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Regulation of Hsp27 and Hsp70 expression in human and mouse skin construct models by caveolae following exposure to the model sulfur mustard vesicant, 2-chloroethyl ethyl sulfide

Adrienne T. Black¹, Patrick J. Hayden², Robert P. Casillas³, Diane E. Heck⁴, Donald R. Gerecke¹, Patrick J. Sinko⁵, Debra L. Laskin¹, and Jeffrey D. Laskin^{6,*}

¹Pharmacology and Toxicology, Rutgers University, Piscataway, NJ

²MatTek Corporation, Ashland, MA

³Battelle Memorial Institute, Columbus, OH

⁴Environmental Health, New York Medical College, Valhalla, NY

⁵Pharmaceutics, Rutgers University, Piscataway, NJ

⁶Environmental and Occupational Medicine, UMDNJ-Robert Wood Johnson Medical School, Piscataway, NJ, USA

Abstract

Dermal exposure to the vesicant sulfur mustard causes marked inflammation and tissue damage. Basal keratinocytes appear to be a major target of sulfur mustard. In the present studies, mechanisms mediating skin toxicity were examined using a mouse skin construct model and a full-thickness human skin equivalent (EpiDerm-FT™). In both systems, administration of the model sulfur mustard vesicant, 2-chloroethyl ethyl sulfide (CEES, 100–1000 μM) at the air surface induced mRNA and protein expression of heat shock proteins 27 and 70 (Hsp27 and Hsp70). CEES treatment also resulted in increased expression of caveolin-1, the major structural component of caveolae. Immunohistochemistry revealed that Hsp27, Hsp70 and caveolin-1 were localized in basal and suprabasal layers of the epidermis. Caveolin-1 was also detected in fibroblasts in the dermal component of the full thickness human skin equivalent. Western blot analysis of caveolar membrane fractions isolated by sucrose density centrifugation demonstrated that Hsp27 and Hsp70 were localized in caveolae. Treatment of mouse keratinocytes with filipin III or methyl-β-cyclodextrin, which disrupt caveolar structure, markedly suppressed CEES-induced Hsp27 and Hsp70 mRNA and protein expression. CEES treatment is known to activate JNK and p38 MAP kinases; in mouse keratinocytes, inhibition of these enzymes suppressed CEES-induced expression of Hsp27 and Hsp70. These data suggest that MAP kinases regulate Hsp 27 and Hsp70; moreover, caveolae-mediated regulation of heat shock protein expression may be important in the pathophysiology of vesicant-induced skin toxicity.

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*To whom correspondence should be addressed. Jeffrey D. Laskin, Ph.D., Department of Environmental and Occupational Medicine, UMDNJ-Robert Wood Johnson Medical School, 170 Frelinghuysen Road, Piscataway, NJ 08854, jlaskin@ehsi.rutgers.edu.

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Conflict of Interest statement: One of the authors (P.J. Hayden) is employed by MatTek Corporation, manufacturer of the EpiDerm-FT™ full-thickness skin equivalent used in the experiments. The other authors have no conflicts of interest to declare.

Keywords

caveolin-1; sulfur mustard; MAP kinase; skin; heat shock proteins; caveolae

Introduction

Exposure to the chemical warfare agent sulfur mustard, a highly reactive vesicant, and related analogs including, 2-chloroethyl ethyl sulfide (CEES) or half mustard, is known to result in persistent skin damage that includes keratinocyte necrosis, inflammation and blistering (Papirmeister *et al.*, 1991). Toxicity is thought to be mediated by alkylation of cellular macromolecules including proteins, lipids and nucleic acids (Shakarjian *et al.*, 2010; Laskin *et al.*, 2010). This can result in activation of signal transduction pathways regulating cell growth, differentiation and apoptosis contributing to toxicity (Dillman *et al.*, 2004; Rebholz *et al.*, 2008; Kehe *et al.*, 2009; Pal *et al.*, 2009; Black *et al.*, 2010b).

