RESEARCH ARTICLE

Cobalt Chloride Induces Necroptosis in Human Colon Cancer HT-29 Cells

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Abstract

Necroptosis, also known as “programmed necrosis”, has emerged as a critical factor in a variety of pathological and physiological processes and is considered a cell type-specific tightly regulated process with mechanisms that may vary rather greatly due to the change of cell line. Here we used HT-29, a human colon cancer cell line, to establish a necroptosis model and elucidate associated mechanisms. We discovered that cobalt chloride, a reagent that could induce hypoxia-inducible factor-1α (HIF1α) expression and therefore mimic the hypoxic microenvironment of tumor tissue in some aspects induces necroptosis in HT-29 cells when caspase activity is compromised. On the other hand, apoptosis appears to be the predominant death form when caspases are functioning normally. HT-29 cells demonstrated significantly increased RIPK1, RIPK3 and MLKL expression in response to cobalt chloride plus z-VAD treatment, which was accompanied by drastically increased IL1α and IL6 expression, substantiating the notion that necrosis can induce profound immune reactions. The RIPK1 kinase inhibitor necrostatin-1 and the ROS scavenger NAC each could prevent necrosis in HT-29 cells and the efficiency was enhanced by combined treatment. Thus by building up a necroptosis model in human colon cancer cells, we uncovered that mechanically RIP kinases collaborate with ROS during necrosis promoted by cobalt chloride plus z-VAD, which leads to inflammation. Necroptosis may present a new target for therapeutic intervention in cancer cells that are resistant to apoptotic cell death.

Keywords: Cobalt chloride - colon cancer cells - necroptosis - RIPK1 - ROS

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Introduction

Colon cancer is the third most common malignant disease in men and the second in women worldwide and it is seen with substantially increasing mortality in the Asia-Pacific region (Ku et al., 2012). Besides surgical procedures, adjuvant chemotherapy only benefit high-risk stage II and III colon cancer patients slightly with approximately 20 months median overall survival (Ku et al., 2012). Considering its high financial cost and toxic effects, it is critical to discover potent targeted treatment to fight colon cancer.

Understanding cell death mechanisms in cancer cells and how the immune system responds to it is pivotal in unraveling effective cancer treatment. Apoptosis and necrosis are two major forms of cell death (Wang et al., 2008; ME, 2011; Sun et al., 2012). Apoptosis is executed by intracellular cysteine protease family of caspases. Conversely, necrosis has been perceived as a passive and unregulated cellular demise resulted from unmanageable physical damages (Wang et al., 2008; ME, 2011; Sun et al., 2012). However, accumulating evidence indicates that necrosis is involved in a variety of physiological processes including T lymphocyte development, bacterial and viral infection, neurodegenerative diseases, auto-immune disorders, as well as ischemia-reperfusion injury, which underscores the crucial role of programmed necrosis in maintaining metazoans stable (Cho et al., 2009; Welz et al., 2011; Zhang et al., 2011; Narayan et al., 2012; Saddoughi et al., 2013). Furthermore some necrosis can be induced by regulated signal transduction pathways such as receptor interacting protein (RIP) kinase. Researches showed that in certain cells deficient for caspase-8 or treated with the pan-caspase inhibitor z-Vad, stimulation of death domain receptors such as the Fas and TNFR family causes a RIP1 kinase-dependent programmed necrotic cell death instead of apoptosis (Wang et al., 2008; ME, 2011; Sun et al., 2012). Other studies have suggest otherwise (Fu et al., 2013). Programmed necrotic cell death induced by either alkylating or oxidizing agents has been shown to be independent of the RIP kinase-dependent necrototic pathway (Fu et al., 2013).

Cobalt chloride is among the most well-established hypoxia mimics in the generation of ROS and activation of HIF1α, p53, p21 and PCNA, which is widely used in a variety of tissues and cell lines to create anemia and hypoxia environment (Zou, 2001). Here we used cobalt chloride to mimic the hypoxia microenvironment of

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colonic neoplasm thus studying necroptosis in a more physiological context.

