCND1.

Table II. Previous studies examining microRNAs in cholangiocarcinoma.

MicroRNA (promoter)	References	MicroRNA (inhibitor)	References
miR-191	Li <i>et al</i> (2017) (29)	miR-195	Li <i>et al</i> (2017) (30)
miR-205-5p	Kitdumrongthum <i>et al</i> (2018) (31)	miR-410	Palumbo <i>et al</i> (2016) (32)
miR-21	Lampis <i>et al</i> (2018) (33)	miR-203	Li <i>et al</i> (2015) (34)
miR-181c	Wang <i>et al</i> (2016) (35)	miR-15a	Utaijaratrasmi <i>et al</i> (2018) (36)
miR-193-3p	Han <i>et al</i> (2018) (37)	miR-34a	Han <i>et al</i> (2016) (38)
miR-383	Wan <i>et al</i> (2018) (27)	miR-433	Mansini <i>et al</i> (2018) (28)
		miR-22	
		miR-199a-3p	Li <i>et al</i> (2017) (39)
		miR-144	Yang <i>et al</i> (2014) (40)
		miR-590-3p	Zu <i>et al</i> (2017) (41)
		miR-101	Deng <i>et al</i> (2015) (42)
		miR-26b-5p	Fan <i>et al</i> (2018) (43)
		miR-24	Ehrlich <i>et al</i> (2017) (44)
		miR-122	Liu <i>et al</i> (2015) (45)
		miR-26a	Wang and Lv (2016) (46)
		miR-551b-3p	Chang <i>et al</i> (2019) (47)

