

RESEARCH PAPER

Dimethyl fumarate induces necroptosis in colon cancer cells through GSH depletion/ROS increase/MAPKs activation pathway

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BACKGROUND AND PURPOSE

Dimethyl fumarate (DMF) is a newly approved drug for the treatment of relapsing forms of multiple sclerosis and relapsing-remitting multiple sclerosis. Here, we investigated the effects of DMF and its metabolites mono-methylfumarate (MMF and methanol) on different gastrointestinal cancer cell lines and the underlying molecular mechanisms involved.

EXPERIMENTAL APPROACH

Cell viability was measured by the MTT or CCK8 assay. Protein expressions were measured by Western blot analysis. LDH release, live- and dead-cell staining, intracellular GSH levels, and mitochondrial membrane potential were examined by using commercial kits.

KEY RESULTS

DMF but not MMF induced cell necroptosis, as demonstrated by the pharmacological tool necrostatin-1, transmission electron microscopy, LDH and HMGB1 release in CT26 cells. The DMF-induced decrease in cellular GSH levels as well as cell viability and increase in reactive oxygen species (ROS) were inhibited by co-treatment with GSH and N-acetylcysteine (NAC) in CT26 cells. DMF activated JNK, p38 and ERK MAPKs in CT26 cells and JNK, p38 and ERK inhibitors partially reversed the DMF-induced decrease in cell viability. GSH or NAC treatment inhibited DMF-induced JNK, p38, and ERK activation in CT26 cells. DMF but not MMF increased autophagy responses in SGC-7901, HCT116, HT29 and CT26 cancer cells, but autophagy inhibition did not prevent the DMF-induced decrease in cell viability.

CONCLUSION AND IMPLICATIONS

DMF but not its metabolite MMF induced necroptosis in colon cancer cells through a mechanism involving the depletion of GSH, an increase in ROS and activation of MAPKs.

Abbreviations

3-MA, 3-methyladenine; CsA, cyclosporine A; DMF, dimethyl fumarate; DMH1, 4-[6-(4-isopropoxyphenyl)pyrazolo [1,5-a]pyrimidin-3-yl] quinolone; EthD-1, ethidium-homodimer-1; HMGB1, high-mobility group box 1; MMF, mono-methylfumarate; NAC, N-acetylcysteine; Nec, necrostatin-1; ROS, reactive oxygen species; TEM, transmission electron microscopy

Tables of Links

TARGETS	
Caspase 3	JNK
ERK	p38

LIGANDS	
Cyclosporine A (CsA)	PD98059
Dimethyl fumarate (DMF)	SB203580
GSH	SP600125

These tables list key protein targets and ligands in this article that are hyperlinked to corresponding entries in <http://www.guidetopharmacology.org>, the common portal for data from the IUPHAR/BPS Guide to PHARMACOLOGY (Pawson *et al.*, 2014) and are permanently archived in the Concise Guide to PHARMACOLOGY 2013/14 (Alexander *et al.*, 2013).

Introduction

Dimethyl fumarate (DMF), the methyl ester of fumaric acid, was initially recognized as an effective hypoxic cell radiosensitizer (Held *et al.*, 1988). Recently, DMF (BG-12, Tecfidera®, BIOGEN IDEC Inc., Weston, MA, USA) was approved by the FDA and European Medicines Agency as a new oral drug for the treatment of relapsing forms of multiple sclerosis and relapsing-remitting multiple sclerosis (Burness and Deeks, 2014; Fox *et al.*, 2014). DMF suppresses the expression of inflammatory cytokines such as inducible NOS, IL-1 β , IL-6 in glial cells (Wierinckx *et al.*, 2005; Wilms *et al.*, 2010), and exerts neuroprotective effects via the antioxidant pathway in the CNS (Linker *et al.*, 2011; Scannevin *et al.*, 2012). All of these effects can account for the efficacy of DMF at treating multiple sclerosis.

Furthermore, in the field of multiple sclerosis, DMF has also been found to exhibit anti-tumour effects. Loewe *et al.* reported that DMF inhibited melanoma growth and metastasis both *in vitro* and *vivo* (Loewe *et al.*, 2006); Kirilin *et al.* reported that DMF induced apoptosis in HT29 colon carcinoma cells (Kirilin *et al.*, 1999), and others have shown that DMF enhances the anti-tumour effects of some chemotherapy drugs (Gu and DeAngelis, 2005; Valero *et al.*, 2010). However, it should be noted that, recent studies revealed that DMF could not be detected in plasma because it was immediately and completely hydrolyzed by esterases to mono-methylfumarate (MMF) (Nibbering *et al.*, 1993; Werdenberg *et al.*, 2003), and during this process, methanol was also generated (Rostami-Yazdi *et al.*, 2010). Therefore, it is essential to identify the effects of DMF and its metabolites (MMF and methanol) on cancer cells and the underlying molecular mechanisms.

Necroptosis is a necrotic-like, caspase-independent cell death that is characterized by loss of plasma membrane integrity, excessive autophagy, loss of mitochondria membrane potential and the generation of reactive oxygen species (ROS) is enhanced in necrotic cells (Degtarev *et al.*, 2005). Necroptosis is emerging as an important target for anticancer treatment (Fulda, 2014); targeting necroptosis has been shown to overcome drug resistance in cancer chemotherapy (Han *et al.*,

2007; Steinhart *et al.*, 2013). Many necroptotic stimuli, such as shiconin, smac mimetic and granulysin, have been shown to have antitumour effects or potentiate the effect of chemotherapeutic drugs (Han *et al.*, 2007; Zhang *et al.*, 2009; Steinhart *et al.*, 2013). Here, we used different gastrointestinal cancer cell lines to investigate the effects of DMF and its metabolites (MMF and methanol), and the underlying molecular mechanisms involved. We found that DMF but not MMF induced necroptosis in colon cancer cells by depleting cellular GSH and subsequently activating MAPKs. We propose that DMF has the potential to be an anticancer drug provided an appropriate drug delivery system can be devised for it.

Methods

Cell culture

CT26, HT29, HCT116 and SGC-7901 cells were maintained in DMEM containing 25 mmol·L⁻¹ glucose supplemented with 10% FBS, 100 u·mL⁻¹ penicillin and 100 μ g·mL⁻¹ streptomycin at 37°C, 5%CO₂. The time of treatment and concentration of agents were shown in the figures and/or corresponding figure legends.

Western blot analysis

Western blot analysis was performed as described in our previous work (Sheng *et al.*, 2014). Cells were lysed with RIPA buffer containing 1% protease inhibitor and centrifuged at 16 500 \times g for 15 min at 4°C. Then the supernatants were collected and the protein concentrations were determined by BCA assay kit (Beyotime). The protein was applied to 10% to 12% SDS-PAGE gels, transferred to nitrocellulose membranes. After incubation with the appropriate primary and secondary antibodies, Western blot bands were quantified by using Odyssey infrared imaging system (Li-Cor Inc., Lincoln, NE, USA) and Odyssey v3.0 software.

Cell viability measurement

Cell viability was measured by using either the MTT or CCK8 assay, as indicated in Figures and/or corresponding Figure legends.

LDH assay

Cell culture medium was collected for LDH determination according to the manufacturer's instructions (LDH cytotoxicity assay kit; Beyotime Biotech). LDH can catalyze the synthesis of pyruvic acid from lactic acid and then pyruvic acid reacts to form 2,4-dinitrophenyl-hydrazine, which shows up as a brownish red colour in basic solution. After the reaction, the absorbance was read at a wavelength 490 nm. LDH release reflected the cell death. The cell death ratio was calculated by the following formula according to the manufacturer's instructions:

The cell death ratio (%) = $(A_{\text{sample}} - A_{\text{control}}) / (A_{\text{max}} - A_{\text{control}}) \times 100$

A_{sample} = sample absorbance value

A_{control} = the absorbance value of control group

A_{max} = the absorbance value of positive group

Transmission electron microscopy (TEM)

TEM was performed to identify the cells undergoing necroptosis. The cells were fixed with ice-cold 2.5% glutaraldehyde in PBS (pH 7.3) at 4°C for 4 h. Fixed cells were post-fixed in 2% OsO₄, dehydrated in graded alcohol, embedded in Epon 812 (Electron Microscopy Sciences, Fort Washington, PA, USA), sectioned with ultramicrotome, and stained with uranyl acetate and lead citrate. The sections were examined with a TEM (Technai 10; Philips, Eindhoven, The Netherlands).

Live- and dead-cell staining

The LIVE/DEAD® Viability/Cytotoxicity Assay Kit (Invitrogen) was used to detect the live and dead cells. Briefly, cells were grown on coverslips at a density of 3.75×10^4 cells mL⁻¹ and incubated overnight at 37°C in a humidified 5% CO₂ incubator. The cells were washed with PBS and dyed according to the manufacturer's instructions. The labelled cells were photographed under a fluorescence microscope. The live cells fluoresce green and dead cells fluoresce red.

Measurement of intracellular GSH

Intracellular GSH contents were measured using a total GSH assay kit (Beyotime Biotech) according to the manufacturer's instructions. Cells were harvested and lysed in protein removal solution S provided in the kit, then repeatedly (twice) frozen and thawed in liquid nitrogen and 37°C water respectively. After incubation for 5 min at 4°C, the samples were centrifuged at 10000× *g* for 10 min at 4°C. The supernatant was treated with Ellman's reagent (DTNB) in combination with GSH reductase enzyme and NADPH. Finally the absorbance values were measured at a wavelength of 412 nm by a microplate reader. The intracellular GSH content was quantified against the corresponding standard curves.

