Capsaicin-induced TRIB3 upregulation promotes apoptosis in cancer cells

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Background: Capsaicin (8-methyl-N-vanillyl-6-nonenamide) is one of the main pungent components of chili peppers and has been shown to exert various effects on numerous physiological processes. Recent studies have focused on the chemopreventive effects of capsaicin, which can combat growth in various human cancer cell systems. The tribbles-related protein 3 (TRIB3) is evolutionarily conserved from Drosophila to humans. In the latter, TRIB3 is a key determinant in numerous cellular processes, including apoptosis.

Purpose: The aim of this study was to examine the importance of TRIB3 in the antitumor efficacy of capsaicin in human cancer cells, and further assess potential mechanism(s) underlying the capsaicin-induced upregulation of TRIB3.

Methods: Human cancer cell lines were treated with capsaicin, then evaluated for levels of TRIB3 and molecules related to apoptosis or signaling pathways. The impact of TRIB3 on capsaicin-induced apoptosis was investigated using siRNA or overexpression of TRIB3.

Results: It is the first time to show that TRIB3 is targeted by capsaicin to promote apoptosis. Capsaicin promotes apoptotic cell death by upregulating TRIB3 expression in cancer cells. Overexpression of TRIB3 enhances capsaicin-induced apoptosis, and TRIB3 knockdown experiments demonstrate that the effect of capsaicin in apoptotic cell death is correlated with the induction of TRIB3 in cancer cells. Finally, enhancements in gene expression and protein stability are involved in the capsaicin-induced upregulation of TRIB3.

Conclusion: Our results show that the capsaicin-induced upregulation of TRIB3 triggers apoptosis and thereby contributes to the suppression of cell growth in cancer cell lines.

Keywords: capsaicin, TRIB3, apoptosis, cancer, Protein stability

Introduction
Capsaicin (8-methyl-N-vanillyl-6-nonenamide) is one of the main pungent components of chili peppers and has been shown to exert various effects on numerous physiological processes. It is commonly used as an anticancer compound owing to its inhibitory effects on cell proliferation. Recent studies focusing on the chemopreventive and chemotherapeutic effects of capsaicin have demonstrated that it can decrease growth in various human cancer cell systems (eg, liver, lung, colon, pancreas, stomach, prostate, leukocyte, breast, esophagus, and tongue) through its effects on cell cycle arrest and apoptosis.1–12 As a principal natural chemopreventive compound, capsaicin has been shown to have a chemosensitizing activity that improves cytotoxicity among 5-fluorouracil-treated gastric cancer cells.13 In the cisplatin-resistant Korean human gastric cancer cell line, SNU-668, co-treatment of cells with capsaicin plus cisplatin induced higher apoptotic cell death than treatment with either agent alone, suggesting that this strategy could potentially overcome cisplatin resistance.14
Cancer cells show differential susceptibility to capsaicin, perhaps reflecting the ability of this agent to modulate diverse signaling pathways that contribute to cell death or survival. However, we do not yet fully understand the molecular mechanism(s) underlying capsaicin-induced growth inhibition and apoptosis, nor have we fully elucidated the complicated cross-talk among the involved signaling pathways. Thus, sustained research efforts are being made to understand the action mechanisms of capsaicin and explore its potential therapeutic use in cancer.

Several mechanisms are thought to be involved in capsaicin-induced apoptosis. For example, capsaicin has been shown to exert cytotoxic effects in an array of mammalian cell lines via various signaling pathways, including the AMPK,15,16 c-Jun N-terminal kinase 1 (JNK),17 p38 MAPK,17 extracellular signal-regulation kinase (ERK),18 phosphoinositide 3-kinase (PI3K)/Akt,19 and interleukin (IL)-620 pathways. Capsaicin has also been shown to exert its chemopreventive activities via its multitargeting property. For instance, capsaicin enhances p53 gene expression in SNU-1 stomach cancer cells21 and stabilizes p53 protein stability in human colon cancer cells,22 triggering apoptosis in both growth factor deprivation,37 and several antitumor drugs such as tetrahydrocannabinoids, salinomycin, or the lipid derivative ABTL0812.38–40 Interestingly, several studies have shown that capsaicin-enhanced upregulation of TRIB3. Our data show that capsaicin enhances the protein expression of TRIB3 in various human cancer cells and significantly increases the mRNA and protein stability of TRIB3, and that these effects are accompanied by increased apoptotic cell death. TRIB3 knockdown experiments further demonstrated that capsaicin-induced apoptotic cell death is correlated with the induction of TRIB3 in cancer cells. We also report that the apoptosis associated with capsaicin-mediated induction of TRIB3 suppresses cell growth in cancer cell lines in vitro. It is clear that TRIB3 acts as a critical factor for capsaicin-promoted apoptosis in cancer cells; however, JNK, p38 and PI3K–AKT signaling pathways are not associated with this capsaicin-enhanced upregulation of TRIB3.

