Inhibition of Transient Receptor Potential Channel 5 Reverses 5-Fluorouracil Resistance in Human Colorectal Cancer Cells*

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Background: Resistance to 5-fluorouracil leads to the failure of chemotherapy for colorectal cancer. Results: Suppressing TrpC5 expression decreased nuclear β-catenin accumulation, reduced the induction of ABCB1, and reversed 5-fluorouracil resistance. Conclusion: TrpC5 is essential in ABCB1 induction and drug resistance in human colorectal cancer cells. Significance: These findings may help develop a novel target for overcoming resistance to chemotherapy in colorectal cancer.

5-Fluorouracil (5-Fu) is commonly used in the chemotherapy of colorectal cancer (CRC), but resistance to 5-Fu occurs in most cases, allowing cancer progression. Suppressing ABCB1 (ATP-binding cassette, subfamily B, member 1), which is a pump overproduced in cancer cells to export cytotoxic drugs, is an attractive strategy to overcome drug resistance. In the present study, transient receptor potential channel TrpC5 was found to be overproduced at the mRNA and protein levels together with ABCB1 in 5-Fu-resistant human CRC HCT-8 (HCT-8/5-Fu) and LoVo (LoVo/5-Fu) cells. More nuclear-stabilized β-catenin accumulation was found in HCT-8/5-Fu and LoVo/5-Fu cells than in HCT-8 and LoVo cells. Suppressing TrpC5 expression with TrpC5-specific siRNA inhibited the canonical Wnt/β-catenin signal pathway, reduced the induction of ABCB1, weakened the ABCB1 efflux pump, and caused a remarkable reversal of 5-Fu resistance in HCT-8/5-Fu and LoVo/5-Fu cells. On the contrary, enforcing TrpC5 expression resulted in an activated Wnt/β-catenin signal pathway and up-regulation of ABCB1. Taken together, we demonstrated an essential role of TrpC5 in ABCB1 induction and drug resistance in human CRC cells via promoting nuclear β-catenin accumulation.

Colorectal cancer (CRC) is a major cause of mortality and morbidity throughout the world (1), the third most common malignant tumor, and the fifth cause of cancer-related death in China. 5-Fluorouracil (5-Fu) is one of the most commonly used cytotoxic drugs in the chemotherapy of CRC. Resistance to 5-Fu occurs in most cases, and this results in cancer progression. High expression of ABCB1 (ATP-binding cassette, subfamily B, member 1) is the most important mechanism in drug resistance and has been considered as a target for its reversal (2). Although many ABCB1 modulators have been developed to inhibit its activity, most are either too toxic or induce intolerable pharmacokinetic interactions (2).

Intracellular Ca^{2+} ([Ca^{2+}]) is one of the crucial signals that modulate cellular functions, and some members of the Ca^{2+}-permeable transient receptor potential canonical (TrpC) family of channel proteins are known to play roles in cancer progression. For example, TrpC1, TrpC3, and TrpC6 play roles in tumor cell proliferation. TrpC1 and TrpC3 are involved in the proliferation of breast and ovarian cancer cells, respectively (3, 4), whereas TrpC6 is involved in the proliferation of liver, prostate, breast, and brain tumor cells (7–10). The roles of these channels in cancer may involve changes in [Ca^{2+}], (3). Previously, we found that TrpC5-mediated Ca^{2+} entry stimulates ABCB1 overproduction in drug-resistant human breast cancer cells, and inhibiting/suppressing TrpC5 reverses the drug resistance through the TrpC5-NFATC3-Ca^{2+}-ABCB1 signal pathway (11). However, to date there is still no report on TrpC5 channel involvement in drug resistance and/or ABCB1 production in human CRC cells. The detailed mechanism by which TrpC5 regulates ABCB1 expression also deserves exploration.

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The abbreviations used are: CRC, colorectal cancer; 5-Fu, 5-fluorouracil; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide; RFI, relative fluorescence intensity; CI, confidence interval; MDR, multidrug resistance.
Activation of the Wnt/Ca\(^{2+}\) pathway results in the mobilization of [Ca\(^{2+}\)]\(_i\) (12), which was recently found to prompt the nuclear accumulation of \(\beta\)-catenin (13), followed by activation of the transcription of T-cell factor/lymphoid enhancer factor-responsive genes (14), including \(\text{ABCB1}\) (15). Therefore, we designed experiments to explore the possible roles of \(\text{TrpC5}\) in regulating the nuclear translocation of \(\beta\)-catenin and \(\text{ABCB1}\) expression in the context of drug resistance.

