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Myeloperoxidase deficiency attenuates nitrogen mustardinduced skin injuries

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Abstract

The pathologic mechanisms of skin injuries, following the acute inflammatory response induced by vesicating agents sulfur mustard (SM) and nitrogen mustard (NM) exposure, are poorly understood. Neutrophils which accumulate at the site of injury, abundantly express myeloperoxidase (MPO), a heme protein that is implicated in oxidant-related antimicrobial and cytotoxic responses. Our previous studies have shown that exposure to SM analog 2-chloroethyl ethyl sulfide (CEES) or NM results in an inflammatory response including increased neutrophilic infiltration and MPO activity. To further define the role of neutrophil-derived MPO in NMinduced skin injury, here we used a genetic approach and examined the effect of NM exposure (12 h and 24 h) on previously established injury endpoints in C57BL/6J wild type (WT) and B6.129X1-MPOtm1Lus/J mice (MPO KO), homozygous null for MPO gene. NM exposure caused a significant increase in skin bi-fold thickness, epidermal thickness, microvesication, DNA damage and apoptosis in WT mice compared to MPO KO mice. MPO KO mice showed relatively insignificant effect. Similarly, NM-induced increases in the expression of inflammatory and proteolytic mediators, including COX-2, iNOS and MMP-9 in WT mice, while having a significantly lower effect in MPO KO mice. Collectively, these results show that MPO, which generates microbicidal oxidants, plays an important role in NM-induced skin injuries. This suggests the development of mechanism-based treatments against NM- and SM-induced skin injuries that inhibit MPO activity and attenuate MPO-derived oxidants.

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Keywords

Sulfur mustard; nitrogen mustard; myeloperoxidase; inflammation; apoptosis; DNA damage

1. Introduction

Sulfur mustard [SM; bis (2-chloroethyl) sulfide] and nitrogen mustard (NM; mechlorethamine) are primary vesicating agents, which pose potential threat for use in chemical warfare and terrorist attack (Saladi et al., 2006; Sharma et al., 2010; Smith et al., 1995). Both of these mustard agents are highly toxic bifunctional alkylating agents, which upon exposure cause severe skin injury including inflammation and disruption of epidermaldermal layers leading to vesication (Graham et al., 2005; Hayden et al., 2009; Jain et al., 2011b; Joseph et al., 2011; Laskin et al., 2010; McManus and Huebner, 2005). In the past few years, numerous studies have been carried out to explore the mechanisms of cell death, inflammation and vesication caused by warfare and alkylating agent SM exposure (Casillas et al., 2000; Inturi et al., 2011; Jain et al., 2011b; Kehe et al., 2009; Kehe and Szinicz, 2005; Pal et al., 2009; Rice, 2003; Shakarjian et al., 2010; Tewari-Singh et al., 2010). However, further investigations are needed to define the mechanism of skin injury and healing processes after vesicant exposure, especially pathogenesis involving the inflammatory response where infiltration of polymorphonuclear lymphocytes (PMNs) is reported to play an important role (Dacre and Goldman, 1996; Graham et al., 2005; Millard et al., 1997; Wormser, 1991).

SM-induced PMN infiltration, mainly neutrophils, has been shown as an early event occurring within 30 min and peaking at 6-12 h post-exposure in human skin (Lindsay and Rice, 1996; Millard et al., 1997; Wormser et al., 2005). Neutrophil infiltration plays a key role in host defense. However, its excessive stimulation induces the release of cytokines and growth factors which contribute to the healing process (Segal, 2005; Weiss, 1989). Conversely, neutrophils and to some extent monocytes and macrophages, secrete myeloperoxidase (MPO). MPO is a member of heme peroxidase-cyclooxygenase family, at the site of tissue injury generating powerful oxidants, free radicals and hypochlorous acid, which can cause tissue injury (Brennan et al., 2001; Klebanoff, 2005; Shiba et al., 2008). Infiltration of neutrophils also plays an important role in blister formation by secreting a variety of proteolytic enzymes including collagenase, gelatinase and elastase. These enzymesare able to degrade specific elements in the extracellular matrix leading to epidermal-dermal separation as observed in skin diseases like psoriasis (Glinski et al., 1990) and bullous pemphigoid (Liu et al., 1997; Liu et al., 2000). SM-induced biochemical changes include high MPO activity, free radical generation and cytokine production (Ham et al., 2012). Our previous studies with SM analog 2, chloroethyl ethyl sulfide (CEES) and NM have shown an increase in neutrophil infiltration and MPO (a marker indicating neutrophil infiltration) activity in mouse skin (Jain et al., 2011b; Tewari-Singh et al., 2009). We have also shown that both CEES and NM exposure induces DNA damage, cell death, oxidative stress, microvesication, activation of proteolytic and inflammatory mediators as well as related signaling pathways. These previous studies indicated that neutrophil-derived MPO could play a vital role in vesicant-induced skin lesions (Jain et al., 2011a; Pal et al., 2009;

