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Flightless-1 inhibits ER stress-induced apoptosis in colorectal cancer cells by regulating Ca²⁺ homeostasis

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Abstract

The endoplasmic reticulum (ER) stress response is an adaptive mechanism that is activated upon disruption of ER homeostasis and protects the cells against certain harmful environmental stimuli. However, critical and prolonged cell stress triggers cell death. In this study, we demonstrate that Flightless-1 (Flil) regulates ER stress-induced apoptosis in colon cancer cells by modulating Ca²⁺ homeostasis. Flil was highly expressed in both colon cell lines and colorectal cancer mouse models. In a mouse xenograft model using CT26 mouse colorectal cancer cells, tumor formation was slowed due to elevated levels of apoptosis in Flil-knockdown (Flil-KD) cells. Flil-KD cells treated with ER stress inducers, thapsigargin (TG), and tunicamycin exhibited activation of the unfolded protein response (UPR) and induction of UPR-related gene expression, which eventually triggered apoptosis. Flil-KD increased the intracellular Ca²⁺ concentration, and this upregulation was caused by accelerated ER-to-cytosolic efflux of Ca²⁺. The increase in intracellular Ca²⁺ concentration was significantly blocked by dantrolene and tetracaine, inhibitors of ryanodine receptors (RyRs). Dantrolene inhibited TG-induced ER stress and decreased the rate of apoptosis in Flil-KD CT26 cells. Finally, we found that knockdown of Flil decreased the levels of sorcin and ER Ca²⁺ and that TG-induced ER stress was recovered by overexpression of sorcin in Flil-KD cells. Taken together, these results suggest that Flil regulates sorcin expression, which modulates Ca²⁺ homeostasis in the ER through RyRs. Our findings reveal a novel mechanism by which Flil influences Ca²⁺ homeostasis and cell survival during ER stress.

Introduction

Colorectal cancer (CRC) is one of the most common lethal cancers around the world¹. The disease is characterized by local recurrence, distant metastasis, and extremely rapid progression¹. Surgical resection, systemic chemotherapy, and radioembolism are the major

strategies for treating CRC, but high mortality and poor prognosis remain serious clinical problems¹. The main objective of cancer therapy is to destroy all cancer cells while causing minimal damage to normal tissue. Apoptosis, a genetically regulated form of programmed cell death, is an emerging target for anticancer therapy². In general, the activation of apoptosis is an important mechanism for the prevention and treatment of CRC, as well as other cancers². Three different pathways lead to apoptosis: the extrinsic death receptor pathway, the intrinsic mitochondrial pathway, and the endoplasmic reticulum (ER) stress pathway². The mechanisms involved in the intrinsic and ER stress-mediated pathways influence each other. Accordingly, the ER stress pathway

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represents an emerging the rapeutic target for cancer treatment 3 .

The ER is a multifaceted organelle involved in initial protein maturation, lipid synthesis, and maintenance of intracellular calcium homeostasis⁴. Disruption of ER homeostasis following glucose deprivation, hypoxia, disruption of Ca²⁺ homeostasis, or accumulation of misfolded proteins activates the unfolded protein response (UPR) and initiates an intracellular signaling pathway that protects the cell⁴. Under normal conditions, three ER stress sensors, protein kinase-like ER kinase (PERK), activating transcription factor 6 (ATF6), and inositolrequiring kinase-1 (IRE1), are bound to the ER-resident chaperone glucose-regulated protein-78 (GRP78/BiP)⁵. However, under ER stress, these proteins are released from GRP78/BiP and initiate their downstream cascades. IRE1α initiates the splicing of X-box binding protein-1 (XBP-1) mRNA. Spliced XBP-1 functions as a transcriptional activator of UPR target genes, including GRP78/BiP and calreticulin⁵. PERK is a transmembrane kinase that phosphorylates and inactivates eukaryotic translation initiation factor 2 subunit α , leading to reduced protein synthesis⁵. The PERK/EIF1α pathway also activates ATF4, which upregulates CCAAT/enhancer-binding proteinhomologous protein (CHOP). Concomitantly, during ER stress, cleaved and activated ATF6α translocates to the nucleus and transactivates the genes encoding various chaperones and ER stress markers, including CHOP and GRP78/BiP itself. Activation of the UPR promotes cell survival by decreasing protein synthesis and increasing protein folding capacity. However, if ER stress persists and homeostasis cannot be restored, the UPR triggers cell death in cells that are beyond repair⁶.

Ca²⁺ is a ubiquitous intracellular messenger that controls multiple cellular functions, including transcription, exocytosis, apoptosis, and proliferation. Ca2+ is mainly stored in the ER, and its levels are tightly regulated by multiple pumps, channels, and binding proteins in that organelle⁷. Ca²⁺ movement across the ER membrane is mediated by Ca²⁺ release channels, ryanodine receptors (RyRs), ER-resident inositol 1,4,5-trisphosphate receptor (IP₃Rs), and the Ca²⁺ uptake pump sarco-ER Ca²⁺-ATPase (SERCA)⁸⁻¹⁰. However, despite tight regulation of ER Ca2+, alterations in ER Ca2+ homeostasis due to dysfunction of these proteins or critical and continuous ER stress provoke ER Ca²⁺ depletion and an overload of intracellular Ca2+, resulting in excessive Ca²⁺ accumulation in the mitochondria. This, in turn, causes apoptosis by increasing mitochondrial membrane permeabilization and promoting the release of cytochrome c^{11} .