By establishing a human colon cancer model of necroptosis with cobalt chloride, we have shown that in the presence of pan-caspase inhibitor z-Vad, necrosis becomes the predominant form of tumor cell loss. Using a combination of morphological, biochemical, genetic, and pharmacological investigations, we further identified RIP kinases and ROS as essential mediators during this sophisticated process.

Materials and Methods

Cell culture

The human colon cancer cell line HT-29 was obtained from American Type Culture Collection (Manassas, VA). Cells were grown in Maccoy’s 5A medium supplemented with 10% fetal bovine serum (FBS, Hyclone Laboratories Inc., Logan, UT) and penicillin (100U/ml) and streptomycin (100mg/ml). Cultures were maintained at 37°C in a humidified atmosphere of 5% CO2.

Antibodies and reagents

TNF-α, z-VAD were obtained from R&D systems (R&D,MN). Necrostatin-1 was purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Propidium iodide and DCFHDA were from Sigma-Aldridge (Sigma-Aldridge, MO). Cobalt Chloride was obtained from BD Biosciences. Anti-rabbit IgG-HRP and GDPDH antibody were purchased from Sangon (Sangon, Shanghai). Anti-mouse IgG-HRP was purchased from Daco (Glostrup, Denmark).

Fluorescence microscopy

For detection of necrosis and apoptosis, 1.5x10^4 cells were seeded per well in a 24-well plate. Following 12 hours incubation for adherence, cells were stimulated as previously described for 48 hrs. Subsequently, cells were incubated with propidium iodide (50μg/ml) and annexin V-FITC (5μl,BD Biosciences) for 15 min at room temperature immediately followed by fluorescence microscopy. Digital images were processed in an identical manner using the advanced DP controller software.

Quantitative real-time PCR analysis

Total RNA was extracted from cultured cells using Trizol reagent (Invitrogen, Carlsbad, California, USA) and dissolved in diethylpyrocarbonate treated (DEPC) water. cDNA was synthesized using the Takara Reverse Transcription System Kit (Takara Biotechnology Co. Ltd., Japan) according to the manufacturer’s instruction. Real time quantitative RT-PCR was performed using the Sybr green premix kit (Takara). The PCR conditions were as following: 95°C for 3 min followed by 40 cycles at 95°C for 10 s and 55°C for 30 s. All reactions were performed in triplicate both biologically and technically.

Flow cytometry analysis

Enumeration of apoptotic cells was done by using FITC conjugated Annexin V and PI (BD Pharmingen, San Jose, CA). Cells were washed twice in cold PBS and resuspended in Annexin V-binding buffer (BD Pharmingen) at a concentration of 1x10^6 per ml. This suspension (100μl) was stained with 5μl of Annexin V-FITC and 5μl PI. These cells were gently vortexed and incubated for 15 min at room temperature in the dark. After addition of 400μl of binding buffer to each tube, cells were analyzed by flow cytometry.

ROS measurement

The reactive oxygen species (ROS) assay was performed according to the manufacturer’s instructions. Briefly, HT-29 cells were seeded in 96-well plates at a density of 2.5x10^4 cells per well and incubated at 37°C overnight for attachment. Change the culture medium to stimuli and incubated for an hour. Cells were then washed with PBS twice followed by the addition of 100μl of fresh culture medium without phenol red or FBS containing 25μM DCFHDA (Sigma) dissolved in dimethylsulfoxide (DMSO). After the cells were incubated for 30 min, they were washed with PBS twice and stayed in medium without phenol red and FBS. ROS absorbances were measured at 525 nm using a fluorescence microplate reader.