Tewari-Singh et al., 2010). Though some studies have examined the role of neutrophils in injury with SM (Ham et al., 2012; Levitt et al., 2003; Shakarjian et al., 2010; Vavra et al., 2004), little is known about the mechanism and role of neutrophil-derived MPO-mediated inflammation and injury in vesicant-exposed skin tissue. Accordingly, the current study was designed to evaluate the role of neutrophil derived MPO in NM-induced skin injury by utilizing C57BL/6J wild type and MPO knockout (B6.129X1-Mpotm1Lus/J) mice. Our results showed an attenuation of NM-induced skin injury biomarkers in MPO KO mice, indicating an important role of neutrophil-derived MPO in vesicant-induced skin injury and inflammation.

2. Materials and methods

2.1 Materials

Hematoxylin and eosin stains and NM (mechlorethamine hydrochloride; purity, 98%) were obtained from Sigma-Aldrich Chemicals Co. (St. Louis, MO). Paraffin wax for tissue block preparation was obtained from Triangle Biomedical Sciences (Durham, NC). DeadEndTM Colorimetric TUNEL (TdT-mediated dUTP Nick-End Labeling) system was purchased from Promega (Madison, WI) and Fluoro MPOTM Fluorescent MPO Detection Kit was obtained from Cell Technology (Mountain View, CA). Primary antibodies for H2A.X and MMP-9, COX-2, and iNOS were obtained from Cell Signaling (Beverly, MA, USA), Cayman Chemicals (Ann Arbor, MI, USA), and Abcam (Cambridge, MA), respectively. We used β actin antibody from Sigma-Aldrich (St. Louis, MO).

2.2 Animal exposure with NM

Male C57BL/6J wild type (hereafter refereed as WT mice) and B6.129X1-MpotmlLus/J mice, homozygous null for MPO gene where no MPO gene product is detected (hereafter refereed as MPO KO mice), were purchased from Jackson Laboratory (Bar Harbor, Maine). Before starting the experiments, mice were acclimatized for one week under standard housing and feeding conditions and all experiments were performed according to approved IACUC protocols of the University of Colorado Denver, CO (Protocol No – 57912(08)1E). Prior to NM exposure, mice were shaved using clippers to remove hair from their dorsal skin. Both wild type and MPO KO mice (n=3) were exposed with 6 mg NM in 200 µL of acetone/mouse applied topically to the dorsal shaved skin. Mice were 5-6 weeks of age with mean weight of 25g at the start of the experiment. Acetone (200 µL) alone was applied as vehicle control for NM and another group of mice were kept without any exposure, and used as untreated control. All the results following NM exposure were compared with vehicle control group and each study group consisted of three mice. Since MPO plays an important role in SM-induced skin injury and inflammation, we employed a high dose of NM to assess the effect of MPO deletion on skin injuries in both MPO KO and WT mice. Skin bi-fold thickness was measured at 12 h and 24 h post-exposure using digital calipers as reported earlier (Tewari-Singh et al. 2009). Immediately following measurement, at 12 h and 24 h post-exposure, mice were euthanized, dorsal skin collected and skin punches fixed for histology and immunohistochemical (IHC) analyses. The rest of the skin tissue was snap frozen in liquid nitrogen for molecular studies.

2.3 Histopathological analysis

Skin tissues were fixed O/N in 10% (v/v) phosphate buffered formalin and dehydrated in ascending concentrations of ethanol, cleared with xylene and embedded in paraffin. 5μ m thick skin sections were processed for H&E staining as reported earlier (Jain et al., 2011b; Tewari-Singh et al., 2009). H&E slides were microscopically evaluated for measurement of epidermal thickness, neutrophil infiltration, dermal injury and the incidences of microvesication as reported earlier (Jain et al., 2011b).

2.4 TUNEL assay

Apoptotic cell death was detected using DeadEndTM Colorimetric TUNEL assay according to the vendor's protocol as detailed earlier (Jain et al., 2011b; Tewari-Singh et al., 2010). At the end of the assay, TUNEL positive cells (brown colored) were quantified in 15 randomly selected fields using 400 magnification. Apoptotic cell index (% apoptotic cell death) was calculated as the number of apoptotic cells \times 100 divided by total number of cells.