Measurement of mitochondrial membrane potential (MMP)

The MMP was measured by using a MMP assay kit (Beyotime Biotech) according to the manufacturer's instructions. Cells were washed with PBS and then incubated in the culture medium containing 1 µg·mL⁻¹ JC-1 (5,5',6,6'-tetrachloro-

1,1',3,3'-tetraethyl-imidacarbocyanine iodide) for 20 min at 37°C. Then, the supernatant was removed and the cells were washed with JC-1 buffer solution twice. The fluorescence intensity was detected by a fluorescence microscope. JC-1 monomer fluorescence (green), which indicates a low MMP, was observed at the wavelength of excitation/emission = 485/530 nm. JC-1 aggregate fluorescence (red), which indicates a high MMP, was detected at the wavelength of excitation/emission = 485/590 nm.

Measurement of intracellular ROS accumulation

Intracellular ROS accumulation was determined by an ROS assay kit (Beyotime Biotech) that utilizes 2',7'-dichlorofluorescein diacetate (DCFH-DA) as a fluorescent probe. In principle, DCFH-DA penetrates the cell membrane freely and is hydrolysed by esterase into DCFH, which cannot escape from the cell, then ROS further oxidizes DCFH to DCF, which fluoresces bright green. After drug treatments, cells were incubated with 10 µM DCFH-DA for 20 min at 37°C and then were washed with DMEM three times. Intracellular ROS were determined by a fluorescence microscope at an excitation wavelength of 488 nm and an emission wavelength of 525 nm. The fluorescence intensity reflected the amount of ROS generated, an increase in fluorescence intensity represents an enhanced generation of ROS.

Detection for HMGB1 release

CT26 cells were seeded in 6-well plates, and then incubated at 37°C, 5%CO₂ overnight. The culture medium was collected after different durations of DMF treatment. The samples were centrifuged at 1000× *g* for 20 min, then the HMGB1 protein in supernatants was detected by using an HMGB1 ELISA kit (Cloud-Clone Corporation, Houston, TX, USA) according to the manufacturer's instructions.

PARP-1 activity assay

PARP-1 activity in HCT116 cells was examined by using a universal colorimetric PARP-1 assay kit (Trevigen, Gaithersburg, MD, USA). After drug treatment, cells were collected and suspended in PARP lysis buffer, which contained protease inhibitor. Then, the samples were centrifuged at 16 500× *g* for 15 min at 4°C, and the protein concentration was assayed using a BCA protein assay kit (Beyotime); 50 µg of protein from each sample were loaded into a 96-well plate coated with histones and biotinylated poly ADP-ribose, and PARP cocktail was also added to each well and then incubated for 60 min at room temperature. Next, the supernatant was aspirated and then the diluted Strep-HRP was added and the samples were incubated for another 60 min at room temperature followed by TACS-Sapphire treatment for 15 min in the dark. To stop the reaction, 0.2 M HCl was added to each well and the plate was read at 450 nm.

Data analysis

Data are presented as mean ± SEM. Significance was determined by using ANOVA, followed by Holm–Sidak. *P* < 0.05 was considered significant.

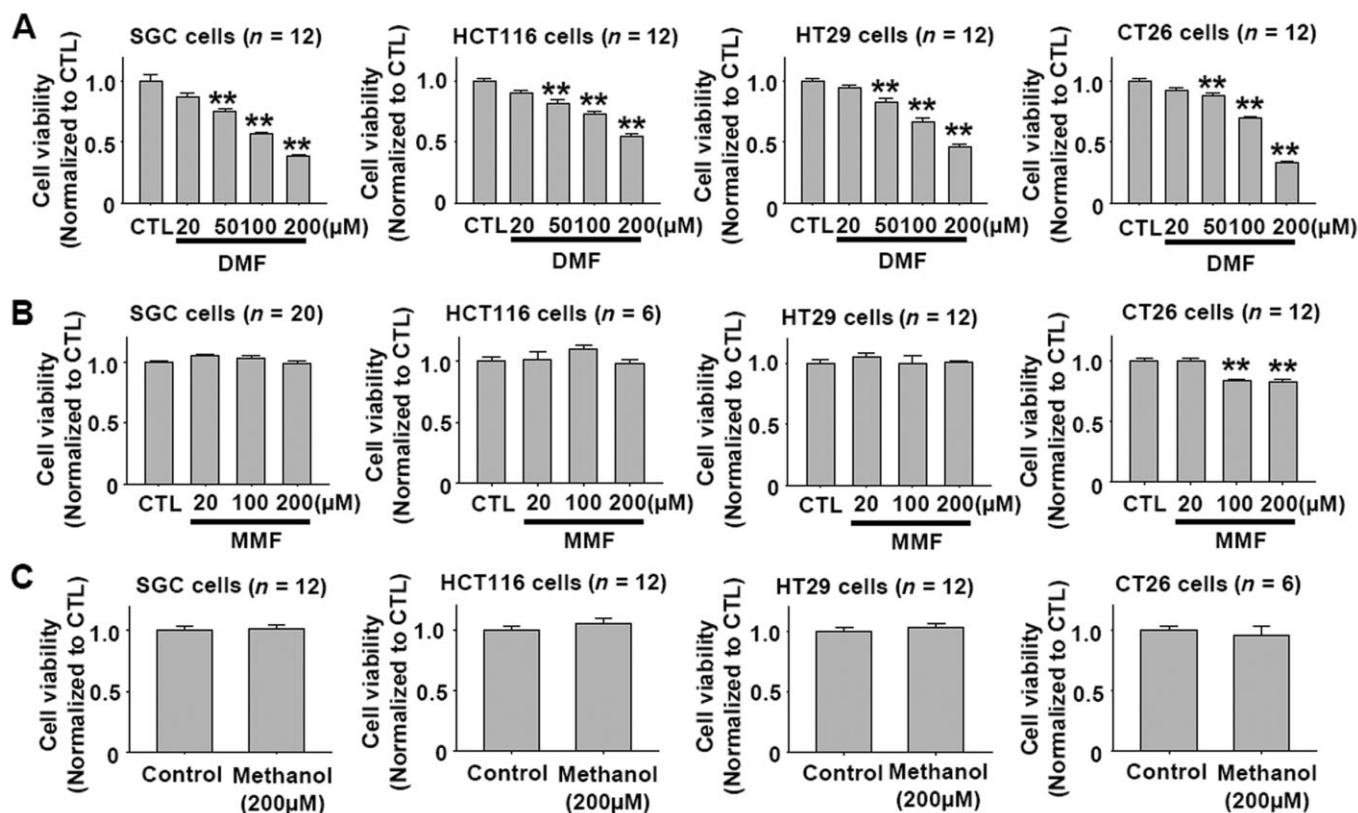


Figure 1

DMF, but not MMF or methanol, reduces the viability of SGC-7901, HT29, HCT116 and CT26 cancer cells after a 24 h treatment. (A) DMF dose-dependently reduced the viability of SGC-7901, HCT116, HT29, and CT26 cancer cells. $**P < 0.01$ versus CTL. CTL, control. (B–C) MMF and methanol showed no significant effect on the viability of SGC-7901, HCT116, HT29 and CT26 cancer cells, except that MMF at 100 and 200 μM slightly reduced CT26 cell viability. $**P < 0.01$ versus CTL. CTL, control.

Cell lines and agents

The murine colon adenocarcinoma cell line (CT26), human colon adenocarcinoma cell line (HT29), human colon cancer cell line (HCT116) and human gastric cancer cell line (SGC-7901) were purchased from the Type Culture Collection of the Chinese Academy of Sciences (Shanghai, China). DMEM was purchased from Hyclone (Logan, UT, USA). FBS was purchased from Biological Industries (Beit Haemek, Israel). The LIVE/DEAD® Viability/Cytotoxicity Assay Kit was obtained from Invitrogen (Eugene, OR, USA). Cyclosporine A (CsA), Ac-DEVD-CHO and LDH assay kits were purchased from Beyotime Institute of Biotechnology (Shanghai, China). Methanol was purchased from Tianjin Yongda Chemical Reagent Company (Tianjin, China), DMF, mono-methyl fumarate, DMH1(4-[6-(4-isopropoxyphenyl)-pyrazolo [1,5-a]pyrimidin-3-yl] quinoline) and 3-methyladenine (3-MA) were purchased from Sigma-Aldrich (St. Louis, MO, USA). Necrostatin-1 (Nec) and PARP-1 antibody were purchased from Santa Cruz (Eugene, OR, USA). Anti-LC3B antibody was purchased from Sigma-Aldrich. Anti-bax, Anti-bcl-2, Anti-gapdh and Anti-actin antibodies were purchased from Sangon Biotech (Shanghai, China). Anti-cleaved caspase-3 antibody was obtained from Cell Signaling Technology (Beverly, MA, USA). MTT and DMSO were obtained from Amresco (Solon, OH, USA). Cell counting kit-8 (CCK-8) was

obtained from Dojindo (Kumamoto, Japan). DMSO was used as a solvent for stock solutions and its final concentration was limited to 4%, which showed no effect on cell viability. The positive inducers of apoptosis, apopida and apobid, were purchased from the Beyotime Institute of Biotechnology.