**EXPERIMENTAL PROCEDURES**

**Cells and Cell Culture**—The human CRC cell line HCT-8 and 5-Fu-resistant HCT-8 cells (HCT-8/5-Fu) were from Keygen Biotech Co. Ltd (Nanjing, China). The human CRC cell line LoVo was from the Cell Resource Center of Shanghai Institutes for Biological Sciences, Type Culture Collection of the Chinese Academy of Sciences (Shanghai, China). 5-Fu-resistant LoVo cells (LoVo/5-Fu) were obtained by treating LoVo cells with stepwise increasing concentrations of 5-Fu over 6 months. HCT-8 and LoVo cells were cultured in RPMI 1640 supplemented with 10% (v/v) inactivated fetal calf serum, penicillin (100 units/ml), and streptomycin (100 \(\mu\)g/ml). HCT-8/5-Fu cells were cultured in RPMI 1640 supplemented with 5 mg/liter 5-Fu, and LoVo/5-Fu cells were cultured in RPMI 1640 supplemented with 5 mg/liter 5-Fu.

**Antibodies, siRNA, and Reagents**—The primary antibodies anti-\(\text{TrpC5}\) (ab63151), anti-\(\text{ABCB1}\) (ab10333), anti-\(\beta\)-catenin (ab6301), and anti-cyclin D1 (ab40754) were from Abcam Biotechnology (Cambridge, MA), anti-histone (AH433), and anti-\(\text{cyclin D1}\) (ab40754) were from Abcam Biotechnology. RIPA (P0013B) and the nuclear and cytoplasmic protein extraction kit. The same quality of total protein was loaded onto each lane. The proteins were electrophoresed on 8% polyacrylamide gel containing 0.1% SDS. The resolved proteins were semi-dry transferred to PVDF membrane. The primary antibodies anti-\(\text{TrpC5}\) (1:1000), anti-\(\text{ABCB1}\) (1:10), and anti-\(\beta\)-catenin (1:200) were used to detect expression of the \(\text{TrpC5}\), \(\text{ABCB1}\), and \(\beta\)-catenin proteins. The cells were visualized under the confocal laser scanning microscope (Leica TCS SP8, Wetzlar, Germany).

**Western Blot**—Whole cell protein was obtained using RIPA containing 1 mM PMSF. Nuclear protein was isolated using a nuclear and cytoplasmic protein extraction kit. The same quality of total protein was loaded onto each lane. The proteins were electrophoresed on 8% polyacrylamide gel containing 0.1% SDS. The resolved proteins were semi-dry transferred to PVDF membrane. The primary antibodies anti-\(\text{TrpC5}\) (1:500), anti-\(\text{ABCB1}\) (1:500), anti-\(\beta\)-catenin (1:1000), and anti-cyclin D1 (1:200) were used to detect the expression of the proteins of interest. \(\beta\)-Actin and histone were the internal references. The antigen-antibody complexes were visualized by an enhanced chemiluminescent reaction. Protein bands were analyzed using ImageJ software (National Institutes of Health, Bethesda, MD).

**Real Time PCR**—Total RNA was extracted from cells using TRIzol. cDNA was synthesized from RNA with oligo(dT) as primer and AMV reverse transcriptase. Ten microliters of 2X Sybrgreen PCR master mix. 0.4 \(\mu\)l of 50X 6-carboxy-X-rhodamine, 0.4 \(\mu\)l of forward and reverse primers, 2 \(\mu\)l of cDNA template, and 6.8 \(\mu\)l of double-distilled \(\text{H}_2\text{O}\) made up the quantitative PCR amplification system. The primer pairs are listed in Table 1. Samples were assayed in triplicate. The relative mRNA

**TABLE 1**

| Gene   | Forward primer sequence (5’-3’) | Reverse primer sequence (5’-3’) |
|--------|---------------------------------|---------------------------------|
| \(\beta\)-Actin | GCCCTTGCTCCCCCTCCCATTC | CGGAATCTCTGCTGACCCCATCT |
| \(\text{TrpC5}\) | CACCCAGCTACGATAGTGAAG | CGAAACAGCCAATTTACCC |
| \(\text{ABCB1}\) | ATTCCCTGATCAATGGTC | TAACAAAGGCACAGCTTAKG |