Flightless-1 (FliI), originally identified as a *Drosophila melanogaster* mutant, is a member of the gelsolin superfamily with an N-terminal leucine-rich repeat domain and

a C-terminal gelsolin-like domain¹². Through its bipartite domain structure, FliI can bind to numerous structural and signaling proteins and thus regulate cell migration, wound healing, and inflammation 13-17. The main roles of the gelsolin family are Ca²⁺- and phosphatidylinositol 4,5bisphosphate-regulated actin binding 18. However, FliI is more divergent from gelsolin than other family members, and its actin-binding and actin-severing activities are Ca²⁺ independent¹⁹. By contrast, FliI interacts in a Ca²⁺-dependent manner with nonmuscle myosin IIA, which plays an essential role in cell extension by activating transient receptor potential cation channel subfamily V member 4²⁰. Furthermore, FliI modulates cell proliferation and survival in cancer cells by interacting with transcription factors such as androgen receptor, estrogen receptor (ER), and carbohydrate responsive elementbinding protein, which regulate tumor progression in prostate cancer and CRC cells^{21–23}. Recently, FliI was shown to promote breast cancer progression by impeding selective autophagy through an interaction with $p62^{24}$. Here, we report a novel function of FliI: FliI suppresses ER stress-induced UPR signaling and apoptosis in colon cancer by regulating Ca²⁺ homeostasis through modulation of RyR activity.

Materials and methods

Cell culture, stable cell line generation

CT26 (ATCC: CRL-2368) cells were cultured in RPMI 1640 medium supplemented with 10% fetal bovine serum (GIBCO BRL, Grand Island, NY, USA) and antibiotics (100 U/ml penicillin and 100 ug/ml streptomycin). Cells were grown at 37 °C under a humidified 5% CO₂ atmosphere. The sequence used for the lentiviral shRNA expression vector (pLKO.1; Open Biosystems, Huntsville, AL, USA) targeting FliI was 5′-TTCTAGGTTGTT GTTGGCAGC-3′. For lentivirus production, HEK-293T cells (ATCC; Manassas, VA, USA) were transfected with 10 μg lentiviral vectors. Following infection with lentivirus, cells were selected with 1 μg/ml puromycin.

Live-cell imaging for intracellular calcium

shRNA-Ctrl or shRNA-FliI CT26 cells were incubated at 37 °C for 30 min in media containing 1 μ M Fluo-4AM (Invitrogen, Carlsbad, CA, USA). After washing with Hank's buffer, cells were analyzed by flow cytometry or imaged on a fluorescence microscope (Olympus, 20×), with excitation and emission wavelengths of 488 and 505 nm, respectively. For analysis using GCaMP6s, shRNA-Ctrl or shRNA-FliI CT26 cells were seeded on a cover glass and transfected with pDEST-mCherry-GCaMP6s using Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA). After 24 h, intracellular Ca²⁺ was imaged on an IX83 microscope (Olympus) equipped with an Olympus 40× objective lens (oil, NA 1.30), a fluorescent

lamp (Olympus), a stage controller (LEP), and a CCD camera (Andor, Concord, MA, USA). Images were processed with MetaMorph software (Molecular Devices, San Jose, CA, USA). For ratiometric Ca²⁺ imaging, cells were pretreated for 2 h with dantrolene (50 µM; Sigma-Aldrich, St. Louis, MO, USA), 2-APB (50 µM; Sigma-Aldrich, St. Louis, MO, USA), tetracaine (50 µM; Sigma-Aldrich), DBHO (2,5-di-tert-butylhydroguinone; 1 µM; Santa Cruz Biotechnologies, Santa Cruz, CA, USA), and loaded with 1 μM Fura-2AM (Molecular Probes) for 30 min. Ratiometric Ca²⁺ imaging at 340 and 380 nm was performed at room temperature in calcium-free Tyrode's solution (129 mM NaCl, 5 mM KCl, 3 mM MgCl₂, 30 mM glucose, and 25 mM HEPES [pH 7.4]) with or without 5 µM ionomycin, 1 μM thapsigargin (TG), and 50 μM DBHQ on an IX81 microscope equipped with an Olympus 40× objective lens (oil, NA 1.30), a fluorescent arc lamp (Lambda LS), an excitation filter wheel (Sutter, Lambda 10-2), a stage controller (ASI, MS-2000), and a CCD camera (Hamamatsu, C10600). Images were processed using MetaMorph software (Molecular Devices) and analyzed using Igor Pro software (WaveMetrics, Portland, OR, USA).