2.5 Immunohistochemical staining

Phospho-H2A.X ser139 (a marker of DNA damage) was analyzed by IHC as described previously (Jain et al., 2011a). Briefly, paraffin embedded skin sections were deparaffinized, rehydrated and treated for antigen retrieval by heat treatment in sodium citrate buffer (pH-6). The endogenous peroxidase activity was blocked and sections were incubated with rabbit polyclonal phospho-H2A.X ser139 antibody (1:100) O/N in humidified chamber. After washing with PBS, sections were incubated with appropriate biotin conjugated secondary antibody and then with streptavidin-HRP conjugates. Following another wash step, sections were incubated with DAB (3,3-diaminobenzidine) for immunopositivity reaction and color development. Thereafter, slides were mounted for microscopic observation. After IHC staining, cells in the skin epidermis were counted in 15 randomly selected fields using 400 magnification and the percent of phospho-H2A.X ser139 positive cells were quantified according to the method described above. All the H&E and IHC stained slides were observed using a Zeiss Axioscope 2 microscope (Carl Zeiss, Inc., Germany) and image analysis and representative pictures were taken by using Carl Zeiss Axiovision Rel 4.5 software.

2.6 MPO activity assay

MPO activity was measured in skin tissue samples through use of the Fluorescent Myeloperoxidase Detection Kit from Cell Technology as reported earlier (Tewari-Singh et al. 2009). In brief, approximately 100 mg tissue samples were homogenized in homogenization buffer (provided in kit) and isolated protein was measured by Lowry's method (Bio-Rad DC protein assay kit, Bio-Rad Laboratories, Hercules, CA). A 50 μg of protein/tissue sample was used to determine MPO activity from three separate animals as described previously (Jain et al., 2011b). MPO activity was expressed as mU/ μg of protein using MPO standard curve.

2.7 Western Blot analysis

COX-2, iNOS and MMP-9 were detected in frozen skin tissue samples by Western blotting. Approximately 100 mg of tissue was taken from the skin samples and homogenized in lysis buffer as published earlier (Jain et al., 2011a) and protein content was measured using Lowry's method (Lowry et al., 1951). For western blotting, 50 μ g of protein from three animals per group was subjected to SDS-PAGE and separated proteins were transferred onto nitrocellulose membrane by western blotting. After blocking, membranes were probed with primary antibodies against COX-2 (1:500), iNOS (1:500), and MMP-9 (1:500) for overnight at 4°C, followed by peroxidase-conjugated appropriate secondary antibodies for 1 h at room temperature. Protein bands were visualized by an enhanced chemiluminescence detection system (GE Healthcare Life Sciences, Pittsburgh, PA). In each case, protein loading was monitored by stripping and re-probing the blots with β -actin.

2.8 Statistical analysis

Statistically significant differences between groups was determined by one-way ANOVA using SigmaStat 3.5 software (Jandel scientific, San Rafael, CA) and the Tukey test for multiple comparisons, with p-value of < 0.05 considered as significant. Data is represented in \pm SEM from three replicate animals.

3. Results

3.1 MPO deletion in mice attenuated NM-induced tissue MPO activity

Our previous studies in mice have shown a profound increase in neutrophil infiltration and MPO activity in skin after dorsal exposure with CEES and NM at 12 h and 24 h time points (Inturi et al., 2013; Jain et al., 2011b; Tewari-Singh et al., 2009). To further investigate the role of the neutrophil-derived MPO enzyme in NM-induced skin injury, we employed a genetic approach using MPO KO mice where neutrophil infiltration and MPO activity were determined following NM exposure. H&E staining of the skin tissue sections from wild type and MPO KO mice 12 h and 24 h after NM exposure showed high neutrophil infiltration and dermal edema in NM-exposed wild type mice as compared with an untreated control (Fig. 1A). However, the NM-induced increase in neutrophil infiltration or edema was not evident in MPO KO mice (Fig. 1C). To confirm the absence of MPO activity and the markedly diminished neutrophil infiltration in MPO KO mice, MPO activity was measured in wild type and MPO KO mice (Fig. 1B and D). In comparison with control wild type mice where MPO activity was 2.4 mU/µg protein, high MPO activity was observed after 12 h (16.3 mU/µg protein) of NM exposure. This was maximal at 24 h (116.9 mU/µg protein) postexposure (Fig. 1B). Conversely, very low MPO activity was observed in MPO KO mice at 12 h (2.14 mU/µg protein) and 24 h (1.40 mU/µg protein) post-NM exposure, which was comparable to MPO activity in control mice (1.48 mU/µg protein; Fig. 1D).