Results

DMF but not MMF or methanol reduces cell viability in SGC-7901, HT29, HCT116 and CT26 cancer cells

DMF was reported to inhibit melanoma growth and metastasis *in vivo* and *in vitro* (Loewe *et al.*, 2006; Yamazoe *et al.*, 2009). The concentration of DMF commonly used in these *in vitro* experiments is in the range 50 to 200 μM (Loewe *et al.*, 2006; Ghods *et al.*, 2013). Hence, we initially examined the effects of DMF 20–200 μM on cell viability in four types of gastrointestinal cancer cell lines, SGC-7901, HT29, HCT116 and CT26. Because DMF is metabolized into MMF and methanol in the body (Werdenberg *et al.*, 2003; Rostami-Yazdi *et al.*, 2010), we further tested the effects of MMF and methanol on the cell viability in these cell lines. As shown in Figure 1A, DMF dose-dependently reduced the viability of SGC-7901, HT29, HCT116 and CT26 cancer cells in the range

20–200 μM . However, MMF and methanol at the same concentrations showed no significant effects on cell viability, with the exception that MMF at 100 and 200 μM slightly reduced the viability of CT26 cells (Figure 2A and B).

DMF does not induce cell apoptosis in HCT116 and CT26 cancer cells

DMF was reported to induce cell apoptosis in human mast cells (Forster *et al.*, 2013). Since DMF reduced the viability of SGC-7901, HT29, HCT116 and CT26 cells, we further investigated whether the reduced cell viability was due to the

induction of apoptosis in HCT116 and CT26 cells by measuring different apoptosis-related factors, bcl-2, bax, caspase-3, in these cells after DMF treatment. As shown in Figure 2A and B, DMF in the range 20 to 200 μM showed no effect on the protein expressions of bcl-2, bax, cleaved caspase-3 in HCT116 and CT26 cells after a 24 h treatment. DMF at 100 μM also showed no effect on bax, cleaved caspase-3 in HCT116 and CT26 cells after treatment from 3 to 24 h (Figure 2C and D). Furthermore, DMF-induced decrease in cell viability in CT26 cells was not affected by the caspase-3 inhibitor Ac-DEVD-HO (Figure 2E). The above data indicated that DMF-induced decrease in cell viability was not due to the

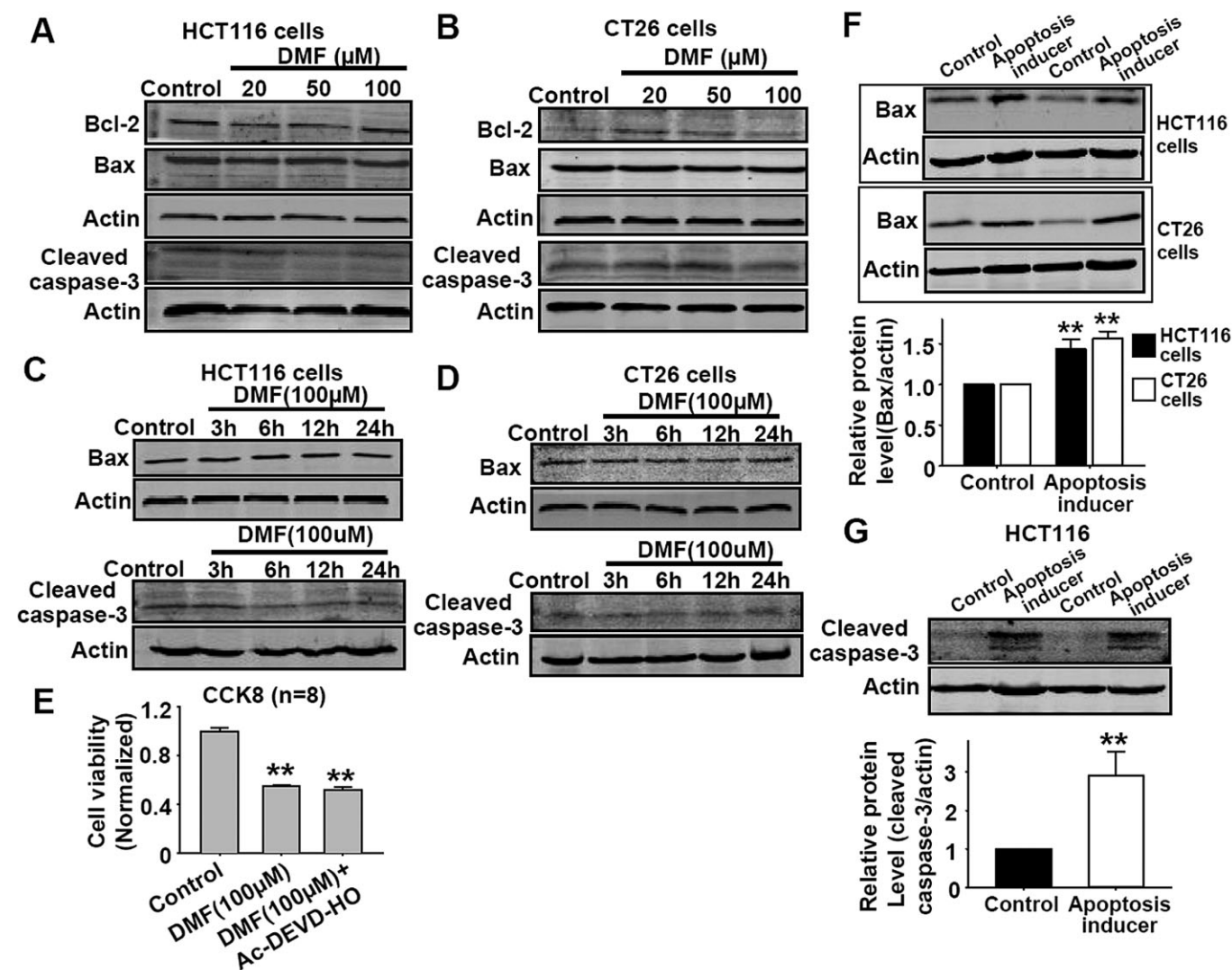


Figure 2

DMF does not affect apoptotic parameters in HCT116 and CT26 cancer cells. (A–B) DMF in the range from 20 μM to 200 μM showed no effect on bcl-2, bax, cleaved caspase-3 protein expressions in HCT116 and CT26 cells after 24 h treatment. (C–D) DMF (100 μM) showed no effect on bax, cleaved caspase-3 protein expressions in HCT116 and CT26 cells after different treatment durations (3, 6, 12 and 24 h). (E) DMF-induced reduction in cell viability of CT26 cells was not affected by pretreatment with the caspase-3 inhibitor Ac-DEVD-HO (100 μM) for 30 min. The treatment time of DMF was 24 h. ** $P < 0.01$ versus CTL. CTL, control. (F) The positive apoptosis inducer increased bax protein expression in HCT116 and CT26 cells; $n = 6$, ** $P < 0.01$ versus control. (G) The positive apoptosis inducer increased the cleaved caspase-3 protein expression in HCT116 cells; $n = 5$, ** $P < 0.01$ versus control. The positive apoptosis inducer was apopida and apobid at a concentration of 1:1000 in the medium.

induction of apoptosis. In order to confirm our experimental conditions, we treated these cells with a positive apoptosis inducer and measured the protein expressions of bax and cleaved caspase-3 protein; this positive apoptosis inducer increased the expression of bax and cleaved caspase-3 (Figure 2F and G).

DMF induces cell necroptosis

LDH release is a typical property of cell necrosis (Huang *et al.*, 2013). The LDH release from CT26 cells treated with 100 μ M DMF for different time was measured to determine whether necrosis was involved in DMF-induced decrease in cell viability. As shown in Figure 3A, DMF (100 μ M) time-dependently increased the release of LDH, which reflected necrotic cell death. We further examined the release of HMGB1 and the activation of PARP-1, which reflect cell necroptosis. As shown in Figure 3B, DMF induced HMGB1 release in CT26 cells. It had no significant effect on the protein expression of PARP-1 (Figure 3C), but DMF treatment significantly increased PARP-1 activity (Figure 3D). Nec, an inhibitor of receptor interacting protein-1 kinase, is a pharmacological tool used to identify cell necroptosis (Degterev *et al.*, 2008); the dose of Nec needed to have this effect has been shown to range from 30 to 100 μ M (Degterev *et al.*, 2008; El-Mesery *et al.*, 2015). We found that Nec at 20 μ M showed no significant effect on DMF-induced decrease in cell viability, but it significantly attenuated DMF-induced decrease of cell viability when the dose was increased to 60 and 100 μ M (Figure 3E). Optical photography also showed that DMF (100 μ M) reduced the cell density and Nec treatment partially reversed this decrease in cell density (Figure 3F). We further tested the effect of Nec using the LIVE/DEAD® Viability/Cytotoxicity Assay. As shown in Figure 3G, DMF (100 μ M) significantly reduced the number of green cells and increased the number of red cells, and these changes were partially reversed by Nec treatment. In principal, the red staining reflects the cell membrane integrity. Therefore, we speculated that DMF disrupted the cell membrane integrity. We further used TEM to observe the DMF-induced cell death in CT26 cells. As shown in Figure 3H, DMF (100 μ M) increased cell death, which was characterized by mitochondrial swelling, reticulum expansion and autolysosome formation. Pretreatment with Nec (100 μ M) for 1 h significantly attenuated DMF-induced morphological changes in CT26 cells. Taken together, the above results indicate that DMF induced cell necroptosis.