Analysis of cell viability and apoptosis induction

shRNA-Ctrl or shRNA-FliI CT26 cells (1 × 10⁴ cells/ well) were cultured in 96-well plates overnight and then treated with TG for 48 h. To determine cell viability, cells were exposed to 3-(4,5-dimethyl-thiazol-2-yl)2,5 diphenyltetrazolium bromide (MTT; Sigma-Aldrich), and crystallized formazan was quantified by measuring absorbance at 595 nm on an Infinity M200 microplate reader (Tecan, Männedorf, Switzerland). Absorbance data were normalized against the vehicle control and expressed as percent viability. Alternatively, after treatment with TG, shRNA-Ctrl or shRNA-FliI CT26 cells were stained with annexin V-FITC (Invitrogen) and propidium iodide (PI; Invitrogen) and quantified on a FACSCalibur system (BD Biosciences, San Jose, CA, USA). Data were processed using FlowJo software (BD Biosciences). For colony formation assays, shRNA-Ctrl or shRNA-FliI CT26 cells $(1 \times 10^3 \text{ cells/well})$ were treated with TG for 48 h, after which the medium was replaced with drug-free medium, and the cells were incubated for 14 days. The colonies were washed with PBS, fixed with methanol, and stained with crystal violet.

Transient transfection of plasmid DNA

Mouse sorcin expression plasmids were purchased from Sino Biological (BDA, Beijing, China). Plasmid DNA was transiently transfected into FliI-KD CT26 cell lines using jetOptimus (Polyplus-transfection, Illkirch-Graffenstaden, France). Cells were treated with TG at the indicated concentration to induce ER stress.

ER Ca²⁺ measurement

For ER calcium measurements, cells were transiently transfected with a plasmid encoding the D1ER Cameleon. pcDNA-D1ER was a gift from Amy Palmer & Roger Tsien (Addgene plasmid #36325)²⁵. ZEN (Zeiss, Oberkochen, Germany) was used to calculate fluorescence intensities from images taken under three conditions: IDD, donor fluorescence intensity excited by an excitation laser (458 nm); IDA, acceptor intensity under a donor excitation laser; and IAA, acceptor intensity excited by an acceptor excitation laser (514 nm). The FRET efficiency is defined as follows: ²⁶ FRET efficiency = $(I_{DA} - \beta \times I_{DD} - \beta)$ $\gamma \times I_{AA}$)/ I_{AA} , where β is a bleeding-through coefficient from donor to acceptor channels and γ is the ratio of fluorescence intensity of acceptor molecules directly excited by a donor excitation laser to fluorescence intensity of acceptor molecules excited by an acceptor excitation laser.

Immunoblotting

Samples were lysed in RIPA lysis buffer (50 mM Tris, pH 8.0, 150 mM NaCl, 1% NP-40, 0.1% SDS, and 0.25% N-deoxycholate) containing protease and phosphatase inhibitors (Sigma-Aldrich). Equal amounts of protein were separated by SDS-PAGE and transferred onto nitrocellulose membranes (GE Healthcare, MA, USA). Membranes were blocked in 5% bovine serum albumin blocking buffer and incubated at 4°C overnight with specific primary antibodies against phospho-PERK, phospho-IREα, GRP78/BiP, CHOP (Cell Signaling Technology, Danvers, MA, USA), sorcin, and FliI (Abcam, Cambridge, MA, USA). Signals were detected using an ECL detection kit (GE Healthcare, Chicago, IL, USA) and subsequently incubated with horseradish peroxidaseconjugated secondary antibodies (Thermo Fisher Scientific, Waltham, MA, USA).

Gene expression analysis

Total RNA was isolated from cells or tissues using TRIzol reagent (Invitrogen). RNA was reverse transcribed using the ABI Reverse Transcription kit. Quantitative PCR was performed with SYBR green fluorescent dye on an ABI 9300 PCR instrument. Relative mRNA expression was determined by the $\Delta\Delta$ Ct method and normalized against TBP mRNA expression.

Colon cancer models

Colon tissues from *APC*^{min+}- and colitis-induced mouse models of CRC were prepared as previously described^{27,28}. Briefly, for colitis-induced CRC, 7- to 8-week-old male mice were intraperitoneally injected with azoxymethane (AOM, 10 mg/kg body weight) and maintained on a regular diet and water for 7 days. Mice were then subjected to five cycles of dextran sodium sulfate (DSS) treatment, in

which each cycle consisted of 1.5% DSS for 7 days followed by a 7-day recovery period with regular water.

Xenograft tumor models

All experiments involving animals were approved by UNIST (IACUC-12-003-A). shRNA-Ctrl or shRNA-FliI CT26 cells (1×10^6) suspended in sterile PBS ($200~\mu$ l) were injected subcutaneously into the left flanks of 6-week-old BALB/c *nu/nu* male mice (SLC Inc., Hamamatsu, Shizuoka, Japan). Tumor sizes were measured using calipers every 2 days for 17 days. Tumor volume was calculated using the following formula: $V=1/2\times(\text{width})^2\times\text{length}$. At the end of the experiment, the animals were euthanized, and tumors were collected for western blotting.

Statistical analysis

Data are presented as the mean \pm standard errors of the mean as indicated in the figure legends. Comparisons between two groups were made by unpaired two-tailed Student's t tests. p values < 0.05 were considered statistically significant. Microsoft Excel was used for statistical calculations.