**TABLE 4**

| Gene     | Forward primer sequence (5’-3’) | Reverse primer sequence (5’-3’) |
|----------|---------------------------------|---------------------------------|
| \(\text{TrpC5}\) | ATGCTGGGACCTCCAGCTA | CCAAAACAGCCAATTTACCC |
| \(\text{ABCB1}\) | GGCTTGCTCCCCCTCCCATTC | CGGAATCTCTGCTGACCCCATCT |

**Amino Acid Co. Ltd (Tianjin, China).** The pCI-\(\text{TrpC5}\) plasmid was constructed in our laboratory (16).
levels of TrpC5 and ABCB1 were compared using the delta-delta cycle threshold (ΔΔCT) method (17). The β-actin ampli-
con was used as internal control for normalization.

Rhodamine 123 Efflux Assay—Confocal microscopy and flow cytometry (FACS) were used to evaluate the efflux activity of ABCB1. For indirect measurement of rhodamine 123 efflux from cells, HCT-8, HCT-8/5-Fu/Scrambled, and HCT-8/5-Fu/ RNAi cells were seeded in culture dishes. When the cells grew to 30–50% confluence, the medium was replaced with RMPI 1640 supplemented with 1 μM rhodamine 123. After 30 min of culture at 37 °C, the cells were fixed with 4.0% paraformalde-
hyde, washed twice with PBS, and visualized under a confocal laser scanning microscope (Leica TCS SP8). The rhodamine 123 signal was excited by 488-nm laser light, and emission was captured at 530 nm. Each field of cells was photographed to calculate the relative fluorescence intensity (RFI). For FACS analysis, HCT-8, HCT-8/5-Fu/Scrambled, and HCT-8/5-Fu/ RNAi cells were resuspended in fresh culture medium containing rhodamine 123 (final concentration, 1 μM). After incubation at 37 °C in the dark for 30 min, the cells were resuspended in cold PBS and analyzed by FACS using a flow cytometer (BD FACSCalibur). Data analysis was carried out with FLOWJO 7.6 software. At least three experiments were performed with each cell type.

[Ca^{2+}], Measurement—La^{3+} is known to potentiate TrpC5 activity but inhibit many other Ca^{2+}-permeable channels (11, 18). We used a Ca^{2+}-sensitive molecular construct, GECO1.2 (19). Normal physiological saline solution contained 140 mM NaCl, 5 mM KCl, 1 mM CaCl_2, 1 mM MgCl_2, 10 mM glucose, and 5 mM Hepes (pH 7.4) (15). GECO1.2 fluorescence signals were measured at room temperature using an Olympus fluorescence imaging system. Changes in [Ca^{2+}], were displayed as the fluorescence intensity relative to the value before La^{3+} application (100 μM, F_1/F_0). Genistein (50 μM) was included in the bath solution to potentiate TrpC5.

MTT Assay—The cytotoxic effect of 5-Fu was evaluated at 48 h with the MTT assay. Cells (1 × 10^4 cells/200 μl) were seeded in 96-well plates and placed in an incubator at 37 °C for 12 h. The medium was then replaced with medium containing different concentrations of 5-Fu. After 48 h, the medium was discarded, and each well received 200 μl of fresh medium containing 5 mg/ml MTT for 4 h. DMSO (150 μl) was added to each well, and then the absorbance was determined at 490 nm.

Statistical Analysis—The results are presented as means ± S.E. Statistical significance was determined by Student’s t test. A value of p < 0.05 was considered statistically significant.

RESULTS

Up-regulated Expression of TrpC5 and ABCB1 in 5-Fu Chemoresistant Human CRC Cells—Real time PCR, Western blot, and immunostaining were analyzed to determine TrpC5 and ABCB1 expression at the mRNA and protein levels in

FIGURE 1. A and B, real time PCR (A) and Western blot (B) showed much higher expression of both TrpC5 and ABCB1 at the mRNA and protein levels in HCT-8/5-Fu and LoVo/5-Fu cells than in their parental lines (HCT-8 and LoVo cells). *, p < 0.05 (Student’s t test). B, Western blot showed more nuclear-stabilized β-catenin accumulation in HCT-8/5-Fu and LoVo/5-Fu cells than in HCT-8 and LoVo cells. *, p < 0.05, (Student’s t test). C, immunofluorescent staining showed higher expression of TrpC5 and ABCB1 proteins and more nuclear-stabilized β-catenin accumulation in HCT-8/5-Fu cells than in HCT-8 cells. TrpC5 and ABCB1 proteins were mostly localized near the cell surface. Scale bars, 20 μm.
TrpC5 Regulates ABCB1 Production in Colorectal Cancer