Necroptosis involves the active disintegration of mitochondrial membranes (Han *et al.*, 2007; Vandenabeele *et al.*, 2010). CsA inhibits mitochondrial permeability transition pore (MPTP) opening and attenuates necrotic damage (Petrosillo *et al.*, 2009; Vaseva *et al.*, 2012; Zhang *et al.*, 2014). Hence, we investigated the effect of CsA on DMF-induced cell death in CT26 and HT29 cells using the LIVE/DEAD® Viability/Cytotoxicity Assay. As shown in Figure 4A and B, CsA treatment significantly inhibited DMF-induced cell death in CT26 and HT29 cells. MMF and methanol treatments were investigated as parallel controls and showed no effects. Indeed, CsA treatment also inhibited DMF-induced mitochondrial membrane depolarization (Figure 4C). These data indicate that MPTP opening was involved in DMF-induced cell necroptosis.

DMF induces cell necroptosis by depleting GSH

GSH is the major redox buffer in cells (Mailloux *et al.*, 2013). Supplements of exogenous GSH or the GSH precursor N-acetylcysteine (NAC) can attenuate hemin-induced astrocyte and oridonin-induced hepatic stellate cell death (Laird *et al.*, 2008; Kuo *et al.*, 2014). DMF has been shown to deplete cellular GSH (Lehmann *et al.*, 2007; Schmidt *et al.*, 2007), indicating that DMF-induced GSH depletion might contribute to DMF-induced cell necroptosis. As shown in Figure 5A and B, the DMF-induced decrease in cellular GSH level and cell viability was inhibited by co-treatment with GSH and NAC in CT26 cells, but Nec and CsA treatments showed no effects on DMF-induced decrease of cellular GSH (Figure 5C). The protective effects of GSH and NAC against DMF-induced cell death were further confirmed by using the LIVE/DEAD® Viability/Cytotoxicity Assay (Figure 5D). The DMF-induced increase in ROS generation was also inhibited by the co-treatment with GSH and NAC (Figure 5F), and DMF-induced mitochondrial depolarization was also inhibited by co-treatment with GSH, NAC and Nec in CT26 cells (Figure 5F).

MAPKs activation is involved in DMF-induced cell necroptosis

MAPKs are the important downstream signals of ROS. Since DMF induced ROS generation, MAPKs would be subsequently activated. As shown in Figure 6A, DMF (100 μ M) significantly activated JNK, p38 and ERK in CT26 cells after treatment from 3 to 24 h, and the maximal activation appeared at the 6 h time point. Treatment with the JNK inhibitor SP6001257, p38 inhibitor SB202190 or ERK inhibitor PD98059 partially reversed DMF-induced decrease in cell viability (Figure 6B).

Previous data showed that co-treatment with GSH or NAC inhibited DMF-induced necroptosis (Figure 5B). Similarly, co-treatment with GSH or NAC inhibited DMF-induced JNK, p38 and ERK activation in CT26 cells (Figure 7). Although Nec restored DMF-induced necroptosis, Nec did not inhibit DMF-induced JNK, p38 and ERK activation (Figure 7), indicating that the target of Nec was downstream of the MAPKs signals.

DMF induces cell autophagy

Autophagy is a natural cellular response to many stimuli, and is usually represented by LC3-II protein expression. Since DMF induced cell necroptosis, the effects of DMF on the levels of LC3-II in CT26, HT29, HCT116 and SGC-7901 cells were examined. As shown in Figure 8A–8D, DMF, but not MMF, increased autophagy responses in SGC-7901, HCT116, HT29 and CT26 cells. DMH1, an inhibitor of autophagy (Sheng *et al.*, 2014), inhibited the DMF-induced increased autophagy response in CT26 cells (Figure 8E). Although DMH1 inhibited the DMF-induced increase in autophagy in CT26 cells, it did not prevent the DMF-induced decrease in cell viability (Figure 8F).

In order to identify the role of autophagy in DMF-induced cell necroptosis, we further studied the effects of Nec and CsA on basal and DMF-induced autophagy responses. As shown in Figure 9, Nec or CsA treatment attenuated DMF-induced autophagy responses, but Nec or CsA alone did not affect the

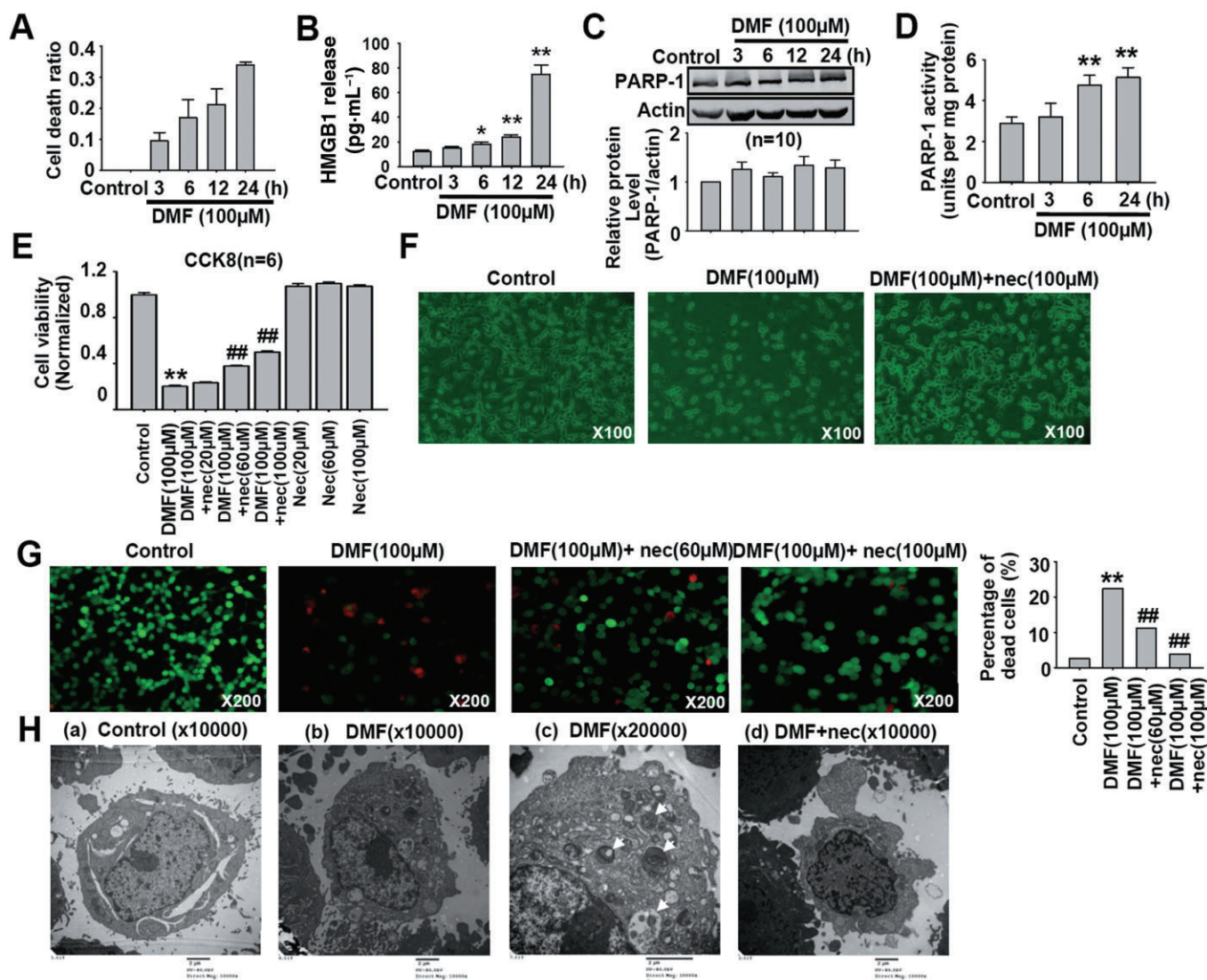


Figure 3

Nec inhibits DMF-induced cell necroptosis. (A) DMF (100 μM) time-dependently increased the cell death ratio in CT26 cells. The treatment time of DMF was set as 3, 6, 12 and 24 h. (B) DMF induced HMGB1 release in CT26 cells. The treatment time of DMF was set as 3, 6, 12 and 24 h. * $P < 0.05$, ** $P < 0.01$ versus control; $n = 4$. (C) DMF treatment showed no significant effect on PARP-1 protein expression in HCT116 cells. The treatment time of DMF was set as 3, 6, 12 and 24 h. (D) DMF treatment significantly increased PARP1 activity in HCT116 cells. ** $P < 0.01$ versus control; $n = 6$. (E) CCK8 cell viability assay showed that Nec inhibited DMF-induced decrease in CT26 cell viability. Nec alone had no effect on CT26 cell viability. Cells were pretreated with Nec for 1 h, then treated with both DMF and Nec for 24 h. ** $P < 0.01$ versus control. ## $P < 0.01$ versus DMF. (F) Representative photos of CT26 cells treated with DMF (100 μM) and DMF (100 μM) plus Nec (100 μM). Cells were pretreated with Nec for 1 h, then treated with both DMF and Nec for 24 h. (G) The LIVE/DEAD® Viability/Cytotoxicity assay showed that DMF(100 μM) induced CT26 cell death, which was partially reversed by Nec (100 μM) treatment. The live cells appear as green and dead cells as red. The number of cells counted: 4136, 5874, 2276 and 2122 in control, DMF, DMF plus Nec (60 μM), and DMF plus Nec (100 μM) respectively. Cells were pretreated with Nec for 1 h, then treated with both DMF and Nec for 24 h. χ^2 test was used, ** $P < 0.01$ versus control; ## $P < 0.01$ versus DMF. (H) Representative electron microscope photographs showed that Nec attenuated DMF-induced cell necroptosis. (a) CT26 cells treated with DMSO were considered as control. (b, c) DMF (100 μM) treatment led to cell necrosis, which is characterized by a disrupted cell membrane, mitochondrial swelling, reticulum expansion and autolysosome formation in CT26 cells as indicated by arrows. (d) Pretreatment with Nec(100 μM) for 1 h attenuated DMF-induced necroptosis in CT26 cells with intact cell membrane and reduced autolysosome formation.

