Genotoxicity of chloroacetamide herbicides and their metabolites in vitro and in vivo

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Abstract. The toxicity of chloroacetamide herbicide in embryo development remains unclear. Acetochlor (AC) is a chloroacetamide that metabolizes into 2-ethyl-6-methyl-2-chloroacetanilide (CMEPA) and 6-ethyl-o-toluidine (MEA). The present study determined the potential effect of AC and its metabolites on embryo development. Both HepG2 cells and zebrafish embryos were exposed to AC, CMEPA and MEA in the presence or absence of co-treatment with anti-reactive oxygen species (ROS) reagent N-acetylcysteine. The generation of ROS, levels of superoxide dismutase (SOD) and glutathione (GSH) in HepG2 cells and lactate dehydrogenase (LDH) leakage from HepG2 cells were investigated. The effects of AC, CMEPA and MEA on DNA breakage, MAPK/ERK pathway activity, viability and apoptosis of HepG2 cells were examined by comet assay, western blotting, MTT assay and flow cytometry, respectively. Levels of LDH, SOD and GSH in zebrafish embryos exposed to AC, CMEPA and MEA were measured. The hatching and survival rates of zebrafish embryos exposed to AC, CMEPA and MEA, were determined, and apoptosis of hatched fish was investigated using acridine orange staining. The present data showed AC, CMEPA and MEA induced generation of ROS and decreased levels of SOD and GSH in HepG2 cells, which in turn promoted DNA breakage and LDH leakage from cells, ultimately inhibiting cell viability and inducing apoptosis, as well as phosphorylation of JNK and P38. However, co-treatment with N-acetylcysteine alleviated the pro-apoptosis effect of AC and its metabolites. Moreover, exposure to AC, CMEPA and MEA lead to toxicity of zebrafish embryos with decreased SOD and GSH and increased LDH levels and cell apoptosis, ultimately decreasing the hatching and survival rates of zebrafish, all of which was attenuated by treatment with N-acetylcysteine. Therefore, AC and its metabolites (CMEPA and MEA) showed cytotoxicity and embryo development toxicity.

Introduction

Chloroacetamide herbicides are commonly used worldwide in agriculture to control annual grasses and broadleaf weeds (1). China uses >10 million kg chloroacetamide annually (2). The effect of chloroacetamide herbicides on wildlife and humans has drawn attention (3,4). These herbicides enter the environment or biological systems by a variety of routes, such as water contamination, and persist from a few months to several years (5). Studies have detected 17.9-1,054.9 ng/l acetochlor (AC, a chloroacetamide) in surface water in China (6,7), which can be absorbed by aquatic organisms and induce adverse events. For example, juvenile red swamp crayfish treated with AC exhibit belly arch, equilibrium loss and lethargy (8). Moreover, through the food chain, chloroacetamide can spread to the whole ecosystem, including humans.

The US Environmental Protection Agency has identified AC as a potential B-2 carcinogen (9). AC induces oxidative stress and regulates expression levels of genes both in vitro and in vivo (10,11). Li et al (12) suggested that AC affects larval development and adult brain of rare minnow and Yang et al (13) found that AC alters gene expression levels of the hypothalamic-pituitary-thyroid axis, which changes thyroid hormone levels in zebrafish. Furthermore, exposure to chloroacetamide has been found to increase the risk of certain diseases in humans, such as cancer (14) and Parkinson's disease (15).

Although potential toxicity has been addressed by researchers and adverse effects of chloroacetamide in vivo have been reported, such as cardiovascular toxicity in zebrafish larvae (16), evidence is still lacking on whether chloroacetamide affects embryo development and the potential underlying mechanism. Moreover, most research (13,16,17) on chloroacetamide has focused on AC, which is metabolized once absorbed in the body. AC and metolachlor, another chloroacetamide herbicide, are metabolized into 2-ethyl-6-methyl-2-chloroacetanilide (CMEPA) in the liver and then into 6-ethyl-o-toluidine (MEA; Fig. S1) (18). Both CMEPA and MEA are bioactivated by para-hydroxylation (18).
There is a lack of data on the toxicity of CMEPA and MEA. Therefore, in the present study, HepG2 cells (a liver cancer cell line) and zebrafish embryos were used to assess the potential toxicity effect of AC, one of most common chloroacetamides, and its metabolites (CMEPA and MEA) by investigating cell viability, oxidative stress, DNA damage and cell apoptosis.