HCT-8, HCT-8/5-Fu, LoVo, and LoVo/5-Fu cells. A much higher expression of both TrpC5 and ABCB1 at the mRNA and protein levels was found in HCT-8/5-Fu and LoVo/5-Fu cells, whereas only a low level of TrpC5 and ABCB1 expression was detected in their parental lines HCT-8 and LoVo (Fig. 1, A–C). Immunostaining showed that TrpC5 and ABCB1 were mostly localized near the cell surface (Fig. 1C).

Expression of TrpC5 Was Required for the Increase of $[\text{Ca}^{2+}]_i$ in HCT-8/5-Fu and LoVo/5-Fu Cells—Application of La$^{3+}$ (100 μM) elicited a rise in $[\text{Ca}^{2+}]_i$ in HCT-8/5-Fu (A) and LoVo/5-Fu (B) but not in HCT-8 and LoVo cells. The La$^{3+}$-elicited $[\text{Ca}^{2+}]_i$, rise in HCT-8/5-Fu and LoVo/5-Fu cells was inhibited by TSE3, a TrpC5-specific blocking antibody, pre IgG, preimmune IgG.

We next explored the possible mechanism by which $\beta$-catenin involved in the regulation of ABCB1 by TrpC5—TrpC5 is considered to increase $[\text{Ca}^{2+}]_i$, and this has been reported to facilitate $\beta$-catenin translocation into the nucleus (13). In HCT-8/5-Fu and LoVo/5-Fu cells, together with higher membrane expression of TrpC5, $\beta$-catenin showed more stable accumulation in the nucleus than in HCT-8 and LoVo cells (Fig. 1, B and C). Suppressing TrpC5 expression with TrpC5-specific siRNA in HCT-8/5-Fu cells resulted in decreased nuclear accumulation of $\beta$-catenin, and on the contrary, enforcing TrpC5 expression in HCT-8 (HCT-8/TrpC5) cells increased the nuclear accumulation of $\beta$-catenin (Fig. 4A).

We next explored the possible mechanism by which $\beta$-catenin involved in the regulation of ABCB1 by TrpC5. Cyclin D1 and ABCB1 are considered to be the target genes of $\beta$-catenin in the activated canonical Wnt/$\beta$-catenin signal pathway (15, 20–22). Western blot showed more cyclin D1 protein expression in the nucleus of HCT-8/5-Fu/DMSO than HCT-8 cells (Fig. 4B). Administration of XAV939 decreased the nuclear accumulation of $\beta$-catenin in HCT-8/5-Fu cells, decreased ABCB1 expression, and decreased the nuclear expression of cyclin D1. Also, the 5-Fu IC$_{50}$ decreased to 35.34 ng/ml (95% CI: 31.73–39.36 mg/liter). On the contrary, exposure to LiCl increased the nuclear accumulation of $\beta$-catenin, as well as the nuclear expression of cyclin D1 and ABCB1 expression in HCT-8 cells (Fig. 4B). Furthermore, up-regulation of TrpC5 expression in HCT-8 cells led to translocation of $\beta$-catenin to the nucleus and up-regulated expression of cyclin D1 and ABCB1 (Fig. 4C).

DISCUSSION

Resistance to chemotherapy occurs universally during malignant tumor treatment and contributes to most cancer-related deaths. The most common and standard strategy to overcome resistance is to use other cytotoxic agents, but this does not work in most cases because of multidrug resistance (MDR).

MDR tumor cells can become resistant to the drugs originally used to treat them and cross-resistant to other drugs with different mechanisms of action. MDR is believed to cause treatment failure in >90% of patients with metastatic cancer (23). A typical example is the outcome of the randomized GERCOR study (24), in which previously untreated advanced CRC patients received FOLFIRI (Irinotecan, 5-Fu, and leucovorin).
followed by FOLFOX (oxaliplatin, 5-Fu, and leucovorin) or the reverse sequence. Although both regimes achieved response rates of \( \geq 50\% \) in first line therapy, considerably fewer patients (4 and 15\%) responded to second line therapy with both regimes. A reasonable explanation for this is MDR.