Western blotting

Equal amounts of protein were separated by 10% SDS polyacrylamide gel electrophoresis and transferred to PVDF membranes. After blocking with 5% non-fat milk in TBST for 1 h, membranes were incubated overnight at 4°C with Rip1, GAPDH primary antibodies followed by incubation with HRP-conjugated secondary antibodies. Immunoreactive bands were detected with enhanced chemiluminescent HRP substrate (Millipore, Bolleria, MA).

Statistical analysis

Data are presented as the mean±standard error of the mean (SEM). Values were compared and analyzed using Student’s t-test. We considered P-values less than 0.05 to be statistically significant.

Results

Cobalt chloride induces necrosis in HT-29 cells

It has been established that death-inducing cytokines as TNF-α causes necrosis in HT-29 cell line when caspases are inhibited (Wang et al., 2008; ME, 2011; Sun et al., 2012). We discovered that necrosis also occurred...
when cells were treated with cobalt chloride and even exacerbated necrosis emerged when cells were treated with cobalt chloride plus z-VAD (Figures 1A and 1B) while no major differences have been observed in apoptosis (Figure S1). We also noticed that cobalt chloride alone has greater necrosis-inducing effect than TNF-α (data not shown). To verify this cell death was indeed necrosis, cells treated with either cobalt chloride or cobalt chloride plus z-VAD were analyzed by transmission electron microscopy (TEM). Surprisingly, most cells treated with cobalt chloride exhibited apoptotic morphology changes as condensed chromatin and plasma membrane blebbing while a significant percentage of cells treated with cobalt chloride plus z-VAD displayed mitochondria swelling and discontinuous cytoplasmic membrane which are typical necrosis morphology (Figure 1C). We suspected that the lack of necrosis morphology in cells treated with cobalt chloride may attribute to the undistinguishable essence between late stage apoptosis and necrosis with propidium iodide staining.

**RIPK1-RIPK3 axis is required for necrosis induced by cobalt chloride**

Since RIPK1/RIPK3/MLKL complex is the best-characterized complex during necroptosis which is fundamental to cytokines as TNF-α, TRAIL, CD95 (Wang et al., 2008; ME, 2011; Sun et al., 2012) but is dispensable to either alkylating or oxidizing agents (Fu et al., 2013), we checked whether it is required for the observed necrosis caused by cobalt chloride. HT-29 cells under cobalt chloride plus z-VAD treatment expressed increased level of RIPK1, RIPK3 which expression steadily augmented over time while in cells treated with cobalt chloride alone RIPK1 RIPK3 had higher expression.
during earlier hours and then ameliorated (Figure 2A). MLKL expression peaked at 2 hours during both treatment (Figure 2A). RIPK1 expression was further confirmed by Western blotting (Figure 2B) after 12 hours of stimulation and we ruled out the possible effects of z-V AD alone (Figure S2 and data not shown). As a validation to our finding, necrostatin-1, an RIPK1 kinase inhibitor rendered HT-29 cells resistant to this type of cell death (Figure 2C). Furthermore necrosis is always associated with ensuing inflammation reaction (Zheng et al., 2013). In consistence with this notion, we find increasing levels of IL1-α and IL6 (Figure 2D) overtime. We also noticed that there was certain basal level of endogenous RIPK1 in HT-29 cells which may accordingly render HT-29 susceptible to programmed necrotic death in the first place.

**ROS is indispensable for Cobalt chloride-induced necrosis**

RIPK1 is a key regulator of necroptosis and its expression level has been shown to correlate with responsiveness to programmed necrosis (Wang et al., 2008; ME, 2011; Sun et al., 2012). We monitored RIPK1 expression level during early hours of necrosis and noticed a different trend between cobalt chloride and typical TNF-α treatment indicating that there may be different mechanisms lying here (Figure 3A) meanwhile z-V AD alone didn’t impose major effects (Figure S3). Since cobalt chloride alone could generate ROS (Zou W, 2001) we tested if ROS played a significant role under