Materials and methods

Cell culture. Human HepG2 cells (a liver cancer cell line, HB-8065; American Type Culture Collection, authenticated via STR profiling) were cultured in Eagle's Minimum Essential Medium (Sigma-Aldrich; Merck KGaA) supplemented with 10% FBS (HyClone; Cytiva), penicillin, and streptomycin (Beyotime Institute of Biotechnology) at 37˚C with 5% CO₂.

AC was purchased from Shanghai Aladdin Biochemical Technology Co., Ltd. CMEPA was purchased from Tokyo Chemical Industry Co., Ltd. and MEA was purchased from Dr. Ehrenstorfer GmbH (LGC Ltd.). Cells were treated with 5 mM N-acetylcyesteine (NAC; cat. no. ST1546; Beyotime Institute of Biotechnology) for 72 h at 37˚C or ERK1/2 inhibitor PD98059 (Selleck Chemicals) at 20 µM, for 72 h at 37˚C as previously described (19). All reagents were dissolved in 40% (v/v) methanol. Controls were treated with 40% (v/v) methanol for 72 h at 37˚C.

In order to generate the dose-effect curves of AC, CMEPA and MEA, HepG2 cells and zebrafish embryos were exposed to each drug at 1.95-500.00 µM (with 2-fold interval; Fig. S2). For HepG2 cells, treatment lasted for 72 h at 37˚C, and then cell viability was measured via MTT assay (Beyotime Institute of Biotechnology) in accordance with the manufacturer's instructions, to calculate IC₅₀ of AC, CMEPA and MEA. For zebrafish (20 embryos/group), treatment started after fertilization, and lasted for 48-120 h (h post-fertilization, hpf) at 28.5±0.5˚C, and then the mortality was evaluated.

Thereafter, cells or zebrafish embryos were exposed to AC, CMEPA and MEA at a concentration of 10-100 µM to investigate the potential toxic effect based on IC₅₀ for cells or LC₅₀ for zebrafish (Fig. S2).

Cell viability assay. The cells were seeded (5x10³ cells/well) in 96-well plates and exposed to AC, CMEPA, and MEA for 72 h at 37˚C. Thereafter, the cell viability was assessed via MTT assay (Beyotime Institute of Biotechnology) according to the manufacturer's protocol. DMSO was used to dissolve the formazan. The absorbance of each well was measured using a multiple plate reader, with excitation at 502 nm and emission at 525 nm.

Reactive oxygen species (ROS) assay. An ROS Assay kit (Elabscience, Inc.) was applied to investigate ROS generation in cells. In brief, cells were collected and suspended in 2′,7′-dichlorofluorescein diacetate (DCFH-DA; 10 mM) at 5x10⁵ cells/ml at 37˚C for 30 min. Then, the fluorescent activity was measured by a multiple plate reader with excitation at 502 nm and emission at 525 nm.

Assay of cellular superoxidase dismutase (SOD) and glutathione (GSH) levels and lactate dehydrogenase (LDH) leakage. In order to evaluate the enzyme activity of cells or embryos exposed to AC, CMEPA and MEA, assay kits for LDH, SOD and GSH were used according to the manufacturer's instructions (all Beyotime Institute of Biotechnology).

Briefly, the cells were seeded in 96-well plates at 5,000 cells/well and exposed to AC, CMEPA and MEA for 72 h at 37˚C. In order to evaluate LDH leakage from cells, culture medium was collected. Samples were prepared using the kit and measured by a multiple plate reader at 490 nm. In order to determine the activity of SOD and GSH, the cells were collected after culture medium was removed. Then, the samples were prepared and measured by a multiple plate reader at 450 nm (for SOD) and emission at 340 nm (for GSH).

In order to measure the activity of SOD, GSH and LDH in embryos, tissue homogenates were diluted in the buffer provided by each kit at a concentration of 10 mg tissue/ml. The samples were prepared according to the manufacturer's instructions and measured using a multiple plate reader, as aforementioned.