Results

Flil is upregulated in CRCs

To investigate the clinical relevance of FliI in CRC, we first examined FliI expression in both CRC cell lines and cancer tissue in the $\widehat{APC}^{\min+}$ - and colitis-induced mouse model of CRC. FliI expression in CRC cell lines and cancer tissues was significantly higher than that in the corresponding normal cell line and nontumor tissues (Fig. 1a, b). To directly assess the cell-autonomous function of FliI in mouse CRC, CT26 cells stably transduced with either shRNA-Ctrl or shRNA-FliI lentiviral shRNAs were implanted subcutaneously into immunodeficient mice to form xenograft tumors. In accordance with the results shown in Fig. 1b, knockdown (KD) of FliI significantly repressed xenograft tumor growth (Fig. 1c). Moreover, FliI-KD tumors had elevated levels of cleaved PARP-1 and caspase-3 (Fig. 1d), suggesting that the reduced level of FliI promoted apoptosis in CT26 cells.

Flil inhibits ER stress-induced apoptosis in CT26 cells

The microenvironment within a solid tumor differs from that of normal tissues; the former is characterized by glucose deprivation, low pH, hypoxia, and an imbalance between production and removal of reactive oxidative stress²⁹. All of these environmental factors contribute to ER stress; therefore, cancer cells must find effective ways to adapt and prevent ER stress-induced apoptosis³. To elucidate the molecular mechanisms by which FliI-KD promotes apoptosis in the xenograft mouse model, we first determined the effect of FliI on cell survival after

treatment with the ER stress-inducing drug TG. Cell viability was assessed by MTT and cell counting assays, which revealed that depletion of FliI decreased cell survival following TG treatment (Fig. 2a, b). Colony forming assays confirmed that FliI was required for cell survival following TG treatment (Fig. 2c). Furthermore, FACS analysis using annexin V-PI labeling revealed that depletion of FliI increased ER stress-induced apoptosis (Fig. 2d). Consistent with the FACS analysis, depletion of FliI increased the levels of cleaved PARP-1 and caspase-3 following TG treatment (Fig. 2e). Another ER stress-inducing drug, tunicamycin (TM), significantly increased apoptosis in FliI-KD cells (Supplementary Fig. 1). Together, these data indicate that FliI inhibits ER stress-induced apoptosis.

Knockdown of Flil sensitizes cells to ER stress

Disturbances in normal ER functions lead to accumulation and aggregation of unfolded proteins, which initiates an adaptive response, the UPR, to restore normal ER function⁶. Failure to activate the adaptive response results in apoptosis³⁰. To obtain insights into the molecular mechanism of action of FliI in ER stress-induced apoptosis, we first evaluated the effect of FliI on the ER stress response. As shown in Fig. 3a, within 3 h of TG treatment, phosphorylation of PERK and IREα was triggered more strongly in FliI-KD cells than Ctrl cells. In addition to UPR sensor protein activation, GRP78/BiP and CHOP levels were also elevated in FliI-KD cells. Similar results were observed after treatment of FliI-KD cells with TM (Supplementary Fig. 2a). Furthermore, the expression of UPR genes, including GRP78/BiP, ATF4, ATF6, and CHOP, was significantly elevated in FliI-KD cells after treatment with TG or TM (Fig. 3b and Supplementary Fig. 2b). These results indicate that KD of FliI made CT26 cells more sensitive to ER stress.

Knockdown of Flil promotes ER stress-induced apoptosis by disrupting intracellular Ca²⁺ homeostasis

Ca²⁺ is a major player in the regulation of apoptosis, both at the early and late stages³¹. Disruption of intracellular Ca²⁺ homeostasis can induce ER stress-mediated apoptosis in response to various pathological conditions^{32–34}. Therefore, we next investigated whether ER stress-induced apoptosis sensitized by KD of FliI influence the intracellular Ca²⁺ concentration. Using Fluo-4AM, a calcium-sensing dye, we observed that depletion of FliI triggered upregulation of intracellular Ca²⁺ (Fig. 4a, b). Similarly, using GCaMP6s, the Ca²⁺ indicator, we found that intracellular Ca²⁺ was significantly increased in FliI-KD cells compared with Ctrl cells (Fig. 4c). Furthermore, 1,2-bis(o-aminophenoxy)ethane-N,N,-N',N'-tetraacetic acid-acetoxymethyl ester (BAPTA-AM), a highly selective Ca²⁺ chelator, abrogated TG-induced apoptosis in FliI-

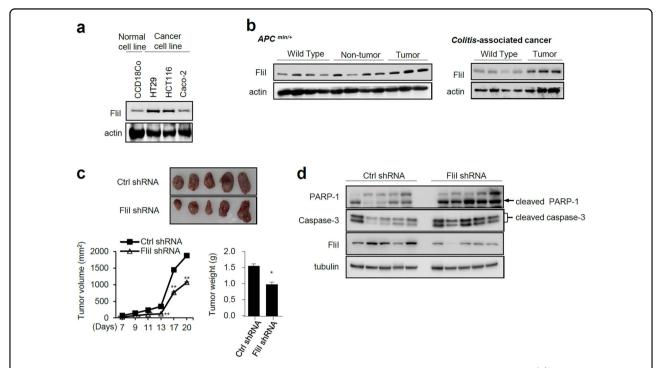


Fig. 1 Flil is overexpressed in colorectal cancer. Flil expression in a colon cancer cell line (**a**) and colon tissue from $APC^{\min/+}$ and colitis-associated cancer mouse models (**b**) were analyzed by western blotting. **c** Orthotopic xenografts of Ctrl or Flil-knockdown CT26 cells were detected (n = 5). Images show tumors dissected from the mice. Tumor volume was measured every 2 days, and tumor weight was measured after resection at the end of the experiment. Data are shown as the means \pm SEM. *p < 0.05 compared with shRNA-Ctrl cells. **d** Protein levels of cleaved PARP-1, cleaved caspase-3, and Flil were detected in mouse tumor tissue derived by western blotting.