**Figure 3.** ROS is Indispensable for Cobalt Chloride-Induced Necrosis. (A) HT-29 cells were treated with DMSO, cobalt chloride and/or z-V AD, TNFα and/or z-V AD for indicated time. Cell lysates were collected and subjected to western blot analysis of RIPK1 and GAPDH levels. (B) HT-29 cells were treated with DMSO, cobalt chloride and/or z-V AD for 2 hr. Cells were incubated with 25 μM DCFHDA. Intracellular ROS levels were quantified with a fluorescence microplate reader. Data are mean ± standard deviation. P<0.05 (C and D) Representative fluorescence microscopy and FACS analysis of annexin V and propidium iodide (PI) staining 2 h after stimulation with cobalt chloride plus z-V AD and cobalt chloride plus z-V AD plus NAC (1 mM). All experiments were at least repeated three times with similar results.

**Figure 4.** ROS Pathway and RIPK1-RIPK3 Axis Collaborate During Necroptosis. (A) After indicated treatment for 2 hr, cells were incubated with 25 μM DCFHDA. Intracellular ROS levels were quantified with a fluorescence microplate reader. Data are mean ± standard deviation. P<0.05. (B) HT-29 cells were treated as indicated for 12 hr. Cell lysates were collected and subjected to western blot analysis of RIPK1 and GAPDH levels. (C) Representative FACS analysis of annexin V and propidium iodide (PI) staining 2 h after stimulation as indicated. All experiments were at least repeated three times with similar results.
ROS pathway and RIPK1-RIPK3 axis collaborate during necroptosis

Because RIPK1-RIPK3 signaling can promote mitochondrial ROS production (Sakurai et al., 2008), necrostatin-1 was used to eliminate the possibility that ROS required for necrosis was induced by RIPK1-RIPK3 pathway. Indeed, ROS production increase was hindered but not completely obstructed by necrostatin-1 (Figure 4A), implying that RIP kinase did regulate downstream ROS production but there were more pathways responsible for ROS accumulation. We next sought to determine if RIPK1-RIPK3 signaling and ROS functions cohesively. As shown in Figure 4B, RIPK1 expression was elevated in cobalt chloride plus z-VAD treated cells and was substantially inhibited with the combination of RIPK1 kinase inhibitor necrostatin-1 and anti-oxidant NAC. Consistent with this, flow cytometry analysis showed that the presence of both necrostatin-1 and NAC significantly reduced PI positive cells (Figure 4C). Collectively these data demonstrate that ROS and RIPK1-RIPK3 cooperated during HT-29 human colon cancer cell necroptosis induced by cobalt chloride and z-VAD.

Discussion

Colon cancer is seen with increasing frequency in the Asia-Pacific region and it is one of the most important causes of cancer mortality worldwide. A great comprehension of tumor cell loss and its ensuing immune responses is important to elicit a successful anti-cancer treatment (Ullrich et al., 2008). While many of the cellular aspects of apoptosis have been characterized, the initiating factors and regulating mechanisms of programmed necrotic tumor cell death are only beginning to be found. In this study we established a programmed necrosis model in HT-29 human colon cancer cell line with cobalt chloride, a hypoxia-inducing agent mimicking tumor microenvironment. Furthermore, we found that additional pan-caspase inhibitor z-VAD treatment increases necrotic cell death during tumorigenesis but doesn’t affect apoptosis (Figure 1). These findings clearly demonstrate programmed necrosis occurs during tumorigenesis which is further revealed when caspases are inhibited. We then utilized this model to investigate the essential mediators of this specific process.