Comet assay. The cells were exposed to AC, CMEPA or MEA at 100 µM for 72 h at 37˚C before they were collected. The cells were suspended in PBS buffer at 1x10⁵/ml. The heated agarose (0.75%) was pre-coated on the slides at 37˚C, and immediately placed at 4˚C for 1 h. The cell suspension was mixed with 0.5% agarose at 37˚C and 30 µl was immediately added onto slides. Subsequently, the slides were placed at 4˚C in the dark for 10 min and then immersed in a 4˚C lysis solution in the dark overnight. Then, the slides were incubated with alkaline electrophoresis solution for 1 h at 4˚C in a dark environment, followed by electrophoresis for 30 min (25 V; 1 V/cm). The slides were neutralized in 0.4 M Tris-HCl for 5 min at room temperature and then washed with deionized water. The slides were fixed with 70% ethanol for 5 min at room temperature and then left to dry at 37˚C for 10 min in the dark. They were stained with propidium iodide (10 µl; Beyotime Institute of Biotechnology) at room temperature for 10 min. Finally, slides were observed under a fluorescence microscope (Nikon Corporation) under 400x magnification. In each sample, 50 cells were scored based on a five-grade scale (0-4). The entire DNA is in the ‘head’ at grade 0 and in the ‘tail’ at grade 4. Grade 1, 2 and 3 were defined as ~25, 50 and 75% DNA in the tail, respectively, by visual scoring (20,21).

Cell apoptosis assay. The cells were collected following exposure to AC, CMEPA or MEA and resuspended in 500 µl binding buffer at 1x10⁶ cells/ml. Then 5 µl Annexin V-fluorescein isothiocyanate (Beyotime Institute of Biotechnology) was added into 100 µl cell suspension, which then was incubated for 10 min at room temperature in the dark. Next, the cell suspension was mixed with 2 ml binding buffer and centrifuged at 500 x g for 5 min at room temperature. Thereafter, the cell pellets were collected and resuspended in 500 µl binding buffer. A total of 10 µl propidium iodide was added into the cell suspension, which then was incubated for 15 min at room temperature in the dark. Following incubation, a FACScan flow cytometer (BD Biosciences) was used to evaluate cell apoptosis. The results analyzed by FlowJo software (V10.6; BD Biosciences) to determine the apoptosis rate.

Zebrafish experiments. AB-type zebrafish (Danio rerio) were obtained from the Animal Experimental Center of...
Hebei Medical University and the procedures and experiments were approved by the Committee on Ethics of Animal Experiments of Hebei Medical University. Zebrafish were maintained at 28.5±0.5˚C, SaO₂>80%, pH, 7.5±0.5, with a 14-h light and 10-h dark cycle. Normally, a zebrafish embryo has a hatching period of 48‑72 hpf (22). At 2 hpf, the embryos were examined under a stereomicroscope to ascertain whether normal embryos had grown. Then, the normal embryos were randomly distributed (30 embryos per group) and exposed to AC, CMEPA or MEA at 10‑100 µM for 120 hpf as follows: i) Treatment with AC at 10, 50, 100 or 100 µM AC + 5 mM NAC; ii) treatment with CMEPA at 10, 50, 100 or 100 µM CMEPA +5 mM NAC; iii) treatment with MEA at 10, 50, 100 or 100 µM MEA +5 mM NAC. Controls for each group were treated were methanol. The hatching status was determined by calculating the hatching and survival rates.

In order to analyze the apoptotic status of embryos following exposure to AC, CMEPA or MEA at 100 µM for 120 hpf as follows: i) Treatment with AC at 10, 50, 100 or 100 µM AC + 5 mM NAC; ii) treatment with CMEPA at 10, 50, 100 or 100 µM CMEPA +5 mM NAC; iii) treatment with MEA at 10, 50, 100 or 100 µM MEA +5 mM NAC. Controls for each group were treated were methanol. The hatching status was determined by calculating the hatching and survival rates.

Western blotting. In order to collect protein from cells, cold RIPA Buffer (Sigma-Aldrich; Merck KGaA) containing protease inhibitors was added to the cells for 5 min on ice, then the lysate was collected using a cell scraper. Samples were centrifuged at 14,000 x g for 15 min at 4°C to collect protein in the supernatant. The protein levels were determined by BCA method. Equal amounts of protein (20 µg/lane) were separated on 8-10% SDS-polyacrylamide gels for electrophoresis