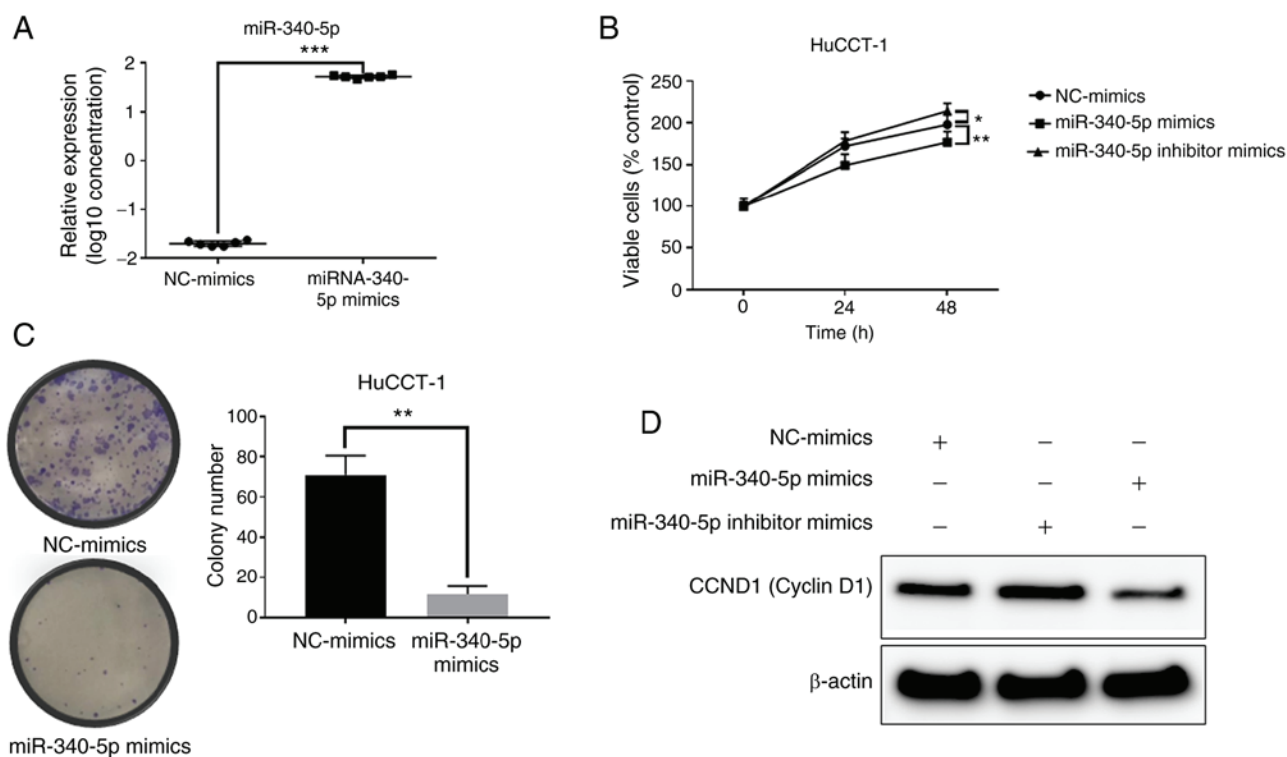


Figure 6. Role of miR-340-5p in HuCCT-1 cells. (A) Relative quantification of miR-340-5p following transfection of miR-340-5p mimics or NC. (B) Cell proliferation in HuCCT-1 cells after transfection of NC-mimics, miR-340-5p mimics, or an miR-340-5p inhibitor. (C) miR-340-5p overexpression inhibited colony formation. (D) The levels of cyclin D1 after transfection of NC mimics, miR-340-5p mimics, or an miR-340-5p inhibitor. * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$, vs. the control. NC, negative control.

miR-340-5p inhibits the proliferation of HuCCT-1 cells and decreases the expression levels of cyclin D1. After transfection of miR-340-5p mimics, miR-340-5p expression was significantly increased in the HuCCT-1 cells (Fig. 6A). Transfection of miR-340-5p mimics decreased proliferation in HuCCT-1 cells (Fig. 6B). Moreover, overexpression of miR-340-5p decreased

the levels of cyclin D1, whereas inhibition induced increased cyclin D1 levels (Fig. 6D). The colony formation assay indicated that overexpression of miR-340-5p decreased the cell proliferation ability of HuCCT-1 cells (Fig. 6C). Therefore, increasing the levels of miR-340-5p inhibited cyclin D1 expression and decreased the cell proliferation ability.