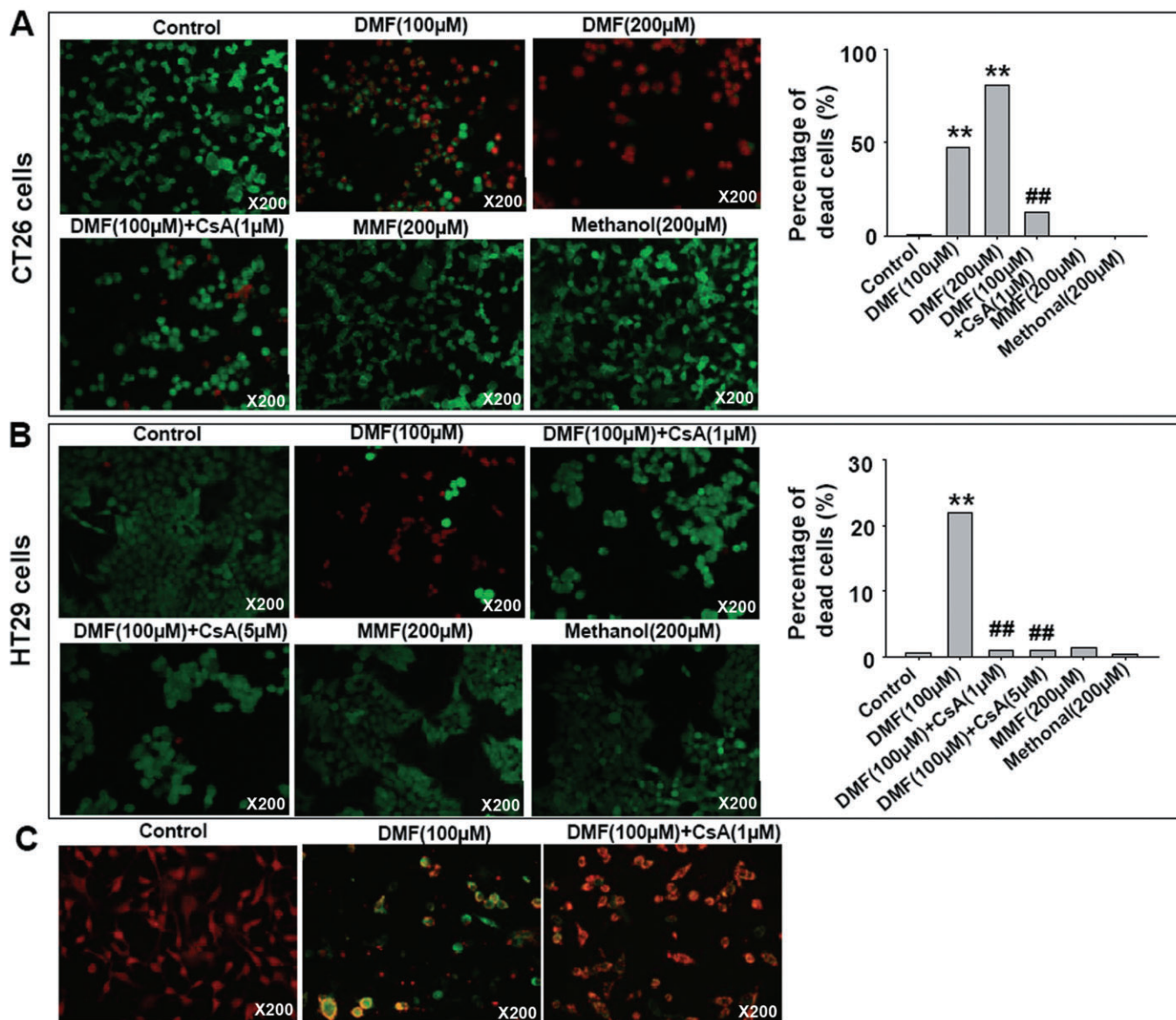


Figure 4

CsA inhibited DMF-induced cell death. (A, B) CsA inhibited DMF-induced cell death in CT26 and HT29 cells as indicated by using live/dead assay. MMF and methanol did not induce cell death. The live cells appear as green and dead cells as red. (A) The number of cells counted: 7833, 1921, 6826, 1123, 14381 and 12381 in control, DMF (100 μM), DMF (200 μM), DMF (100 μM) + CsA (1 μM), MMF (200 μM) and methanol (200 μM) respectively. (B) The number of cells counted: 4592, 4038, 5429, 1189, 4756 and 3568 in control, DMF (100 μM), DMF (100 μM) + CsA (1 μM), DMF (100 μM) + CsA (5 μM), MMF (200 μM) and methanol (200 μM) respectively. Cells were pretreated with CsA for 30 min, then treated with both DMF and CsA for 24 h. χ^2 test was used, $**P < 0.01$ versus control; $##P < 0.01$ versus DMF. (C) CsA treatment attenuated DMF-induced mitochondrial membrane depolarization in CT26 cells, evaluated by JC-1 staining. In normal conditions, JC-1 accumulated in the mitochondrial matrix to form J-aggregates, which fluoresced red and indicated a high MMP. When cell MMP was low, JC-1 would not accumulate in the mitochondrial matrix and exists in its monomer form, which fluoresced green.

basal or starvation-induced autophagy responses in CT26 cells. These data indicate that DMF-induced autophagy was a secondary response in these cells.

Discussion

DMF, the methyl ester of fumaric acid, has been approved for the treatment of relapsing forms of multiple sclerosis and

relapsing-remitting multiple sclerosis clinically (Burness and Deeks, 2014; Fox *et al.*, 2014). In the present study, we found that DMF but not its metabolite MMF induces necroptosis in colon cancer cells and the mechanism involves GSH depletion, an increase in ROS and activation of MAPKs-mediated signalling. The mechanism of DMF-induced cell necroptosis is summarized in Figure 10. In the present study, we examined the effects of DMF on cell viability in four types of cancer cell lines, SGC-7901, HT29, HCT116 and CT26, and