Caspase-3/8 activity. The caspase-3/8 activity was assayed using Caspase-3 and Caspase-8 Assay kits (both Beyotime Institute of Biotechnology), according to the manufacturer's instruction. The fresh protein lysates from cells were prepared using cell lysis buffer (Beyotime Institute of Biotechnology). Then, 85 µl reaction buffer and 5 µl Leu-Glu-His-Asp-p-nitroanilide were added to each sample and incubated at 37°C for 2 h. The absorbance was measured using a multplate reader at 450 nm.
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and then transferred to nitrocellulose membranes. Next, the membranes were blocked with 5% non-fat milk in Tris-Cl-buffered saline (TBS-T, 0.1% Tween-20) at room temperature for 2 h. The membranes were incubated with primary antibodies at 4˚C overnight. Thereafter, the membranes were washed with TBS-T and incubated with horseradish peroxidase-conjugated anti-mouse or anti-rabbit (both 1:5,000; cat. nos. sc-516102 and sc-2357, respectively; both Santa cruz Biotechnology, Inc.) secondary antibody at room temperature for 1 h and subsequently processed for enhanced chemiluminescence detection (Pierce; Thermo Fisher Scientific, Inc.). The signals were detected via a chemiluminescence detection system (Bio‑Rad Laboratories, Inc.). The protein expression levels were quantified with ImageJ software (version 1.3.2; National Institutes of Health). The following monoclonal primary antibodies were used: Mouse P38 (1:3,000; cat. no. sc-7972; Santa Cruz Biotechnology, Inc.), phosphorylated (Pho)-P38 (1:3,000; cat. no. sc-166182; Santa Cruz Biotechnology, Inc.), JNK (1:3,000; cat. no. sc-7345; Santa cruz Biotechnology, Inc.), rabbit Pho-JNK (1:3,000; Thr183/Tyr185; 1:3,000; cat. no. 4668; Cell Signaling Technology, Inc.), caspase-3 (1:3,000; cat. no. 9664S; Cell Signaling Technology, Inc.) and mouse GAPdH (1:3,000; cat. no. sc-32233; Santa Cruz Biotechnology, Inc.).

**Results**

ROS generation and SOD and GSH activity in cells and zebrafish embryos exposed to AC, CMEPA and MEA. Intracellular ROS was measured by DCF fluorescence intensity. Treatment with AC, CMEPA and MEA increased DCFH-dA intensity, indicating elevated ROS production after cells were exposed to AC, CMEPA and MEA (Fig. 1). In order to analyze the oxidative stress response, SOD activity and GSH content of cells were determined following exposure to various concentrations of AC, CMEPA, and MEA. Following 72 h exposure at 10-100 µM, all treated groups (AC, CMEPA and MEA) showed a decrease in SOD activity and GSH levels (Fig. 1). These data showed that exposure to AC, CMEPA and MEA decreased GSH and SOD levels in cells (Fig. 1).

In order to evaluate the toxicity of AC, CMEPA and MEA in vivo, the effects of AC, CMEPA and MEA on ROS generation were investigated using zebrafish embryos. After zebrafish embryos were exposed to AC, CMEPA and MEA for 120 hpf at 10-100 µM, SOD and GSH levels decreased (Fig. 2).

**Cell injury assay by chloroacetamide in vitro.** In order to address whether AC, MEPA, and MEA induce cell injury, leakage of cell injury biomarker LDH from cells exposed to...
The data showed that exposure to AC, CMEPA, and MEA increased LDH leakage from cells in a dose-dependent manner (Fig. 3). These results suggested that these chemicals increased the permeability of the cell membrane. Moreover, the LDH levels in the embryos increased following exposure to AC, CMEPA, or MEA (Fig. 2). The comet assay was used to measure the DNA strand breaks in HepG2 cells following exposure to AC, CMEPA, and MEA. As observed under a fluorescence microscope, DNA strand breaks were identified in cells following exposure to Ac, cMEPA, and MEA for 72 h. The olive tail moments (OTMs) significantly increased following cell exposure to AC, CMEPA, and MEA by 4-8-fold compared with the control (Fig. 4).

Figure 3. LDH leakage from HepG2 cells exposed to AC, CMEPA and MEA. HepG2 cells were exposed to AC, CMEPA and MEA at 10-100 µM for 72 h; control cells were treated with methanol. The cell culture medium was collected and LDH leakage from the cells was measured. Exposure to AC, CMEPA and MEA caused LDH leakage from cells in a dose-dependent manner. The data are presented as the mean ± SEM (n=3). *P<0.05, **P<0.01. LDH, lactate dehydrogenase; AC, acetochlor; CMEPA, 2-ethyl-6-methyl-2-chloroacetanilide; MEA, 6-ethyl-o-toluidine.