Heat shock proteins (Hsp's) are molecular chaperones important in the folding, assembly and degradation of proteins; they are upregulated in response to cellular stresses such as heat shock, inflammation, and electrophilic and oxidative stress (Macario and Conway de Macario, 2007; Liberek *et al.*, 2008). Hsp's also play critical roles in regulating signal transduction, protein trafficking, cellular proliferation and differentiation (Helmbrecht *et al.*, 2000). Two prominent Hsp's are 27 kDa Hsp (Hsp27) and 70 kDa Hsp (Hsp70). While Hsp27 is an ATP-independent molecular chaperone that protects against stress-induced aggregation of proteins, Hsp70 is ATP-dependent and functions in protein folding (Macario and Conway de Macario, 2007). In intact skin, Hsp70 expression is observed throughout the epidermis, while Hsp27 is localized primarily in the granular and spinous layers, indicating that this protein is also involved in keratinocyte differentiation (Robitaille *et al.*; Gandour-Edwards *et al.*, 1994; Trautinger *et al.*, 1995; Trautinger, 2001). Increases in both Hsp27 and Hsp70 have been reported in human and mouse skin, as well as in cultured keratinocytes and skin explants in response to heat shock, UVB light, heavy metals, oxidative stress, and inflammatory mediators (Maytin, 1995; Trautinger, 2001). Hsp's are thought to protect against cellular stress and toxicity by modulating intracellular signal transduction pathways such as MAP kinase and Nrf2 (Mosser *et al.*, 2000). Hsp27 and Hsp70 are also known to interact with the cytoskeleton and are involved in the regulation of actin and microtubulin polymerization and reorganization (Liang and MacRae, 1997).

Caveolae comprise a subset of membrane lipid rafts important in intracellular signaling, lipid transport and protein trafficking (Cohen *et al.*, 2004). Caveolin-1 is the primary structural protein in caveolae (Liu *et al.*, 2002). Although caveolae play an important role in regulating growth and differentiation in keratinocytes, they do not appear to be required for skin development (Roelandt *et al.*, 2009). Thus, while caveolin-1 knockout mice can generate functional epidermis, aberrant keratinocyte proliferation and differentiation are observed (Capozza *et al.*, 2003; Roelandt *et al.*, 2009). Previous studies have shown that Hsp27 and Hsp70 are present in lipid rafts and co-localize with caveolar proteins (Broquet *et al.*, 2003; Chen *et al.*, 2005; Staubach *et al.*, 2009). Moreover, there appears to be a link between caveolae-mediated cell signaling and cytoskeletal organization which may be important in regulating keratinocyte responses to stress (Head *et al.*, 2006).

In the present studies, we analyzed the effects of CEES on expression of Hsp27 and Hsp70 in a full-thickness human skin equivalent and a mouse keratinocyte skin construct model (Black *et al.*, 2010a) and investigated the role of caveolae in regulating expression of these proteins. CEES was found to markedly upregulate expression of Hsp 27 and Hsp70, a MAP kinase-dependent process in which these Hsp's were selectively localized in caveolae.

Disruption of caveolae blocked CEES-induced changes in expression of the Hsp's, demonstrating a direct link between sensitivity to CEES and caveolae expression. Hsp expression via the formation of caveolae may be an adaptive response important in protecting cells during stress. Thus, caveolae may play a key role in activation of signaling pathways and protein trafficking in mediating the keratinocytes response to sulfur mustard.

Materials and Methods

Chemicals and reagents

All primary antibodies were affinity-purified. Rabbit polyclonal antibodies to p38, phospho-p38, JNK, phospho-JNK, ERK1/2, and phospho-ERK1/2 were from Cell Signaling Technology (Beverly, MA). Rabbit monoclonal or polyclonal antibodies to caveolin-1 were from Cell Signaling Technology or Santa Cruz Biotechnology (Santa Cruz, CA). Rabbit polyclonal antibodies to Hsp70, goat polyclonal antibodies to Hsp27 and β -actin, and horseradish peroxidase-labeled donkey anti-goat secondary antibodies were from Santa Cruz Biotechnology (Santa Cruz, CA). Horseradish peroxidase-labeled goat anti-rabbit secondary antibodies and detergent-compatible protein assay reagents were obtained from Bio-Rad Laboratories (Hercules, CA). MultiScribe Reverse Transcriptase was from Promega (Madison, WI), the RNeasy purification kit from Qiagen (Minneapolis, MN), and the Western Lightning enhanced chemiluminescence kit (ECL) from Perkin Elmer Life Sciences, Inc. (Boston, MA). Rabbit and goat IgG were from ProSci (Poway, CA) and the Vectastain Rabbit and Goat Kits and the Peroxidase Substrate Kit DAB from Vector Labs (Burlingame, CA). SYBR Green Master Mix and other PCR reagents were purchased from Applied Biosystems (Foster City, CA) and precast gradient polyacrylamide gels from Pierce Biotechnology, Inc. (Rockford, IL). Cell culture reagents were from Invitrogen Corp (Carlsbad, CA) and polyester Transwell permeable supports from Corning Life Sciences Inc (Acton, MA). SP600125 and filipin III were from EMD Biosciences (La Jolla, CA) and CEES, SB203580, methyl- β -cyclodextrin (MBCD), protease inhibitor cocktail and all other chemicals from Sigma-Aldrich (St. Louis, MO). CEES (100 mM stock solution) was prepared fresh in absolute ethanol immediately before use and diluted to the appropriate concentrations in PBS.