3.2 MPO deletion in mice attenuated NM-induced skin bi-fold thickness and epidermal thickness increases

Our previous studies have shown that CEES and NM caused an increase in skin bi-fold thickness and epidermal thickness in mice (Jain et al., 2011b; Tewari-Singh et al., 2013; Tewari-Singh et al., 2009). To examine whether similar effects were present in NM-treated

MPO KO mice, we first measured the skin bi-fold thickness at 12 h and 24 h post-exposure. In WT mice, NM caused a significant (p<0.05) increase in skin bi-fold thickness after 24 h (1.46 fold) in comparison with the untreated control. On the other hand, a significant increase in bi-fold thickness was not observed in MPO KO mice at 12 h (1.3 fold) or 24 h (1.04 fold) post-NM exposure in comparison with control (Fig. 2A). H&E stained tissue sections from both types of mice were next analyzed for epidermal thickness at 12 h and 24 h after NM exposure. In wild type mice, NM caused a significant (P<0.05) 2.2-fold increase in epidermal thickness at 12 h following exposure (Fig. 2C). On the other hand, there was no significant effect of NM on epidermal thickness in MPO KO mice at 12 h or 24 h post-exposure (Fig. 2E).

3.3 MPO deletion in mice attenuated NM-induced microvesication

Our previous studies have reported that CEES and NM cause microvesication in mice, and established this lesion as an important biomarker of vesicant skin injury (Jain et al., 2011b; Tewari-Singh et al., 2013; Tewari-Singh et al., 2009). Similar to our previous reports, microvesication (epidermal-dermal separation) was observed 12 h and 24 h post-NM exposure in WT mice. However, these events were reduced in MPO KO mice (Fig. 3A and C). The incidence of microvesication was quantified according to their size; small ($<100\mu m^2$), medium ($100-500~\mu m^2$) and large ($>500~\mu m^2$) per skin section of mice. In WT mice, small NM-induced lesions of microvesication were 5.0 ± 0.70 at 12 h and 4.05 ± 0.65 at 24 h post-exposure (Fig. 3B). NM-induced lesions of medium and large sized microvesicles gradually increased and were maximal at 24 h post-exposure (Fig. 3A and B) In MPO KO mice, lesser incidences of all sizes of microvesication lesions were observed in comparison with WT mice at both 12 h and 24 h post-NM exposure (Fig. 3 C and D). Maximal occurrence of small sized NM-induced lesions of microvesication in WT mice, were 2.5 ± 0.71 and 3.15 ± 0.70 at 12 h and 24 h post-exposure, respectively, in MPO KO mice (Fig. 3D).

3.4 MPO deletion in mice attenuated NM-induced DNA damage and apoptotic cell death

Phospho-H2A.X Ser139 is a marker of DNA damage (Rogakou et al., 1998), that is shown to increase following vesicant exposure indicating DNA damage (Inturi et al., 2011; Jain et al., 2011a; Joseph et al., 2011; Tewari-Singh et al., 2010). Previously, our studies have shown that CEES or NM exposure causes DNA damage leading to cell death of skin keratinocytes, and this could be a major contributor of skin injury (Inturi et al., 2013; Jain et al., 2011a; Pal et al., 2009; Tewari-Singh et al., 2010). Since MPO takes part in the production of free radicals which may cause DNA damage and apoptotic cell death, we examined the hypothesis that ablation of PMN and MPO activity would decrease the DNA damage and cell death following NM treatment. In WT mice, NM exposure resulted in a significant (p<0.05) increase in the number of phospho-H2A.X Ser139 positive cells (Fig. 4A). After 12 h and 24 h of NM exposure in WT mice, $48.2 \pm 3.5\%$ and $47.6 \pm 1.4\%$ H2A.X Ser139 positive cells, respectively, were observed in NM-exposed mice compared with only 20 ± 1.0% phospho-H2A.X Ser139 positive cells in control mice skin (Fig. 4B). In skin tissue from MPO KO mice, $33 \pm 4.8\%$ and $37 \pm 1.3\%$ phospho-H2A.X Ser139 positive cells were observed at 12 h and 24 h post-NM exposure compared with $18 \pm 8.2\%$ positive cells in control. However, this increase was not significant (Fig. 4C and D). Apoptotic cell death,

which is one of the early responses in NM-induced skin injury, was analyzed by TUNEL staining. Quantification of data from TUNEL-stained skin sections from WT mice showed a significantly (P<0.05) increased percent of apoptotic cell death at 12 h (49 \pm 2.5%) and 24 h (42 \pm 0.68%) after NM exposure in comparison with the untreated control (26.%; Fig. 4E and F). However, in skin tissues from MPO KO mice, no significant increase in apoptotic cell death was observed at 12 h (19 \pm 3.4%) and 24 h (25 \pm 2.2%) after NM exposure, again when compared with control (16 \pm 6.8%; Fig. 4G and H).