Figure 4. Comet assay of DNA damage in HepG2 cells exposed to AC, CMEPA and MEA. HepG2 cells were exposed to AC, CMEPA and MEA at 100 µM for 72 h; control cells were treated with methanol. Exposure to AC, CMEPA and MEA induced DNA breakage in HepG2 cells and a significant increase in OTM was observed compared with the control. Representative images of tailing by damaged DNA in HepG2 cells are shown. Magnification, x200. The data are presented as the mean ± SEM (n=3). *P<0.05, **P<0.01. AC, acetochlor; CMEPA, 2-ethyl-6-methyl-2-chloroacetanilide; MEA, 6-ethyl-o-toluidine; OTM, olive tail moment.

Cytotoxic effects of chloroacetamide in vitro. Cell viability assay was used to determine the viability of HepG2 cells exposed to AC, CMEPA or MEA. Following 72 h incubation with AC at 10-100 µM, cell viability decreased significantly compared with the controls. The inhibitory effect of CMEPA and MEA on cell viability was also observed in a dose-dependent manner (Fig. 5). However, co-treatment with anti-ROS reagent NAC alleviated the inhibitory effect of AC and its metabolites.

AC, CMEPA and MEA induce cell apoptosis both in vitro and in vivo. HepG2 cells were exposed to AC, CMEPA and MEA for 72 h at 10-100 µM before Annexin V and PI staining and flow cytometry analysis. AC and its metabolites (CMEPA and MEA) significantly increased the percentage of apoptotic cells. The pro-apoptosis effects of AC, CMEPA, and MEA were dose-dependent (Fig. 6). Moreover, treatment with AC, CMEPA and MEA increased the apoptotic protein activity (Caspase3 and Caspase8), which was consistent with the results of flow cytometry (Fig. S3). However, the co-treatment with anti-ROS reagent NAC alleviated the pro-apoptosis effect of AC and its metabolites.
Figure 6. Exposure to AC, CMEPA and MEA promotes apoptosis of HepG2 cells in vitro. HepG2 cells were exposed to AC, CMEPA and MEA at 10-100 µM for 72 h; control cells were treated with methanol. The cells were collected, stained with Annexin V and PI and analyzed by FACScan. The number of early (Annexin V-positive) and late apoptotic (Annexin V- and PI-positive) cells indicates the total percentage of gated cells. Representative images are shown. Exposure to AC, CMEPA and MEA promoted cell apoptosis in a dose-dependent manner. However, the pro-apoptosis effects of AC, CMEPA and MEA were alleviated by NAC (5 mM). The co-treatment with NAC and AC, CMEPA and MEA resulted in a lower apoptosis rate than treatment with AC-, CMEPA- and MEA-alone. The data are presented as the mean ± SEM (n=3). *P<0.05, **P<0.01. Ac, acetochlor; CMEPA, 2-ethyl-6-methyl-2-chloroacetanilide; MEA, 6-ethyl-o-toluidine; NAC, N-acetylcysteine.

Figure 7. Cell apoptosis in zebrafish increases following exposure to AC, CMEPA and MEA. Embryos were treated with AC, CMEPA and MEA at 100 µM for 120 hpf (30/group) and then stained with acridine orange; control embryos were treated with methanol. The zebrafish were observed under a fluorescence microscope. A significant increase in the number of apoptotic cells was observed compared with the control. Representative images are shown. Scale bar, 200 µm. *P<0.01. AC, acetochlor; CMEPA, 2-ethyl-6-methyl-2-chloroacetanilide; MEA, 6-ethyl-o-toluidine.
Furthermore, to investigate the apoptosis status in zebrafish embryos, the zebrafish were exposed to AC, CMEPA and MEA and then stained with AO. The formation of apoptotic bodies was clearly observed in treated larvae compared with controls (Fig. 7).

Activation of the MAPK/ERK pathway participates in cell apoptosis (26). Thus, it was investigated whether AC, CMEPA and MEA activate the MAPK/ERK pathway in cells. HepG2 were cultured in vitro and treated with AC, CMEPA and MEA, in the presence or absence of ERK1/2 inhibitor PD98059 (20 µM; 2 h); control cells were treated with methanol. Cells were collected and total proteins were extracted from cell pellets. The protein expression was examined via western blot analysis. The representative images are shown. AC, CMEPA, MEA significantly increased levels of ERK1/2 pathway signaling molecules Pho-P38 and Pho-JNK and expression of Caspase3. ERK1/2 inhibitor decreased expression of Caspase3. AC, acetochlor; CMEPA, 2-ethyl-6-methyl-2-chloroacetanilide; MEA, 6-ethyl-o-toluidine; Pho, phosphorylated.