Mouse keratinocyte skin construct

A three dimensional skin-like structure was generated using PAM212 mouse keratinocytes grown in DMEM supplemented with 10% fetal bovine serum as previously described (Black *et al.*, 2010b). Briefly, cells (1×10^6) were grown on 24-mm Transwell semi-permeable polyester membrane supports (0.4 μ m pore size) in 6-well cell culture plates. After reaching confluence, the medium was removed from the upper portion of the Transwell such that the cells were exposed to air at the apical surface and in contact with the medium at the basolateral surface. After 24 hr incubation at 37°C in a humidified CO₂ incubator, cells were treated with 1 ml vehicle control or CEES for 2 hr. The Transwells were removed, washed in PBS and immediately replaced in the same 6-well culture plates. In some experiments, the cells were preincubated at 37°C for 30 min or 3 hr prior to CEES treatment at the apical surface with p38 MAP kinase inhibitor (SB203580, 10 μ M), JNK inhibitor (SP600125, 20 μ M), caveolae inhibitors (filipin, 10 μ M or MBCD, 5 mM), or DMSO control.

Human skin equivalent construct model

EpiDerm-FT™ full-thickness human skin equivalents (EFT-400) and EFT-400-MM medium, supplemented with growth factors, hormones and lipid precursors, were kindly provided by MatTek Corporation (Ashland, MA). The skin equivalents were placed in 6-well plates in 2 ml of EFT-400-MM medium. After overnight incubation at 37°C in a humidified CO₂ incubator, 1 ml of PBS containing vehicle control or CEES was added to

the apical surface of the tissues. After 2 hr, the skin equivalents were removed from the plates, washed in PBS, immediately replaced in the same 6-well culture dishes and incubated for an additional 2–72 hr. For immunohistochemistry, tissues were fixed in 3% paraformaldehyde in PBS supplemented with 2% sucrose at 4°C for 24 hr, and then removed from the supports, transferred to 50% ethanol and paraffin embedded. Tissue sections (5 µm) were deparaffinized, blocked in 100% serum at room temperature for 2 hr, and then incubated overnight at 4°C with rabbit or goat IgG control, anti-Hsp27 antibody (1:400), anti-Hsp70 antibody (1:500) or anti-caveolin-1 antibody (1:800) followed by incubation with biotinylated anti-rabbit or goat antibodies (1:200) for 30 min at room temperature. Binding was visualized using a Peroxidase Substrate Kit. For PCR and Western blot analysis, the epidermis was removed from unfixed skin equivalents by gentle peeling and immediately analyzed for mRNA or protein expression.

Isolation of caveolae

Caveolar fractions of mouse keratinocytes were prepared as described by Smart et al. (Smart et al., 1995). Briefly, after incubation with 300 µM CEES or control, cells were scraped from the Transwell supports in a sucrose buffer (0.25 M sucrose, 20 mM Tricine, 1 mM EDTA, pH 7.8), centrifuged at 1400 × *g* for 5 min at room temperature, and resuspended in 0.5 ml sucrose buffer. The cells were then homogenized using a glass Dounce homogenizer and centrifuged at 1000 × *g* for 10 min at 4°C. The supernatant was collected and the pellet resuspended in an additional 0.5 ml of sucrose buffer, homogenized, and centrifuged. The second supernatant was combined with the first supernatant which was then carefully layered onto 9 ml Percoll solution (30% Percoll in sucrose buffer) and centrifuged at 84,000 × *g* for 30 min in a Ti-70 rotor using a Beckman Coulter L7–55 ultracentrifuge. The caveolar fraction, visible as a white band approximately 2 cm from the top of the ultracentrifuge tube, was collected and protease inhibitor cocktail (1:100) added.