3.5 MPO deletion in mice attenuated NM-induced COX-2, iNOS and MMP-9 levels

It has been previously reported that COX-2 and iNOS play important roles in SM and SM-analog induced skin injury and inflammation (Jain et al., 2011a; Pal et al., 2009). Therefore, levels of COX-2 and iNOS were quantitated by western blotting following NM exposure. These blots clearly showed an NM-induced increase in expression of COX-2 at 12 h (18.3 fold) and 24 h (15.9 fold) post-exposure; this increase was completely attenuated in MPO KO mice (Fig. 5A). Similarly, NM exposure also caused an increase in iNOS levels at 12 h (5.3 fold) and 24 h (7.3 fold) post-NM exposure, when compared to control skin tissue in WT mice (Fig. 5 A). In MPO KO mice, NM-exposure failed to increase the quantity of iNOS at 12 h post-exposure. However, at 24 h post-exposure NM induced an increase in iNOS in the MPO KO mice albeit less than that observed in WT mice (Fig. 5A).

As described above, NM exposure caused an increased number of microvesication lesions in WT mice when compared to MPO KO mice. MMP-9 levels are reported to increase after vesicant exposure and this plays a role in microvesication because it can degrade the components of extracellular matrix and basement membrane allowing epidermal dermal separation (Jain et al., 2011a; Pal et al., 2009; Shakarjian et al., 2010). To elucidate the role of the MPO enzyme in microvesication, MMP-9 expression was measured at 12 h and 24 h following NM exposure in both WT and MPO KO mice (Fig. 5B). NM exposure caused an increase in MMP-9 levels at 12 h (26 fold) and 24 h (10.2 fold) post-exposure in WT mice. However, in MPO KO mice, NM exposure induced only a slight increase in MMP-9 levels at both 12 h (3.3 fold) and 24 h (4.4 fold) post-exposure (Fig. 5 B).

4. Discussion

Although substantial research efforts have focused on understanding the mechanisms of skin injuries from SM exposure, process causing cutaneous lesions via acute inflammatory response after SM exposure is poorly understood. Earlier studies have shown that PMNs, mainly neutrophils, accumulate at the site of injury after SM exposure (Joseph et al., 2011; Shakarjian et al., 2010; Vavra et al., 2004). Similarly, our studies following exposure to SM analogs CEES and NM have also shown infiltration of inflammatory cells including neutrophils with greatly elevated MPO activity in mouse skin (Inturi et al., 2011; Jain et al., 2011a; Jain et al., 2011b; Pal et al., 2009; Tewari-Singh et al., 2009). Vesicant-induced neutrophil infiltration elicits MPO enzyme secretion which could enhance oxidative stress and inflammation in skin, and this plays an important role in resulting skin injuries. Therefore, in the present study we investigated the role of neutrophil derived MPO enzyme in the inflammation-related mechanism of NM-induced skin injuries. Our results from this study using MPO KO mice demonstrate for the first time that MPO plays an important role

in NM-induced skin injury by attenuating DNA damage, cell death, vesication and inflammation.

Due to limited access to SM, we used NM in this study because both SM and NM are vesicating and powerful alkylating and vesicating agents that show toxic effects following cutaneous exposure (Smith et al., 1998; Tewari-Singh et al., 2013; Wormser et al., 2002). We employed established NM-induced endpoints from our previous studies (Inturi et al., 2013 and unpublished data) in this study in C57BL/6J mice, since MPO KO mice were produced on a C57BL/6 background.

Both SM- and NM-induced skin injuries occur mainly due to their alkylating properties and direct DNA damage (Inturi et al., 2011; Kehe and Szinicz, 2005; Walker, 1971). However, generation of reactive oxygen/nitrogen species (ROS or RNS) after SM and NM exposure also have been shown to cause DNA damage leading to injuries (Kehe et al., 2009; Laskin et al., 2010; Naghii, 2002; Shakarjian et al., 2010). These injuries occur largely due to the sensitivity of the basal epidermal keratinocytes to SM where DNA damage is a major element (Dacre and Goldman, 1996; Graham et al., 2005; Kehe and Szinicz, 2005; Papirmeister et al., 1985). The Inflammatory response is considered to play a vital role in SM-induced pathology; neutrophil infiltration is a major early event in the inflammatory response (Lindsay and Rice, 1996; Millard et al., 1997). The neutrophil infiltration could contribute to oxidative stress and DNA damage, and could cause secondary injuries and inflammation. This may be attributed to the secretion of MPO by azurophilic granules of activated neutrophils at the site of inflammation which oxidizes Cl⁻ to hypoclorus acid (HOCl), a reactive oxidant, and causes indirect DNA damage (El Kebir et al., 2008; Loria et al., 2008; Miyamoto et al., 2006; Mutze et al., 2003). In the present study therefore, the role of MPO in these important events leading to NM-induced skin pathology was analyzed using MPO KO mice. Similar to earlier reports with SM and to our previous studies with CEES and NM, NM exposure in WT mice caused a robust NM-induced increase in neutrophil infiltration and MPO activity (Blank et al., 2000; Inturi et al., 2013; Jain et al., 2011b; Millard et al., 1997; Tewari-Singh et al., 2009; Vavra et al., 2004). However, this NM-induced increase in MPO activity was not observed in MPO KO mice confirming these mice were suitable for this study. Skin edema and hyperproliferation are important events of vesicant exposure that are referred to as injury and healing markers, respectively, as reported in our earlier studies (Jain et al., 2011b; Tewari-Singh et al., 2009). NM caused an increase in skin bi-fold and epidermal thickness of WT mice. However, no significant NM-induced increment in either of these markers was observed in MPO KO mice, indicating less injury and proliferation of cells to compensate for the cell death of effected epidermal keratinocytes. Our data also showed much lower NM-induced DNA damage (Phospho-H2A.X Ser139 IHC staining) and apoptotic cell death (TUNEL staining) in epidermal keratinocytes of MPO KO mice compared to WT mice. Since, DNA damage and apoptotic cell death are main consequences of SM exposure leading to skin injuries (Kehe and Szinicz, 2005; Mellor et al., 1991; Papirmeister et al., 1985; Paromov et al., 2007), a decrease in these effects evidenced in MPO KO mice suggests a vital role of MPO in NM-induced DNA damage and cell death. These are probably due to MPO's ability to induce oxidative stress.