Materials and methods

Chemicals

The MAPK inhibitors, U0126, SB203580, and SP600125, and the proteasome inhibitor, MG132, were purchased from TOCRIS Bioscience (Bristol, UK). The protease inhibitor cocktail was a product of Roche Applied Science (Mannheim, Germany). Cycloheximide was obtained from Sigma-Aldrich Co. (St Louis, MO, USA). Capsaicin (8-methyl-N-vanillyl-6-nonenamide) was purchased from Enzo Life Sciences (Farmingdale, NY, USA). All other chemicals were purchased from Sigma-Aldrich Co. (St Louis, MO, USA) or Amresco (Solon, OH, USA).

Antibodies

The antibody against TRIB3 was obtained from Novus Biologicals (Littleton, CO, USA). The antibodies against PARP, caspase 3, caspase 8, caspase 9, caspase 12, phospho-p38, phospho-ERK1/2, phospho-JNK, and phosphor-AKT were products of Cell Signaling Technology (Beverly, MA, USA). The antibodies against ERK2, JNK, p38, and AKT, beta-actin, and the peroxidase-conjugated secondary antibodies against mouse and rabbit IgG were purchased from Santa Cruz.
Biotechnology Inc. (Santa Cruz, CA, USA). The antibodies against phospho-p90RSK (Thr573), Bid and Bax were from Abcam (Cambridge, UK). The antibody against the Myc tag was from EMD Millipore (Billerica, MA, USA).

Cell culture
The human gastric carcinoma cell lines, AGS, SNU-1, and KATO III, were obtained from Bioresource Collection and Research Center (BCRC; Hsinchu, Taiwan). MKN-45 was purchased from Japanese Collection of Research Bioresources (JCRB) Cell Bank (Osaka, Japan). Human gastric cancer cell line TMK-1, human cervical squamous cell carcinoma SiHa, and human colorectal adenocarcinoma cell line SW480 were gifted by Dr Wu (Department of Education and Research and Division of Gastroenterology, Taichung Veterans General Hospital, Taichung, Taiwan; current address is Taipei Veterans General Hospital and Faculty of Medicine, School of Medicine, National Yang-Ming University, Taipei, Taiwan) and their DNA short tandem repeats (STR) profile was confirmed at BCRC. Cells were cultured in RPMI-1640 (Thermo Fisher Scientific, Waltham, MA, USA). Cultures were supplemented with 10% fetal bovine serum, sodium bicarbonate (2%, w/v), l-glutamine (0.29 mg/mL), penicillin (100 units/ml), and streptomycin (100 µg/mL) (Thermo Fisher Scientific), and grown at 37°C in a humidified 5% CO₂ incubator.

Plasmid
The full-length human TRIB3 cDNA was cloned into the pcDNA3.1-Myc-His vector using primers designed to introduce Hind III and Xho I restriction sites, as follows: sense, 5′-AAC TCG AGG CCA CCA TGC GAG CCA CCC CTC TG-3′ and antisense, 5′-AAA AGC TTG CCA TAC AGA ACC ACT TC-3′. The obtained TRIB3 cDNA was confirmed by sequencing. Cells were transfected with 2 µg TRIB3-Myc plasmid using jetPEI (Polyplus, Illkirch, France) according to the manufacturer’s recommendations.

Cell viability assay
Cells (5×10³/well) were seeded in a 96-well plate. After 24 hours, cells were treated with different concentrations of capsaicin (three wells per concentration) for 48 hours. Cell viability was detected using the WST-1 reagent, according to the manufacturer’s recommendations (Roche Applied Science).