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Activation of the MAPK/ERK pathway participates in cell apoptosis (26). Thus, it was investigated whether AC, CMEPA and MEA activate the MAPK/ERK pathway in cells. HepG2 were cultured in vitro and treated with AC, CMEPA and MEA, in the presence or absence of ERK1/2 inhibitor; AC, CMEPA and MEA promoted phosphorylated levels of both P38 and JNK, and induced apoptotic protein Caspase3 expression, whereas ERK1/2 inhibitor inhibited expression of Caspase3 (Fig. 8).

Survival and hatching of zebrafish embryos exposed to AC, CMEPA and MEA. The zebrafish embryos were exposed to AC, CMEPA and MEA at 10-100 µM for 120 hpf; control embryos were treated with methanol. Exposure to AC, CMEPA and MEA decreased the survival and hatching rate of zebrafish embryos in a dose-dependent manner. However, co-treatment with NAC (5 mM) increased both the hatching and survival rates of zebrafish exposed to AC, CMEPA and MEA. The data are presented as the mean ± SEM. *P<0.05, **P<0.01. AC, acetochlor; CMEPA, 2-ethyl-6-methyl-2-chloroacetanilide; MEA, 6-ethyl-o-toluidine; NAc, N-acetylcysteine.
Discussion

The present study showed the toxicity of chloroacetamide herbicides, including AC, CMEPA and MEA. The toxicity may be caused by excess generation of ROS, which induces cell apoptosis.

To date, it has been difficult to determine the threshold concentration of toxic effects of chloroacetamide herbicides in nature because the sensitivity of different species to chloroacetamide herbicides may be different. For example, Yin et al. (27) suggested the LC50 of AC for Chinese toad is 0.76 mg/l, while Daam et al. (28) suggested that LC50 of AC for Physalaemus cuvieri and Hyperolius pardalis are 4.4 and 7.8 mg/l, respectively. Even in the same species, such as zebrafish, the concentration of AC to induce cytotoxicity is controversial. In zebrafish, studies have reported that AC induces toxicity at 37-74 µM (16,29). Moreover, many studies (19,30,31) have tried to determine the threshold concentration of toxicity of AC in cell lines but the data are inconsistent. Studies have reported that AC inhibits viability of different types of cells in a time- and concentration-dependent manner, including HepG2 cells at 50-130 µM (30,31), A549 cells at 25-200 µM (19) and Veto cells at 130 µM (31). This may be due to varying sensitivity of different cell lines or species, or inconsistent experimental conditions. To the best of our knowledge, previous studies have not investigated the toxicity of CMEPA or MEA in vitro or in vivo. Therefore, the present study determined the IC50 of AC, CMEPA and MEA in HepG2 cells, as well as the LC50 of AC, CMEPA and MEA in zebrafish embryos, which were used to treat cells and zebrafish to investigate the effect of AC, CMEPA and MEA.

Chloroacetamide is absorbed by living creatures, increasing the potential risk of disease, such as Parkinson’s Disease in humans (15). To the best of our knowledge, however, few studies have focused on the mechanism underlying toxicity of chloroacetamide in embryos. In the present study, ROS production in cells and embryos was examined and ROS induction by chloroacetamide and its metabolites was observed. Intracellular ROS are effectively eliminated by the combined action of SOD, GSH and other endogenous antioxidants, providing a repair mechanism for oxidized membrane components (32). SOD serves a critical role in cell defense against the toxic effects of oxygen radicals, which reduce superoxide anions to hydrogen peroxide (33). In the present study, chloroacetamide and its metabolites were found to decrease the SOD level at 72 h post-exposure. A short-term increase in ROS increases SOD levels as an adaptation and defense response against ROS production (34), whereas long-term exposure to toxicity, such as that induced by chloroacetamide, exhausts mobilized SOD, decreasing SOD levels. Similar results were also observed for GSH levels following exposure to chloroacetamide in the present study. GSH is a ubiquitous molecule in the process of antioxidation of ROS and free radicals by scavenging free radicals and other ROS, removing hydrogen and lipid peroxides and preventing the oxidation of biomolecules (35). The present results showed that chloroacetamide and its metabolites decreased GSH levels in a concentration-dependent manner. This may be due to lipid peroxidation caused by a high concentration of chloroacetamide and its metabolites, thereby inducing oxidative stress and exhausting GSH in cells. Due to the decrease in antioxidant enzyme activity, insufficient antioxidant defense leads to accumulation of ROS, which can lead to cell injury, including cell membrane and DNA damage (36).