KD CT26 cells (Fig. 4d). These results strongly suggest that elevation of intracellular Ca²⁺ decreases the cellular threshold for TG-induced apoptosis in FliI-KD cells.

Knockdown of Flil promotes ER Ca²⁺ release through RyR and alteration of intracellular Ca²⁺, resulting in cell death

The ER is the primary site of intracellular Ca²⁺ storage, and ER Ca2+ depletion is the major cause of both ER stress and ER stress-induced apoptosis³⁵. Hence, we postulated that the ER contributes to the increase in intracellular Ca²⁺ caused by FliI-KD. To test this idea, we determined the concentration of free Ca²⁺ within the ER by depleting the ER Ca²⁺ store with the ionophore ionomycin. As shown in Fig. 5a, ionomycin treatment increased intracellular Ca²⁺, and the ionomycin-mediated intracellular Ca²⁺ increase was lower in FliI-KD cells than in control cells. In addition, we measured cytosolic Ca²⁺ released from the ER by treatment with TG or DBHQ, a specific inhibitor of SERCA, to deplete ER Ca2+. Consistent with the results in Fig. 5a, the intracellular Ca²⁺ concentration was lower in TG- or DBHQ-treated FliI-KD cells (Supplementary Fig. 3a, b). Furthermore, when we exposed FliI-KD cells to a low dose (1 μM) of DBHQ for short (30 min) and long periods (3 h), ER Ca²⁺ depletion of FliI-KD was observed to occur in a timedependent manner (Supplementary Fig. 3c), clearly indicating that the elevation in intracellular Ca^{2+} in FliI-KD cells was caused by the release of ER Ca^{2+} . Finally, we directly measured the ER Ca^{2+} concentration using an ER-targeted Cameleon (D1ER) probe. As expected, ER Ca^{2+} was lower in FliI-KD cells than in control cells (Fig. 5b)

Regulation of intracellular Ca²⁺ by ER is mainly mediated by Ca²⁺ reuptake into the ER through SERCA Ca²⁺ pumps and Ca²⁺ release through Ca²⁺ channels, including IP₃Rs or RyRs⁸⁻¹⁰. Therefore, we next investigated whether the increase in intracellular Ca²⁺ in FliI-KD cells was mediated by ER Ca²⁺ channels. For these experiments, we used two different Ca²⁺ channel blockers, dantrolene, tetracaine, and 2-aminoethoxydiphenyl borate (2-APB), which are inhibitors of RyRs and IP₃Rs, respectively. Dantrolene and tetracaine elevated the ionomycininduced increase in intracellular Ca2+ in FliI-KD cells, whereas 2-APB did not (Fig. 5a). Dantrolene and tetracaine also elevated TG-induced and DBHQ-induced intracellular Ca²⁺ increases in FliI-KD cells (Supplementary Fig. 3a, b). Consistent with the results of intracellular Ca²⁺, ER Ca²⁺ was also recovered by treatment with dantrolene and tetracaine in FliI-KD cells (Fig. 5b). Furthermore, dantrolene and tetracaine inhibited TG-

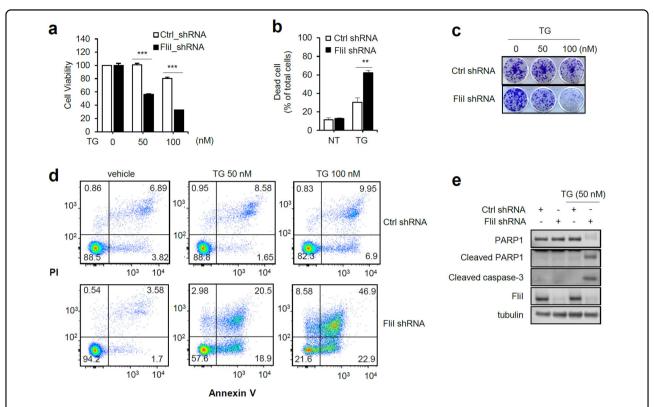


Fig. 2 Knockdown of Flil promotes TG-induced apoptosis in CT26 cells. Cell viability of Flil-KD CT26 cells was measured by MTT assay (**a**), cell counting assay (**b**), and colony assay (**c**) after treatment with TG for 48 h. Data are shown as the mean \pm SEM. **p < 0.01; ****p < 0.001 shRNA-Ctrl cells vs. shRNA-Flil cells. After Ctrl- and Flil-KD cells were treated with TG for 48 h, they were subjected to annexin V-Pl flow cytometry assay (**d**) and western blotting for PARP-1, cleaved caspase-3, Flil, and tubulin (**e**). TG thapsigargin.