Reverse-transcription polymerase chain reaction
Total RNA was isolated from cells using an RNeasy Mini kit (Qiagen Inc. Germantown, MD, USA) and quantified. One microgram of total RNA was quantified and applied to synthesize single-stranded cDNA using an ImProm-II Reverse Transcriptase kit (Promega Corporation, Madison, WI, USA), followed by amplification reactions using primers: sense, 5′-ATG ATT CCC TGT GGG ACA AG-3′ and antisense, 5′-CTG TCT CCC TCC TCT TCC CT-3′. Reaction conditions consisted of 28 cycles of 95°C for 1 minute 30 seconds, 51°C for 1 minute 10 seconds, and 72°C for 2 minutes 10 seconds, followed by a final extension of 5 minutes at 72°C. Polymerase chain reaction (PCR) products were resolved by electrophoresis on 1% agarose gels and visualized by ethidium bromide staining.

RNA interference
For transient siRNA-mediated knockdown, 30 nM double-stranded RNA duplexes targeting the TRIB3 gene (ON-TARGET plus SMARTpool, Cat. No. L-003754) and non-targeting siRNA duplexes (Cat. No. D-001810-10-05, both from Dharmacon Research, Lafayette, CO, USA) were transfected to the indicated cells using Lipofectamine RNAiMAX (Thermo Fisher Scientific) according to the manufacturer’s recommendations. After 24 hours, the cell extracts were prepared and the levels of TRIB3 were determined by Western blot analysis.

Cell lysis and Western blotting
Cell pellets were lysed in Buffer A (50 mM HEPES, pH 7.5, 150 mM NaCl, 5 mM EDTA, 1% Triton X-100, 50 mM NaF, 1 mM Na₂VO₄, and 10% glycerol plus a protease inhibitor cocktail), and extracts were collected by centrifugation at 10,000×g and 4°C for 10 minutes. Protein concentrations were determined by Bradford protein assay (Bio-Rad Laboratories Inc., Hercules, CA, USA). Equal amounts of protein were resolved by sodium dodecyl sulfate-polyacrylamide gel electrophoresis, and the resolved proteins were transferred to a polyvinylidene difluoride (EMD Millipore) membrane. The membrane was blocked with 5% nonfat dry milk and probed with primary antibodies, and the signals were detected using an enhanced chemiluminescence system (EMD Millipore). For the detection of phosphoproteins, membrane was blocked with bovine serum albumin (Sigma-Aldrich) or EasyBlocker (GeneTex, Hsinchu, Taiwan).

Quantitative real-time PCR (qPCR)
Total RNA was isolated using an RNeasy Mini kit (Qiagen) and quantified. Single-stranded cDNA was synthesized from 2 µg of RNA using an ImProm-II Reverse Transcriptase kit (Promega). Specific primers were designed using the Probe Finder software from Roche Applied Science, which is available online at the Universal ProbeLibrary Assay Design Center (Roche Applied Science) and is based on the...
Minimum Information for publication of Quantitative real-time PCR Experiments guidelines. The probes were obtained from the Universal ProbeLibrary collection (Roche Applied Science). The qPCR analysis was performed utilizing a LightCycler Nano instrument (Roche Applied Science), and the results of each experiment were normalized with respect to the expression of the control gene, TBP.

**Real-time cell analysis (RTCA)**

To measure the electrical impedance of touched cell proliferation, cells (7.5×10^5/well) were seeded onto E-plates (Roche Applied Science) and incubated for 30 minutes. The plates were placed onto the RTCA station (xCELLigence Real-Time Cell Analysis System, Roche, Germany) and incubated overnight, and then the cells were exposed to capsaicin. Thereafter, impedance was measured every hour over a period of 99 hours. Impedance is presented as the cell index (CI) over a period of 99 hours. Impedance is presented as the background resistance and Zi is the resistance at an individual time point. A normalized cell index was determined as the cell index at a certain time point (CI_{t}) divided by the cell index at the normalization time point (CI_{nml_time}).