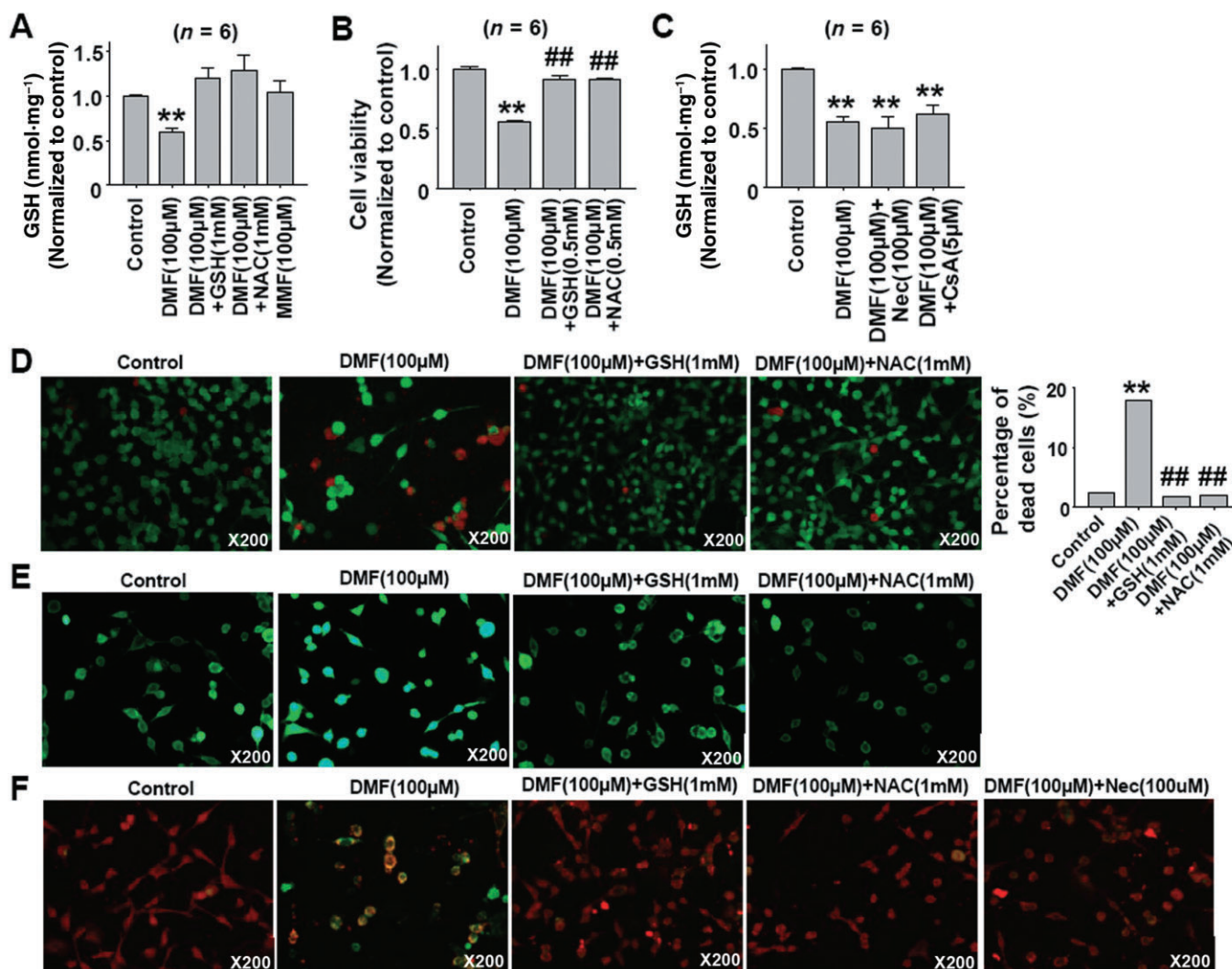


Figure 5

GSH and NAC attenuate DMF-induced CT26 cell necroptosis. (A) DMF induced reduction in cellular GSH levels in CT26 cells and the reversal of this reduction by GSH and NAC treatment. Cells were pretreated with DMF(100 μM) for 1 h, then GSH(1 mM) and NAC(1 mM) were added for 2 h. $**P < 0.01$ versus control. (B) GSH and NAC treatment alleviated DMF-induced the decrease of CT26 cell viability. Cell were pretreated with DMF(100 μM) for 1 h, then treated with both DMF(100 μM) and GSH(0.5 mM) or NAC(0.5 mM) for 24 h. $**P < 0.01$ versus control; $##P < 0.01$ versus DMF(100 μM). (C) Nec and CsA treatment did not affect DMF-induced reduction of cellular GSH levels in CT26 cells. Cells were pretreated with Nec (100 μM) for 1 h and CsA(5 μM) for 30 min, then DMF(100 μM) was added until 3 h. $**P < 0.01$ versus control. (D) GSH and NAC treatment attenuated DMF-induced cell death in CT26 cells, shown using the LIVE/DEAD staining assay. Cells were pretreated with DMF(100 μM) for 1 h, then GSH(1 mM) and NAC(1 mM) were added until 24 h. The live cells appear as green and dead cells as red. The total number of cells counted cells: 4136, 1857, 3988 and 2811 in control, DMF (100 μM), DMF (100 μM) plus GSH (1 mM) and DMF (100 μM) plus NAC (1 mM) respectively. χ^2 test was used, $**P < 0.01$ versus control; $##P < 0.01$ versus DMF. (E) GSH and NAC treatment attenuated the DMF-induced ROS increase in CT26 cells as shown using DCFH-DA staining. Cells were pretreated with DMF(100 μM) for 1 h, then GSH(1 mM) and NAC(1 mM) were added until 3 h. DMF treatment increased the cell fluorescence intensity. (F) DMF-induced mitochondrial depolarization was inhibited by GSH, NAC and Nec treatment in CT26 cells. Cell were pretreated with DMF(100 μM) for 1 h, then GSH (1 mM) and NAC (1 mM) were added for 5 h. For Nec treatment, cells were pretreated with Nec (100 μM) for 1 h, then DMF (100 μM) was added for 6 h. The MMP was evaluated by JC-1 staining. Red fluorescence indicates a high MMP and green fluorescence indicates a low MMP concentration.

then mainly used CT26 cells to elucidate the mechanism of DMF's action.

DMF and its *in vivo* metabolite MMF are fumaric acid esters and an important property of this class of compound is its ability to induce antioxidant effects by up-regulating the expression of nuclear factor (erythroid-derived 2)-like 2

(Nrf2)-driven antioxidant response genes (Gill and Kolson, 2013). In both the nervous and cardiovascular systems, oxidative stress is associated with numerous diseases; thus, fumaric acid esters show protective effects on a wide variety of disorders. DMF and MMF protect neurons and astrocytes against oxidative stress-induced cellular injury and loss

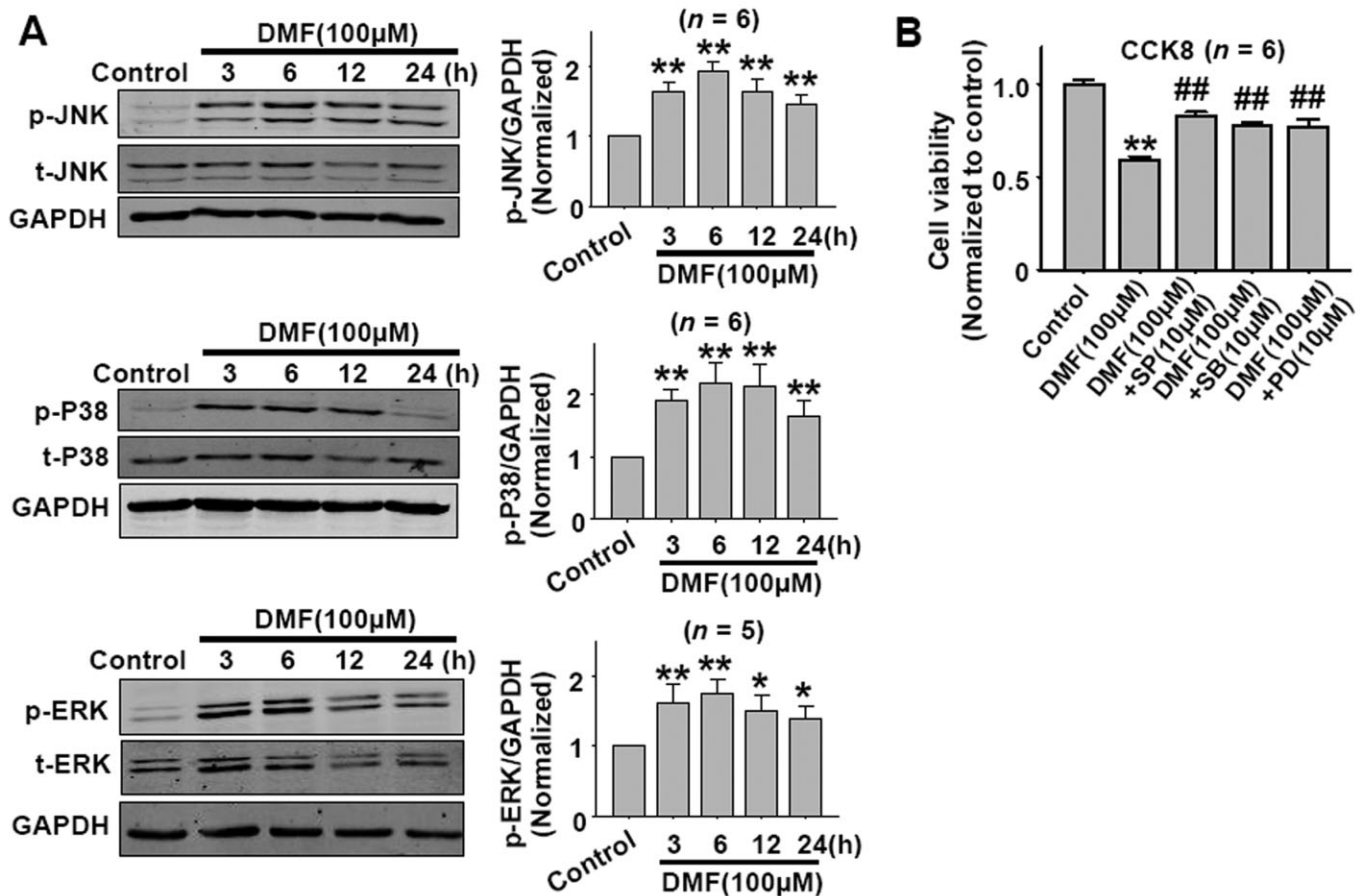


Figure 6

DMF activates JNK, p38 and ERK MAPKs in CT26 cells. (A) The time course of DMF-induced JNK, p38 and ERK activation in CT26 cells. * $P < 0.05$, ** $P < 0.01$ versus control. (B) JNK, p38 and ERK inhibitors attenuated DMF-induced cell death in CT26 cells as shown by the CCK8 assay. SP, JNK inhibitor SP600125; SB, p38 inhibitor SB203580; PD, ERK inhibitor PD98059. ** $P < 0.01$ versus control; ### $P < 0.01$ versus DMF. Cells were pretreated with the JNK inhibitor SP600125, p38 inhibitor SB203580, or ERK inhibitor PD98059 for 1 h, then DMF(100 µM) was added until 24 h.

(Scannevin *et al.*, 2012), and protect neurons and glia from oxidative stress-induced cell death (Linker *et al.*, 2011). DMF treatment attenuates axonal loss and reduces astrocyte activation in wild-type mice, but not in *Nrf2*^{-/-} knockout mice, suggesting a role for Nrf2-dependent antioxidant response element activation in the neuroprotection mediated by fumaric acid esters (Linker *et al.*, 2011). In the cardiovascular system, augmentation of myocardial fumarate, whether through fumarate hydratase depletion or by DMF treatment, was found to be cardioprotective against ischaemia-reperfusion injury (Ashrafi *et al.*, 2012). In spontaneously hypertensive rats that transgenically express human C-reactive protein, fumaric acid esters ameliorated inflammation, oxidative stress and metabolic disturbances (Silhavy *et al.*, 2014). However, from the present data, we speculated that DMF can both induce and inhibit oxidative stress, its effect depending on the dose used.