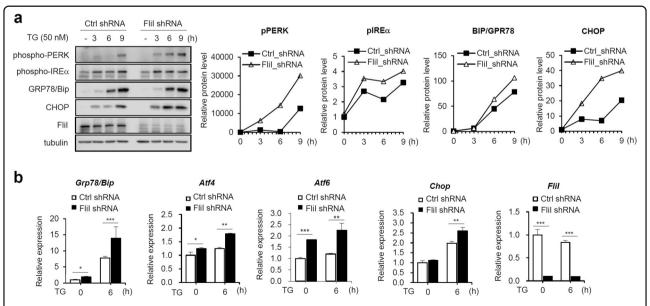


Fig. 3 Knockdown of Flil sensitizes CT26 cells to TG-induced UPR. Ctrl- and Flil-KD cells were treated with 50 nM TG for the indicated times, and the extracts were analyzed by western blotting with antibodies against phospho-PERK, phospho-IREα, GRP78/BiP, CHOP, Flil, and tubulin (a). mRNA expression was analyzed by quantitative real-time PCR (**b**). Data are shown as the mean \pm SEM (n = 3). *p < 0.05; **p < 0.01; ****p < 0.001, shRNA-Ctrl cells vs. shRNA-Flil cells. TG thapsigargin.

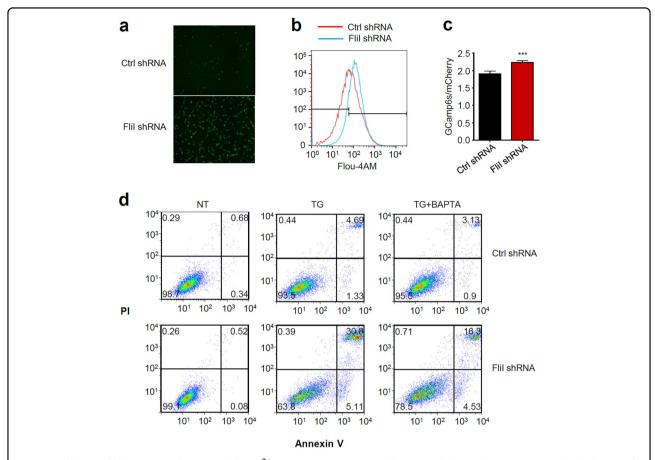


Fig. 4 Knockdown of Flil increases the intracellular Ca²⁺ concentration in CT26 cells. Ctrl- and Flil-KD cells were incubated with Fluo-4AM for 30 min, and intracellular Ca²⁺ was (**a**) imaged on a fluorescence microscope (200×) and (**b**) measured by FACS analysis. **c** Ctrl- and Flil-KD cells were transfected with mCherry-GCaMP6s and imaged on a fluorescence microscope. Intracellular Ca²⁺ levels were monitored based on the ratio of GCaMP6s to mCherry fluorescence intensity. **d** Ctrl- and Flil-KD cells were pretreated with BAPTA (10 μ M) and treated with TG (100 nM) for 48 h, and then apoptosis was detected by annexin V-Pl flow cytometry assay. TG thapsigargin.

induced expression of ER stress marker proteins (Fig. 5c and Supplementary Fig. 4a). We then tested whether RyR-mediated Ca²⁺ release in FliI-KD cells was associated with TG-induced apoptosis. As shown in Fig. 5d and Supplementary Fig. 4b, inhibition of RyR compromised the increase in TG-induced apoptosis caused by FliI-KD. In addition, the TG-induced increases in the levels of cleaved PARP-1 and caspase-3 in FliI-KD cells were rescued by dantrolene treatment (Fig. 5e). Together, these results strongly suggest that RyR channel-dependent ER Ca²⁺ release is an essential upstream event in the sensitization of FliI-KD cells to TG-induced apoptosis.

Knockdown of Flil suppresses the expression of sorcin, which protects against ER stress

To investigate the molecular mechanism by which FliI regulates ${\rm Ca}^{2+}$ release through RyRs, we compared the global gene expression profile using RNA sequencing. We focused on sorcin (soluble resistance-related calciumbinding protein) downregulation by FliI-KD (Fig. 6a)

because sorcin has been reported to negatively regulate ER Ca²⁺ release by mediating RyR and SERCA^{36,37}. Then, we hypothesized that downregulated sorcin contributes to the depletion of ER Ca²⁺ through RyR in FliI-KD cells. To verify this speculation, we overexpressed sorcin in FliI-KD cells and measured ER Ca²⁺. As shown in Fig. 6b, decreased ER Ca²⁺ was recovered by overexpression of sorcin in FliI-KD cells. Furthermore, sorcin overexpression in FliI-KD cells decreased the level of TG-induced ER stress compared with that in vector-expressed FliI-KD cells (Fig. 6c). These results suggest that FliI controls the ER Ca²⁺ pool by regulating sorcin expression.