**Statistical analysis**

All data were obtained from at least three independent experiments. Values are presented as the mean±SD from at least three independent experiments, unless otherwise indicated. The SigmaPlot 10.0 software was used to analyze the data and calculate significant differences between the control and experimental groups, as assessed by the Student’s t-test. The results presented in the graphs are representative of multiple independent experiments, and * indicates *P*<0.05, which was considered statistically significant.

**Results**

**Capsaicin induces the protein expression of TRIB3 in cancer cell lines**

We first examined the expression levels of TRIB3 in seven different cancer cell lines: AGS, SW480, MKN-45, SNU-1, TMK-1, SiHa, and KATO III. As shown in Figure 1A, the TRIB3 protein levels were dramatically different in the various cancer cell lines, with relatively low levels in AGS, SNU-1, MKN-45, SiHa, and KATO III cells and high-level expression in SW480 and TMK-1 cells. Similar expression profiles were obtained from our analysis of TRIB3 mRNA expression (Figure 1A, lower panel). Given that AGS cells possess the lowest malignancy and MKN-45 cells are very malignant, our results suggest that the expression level of TRIB3 may not correlate with the malignancy of cancer cell lines in vitro. Next, to test whether the endogenous level of TRIB3 contributed to the susceptibility of cancer cells to capsaicin, we treated AGS, MKN-45, and TMK-1 cells with the indicated concentrations of capsaicin for 48 hours and then determined cell viability. However, we did not identify any significant difference in the viability of the treated cell lines at 100 µM (*P*>0.05; Figure 1B). To investigate whether capsaicin induces TRIB3 expression in cancer cells, we examined TRIB3 expression in AGS, MKN-45, TMK-1, and SiHa cells exposed to different concentrations of capsaicin for 24 hours, or to a consistent amount of capsaicin for various time periods. Our results revealed that capsaicin dose- (Figure 1C) and time-dependently (Figure 1D) increased TRIB3 expression in different cancer cell lines.

Interestingly, we observed a marked increase in the cleaved forms of caspase 3 and PARP accompanied with TRIB3 upregulation in capsaicin-treated AGS (Figure 2A) and SiHa (Figure 2B), indicating that these cells underwent apoptotic cell death. Consistent with the known ability of TRIB3 to negatively regulate the activation of AKT, cells subjected to the capsaicin-induced upregulation of TRIB3 showed gradual decreases in the activation of AKT (Figure 2A and B). Dose-dependent induction of apoptosis was further confirmed in other cell lines. The increased cleaved form of PARP was also observed in capsaicin-treated TMK-1 (Figure 2C) and MKN-45 (Figure 2D) cells. Further analysis of factors involved in various apoptotic pathways revealed that capsaicin triggered significant increases in molecules involved in both intrinsic mitochondrial pathway (eg, cleaved caspase 9 and Bax) and extrinsic pathway (eg, cleaved caspase 8), as shown in Figure 2D.

**Induction of TRIB3 is essential for capsaicin-induced apoptosis**

Next, we investigated the functional involvement of TRIB3 in capsaicin-mediated apoptosis. TRIB3-Myc was ectopically expressed in AGS cells for 24 hours, and then the cells were treated with capsaicin. As shown in Figure 3A, ectopic expression of Myc-tagged TRIB3 augmented capsaicin-induced cleavage of caspase 3 and PARP (Figure 3A). Monitoring of cell proliferation using a RTCA system revealed that overexpression of TRIB3 attenuated cell proliferation in AGS (Figure 3B). This suggests that TRIB3 contributes to the antiproliferative and proapoptotic effects of capsaicin. Conversely, knockdown of cellular TRIB3 with specific siRNA (si-TRIB3) decreased the levels of cleaved caspase 3 and cleaved PARP in capsaicin-treated AGS and TMK-1 cells.
Upregulation of TRiB3 promote apoptosis compared to the results from cells treated with si-control (Figure 3C). Together, these findings support the hypothesis that capsaicin elevates TRIB3 expression, and that TRIB3 contributes to cell death in capsaicin-treated cancer cells.