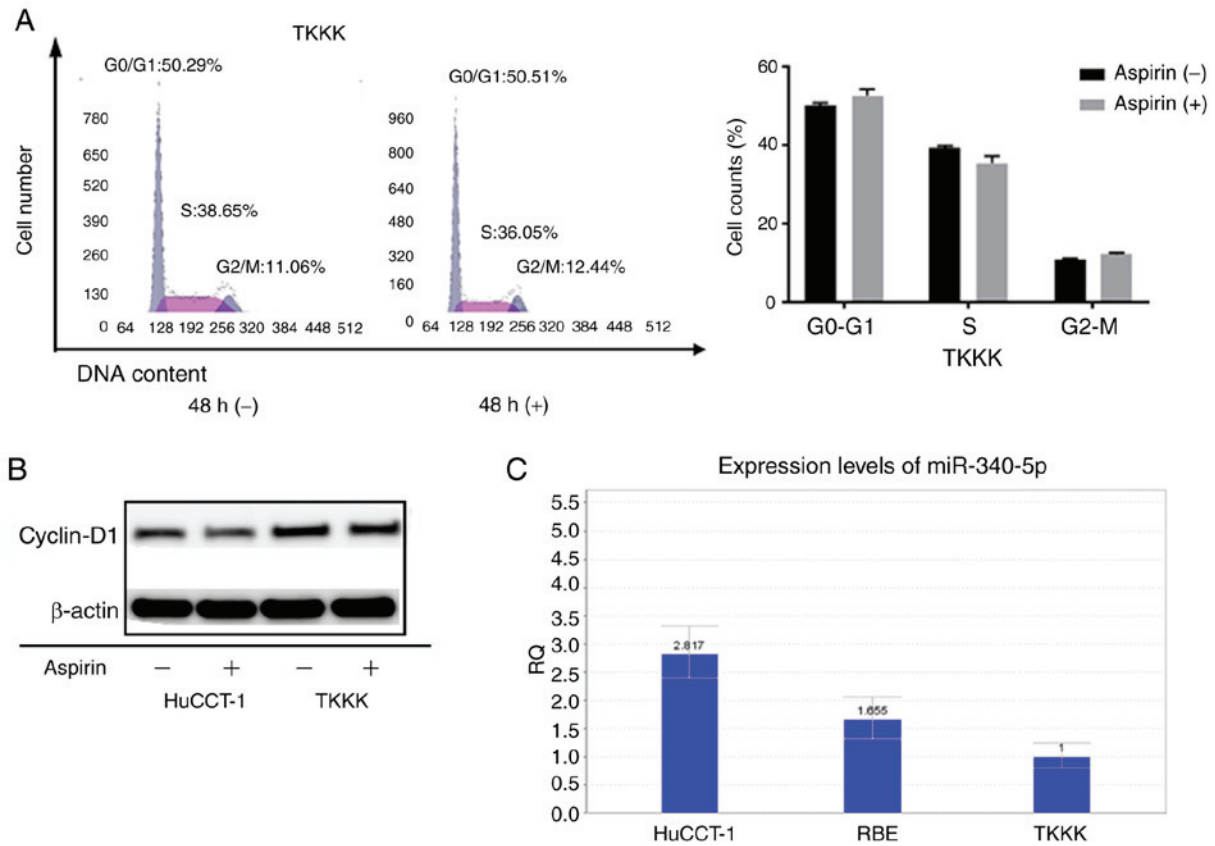


Figure 7. Comparison of the cell cycle and miR-340-5p in aspirin-nonresponsive TKKK cells and aspirin-responsive HuCCT1 cells. (A) Left image: TKKK cells treated with or without 2.5 mmol/l aspirin and analyzed by flow cytometry to estimate the proportion of cells in each phase of the cell cycle. Right image: Graphical representation of the proportion of cells in each phase of the cell cycle. (B) Expression of cyclin D1 in HuCCT-1 cells and TKKK cells after 48 h of aspirin treatment. (C) Relative quantification of miR-340-5p in cholangiocarcinoma cell lines.