Discussion

Globally, CRC is the third most commonly diagnosed malignancy and the second leading cause of cancer-related death¹. In particular, the emergence of refractory disease following chemotherapy continues to be a major contributor to treatment failure and to the high mortality rates observed in advanced CRC. We now know that the

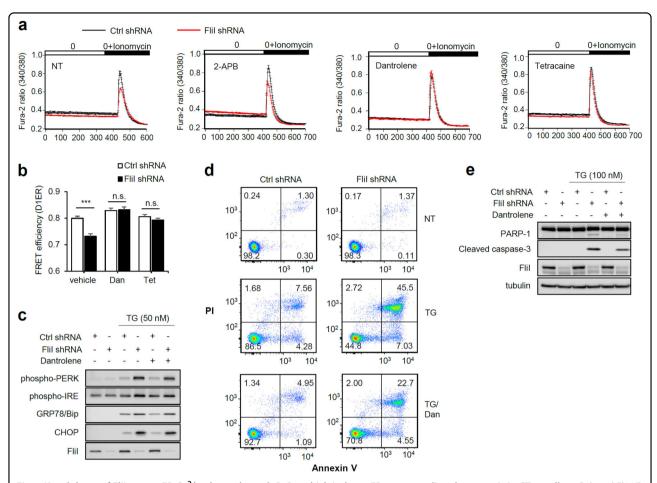


Fig. 5 Knockdown of Flil causes ER Ca²⁺ release through RyRs, which induces ER stress-mediated apoptosis in CT26 cells. a Ctrl- and Flil-KD cells were pretreated for 2 h with 2-APB (50 μM) and dantrolene (50 μM) and tetracaine (50 μM) and then incubated in Fura-2AM for 30 min. Ratiometric Ca²⁺ imaging was performed in 0 mM Ca²⁺ Tyrode's solution with or without 5 μM ionomycin, and Ca²⁺ influx was monitored based on the Fura-2 fluorescence ratio. **b** After 24 h transfection of ctrl- and Flil-KD cells with pcDNA-D1ER, dantrolene (50 μM) and tetracaine (50 μM) were treated for 2 h, and the FRET signal was measured. Data are shown as the mean ± SEM. ***p < 0.001, shRNA-Ctrl cells vs. shRNA-Flil cells. Dan dantrolene, Tet tetracaine. **c** Ctrl- and Flil-KD cells were treated with 50 nM TG for 6 h following pretreatment with 50 μM dantrolene. Cell lysates were analyzed by western blotting for phospho-PERK, phospho-IREα, GRP78/BiP, CHOP, Flil and tubulin. **d**, **e** Ctrl- and Flil-KD cells were pretreated for 2 h with dantrolene (20 μM) and then with TG (100 nM) for 48 h. Apoptosis was detected by annexin V-Pl flow cytometry assay (**d**) and western blotting for PARP-1, cleaved caspase-3, Flil, and tubulin (**e**). TG thapsigargin, Dan dantrolene.

outcome of anticancer treatment can be influenced by the cancer cell response to cellular stress induced by chemotherapeutic agents. The ER stress response is an adaptive mechanism that protects the cell against environmental stimuli⁴. In addition, the ER has been implicated in promoting apoptosis by releasing calcium, which serves as a secondary messenger for mitochondrion-mediated apoptosis³⁸. In a tumor context, high proliferation rates and accelerated metabolism burden cancer cells with cellular and metabolic stress; therefore, as an adaptation, these cells generally activate the ER stress response to support their own survival³⁹. However, this signaling event is a double-edged sword: although it supports cancer cell survival in an adverse environment, harsh and sustained stress conditions, such as those induced by chemopreventive agents, tip the

balance from prosurvival to proapoptosis, culminating in ER stress-mediated cancer cell death³⁹. The mechanisms determining the switch from adaptation to cell death signaling under ER stress have not been clearly defined, and the outcome of ER stress signaling may depend on the type, intensity, and duration of the stimulus in relation to active changes in the cellular environment. Despite this lack of clarity, employing strategies to either sensitize or persuade the cancer cells to ER stress may prove to be a useful tool for overcoming drug resistance and proliferation in cancer cells. Notably, van den Brink et al. reported that ER stressinduced differentiation sensitizes colon cancer stem cells to chemotherapy⁴⁰. Many studies have supported the idea that FliI plays important roles in tumor progression by stimulating cell proliferation, inhibiting apoptosis, and promoting

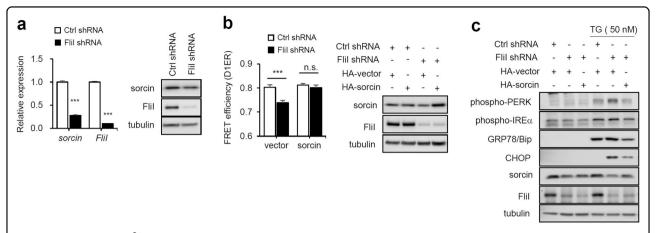


Fig. 6 Flil regulates ER Ca²⁺ through sorcin. a Sorcin expression levels were analyzed by western blot and quantitative real-time PCR. Data are shown as the mean \pm SEM (n=3). ****p<0.001, shRNA-Ctrl cells vs. shRNA-Flil cells. **b** After cotransfection of vector or sorcin with D1ER, FRET signals were measured. Data are shown as the mean \pm SEM. ****p<0.001, shRNA-Ctrl cells vs. shRNA-Flil cells. **c** Ctrl- and Flil-KD cells were treated with 50 nM TG for 6 h following expression of vector or sorcin. Cell lysates were analyzed by western blotting for phospho-PERK, phospho-IREα, GRP78/BiP, CHOP, sorcin, Flil, and tubulin.

invasion^{21–23,41}. In this study, we found that FliI is highly expressed in CRC and that KD of FliI in CT26 cells promoted the induction of ER stress by TG, resulting in apoptosis (Figs. 2, 3). The molecular mechanism by which FliI influences sensitivity to ER stress inducers remains unclear, but we speculate that one important factor is the Ca²⁺ concentration in the ER and cytosol.