SM exposure causes blister formation (vesication) in human skin and microvesication in animals' skin (Cowan et al., 2003; Ghabili et al., 2010; Shakarjian et al., 2010; Smith et al., 1995). Vesication occurs due to apoptosis in basal keratinocytes and degradation of the anchoring filaments. laminin-5, collagen and other extracellular matrix proteins (ECM) after SM exposure (Hess and FitzGerald, 2007). These proteases are either secreted by neutrophils or are directly activated by SM at the time of injury, leading to blister formation (Chang et al., 2009; Mol et al., 2009; Ray et al., 2002; Sabourin et al., 2002). Similar to the microvesication observed following SM exposure as well as our previous studies with CEES and NM, in this study NM induced microvesication in WT mice in this study. However this lesion was attenuated in the NM treated MPO KO mice. Since, vesication is associated with apoptotic cell death, decreased NM-induced apoptotic cell death in keratinocytes of MPO KO mice in comparison with WT mice observed here could contribute to decreased microvesication in MPO KO mice. It has been reported that proteases such as MMP-9, which degrade collagen and other ECM proteins are important mediators in vesicantinduced blister formation (Jain et al., 2011a; Pal et al., 2009; Ries et al., 2009; Shakarjian et al., 2006). Attenuation of the NM-induced MMP-9 expression in MPO KO mice here further suggests a role for MPO in vesicating effects of NM. Our findings support previous studies which show that MPO-derived HOCl modulated the activity of MMPs (Lau et al., 2005).

Many studies have reported an increased expression and importance of inflammatory mediators like COX-2 (the rate limiting enzyme in prostaglandin biosynthesis) in the skin after exposure to SM or its analog CEES (Black et al., 2010; Jain et al., 2011a; Joseph et al., 2011; Pal et al., 2009; Shakarjian et al., 2010). This increase in COX-2 expression could enhance vascular permeability and the further influx of inflammatory cells (Jain et al., 2011a). Our earlier studies also documented that CEES induced an increase in the expression of iNOS, an inducible form of the rate limiting enzyme that makes nitric oxide (NO) in mouse skin. This could be due to neutrophil and macrophage infiltration in the skin (Jain et al., 2011a; Pal et al., 2009). In our study, the observed decrease in NM-induced expression of both COX-2 and iNOS in MPO KO mice compared to WT mice could have further caused decreased inflammatory response in KO mice. Nitric oxide (NO) synthesized by iNOS also plays a potential role in wound healing depending on the production of peroxynitrite (ONOO⁻), which can lead to oxidation of cellular molecules, targets (Korkmaz et al., 2006). MPO-derived secondary oxidants play a role in modulating signaling pathways by oxidizing endothelial-derived NO (Lau et al., 2005). Results here also suggest that MPO deficiency caused a reduction in NM-induced induction of iNOS expression in MPO KO mice as compared to control mice at corresponding time points. This could decrease ONOO by modulating NO-related pathways and subsequent DNA damage as well as cell death.