Previous studies showed that DMF or MMF at 10 µM had protective effects produced through the induction of antioxidant response genes (Ashrafi *et al.*, 2012; Scannevin *et al.*, 2012). We found that DMF at 10 µM showed no cytotoxic effects on colon cancer cells. However, when the concentra-

tion of DMF was increased (50 to 100 µM), significant cytotoxic effects occurred. Furthermore, the rapid cytotoxic effects of DMF (100 µM) appeared 3 h after DMF treatment and the GSH level decreased simultaneously. Thus, even if DMF at this concentration could induce an up-regulation of antioxidant response genes, the cytotoxic effects were so rapid that there is not enough time for the DMF to up-regulate the expression of antioxidant genes, which would protect the cells. GSH scavenges ROS with the aid of GSH peroxidase (Mailloux *et al.*, 2013), and GSH depletion can result in cell death due to increased levels of ROS (You and Park, 2010; Franco and Cidlowski, 2012). As DMF is an α,β -unsaturated carboxylic acid ester, it can react spontaneously with thiols in GSH via a Michael-type reaction, whereas MMF reacts at a much lower rate (Schmidt *et al.*, 2007). Consistent with this, we found that MMF showed no significant effects on cell viability or on the cellular level of GSH.

Apoptosis is defined as a process driven by a set of molecular mechanisms that programmes the cells to die. Apoptotic cell death involves the engagement of pathways that result in the activation of caspase proteases. Necroptosis is defined as a receptor-interacting protein kinase-3 dependent cell death,

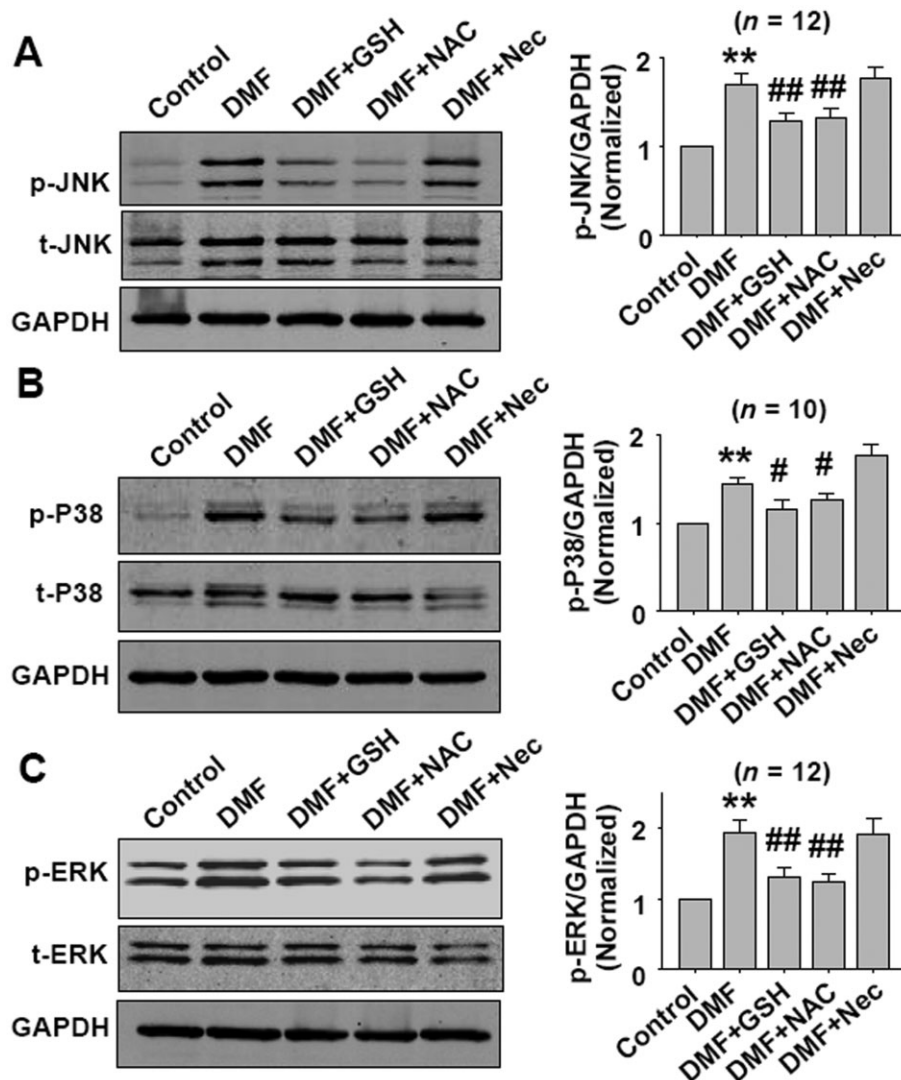


Figure 7

GSH and NAC treatment inhibits DMF-induced JNK, p38 and ERK activation in CT26 cells. Cells were pretreated with DMF (100 μ M) for 1 h, then treated with DMF plus GSH or DMF plus NAC for 5 h. For Nec treatment, cells were pretreated with Nec (100 μ M) for 1 h, then DMF (100 μ M) was added for 6 h. The concentrations of DMF, GSH, NAC and Nec were 100 μ M, 1 mM, 1 mM and 100 μ M respectively. ** P < 0.01 versus control. # P < 0.05, ## P < 0.01 versus DMF.

and is implicated in a variety of disease states, such as stroke, atherosclerosis, ischaemia-reperfusion injury and pancreatitis (Linkermann and Green, 2014). Cell necroptosis has different characteristics compared with cell apoptosis, both biochemically and morphologically, and has been well reviewed (Christofferson and Yuan, 2010; Galluzzi *et al.*, 2012; Linkermann and Green, 2014). It has been proposed that targeting cancer cells by induction of necroptosis may have significant potential in cancer chemotherapy, especially in the apoptotic/drug resistance cancers (Hu *et al.*, 2007). Indeed, recently this hypothesis has been substantiated by the results of several studies (Grassilli *et al.*, 2013; Steinhart *et al.*, 2013; Yan *et al.*, 2013; Chromik *et al.*, 2014). In the present study, we found that DMF did not significantly affect apoptotic parameters, such as the expression of bcl-2, bax, cleaved caspase-3 protein, but it did induce cell necroptosis as

demonstrated by use of the pharmacological tool Nec, TEM and LDH release. We did not exclude the possibility that DMF induces cell apoptosis, but necroptosis was found to be the main method by which DMF induced cell death. Our results showed that DMF increased cellular ROS and induced cell necroptosis, indeed, ROS also induces cell apoptosis. We speculate that DMF depleted GSH and triggered cell necroptosis so rapidly that there was not enough time for the initiation of apoptosis. The ability of DMF to induce necroptosis would be an advantage for DMF as a potential anticancer drug. GSH levels in cancer cells are much higher than that in normal tissues (Perry *et al.*, 1993; Berger *et al.*, 1994) and elevated intracellular GSH is associated with drug resistance in cancer chemotherapy (Berger *et al.*, 1994); therefore, this effect of DMF would be relatively selective for cancer cells because of its ability to deplete GSH.