ABCB1, also known as P-glycoprotein, coded by the ABCB1/MDR1 gene, is an ATP-dependent membrane transport protein, and its high expression in cancer cells is considered to be the major cause of MDR and the failure of chemotherapy. ABCB1 has been a research focus for overcoming resistance, but many ABCB1 antagonists have failed, because of either toxicity or intolerable pharmacokinetic interactions (2).

\( \left[ \text{Ca}^{2+} \right]_{i} \) is involved in the transcriptional regulation of ABCB1 and is a promising target for modulating ABCB1 production (19, 25, 26). \( \text{Ca}^{2+} \) signaling is indispensable for maintaining physiological functions in all living cells. Studies linking \( \text{Ca}^{2+} \) signaling to nonmalignant tumors have emerged (as well as cardiovascular (27) and neurological diseases (28)) over

**FIGURE 3.** A–C, real time PCR (A), Western blot (B), and immunostaining (C) showed that both TrpC5 and ABCB1 expression at the mRNA and protein levels were significantly down-regulated in HCT-8/5-Fu/RNAi and LoVo/5-Fu/RNAi cells compared with scrambled siRNA in HCT-8/5-Fu and LoVo/5-Fu cells. *, \( p < 0.05 \) (Student’s t test). D, confocal laser scanning showed that the fluorescence intensity in HCT-8/5-Fu/Scrambled cells was lower than in HCT-8 cells and increased after inhibition of TrpC5 with siRNA (green fluorescence indicates rhodamine 123, an ABCB1 substrate). E, FACS analysis showed low accumulation of rhodamine 123 immunofluorescence in HCT-8/5-Fu/Scrambled cells compared with HCT-8 cells, and this increased after inhibition of TrpC5 with siRNA. F, MTT assay showed that HCT-8/5-Fu/Scrambled and LoVo/5-Fu/Scrambled cells were much more resistant to 5-Fu-induced cell death than HCT-8 and LoVo/5-Fu cells. Administration of TrpC5-siRNA caused a remarkable reversal of 5-Fu resistance. *, \( p < 0.05 \) (Student’s t test). Scale bars, 20 \( \mu \text{m} \). Scrambled, scrambled siRNA.
many years, whereas only recently, increasing numbers of studies have been concerned with Ca\textsuperscript{2+}/H\textsubscript{11001} signaling in malignant tumors and have demonstrated its involvement in many key aspects of cancer development and progression. Among these, Trp channels have received the most attention. Trp channels are expressed in a wide variety of tissues and have many functions including distinguishing sensations such as pain, temperature, taste, and pressure (29). Recently, abnormal expression of Trp channels has been found in many tumors and is considered to participate in the biological behavior of cancer cells, including invasion (30, 31), proliferation (6), differentiation (32), and tumor vascularization (33).

As Ca\textsuperscript{2+} entry channels, Trp channels have recently been associated with ABCB1 induction. TrpC5 is essential for ABCB1 overproduction in adriamycin-resistant breast cancer cells (MCF-7/ADM) via the TrpC5-Ca\textsuperscript{2+}/NFATc3-ABCB1 signaling cascade (11), and the transfer of TrpC5-containing extracellular vesicles from MCF-7/ADM cells to the parental line MCF-7/WT cells allows recipient cells to acquire TrpC5, consequently stimulating ABCB1 production and thus conferring chemoresistance on nonresistant cells (16).

In this study, ABCB1 expression was higher in HCT-8/5-Fu and LoVo/5-Fu cells than in the parental HCT-8 and LoVo cells, consistent with previous studies (34, 35). Also, TrpC5 was
overexpressed in HCT-8/5-Fu and LoVo/5-Fu cells compared with HCT-8 and LoVo cells. Application of La3+/H11001 elicited a rise in [Ca2+]i in HCT-8/5-Fu and LoVo/5-Fu but not in HCT-8 and LoVo cells, and this was inhibited by T5E3, indicating that TrpC5 in HCT-8/5-Fu and LoVo/5-Fu cells has constitutive activity that is required for the increase of [Ca2+]i. We next knocked down TrpC5 expression in HCT-8/5-Fu and LoVo/5-Fu cells using TrpC5-siRNA. As a result, the mRNA and protein levels of ABCB1 decreased significantly. Rhodamine 123, a fluorescent cationic dye, is an established substrate for ABCB1. It has been used as a molecular probe in studies pertaining to MDR phenotypes (36, 37). We assessed the RFI of rhodamine 123 in cells by confocal microscopy and found that the RFI in HCT-8 cells was markedly higher than in HCT-8/5-Fu cells, and the RFI in HCT-8/5-Fu cells increased after TrpC5 was down-regulated by siRNA. MTT assays showed that inhibition of TrpC5 using TrpC5-siRNA reversed the 5-Fu resistance in HCT-8/5-Fu and LoVo/5-Fu cells. These results showed that overexpression of ABCB1 contributes to the resistance of HCT-8/5-Fu and LoVo/5-Fu cells to 5-Fu and is modulated by Ca2+ entry through TrpC5. In this process, TrpC5 is the key modulator that regulates Ca2+ entry and ABCB1 expression. This is in accord with our prior findings, in which we identified the role of TrpC5 and ABCB1 in breast cancer cell resistance to adriamycin (11).