Earlier reports have shown the involvement of oxidative stress in vesicant-induced activation of DNA damage and MAPK/Akt-NF-kB/AP-1 signaling pathways, and induction of inflammatory mediators in resulting skin injuries (Black et al., 2010; Jain et al., 2011a; Laskin et al., 2010; Pal et al., 2009). The NM-induced WT skin injury could, in part, be due to oxidative stress caused by numerous reactive oxidants from HOCl production or enhanced iNOS expression following the release of MPO from neutrophils. In addition, HOCl generation from MPO can also activate regulatory molecules like p53 and members of

the MAPK pathway (Mutze et al., 2003). Therefore, deletion of MPO in MPO KO mice might be affecting various signaling pathways and oxidative stress, and the injury effects related to DNA damage, cell death and inflammation in these mice. However, these same pro-inflammatory properties of MPO make it a major player in the innate response to foreign invasion and contribute to the microbicidal activity of neutrophils, an important aspect of host defense (Malle et al., 2007). This positive aspect of MPO is important to consider during the development of mechanism-based MPO inhibitors for treatment of injuries. This study also forms the basis of further critical investigation of mechanisms involved in MPO-derived oxidant-induced tissue damage and initiation as well as progression of the SM- and NM-related inflammatory response.

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Highlights

• Myeloperoxidase (MPO) plays an important role in nitrogen mustard (NM)-induced skin injuries.

- Study was carried out in C57BL/6J wild type (WT) and B6.129X1-MPOtm1Lus/J (MPO KO) mice.
- MPO deletion attenuated inflammatory response and microvesication,
- MPO deletion attenuated DNA damage and cell death.
- Deletion of MPO might be affecting various signaling pathways and oxidative stress.

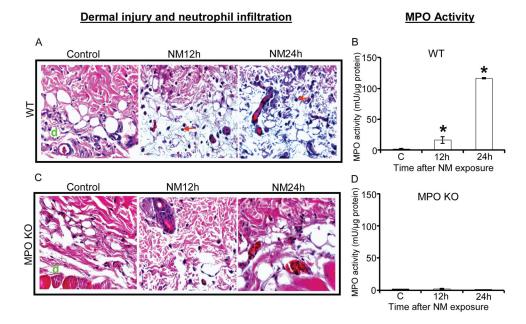
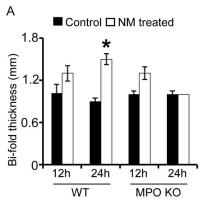


Figure 1. Effect of MPO deletion on MPO activity and neutrophil infiltration after NM exposure Dorsal skin of wild type and MPO KO mice was exposed to 6 mg NM or acetone alone (vehicle control). Following 12 h and 24 h of NM exposure, skin tissue was collected, processed, sectioned and subjected to H&E staining as detailed under Materials and Methods. The H&E stained skin tissue sections from both wild type and MPO KO mice were analyzed microscopically for inflammatory cells including neutrophils in the dermis (A and C). To further assess the MPO activity indicating neutrophil infiltration in the dermis, following similar NM exposure, MPO activity was measured in the skin tissues from wild type and MPO KO mice as detailed under Materials and Methods (B and D). Data presented are mean \pm SEM of three animals. *, p < 0.05 as compared to vehicle control group. D, dermis; WT, wild type mice; MPO KO, MPO knockout mice; red arrows, inflammatory cells.

Skin Bi-fold thickness



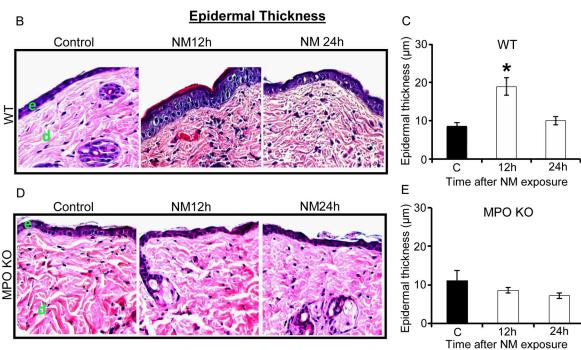


Figure 2. Effect of MPO deletion on skin bi-fo ld thickness and epidermal thickness after NM exposure

Dorsal skin of wild type and MPO KO mice was exposed to 6 mg NM or acetone alone (vehicle control). After 12 h and 24 h of exposure, skin bi-fold thickness was measured using digital calipers (A). Following 12 and 24 h of NM exposure, skin tissue was collected, processed, sectioned and subjected to H&E staining as detailed under Materials and Methods. The H&E stained skin tissue sections from both the wild type and MPO KO mice were analyzed for epidermal thickness (B–E). The NM-induced epidermal thickness in wild type and MPO KO mice shown in representative pictures (B and D) was further quantified (C and D) as detailed under Materials and Methods. Data presented are mean \pm SEM of three animals *, p < 0.05 as compared to vehicle control group. e, epidermis; d, dermis; WT, wild type mice; MPO KO, MPO knockout mice.