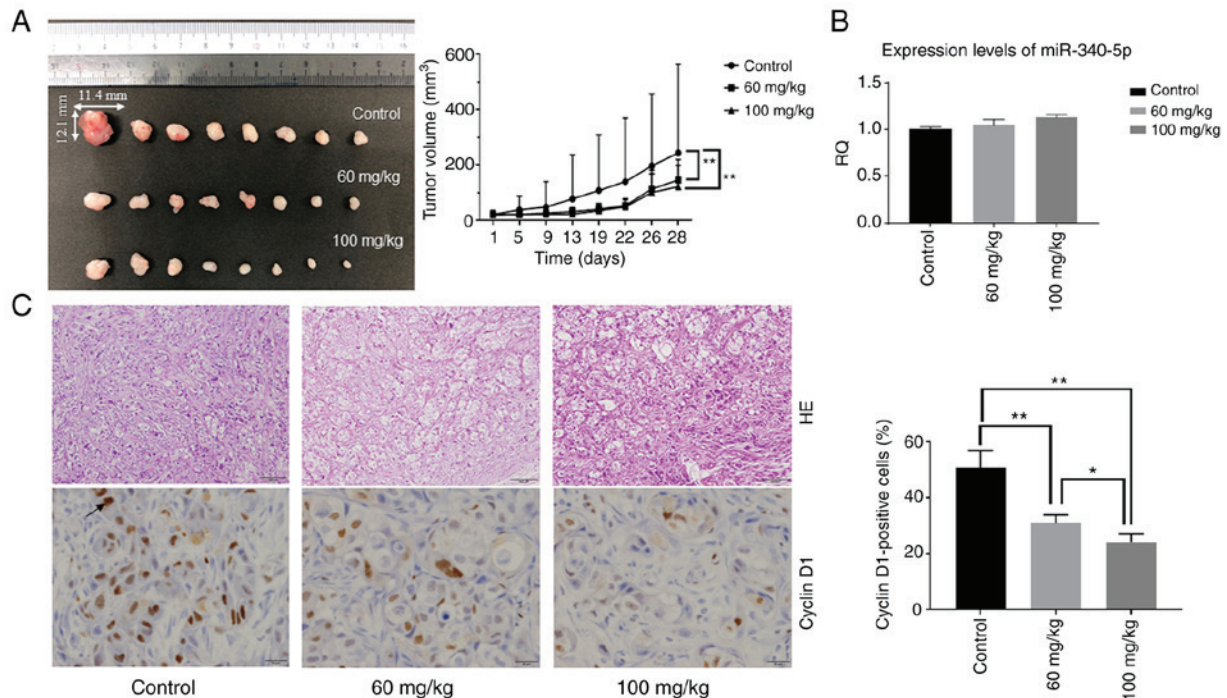


Figure 8. Aspirin inhibits the growth of HuCCT-1 cell xenografts in nude mice. (A) The tumors were significantly smaller in aspirin-treated mice than in vehicle-treated mice. Each point represents the mean \pm standard deviation of 10 animals. ** $P < 0.01$ by mixed analysis of variance. (B) Relative quantification of miR-340-5p in tumor tissue of the three groups of xenografted mice. (C) Left image: H&E-stained images of the xenografted tumor tissues between the aspirin-treated and control mice (scale bar, 100 μ m). Immunohistochemical staining of cyclin D1 between the aspirin-treated and control mice (scale bar, 20 μ m). Right image: Cyclin D1-positive cells in the aspirin-treated groups were reduced in number compared with the control group as determined by one-way analysis of variance. * $P < 0.05$ and ** $P < 0.01$. H&E, hematoxylin and eosin.

Aspirin-nonresponsive TKKK cells express low levels of miR-340-5p. As the CCA cell line TKKK did not exhibit response to aspirin, the cell cycle progression in TKKK cells as compared to HuCCT-1 cells was assessed. Forty-eight hours of aspirin treatment in TKKK cells revealed no obvious difference in the proportion of cells in each phase of the cell cycle (Fig. 7A). In addition, the levels of cyclin D1 were not significantly altered (Fig. 7B). Relative quantification of miR-340-5p was assessed in all cell lines (Fig. 7C). Expression was lowest in TKKK cells, which may indicate that cell lines with high expression of miR-340-5p are more sensitive to cell cycle arrest with aspirin treatment.

Aspirin inhibits tumor proliferation in vivo. Based on the results obtained from *in vitro* studies, the effect of aspirin in an *in vivo* model of CCA was assessed. Nude mice were injected subcutaneously with HuCCT-1 cells followed by intraperitoneal injection of aspirin. The present results revealed that tumor growth was significantly inhibited in mice treated with aspirin compared to untreated mice ($P < 0.05$) (Fig. 8A). No mice succumbed during the observation period. Expression levels of miR-340-5p in tumor tissue were not significantly different, although the RQ was slightly increased in the two aspirin-treated groups (Fig. 8B). H&E-stained images of the xenografted tumor tissues revealed no significant histopathological differences between the aspirin-treated and control mice. Immunohistochemical staining of cyclin D1 indicated that cyclin D1-positive cells in the aspirin-treated groups were reduced in number compared with the control group (Fig. 8C).

Discussion

CCA is an aggressive cancer with high mortality and poor prognosis that accounts for approximately 3% of gastrointestinal malignancies (8). The 5-year survival rate of CCA is only 10%, and the median survival is 24 months (48). Initial analysis revealed that aspirin use was associated with a reduced iCCA risk in men (HR=0.64, 95% CI=0.42-0.98) (49). Subsequent analysis has revealed that aspirin use has a significant inverse association with the development of all three CCA subtypes and leads to an approximately 3-fold reduction in CCA risk (13). This is thought to be due to the anti-inflammatory effect of aspirin. In the present study, we further elucidated the mechanism of the effect of aspirin on CCA. To the best of our knowledge, the present study is the first study revealing that aspirin inhibits the proliferation of CCA cells *in vivo*.