Capsaicin enhances TRIB3 expression at the levels of mRNA expression and protein stability

Given the ability of capsaicin to induce TRIB3, we set out to test whether capsaicin could regulate the expression and function of TRIB3 in cancer cell lines. TRIB3 has been shown as a short-lived protein that is frequently degraded through the ubiquitin–proteasome pathway.44,45 To test whether the differences of TRIB3 in these cancer cell lines were due to differences in protein stability, we examined the levels of TRIB3 in four different cancer cell lines with MG132 treatment. Inhibition of proteasome pathway by MG132 further enhanced the levels of TRIB3 in a time-dependent manner (Figure 4A). This confirmed that the turnover of TRIB3 occurs through 26S proteasome-mediated proteolysis in these cancer cell lines. Furthermore, we treated cells with cycloheximide (a protein synthesis inhibitor) and tested the half-life of TRIB3 by Western blotting at different time points posttreatment (0, 0.5, 1, 2 hours). As shown in Figure 4B, the protein turnover of TRIB3 was not significantly different among the tested cancer lines. These results suggest that the differential expression of TRIB3 might reflect differences in transcriptional regulation among the cancer cell lines. To examine whether

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**Figure 1** Capsaicin induces the upregulation of TRiB3 in different cancer cell lines.

**Notes:** (A) Cancer cell lines vary in their TRiB3 expression levels. The protein (upper panel) and mRNA (lower panel) expression levels of TRiB3 were measured in seven different cancer cell lines. (B) AGS, MKN-45, and TMK-1 cells were seeded in a 96-well plate for 24 hours and then exposed to 0, 50, 100, 200 µM capsaicin for 48 hours. Cell viability was determined using the WST-1 reagent. *P<0.05 as the cell viability of MKN-45 was significantly different from those of AGS and TMK-1. (C) Cells were exposed to different concentrations of capsaicin and the protein levels of TRiB3 were determined. (D) MKN-45 and SiHa cells were treated with 100 µM capsaicin for 6–24 hours and the protein levels of TRiB3 were determined. All data were obtained from at least two to four independent experiments.

**Abbreviation:** n.s., not significant.
Capsaicin induced TRIB3 expression by increasing gene transcription, we treated cells with capsaicin for 24 hours and then determined the TRIB3 mRNA levels. Our qPCR analyses revealed that capsaicin significantly increased the mRNA level of TRIB3 in AGS cells (Figure 4C), indicating that increased gene expression, at least in part, might be involved in the capsaicin-induced upregulation of TRIB3. Since protein turnover also plays a critical role in regulating the protein level and function of TRIB3,44–46 the cycloheximide experiment was performed, which revealed that TRIB3 protein stability was moderately, but significantly enhanced by capsaicin in AGS (Figure 4D) and MKN-45 (Figure 4E) cells. Thus, stabilization of the TRIB3 protein also appears to be an important underlying mechanism of capsaicin-mediated TRIB3 upregulation. Taken together, our data suggest that both transcriptional upregulation and increased protein stability might contribute to capsaicin-mediated TRIB3 expression.

Molecular regulation of TRIB3 expression by capsaicin

Since capsaicin has been reported to modulate various signaling pathways and alter cell growth, we investigated the
Upregulation of TRIB3 promote apoptosis

Indeed, capsaicin was found to dose-dependently enhance the activation of ERK, JNK, and p38 in AGS cells (Figure 5A). However, inhibitor experiments revealed that JNK and p38 MAPK signaling did not involve capsaicin-mediated upregulation of TRIB3 (Figure 5B). Inhibition of MEK-ERK1/2 signaling significantly suppressed the protein level of TRIB3 induced by capsaicin (Figure 5C). To further study whether MEK-ERK1/2 pathway participated in capsaicin-induced TRIB3 gene expression, cells were co-treated with U0126 and capsaicin for 24 hours and then the TRIB3 mRNA levels were determined. The result indicated that MEK1/2 inhibitor U0126 did not interfere with the mRNA expression of TRIB3 in capsaicin-treated AGS cells (Figure 5D), implying that MEK-ERK1/2 signaling may regulate the protein expression of TRIB3 induced by capsaicin at the posttranscriptional level. In addition, knockdown of TRIB3 did not suppress the capsaicin-mediated activation of MAPK signaling in AGS (data not shown). Collectively, our results suggest that capsaicin upregulates TRIB3 expression by increasing its gene transcription and protein stability. Mechanistically, although capsaicin activates three MAPK signaling pathways, JNK and p38 activation does not appear to participate in the

Figure 3 Induction of TRIB3 is essential for capsaicin-induced apoptosis.