Increased LDH leakage from the cells and embryos was observed following exposure to chloroacetamide. LDH is a soluble cytoplasmic enzyme present in almost all cells and is released into extracellular space when the plasma membrane is damaged (37). Therefore, LDH leakage from cells is associated with cell necrosis, which can be considered a biomarker of cell injury. As shown in the present study, chloroacetamide induced excessive production of ROS in cells, which damaged the cell membranes and caused LDH to leak from cells. DNA damage due to exposure to chloroacetamide is caused by excess generation of ROS in the embryo. ROS are continuously generated in cells under normal physiological conditions. These consist of stable oxidants (such as H2O2) and unstable free radicals (such as superoxide anion, nitric oxide, hydroxyl moiety and hypochlorite), which participate in cellular events and regulate cell behavior (38). For example, in rat fibroblasts, hypoxia induces production of nitric oxide, which triggers apoptosis of cells (39). Additionally, superoxide anions induce the migration and recruitment of leukocytes (40). However, excess ROS generation can overwhelm the antioxidant system, thus inducing DNA, lipid and peptide oxidation (19). ROS, such as the highly reactive hydroxyl radical (•OH), interact with the double bonds of the DNA base, which adds to the C4, C5 and C8 positions of purines generating OH adduct radicals, or abstract the H atom from the methyl group of thymine and each of the C-H bonds of 2'-deoxyribose (41). Moreover, the guanine radical cation (guanine•+) is formed by eliminating OH from the C4-OH adduct radical of guanine (42). Reactions of •OH with the sugar moiety of DNA by H abstraction cause sugar modifications and strand breaks (41,43). Thus, modifications of DNA by ROS production induce DNA breakage; DNA damage can trigger apoptosis of cells. ROS can also induce apoptosis through other pathways (44). ROS induce DNA-protein cross-linking and lead to DNA damage by combining with an amino acid radical or adding an aromatic amino acid of proteins (45). ROS can induce expression of apoptosis-associated proteins, such as Fas, caspase3, caspase7 and caspase8 and stimulate cell apoptosis (46,47). Activation of the MAPK/ERK pathway can trigger cell apoptosis (26). In the present study, AC and its metabolites induced cell apoptosis via activation of the MAPK/ERK pathway. Treatment with AC and its metabolites induced phosphorylation of P38 and JNK, which then increased expression levels of apoptotic proteins, such as Caspase3. Consistent with the present findings, Zerin et al. (19) suggested that AC induces lung cancer A549 cell apoptosis via the MAPK/ERK pathway.

Apoptosis serves a key role in the development of embryos. Studies (48-50) using electron microscopy have demonstrated apoptotic changes in the amniotic epithelium and chorionic trophoblast cells, including the condensation of chromatins and nuclear shrinkage. Under normal physiological conditions, cell apoptosis is involved in the formation of vesicles and tubes (51). However, abnormal apoptosis can lead to abnormal development of embryos (52). Ding et al. (53) reported that
Oxidative stress can induce cytotoxicity in vitro [43]. Amide can cause oxidative stress in cells via ROS generation. In vivo [43], promoted generation of ROS and increased cell apoptosis. In the present study, exposure to chloroacetamide and its metabolites promoted generation of ROS and increased cell apoptosis in vitro and in vivo. Exposure to chloroacetamide decreased hatchability and survival rate of embryos, with evidence of excessive apoptosis, indicating toxicity to embryos. Thus, chloroacetamide may be an environmental risk factor during embryo development.

In conclusion, the present study showed that chloroacetamide can cause oxidative stress in cells via ROS generation. Oxidative stress can induce cytotoxicity in vitro by triggering cell apoptosis. The present data may provide insights for better understanding of chloroacetamide toxicity. It is necessary to investigate the potential embryo toxicity of environmental chloroacetamide.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

XM and YZ designed and performed experiments, analyzed the data and wrote the manuscript. MG, WZ, HT, CJ and XT performed the experiments. XM and WK contributed to study design and wrote the manuscript. WK funded the study. XM and WK confirm the authenticity of all the raw data. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Experiments were approved by the Committee on Ethics of Animal Experiments of Hebei Medical University (approval no. SIZCDC2016002).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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