The anti-inflammatory dose of aspirin varies from 0.5-2.5 mM (50); therefore, relying on data from the anti-proliferation assay, 2.5 mM was selected as the concentration of aspirin, which does not have off-target cytotoxicity. It was observed that aspirin inhibited the proliferation of CCA cells (HuCCT-1 and RBE) and induced cell cycle arrest (HuCCT-1) at the G_0/G_1 phase, which was correlated with a marked decrease in the expression of cyclin D1. Aspirin also decreased the phosphorylation of Rb. The expression of cell cycle-related molecules is related to cancer progression and prognosis (51), and aspirin has been revealed to inhibit the expression of cyclin D1 in other types of cancer,

such as oral squamous cell carcinoma (52) and glioblastoma multiforme (53). To determine whether aspirin induced apoptosis, HuCCT-1 cells were treated with 2.5 mM aspirin and analyzed using FCM; the levels of cCK18 were also measured. However, there was no evidence that aspirin induced apoptosis in HuCCT-1 cells. These data indicated that aspirin inhibited CCA cell proliferation mainly through cell cycle arrest. However, aspirin has also been revealed to induce apoptosis in numerous types of cancers (52-55); this discrepancy could be due to differences in the properties of different types of cancers.

miRNAs are short, noncoding, endogenous, single-stranded RNA molecules 19-25 nucleotides in length that regulate target gene expression (56). They are known to regulate the development and progression of various cancers (57). A miRNA expression array was used to identify the miRNAs associated with the antitumor effects of aspirin. miR-340-5p that was significantly upregulated in response to aspirin treatment in cells, was not significantly different in tumor tissue, although the relative quantification (RQ) was slightly increased in the two aspirin-treated groups. Recent studies have indicated that miR-340-5p inhibited non-small cell lung cancer cell growth and metastasis by targeting ZNF503 (58) and suppressed osteosarcoma development via targeting STAT3 (59). In the present study, overexpression of miR-340-5p inhibited proliferation in HuCCT-1 cells. In addition, overexpression of miR-340-5p decreased the levels of cyclin D1. The predicted sequence from the miRDB database and bioinformatics analysis indicated that cyclin D1 may be an miR-340-5p target, and miR-340-5p overexpression decreased cyclin D1 levels in cells; immunohistochemical staining indicated that cyclin D1-positive cells in aspirin-treated mice were reduced compared with control mice. Therefore, it is theorized that aspirin works partially through the miR-340-5p/cyclin D1 axis to inhibit HuCCT-1 cell proliferation.

In the present study, aspirin treatment had no effect on TKKK cells. To investigate the difference between TKKK cells and HuCCT-1 cells, FCM was used to analyze changes in the cell cycle with aspirin treatment. Aspirin did not induce G_0/G_1 arrest in TKKK cells, and there was no difference in the levels of cyclin D1 after 48 h of treatment. In addition, the relative levels of miR-340-5p were assessed in all cell lines, and TKKK cells had the lowest expression. Therefore, aspirin may not suppress proliferation in TKKK cells because it cannot utilize the miR-340-5p/cyclin D1 axis.

In the present *in vivo* model, aspirin inhibited the growth of subcutaneous CCA tumors in athymic nude mice. In accordance with the *in vitro* results and previous studies (23,60), in the *in vivo* experiment, the tumor volumes in both treatment groups (low-dosage and high-dosage) were significantly decreased compared with the tumor volume in the control group. However, there was no obvious difference in tumor volume between the low-dosage and high-dosage groups; long-term treatment may be required to see a difference based on dosage.

In conclusion, the present study indicated that aspirin inhibited cell proliferation and tumor growth in some CCA cell lines by inducing G_0/G_1 phase cell cycle arrest, and the underlying mechanism may partially be through the miR-340-5p/cyclin D1 axis to induce cell cycle arrest.

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Availability of data and materials

All data supporting the conclusions of the present study have been documented in this study.

Authors' contributions

TS, AM, HK and TM conceived and designed the experiments. TS, HI, JG, NN, SL, MN, HY, TN, KO, YS and KF analyzed and interpreted the data. TS performed the experiments. TS wrote, reviewed and edited the manuscript. TM reviewed and edited the manuscript for important intellectual content. All authors reviewed, read and approved the final manuscript.

Ethics approval and consent to participate

All experimental protocols were approved (approval no. 18674) by the Institutional Review Board of the Department of Laboratory Animal Science of Kagawa University (Kawaga, Japan).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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