RIPK1 is a serine/threonine adaptor kinase that acts downstream of downstream of death domain receptors and is essential for both cell survival and death (Green et al., 2011). Although the kinase activity of RIPK1 is not required for NF-κB activation (He et al., 2009), such activity is crucial for both caspases activation and necrosis induction. Consistent with this, RIPK1 kinase inhibitor necrostatin-1 cotreatment effectively suppress necrotic tumor cell death (Figure 2). RIPK1 is reported to form a pro-necrosis complex with RIPK3/FADD/caspase-8 when treated with death cytokines such as TNFα, Fas (Oberst et al., 2011). Since necrosis is differently regulated in this model (Figure 2), the effect protein of RIPK1 under this circumstance is unknown. It should be our priority in the future to identify the unknown components of such pathway. Some study showed that DNA damage triggered by alkylating chemotherapy agents activates a RIPK1-independent necrotic signaling pathway which was carried out with no full consideration of tumor heterogeneity and microenvironment (Junttila and de Sauvage, 2013). By further exploiting this model, we could look into the programmed necrotic death induced by alkylating agents in a more physiological context and thus might uncover potential therapeutic targets.

Reactive oxygen species (ROS) have been implicated in TNF-α induced necrosis. Nox1 NAPDH oxidase has been shown to work downstream of RIPK1 to generate ROS in L929 and MEF cells (Whelan RS, 2012). Alternatively, RIP3 activates metabolic enzymes such as glutamate dehydrogenase 1, and thereby increases mitochondrial ROS production (Tu et al., 2009). However, ROS dependence seems cell-type specific with data showing that ROS quenching failed to prevent necrosis in HT-29 cells treated with TNF-α, Smac mimetic and z-VAD (ME, 2011). Our data indicated that ROS was indispensable during necrosis induced by cobalt chloride plus z-VAD treated HT-29 cells (Figure 3) and only part of which that was generated by RIPK1 (Figure 4).

Since many cancer cell types can evade apoptotic cell death through deficiencies in key apoptotic proteins or by inhibition of apoptotic pathways and evasion of apoptosis is recognized to result in resistance to anti-cancer therapies (de Bruin and Medema, 2008; Krysko and Vandenbergbelee, 2008), elucidation of the necrotic signaling cascades will certainly enable targeted therapeutics in eliminating apoptosis-resistant cancer cells. Indeed recent studies showed that apoptosis is dispensable for complete tumor regression after alkylating agent chemotherapy because other forms of cell death can be activated (Krysko et al., 2012), including programmed necrosis. Moreover, necrosis promotes the clearance of apoptosis-deficient tumors by amplifying the initial damage signal through activating inflammatory response which destroys damaged cells and facilitates the cleaning up of cell debris (Kaczmarek et al., 2013). Cancer cells also exhibit increased ROS generation that may promote cell proliferation and, in many cases, might be coupled with redox adaptation to promote cell survival and drug resistance (Saeidnia and Abdollahi, 2013). Targeting this unique biochemical alteration might be a feasible approach to achieve therapeutic selectivity and to overcome drug resistance. In preclinical models, combination of ROS-generating agents and drugs that inhibit ROS elimination showed great potency to promote...
ROS accumulation and enhance cancer cell cytotoxicity (Trachootham et al., 2009; Sosa et al., 2013). In this work, we identify that during necrosis in HT-29 cells induced by cobalt chloride plus z-VAD treatment RIPK1 kinase acts in concert with ROS. Simultaneous inhibition of both RIP1 kinase and ROS pathways with small molecule compounds provides significant protection of programmed cell death (Figure 4).

There is convincing preclinical and accumulating clinical evidence in support of the notion that successful antineoplastic therapies reinstate immunosurveillance (Motz and Coukos, 2013). Unlike apoptosis, programmed necrosis is highly inflammatory. Extracellular HMGB1 released during necroptosis generally boosts immune responses (Duprez et al., 2011). Therefore programmed necrosis presents potential targets for therapeutic intervention. However, whether the inflammation associated with necrosis is desirable or leads to further tumor growth is under debate (Krysko et al., 2012). Our study demonstrated a crucial link between necroptosis, tumor growth and its regulation by caspases: a mystery in five inflammatory response syndrome. Simultaneous inhibition of RIP kinase-dependent necrosis drives lethal systemic inflammatory response syndrome. Immunity, 35, 908-118.

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