Notes: (A) AGS cells were induced to express 2 µg ectopic TRIB3-Myc for 24 hours and exposed to 200 µM capsaicin for another 24 hours, and then the protein levels of TRIB3, TRIB3-Myc, cleaved caspase 3, and cleaved PARP were determined. (B) 2 µg TRIB3-Myc was expressed in AGS cells for 24 hours before exposure to 200 µM capsaicin for another 24 hours, and cell impedance was determined using an RTCA system. (C) AGS and TMK-1 were subjected to si-RNA–mediated knockdown of endogenous TRIB3 for 24 hours and then exposed to 200 µM capsaicin for another 24 hours. The protein levels of TRIB3, cleaved caspase 3, and cleaved PARP were determined. All data were obtained from at least three to five independent experiments.

Abbreviation: RTCA, real-time cell analysis.
capsaicin-induced upregulation of TRIB3, nor does TRIB3 appear to contribute to the capsaicin-mediated activation of MAPK signaling.

Discussion

Increasing evidence suggests that capsaicin has anticancer effects, but we have not yet fully elucidated the molecular mechanism underlying capsaicin-induced growth inhibition and apoptosis of human cancer cells. In this study, we report that capsaicin dose- and time-dependently induces TRIB3 expression and promotes apoptosis in different cancer cells. Moreover, the antiproliferative and proapoptotic effects of capsaicin were found to correlate with its induction of TRIB3 expression, which occurs via both transcriptional and posttranslational mechanisms (Figure 6). Although the detailed mechanism underlying the capsaicin-induced gene upregulation and protein stabilization of TRIB3 is still poorly understood, our findings clearly indicate that capsaicin-
Figure 5 JNK and p38 MAPK signaling pathways do not appear to participate in the capsaicin-triggered upregulation of TRiB3.

Notes: (A) AGS cells were exposed to different doses of capsaicin for 24 hours, cell lysates were prepared, and the levels of various signaling proteins were determined. (B) AGS cells were pretreated with 10 µM SB203580 or SP600125 for 1 hour and then with 100 µM capsaicin for another 24 hours. The levels of TRiB3 were determined. (C, D) AGS cells were pretreated with 10 µM U0126 for 1 hour and then with 100 µM capsaicin for another 24 hours. The protein (C) and mRNA (D) levels of TRiB3 were determined. n=5. *P<0.05. The mRNA levels of TRiB3 were normalized with respect to the expression of the internal control gene, TBP.

Abbreviation: TBP, TATA-box binding protein.
induced alterations in TRIB3 might be a major event leading to apoptosis in capsaicin-treated cancer cells.

Capsaicin has been shown to exert profound antiproliferative activity in animal/cell models of cancer, mostly by inducing apoptosis. The modulations of signaling pathways and transcription factor functions are major mechanisms contributing to the antiproliferative effect of capsaicin. In addition, capsaicin can induce apoptosis by indirectly inhibiting the ubiquitin–proteasome system, leading to the accumulation of target substrates normally targeted by the proteasome, such as p53, Bax, and p27.47 By inhibiting the interaction between p53 and MDM2, capsaicin has been reported to stabilize the p53 protein and enhance the transcriptional activity of its encoding gene in human colon cancer cells.22 In the present study, we report that TRIB3 is a potent target of capsaicin and its upregulation might play a critical role in the capsaicin-mediated apoptosis of different human cancer cells.

Studies have shown that the protein level of TRIB3 is tightly regulated at several levels, including both transcriptional and translational mechanisms. For example, the TRIB3 mRNA is induced by the unfolded protein response via the transcription factor, ATF4, as well as by the NFκB and PI3K pathways.48 Moreover, the TRIB3 mRNA may be stabilized by the hypoxia-regulated protein, HuR.48 At the protein level, TRIB3 is modulated by ubiquitin-mediated proteolysis. Siah1 interacts with and ubiquitinates TRIB3 in vivo to regulate its turnover via the ubiquitin–proteasome pathway.44 A typical D-box motif localized in the carboxyl-terminal domain of TRIB3 is required for its protein stability, and the APC/C competitor protein is involved in ubiquitinating and destabilizing the TRIB3 protein.45 In the present study, we reveal that capsaicin enhances TRIB3 gene expression and protein stability, thereby augmenting apoptotic cell death in cancer cells. Although JNK and p38 MAPK were not involved in the induction of TRIB3 by capsaicin, MEK-ERK1/2 signaling may regulate capsaicin-induced TRIB3 at the posttranscriptional level. The detailed mechanism should be further investigated in the future.