Microvesication

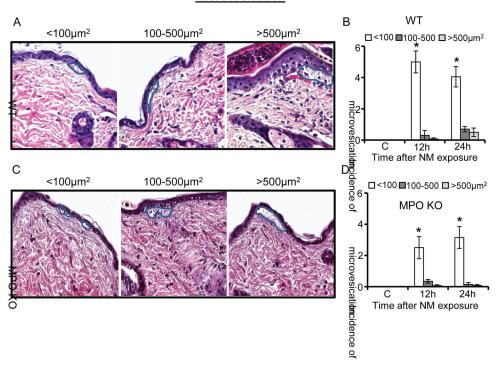


Figure 3. Effect of MPO deletion on microvesication after NM exposure

Dorsal skin of wild type and MPO KO mice was exposed to 6 mg NM or acetone alone (vehicle control). Following 12 h and 24 h of NM exposure, skin tissue was collected, processed, sectioned and subjected to H&E staining as detailed under Materials and Methods. The H&E stained skin tissue sections from both the wild type and MPO KO mice were analyzed for microvesication. The NM-induced incidences of microvesication in wild type and MPO KO mice shown in representative pictures (A and C), were counted according to their sizes and classified into three categories; small (<100 μ m²), medium (100–500 μ m²) and large (>500 μ m²) (B and D). Data presented are mean \pm SEM of three animals *, p < 0.05 as compared to vehicle control group. e, epidermis; d, dermis; WT, wild type mice; MPO KO, MPO knockout mice

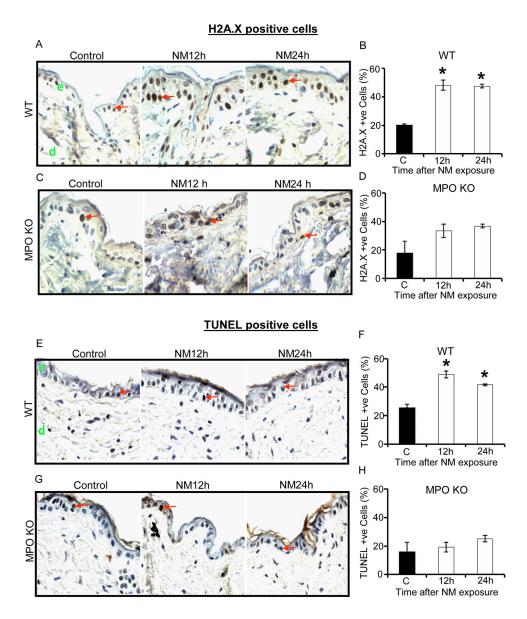


Figure 4. Effect of MPO deletion on DNA damage and apoptotic cell death after NM exposure Dorsal skin of wild type and MPO KO mice was exposed to 6 mg NM or acetone alone (vehicle control). Following 12 h and 24 h of NM exposure, skin tissue was collected, processed, sectioned and analyzed for DNA damage (IHC localization of phospho-H2A.X Ser139, A–D) and cell death (TUNEL assay, E–H) as detailed under Materials and Methods. The NM-induced increase in IHC localization of phospho-H2A.X Ser139 in wild type and MPO KO mice shown in representative pictures (A and C) was quantified (B and D) as detailed under Materials and Methods. The NM induced apoptotic cell death in wild type and MPO KO mice shown in representative pictures (E and G) was further quantified (F and H) as detailed under Materials and Methods. Data presented are mean ± SEM of three animals of each group. *, p<0.001 as compared to vehicle control group. e, epidermis; d, dermis; red arrows, TUNEL positive cells and phosphor-H2A.X Ser139.

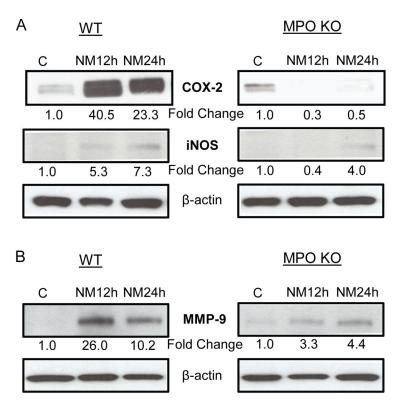


Figure 5. Effect of MPO deletion on the expression of COX2, iNOS, and MMP-9 after NM exposure ${\cal M}_{\rm S}$

Dorsal skin of wild type and MPO KO mice was exposed to 6 mg NM or acetone alone (vehicle control). Following 12 h and 24 h of NM exposure, skin tissue from wild type and MPO KO mice was collected and processed, and subjected to western blot analysis for COX-2 and iNOS (A), and MMP-9 (B) as detailed under Materials and Methods. Data are presented as fold change in comparison with their respective controls. Protein loading was determined by stripping and reprobing of the same membrane for β -actin.