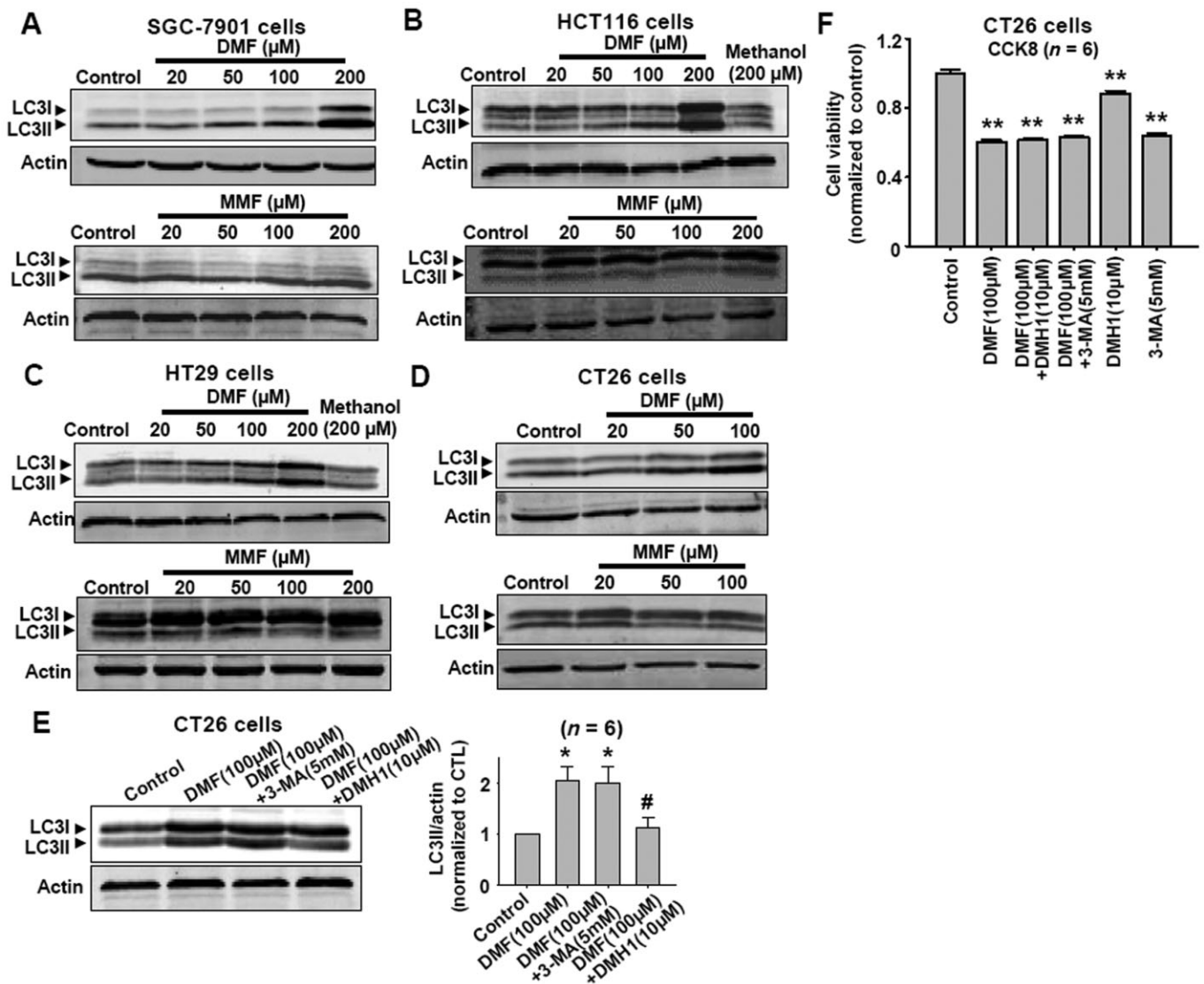


Figure 8

DMF induces autophagy responses in SGC-7901, HCT116, HT29 and CT26 cancer cells. (A–D) Western blot results showing that DMF but not MMF increased LC3-II levels in SGC-7901, HCT116, HT29 and CT26 cancer cells. (E) DMH1 inhibited DMF-induced increase in LC3-II in CT26 cells. * $P < 0.05$ versus control. # $P < 0.05$ versus DMF (100 μM). CTL, control. (F) DMH1 did not affect DMF-induced decrease in CT26 cell viability. ** $P < 0.01$ versus control.

Autophagy is a critical lysosomal process that maintains cellular homeostasis through the degradation of cellular components such as organelles and proteins, but when extensive, the autophagy response leads to autophagic cell death (Yang and Klionsky, 2010). Autophagy plays a complicated role in tumourigenesis and treatment responsiveness. On the one hand, induction of autophagy seems to be beneficial for cancer prevention and induction of autophagic cell death is considered as a therapeutic strategy; on the other hand, inhibiting autophagy can enhance the efficacy of anticancer therapies (Choi, 2012). Since DMF induced cell necroptosis, we further studied the effects of DMF on autophagy responses and the role of autophagy responses in DMF-induced cell death. We found that DMF

increased cell autophagy responses, but inhibiting DMF-induced autophagy by DMH1 did not prevent the DMF-induced decrease in cell viability. Nec or CsA treatment, which showed cytoprotective effects, attenuated DMF-induced autophagy responses, but Nec or CsA alone did not affect the basal and starvation-induced cell autophagy responses. These results indicate that the changes in cell autophagy were secondary responses to DMF stimulation. In the present study, we also used another classical autophagy inhibitor 3-MA. Our previous work and other studies have shown that 3-MA is not a stable autophagy inhibitor and has some non-specific effects (Wu *et al.*, 2010; Sheng *et al.*, 2013). Here, we found that 3-MA did not inhibit the DMF-induced increased autophagy.

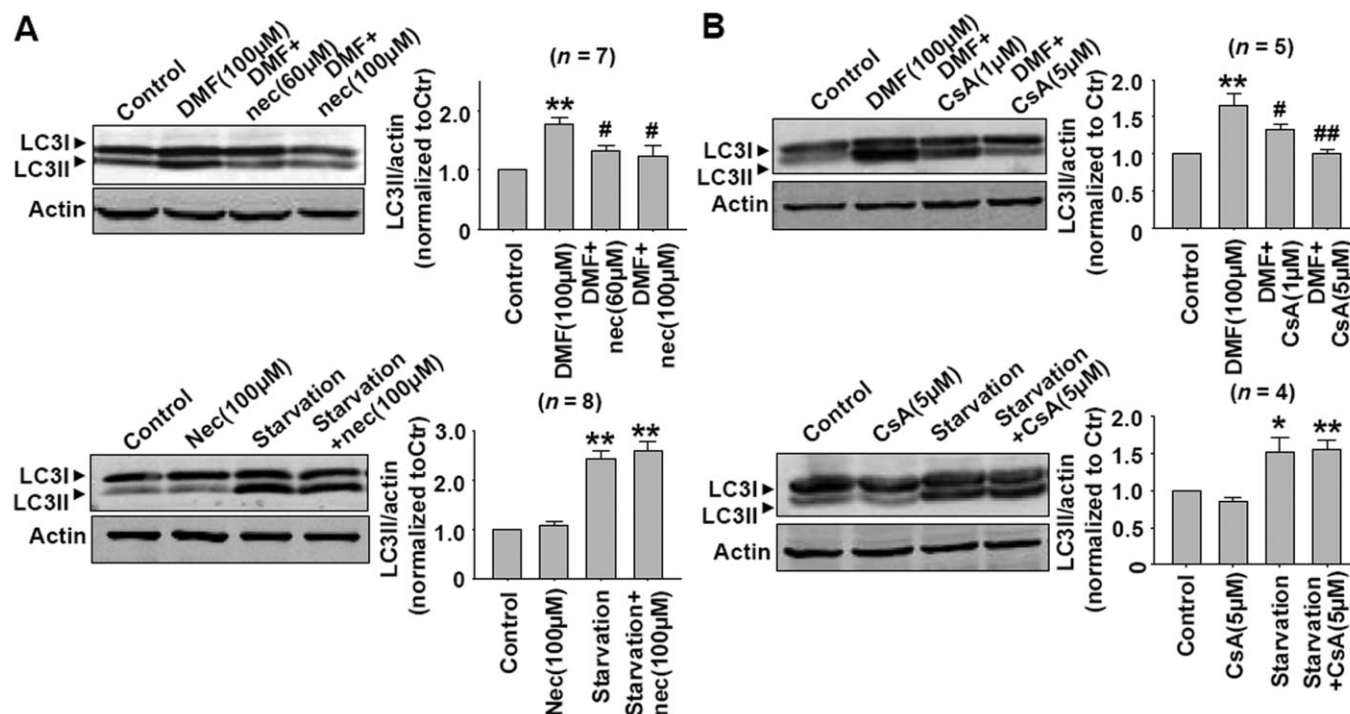


Figure 9

(A) Nec treatment attenuated DMF-induced autophagy responses, but Nec alone did not affect the basal and starvation-induced autophagy in CT26 cells. ** $P < 0.01$ versus control. # $P < 0.05$ versus DMF (100 μM). (B) CsA treatment attenuated DMF-induced autophagy responses, but CsA alone did not affect the basal and starvation-induced autophagy in CT26 cells. * $P < 0.05$, ** $P < 0.01$ versus control. # $P < 0.05$, ## $P < 0.01$ versus DMF (100 μM). The starvation condition was serum-free DMEM (5.5 mM glucose).

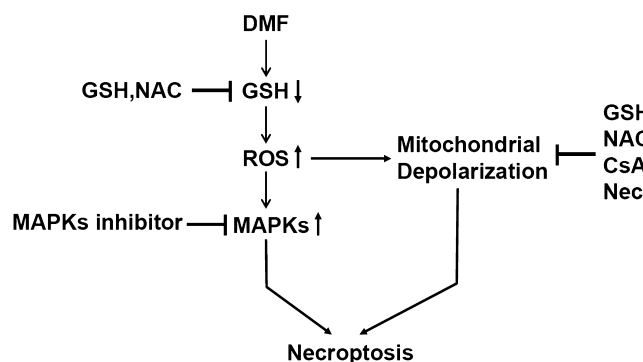


Figure 10

The mechanism of DMF-induced cell necroptosis.

We also measured the effects of DMF on the viability of normal cultured cardiomyocytes and cardiofibroblasts and found that DMF reduced the viability of these cells in a similar manner to the cancer cells (data not shown), indicating that these cytotoxic effects of DMF are non-specific. Currently, no necroptosis responses have been reported as a side effect of the clinical use of DMF. MMF, but not DMF, can be detected in serum after oral DMF ingestion (Werdenberg *et al.*, 2003; Rostami-Yazdi *et al.*, 2010). MMF is biologically active but not cytotoxic, so, the metabolism of DMF in the

intestine might be a process by which the systemic toxicity of DMF is reduced. These properties of DMF indicate that local administration of high concentrations of DMF into the intestine would be needed to directly kill cancer cells, but this route of administration would reduce its systemic toxicity by increasing its degradation to MMF. Therefore, DMF might have potential as a drug for localized colorectal cancer therapy. Colorectal cancer is a common and lethal disease, and tends to have drug resistance (Panczyk, 2014). At present, targeted drug delivery systems are being investigated for the direct administration of chemotherapeutic and chemopreventive agents to the colon and rectum (Krishnaiah and Khan, 2012). DMF is a promising drug candidate for colorectal cancer therapy provided an appropriate local drug delivery system can be devised.

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Author contributions

X X, Y Z, C-Y M, X-M X, Y-Q Z, C-G W, J J, X S, J-L G, N L performed the cell culture, Western blot and parameter detection by commercial kits. D-L D and Z-J S designed the project. D-L D wrote the paper.

Conflict of interest

The authors declare no conflict of interest.

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