Using a functional genetic analysis, we identified the –1561 to +77 bp region upstream of the TRIB3 gene as being relevant to the construction of a luciferase reporter system. However, our results revealed that this upstream region was not responsible for the induction of TRIB3 by capsaicin (data not shown). Future studies are needed to elucidate the detailed mechanisms (eg, TRIB3 phosphorylation, ubiquitin ligase activity, and/or proteasomal function) responsible for the capsaicin-mediated changes in the gene expression and stability of TRIB3.

TRIB3 is often activated by a wide variety of stress signals, including ER stress, hypoxia, and starvation. We previously reported that TRIB3 gene expression is dose-dependently upregulated by gold nanoparticle exposure in a cell model,49 supporting the notion that TRIB3 may act as a marker of stress and increased metabolism. However, the existing results are rather contradictory regarding the potential pro- or antiapoptotic roles of TRIB3 in response to diverse stresses. This suggests that there may be fundamen-

Figure 6 Capsaicin induced apoptosis through upregulating TRIB3 expression in cancer cells.

Notes: The proposed model is consistent with the published studies and the results described here. Capsaicin dose- and time-dependently induces TRIB3 expression via both stimulating gene transcription and stabilizing protein stability mechanisms to promote apoptosis in cancer cells.

Abbreviation: CHX, cycloheximide.
tally different correlations between TRIB3 mRNA/protein changes and the resulting cellular phenotype. Alternatively, the contradictory results may reflect that TRIB3 is responsible for the set points of several regulatory feedback loops.

Interestingly, we have reported that TRIB3 downregulation enhances doxorubicin-induced cytotoxicity in gastric cancer cells, indicating TRIB3 plays a prosurvival role in doxorubicin-induced cytotoxicity in gastric cancer cells. Furthermore, we also suggest that ER stress and extrinsic apoptotic pathways may not be involved in doxorubicin-induced cytotoxicity. Interestingly, in this study, we found that capsaicin-enhanced TRIB3 expression is accompanied by increased levels of Bip (GRP78) and cleaved caspase 12 (data not shown), two well-known markers for ER stress. We anticipate that increased TRIB3 expression plays an important role in capsaicin-induced apoptosis through the induction of ER stress. Consistent with previous reports, it seems like that TRIB3 has the opposite role in apoptosis in gastric cancer cell lines, dependent on apoptotic inducers such as capsaicin or doxorubicin. More specifically, ER stress might be an important switch to determine the pro- or antiapoptotic role of TRIB3. In conclusion, the apparent dual function of TRIB3 may act as a key cellular context-dependent switch between cell survival and apoptosis, and it is not a bona fide member of apoptotic machinery. Therefore, we need to make very careful conclusion for the function of this protein in apoptosis.

Gastric cancer is the fourth most common cancer worldwide, but conventional chemotherapy and radiation are not effective in prolonging the long-term survival of gastric cancer patients. The existing evidence suggests that combinatorial anticancer therapeutic interventions, such as the addition of natural compounds to the conventional strategies, are needed to completely eradicate neoplastic lesions and minimize the side effects. Capsaicin has been demonstrated to negatively regulate cell survival, growth, inflammation, and differentiation. In this study using cell culture model, we presented a novel mechanism whereby capsaicin upregulates TRIB3 to promote apoptosis in gastric cancer cell lines. This suggests that the combination of capsaicin with effectively targeted chemotherapeutic agents might provide an efficient strategy for the treatment of human tumors, such as gastric cancer. However, the effect of capsaicin in cell culture may not be well correlated with the responses and mechanisms in vivo and the concentrations of capsaicin used in this study were not physiologically relevant. The significance of TRIB3 in capsaicin-mediated apoptosis in vivo should be further investigated.

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Disclosure
The authors report no conflicts of interest in this work.

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