Calcium, stored primarily in the ER, is one of the key regulators of cell survival, but it can also induce apoptosis in response to a variety of pathological conditions²⁹. Furthermore, calcium-mediated signaling pathways were implicated in tumorigenesis and tumor progression, including metastasis, invasion, and angiogenesis 42,43. Conversely, several studies showed that sustained intracellular calcium overload can trigger cell death or deregulate calcium-dependent potential tumorigenic pathways²⁹. Recent studies suggest that interfering with the sequestration of calcium into intracellular pools from the ER can trigger apoptosis as part of a stress response 11,38. Thus, the final outcome is determined by the amplitude of the increase and the duration of the change in intracellular calcium level, as well as the nature of the change and the location. In this study, downregulation of FliI in CT26 cells increased intracellular Ca²⁺ but decreased ER Ca²⁺ concentration. This Ca²⁺ depletion was mediated by RyRs (Fig. 5). Thus, depletion of FliI resulted in an increase in the intracellular Ca²⁺ concentration through Ca2+ release via RyRs, which sensitized cells to ER stress inducers. Thereafter, more rapid and marked increases in intracellular Ca²⁺ levels triggered apoptosis.

We next asked how FliI regulates Ca²⁺ release through RyRs. RyRs are modulated directly or indirectly by Ca_V1.1/1.2, various ions, and small molecules and

proteins, including Ca²⁺, Mg²⁺, protein kinase A, FK506 binding proteins, calmodulin, Ca²⁺/calmodulin-dependent protein kinase II (CaMKII), calsequestrin, and triadin³⁷. Most RyR modulators interact with the cytoplasmic region of the channel. Hence, we first hypothesized that FliI stimulates RyR activation because FliI regulates signaling pathways by interacting with signaling proteins 12,13. FliI interacts directly with active CaMKII to inhibit its signaling cascade44, and CaMKII phosphorylates RyRs in the cytosol, altering their gating properties, leading to Ca²⁺ release from the ER⁴⁵. Based on these observations, we postulated that FliI might block the phosphorylation of RyRs by direct interaction with CaMKII, which inhibits RyR activation. However, we did not observe any change in TG-induced UPR signaling or apoptosis when we treated cells with KN-93, a CaMKII inhibitor (data not shown). Therefore, CaMKII activation is not involved in RyR activation by FliI depletion.

Focusing on FliI as a coactivator of transcription factors, we next hypothesized that FliI regulates RyR activation by modulating the expression of genes related to RyRs. Because there was no change in RyR expression in FliI-KD cells (data not shown), we compared global gene expression profiles using RNA sequencing and observed that sorcin (soluble resistance-related calcium-binding protein) was downregulated by depletion of FliI (Fig. 6a). Sorcin is a member of the penta-ERF-hand protein family that localizes to the cytosol, nucleus, plasma membrane, cytoplasmic vesicles, and ER membrane 36,44. In cardiomyocytes, sorcin interacts with RyRs and SERCA; through these interactions, Ca²⁺-bound sorcin negatively regulates the release of Ca²⁺ from the ER and increases the Ca²⁺ load in the ER by inhibiting RyRs and activating SERCA^{36,37}. These changes promote the accumulation of

Ca²⁺ in the ER and decrease the level of intracellular Ca²⁺, thereby preventing ER stress. As expected, overexpression of sorcin in FliI-KD cells increased the ER Ca²⁺ pool, which recovered TG-induced ER stress (Fig. 6). Sorcin is upregulated under ER stress⁴⁶ and overexpressed in many human tumors, including leukemia, gastric, breast, and ovarian cancers ^{47–50}, and in chemoresistant cell lines^{51–55}. Notably, in this regard, overexpression of sorcin is associated with multidrug resistance (MDR), whereas downregulation of the protein increases sensitivity to antitumor drugs^{56,57}. Moreover, sorcin overexpression is correlated with upregulation of MDR1/Pglycoprotein^{54,57}, and the P-glycoprotein-dependent MDR phenotype seems to be related to intracellular Ca²⁺ homeostasis⁵⁸. In fact, we also observed downregulation of MDR1/P-glycoprotein and ABCB1 amplicons, including Dbf4, Crot, and Slc5a40, whose overexpression confers MDR (Supplementary Fig. 5)⁵⁹. Thus, it is intriguing to speculate that sorcin is involved in FliI-mediated regulation of Ca²⁺ homeostasis, possibly by activating RyRs, which may be responsible for sensitizing cells to ER stress inducer-mediated apoptosis. Further studies are needed to show how FliI regulates sorcin expression.

In conclusion, we have uncovered a novel molecular mechanism in which depletion of FliI increases intracellular Ca^{2+} concentration in CRC cells by promoting ER Ca^{2+} release through RyRs, which sensitizes the cells to ER stress inducers and promotes ER stress. Ultimately, this results in ER stress-induced apoptosis. This study provides the first evidence that FliI plays a role in ER stress by controlling Ca^{2+} homeostasis in CRC cells. In addition, FliI expression is correlated with the expression of sorcin as well as ABCBI and its amplicons, which are markers of MDR in CT26 cells. Therefore, targeting FliI represents an innovative and effective strategy for the development of cancer therapeutics.

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Confict of interest

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