Wnt-related signaling includes the canonical Wnt/β-catenin and the noncanonical Wnt/Ca2+ pathways. In the activated canonical Wnt/β-catenin pathway, phosphorylation of glycogen synthase kinase 3β results in the down-regulation of glycogen synthase kinase 3β activity, decreased β-catenin degradation in the cytoplasm, and subsequently the accumulation of β-catenin in the nucleus (38), where it binds to lymphoid enhancer factor/T-cell factor and activates Wnt target genes including cyclin D1 and ABCB1 (15, 20–22). Activation of canonical Wnt/β-catenin signaling has been demonstrated to induce chemoradioresistance in cancer cells (39–41). Recently, it was shown to be associated with up-regulation of ABCB1 and to induce MDR in 5-Fu-resistant human cholangiocarcinoma cells (22). In the present study, we found the same phenomenon in the 5-Fu-resistant human CRC cells HCT-8/5-Fu.

The two Wnt-related signaling pathways were thought to operate independently (42), but recent studies indicated that they act in a coordinated manner (13). Activation of the Wnt/Ca2+ pathway results in an increase of [Ca2+]i and depolarization of both the cell and nuclear membranes, subsequently promoting β-catenin translocation into the nucleus. This indicates a novel pathway through which TrpC5 regulates the induction of ABCB1. As a nonselective Ca2+-permeant cation channel, TrpC5 could modulate Ca2+ entry into the cytoplasm. Thus, overexpression of TrpC5 might increase [Ca2+]i, cause the nuclear translocation of β-catenin, and induce ABCB1 gene production. This hypothesis was validated by our studies. TrpC5 overexpression and greater accumulation of stabilized β-catenin in the nucleus were simultaneously found in 5-Fu-chemoresistant cells (HCT-8/5-Fu and LoVo/5-Fu cells). Furthermore, knockdown of TrpC5 by TrpC5-siRNA caused a remarkable decrease in the nuclear accumulation of β-catenin in HCT-8/5-Fu cells, whereas enforcing TrpC5 expression increased the nuclear accumulation of β-catenin in HCT-8 cells. We carried out additional experiments and demonstrated that the Wnt target genes cyclin D1 and ABCB1 were regulated by the nuclear accumulation of β-catenin in HCT-8 and HCT-
8/5-Fu cells. Importantly, we found that enforcing TrpC5 expression resulted in the increased nuclear accumulation of β-catenin and subsequent up-regulated expression of the Wnt target genes cyclin D1 and ABCB1. Thus, we hypothesize the existence of a TrpC5-Ca^{2+}-β-catenin-ABCB1 signal pathway (Fig. 5).

In our previously study (11), the TrpC5-Ca^{2+}-NFATc3-ABCB1 signal pathway was identified as the mechanism by which TrpC5 regulates ABCB1 production. It has been reported that inhibition of glycerogen synthase kinase 3β promotes the accumulation of both NFATc3 and β-catenin in the nucleus (43, 44), but to date no link between NFATc3 and β-catenin has been demonstrated. Thus, the TrpC5-Ca^{2+}-β-catenin-ABCB1 signal pathway may be a novel means by which TrpC5 regulates ABCB1 production.

It is generally acknowledged that ABCB1 overexpression contributes to most of the chemotherapy resistance. In addition to our previous findings in breast cancer, TrpC5 was demonstrated to be essential in the induction of ABCB1-mediated resistance to 5-Fu in CRC cells. The underlying mechanism involves the translocation of β-catenin into the nucleus facilitated by an increase of [Ca^{2+}], triggered by TrpC5. These findings help to understand the complicated underlying mechanism of MDR. Moreover, our study puts forward the idea that TrpC5 may be a target for the reversal of clinical CRC chemoresistance.

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