Western blotting

Lysates were prepared using buffer containing 50 mM Tris-HCl pH 8.0, 150 mM NaCl and 1% Triton X-100 supplemented with 5 µl protease inhibitor cocktail [4-(2-aminoethyl)benzenesulfonyl fluoride, aprotinin, bestatin hydrochloride, N-(trans-epoxysuccinyl)-L-leucine 4-guanidinobutylamide, EDTA and leupeptin]. Proteins (20 µg) from lysates were separated on 10% SDS-polyacrylamide gels and then transferred to nitrocellulose membranes. After incubation in blocking buffer (5% dry milk Tris-buffered saline containing 0.1% Tween 20) for 1 hr at room temperature, the membranes were incubated overnight at 4°C with anti-Hsp27 (1:400), anti-Hsp70 (1:200) or anti-caveolin-1 (1:1000) antibodies followed by horseradish peroxidase-conjugated secondary antibodies for 1 hr at room temperature. Protein expression was visualized using enhanced chemiluminescence (ECL) reagents.

Real-time polymerase chain reaction

RNA was isolated using the Versagene RNA purification kit following the manufacturer's protocol. RNA was converted to cDNA using Superscript reverse transcriptase. The cDNA was diluted 1:10 in RNase-DNase-free water for PCR analysis. For each gene, a standard curve composed of a serial dilution of pooled cDNA from the samples was used as a reference. All values were normalized to GAPDH (*n* = 3). The control was assigned a value of one and samples calculated relative to control. Real-time PCR was performed on an ABI Prism 7300 Sequence Detection System using 96-well optical reaction plates. SYBR-Green was used for detection of fluorescent signal and the standard curve method was used for relative quantitative analysis. The primer sequences for the genes were generated using Primer Express software (Applied Biosystems) and the oligonucleotides were synthesized

by Integrated DNA Technologies, Inc. (Coralville, IA). The forward and reverse sequences (5' → 3') used are listed in Table 1.

Statistical analysis

All experiments were repeated 3 times with similar results. Data are expressed as mean ± SEM. Statistical differences between the means were determined using two-way ANOVA and considered significant at $p < 0.05$.

Results

Effects of CEES on expression of Hsp's

In initial studies we determined if CEES treatment of human and mouse skin construct models altered expression of Hsp27 and Hsp70. Immunohistochemical analysis of the human skin equivalent revealed low constitutive expression of Hsp27 and Hsp70 in control keratinocytes (Figs. 1–4). Treatment of the human skin equivalent with CEES (100–1000 μM) resulted in a time- and concentration-dependent increase in both Hsp27 and Hsp70 in keratinocytes, as well as fibroblasts which was maximal at 24 hr (Fig. 1 and 2). The Hsp's were identified in both cytosol and nuclei of basal and suprabasal keratinocytes and fibroblasts. CEES-induced upregulation of Hsp27 and Hsp70 was confirmed by western blotting and real time PCR analysis of epidermal sheets isolated from the skin equivalents (Figs. 5 and 6). Maximal effects were noted with 1000 μM CEES. Similar results were observed in the mouse skin construct model. Thus, CEES treatment (100–1000 μM) resulted in concentration-dependent increases in mRNA expression of Hsp27 and Hsp70 24 hr post-treatment (Fig. 6). CEES also upregulated Hsp27 and Hsp70 protein expression with maximal increases detected at 1000 μM CEES (Fig. 5).

Role of caveolae in CEES-induced alterations in Hsp27 and Hsp70 expression

In further studies, we determined if plasma membrane caveolae play a role in mediating CEES-induced increases in expression of Hsp27 and Hsp70. Analysis of tissue sections by immunohistochemistry revealed low constitutive expression of caveolin-1, the major structural component of caveolae, in basal keratinocytes and fibroblasts in human skin equivalents. CEES treatment (100–1000 μM) upregulated expression of this protein in both cell types; maximal expression was evident after 24 hr with 300–1000 μM CEES (Fig. 7). CEES-induced caveolin-1 expression was extranuclear, and predominantly noted as punctate staining of keratinocyte membranes throughout the basal layer (Fig. 7). CEES treatment was also found to up regulate caveolin-1 mRNA (up to 4-fold after 24 hr) and protein expression (Figs. 5 and 6) in isolated epidermal sheets from the human skin equivalents. In the mouse skin construct model, CEES also caused a concentration-dependent increase in caveolin-1 mRNA (up to 8-fold after 24 hr) and protein (Figs. 5 and 6). In these cells, maximal increases were detected at 1000 μM CEES. In further experiments, caveolae were isolated from untreated and CEES (300 μM)-treated keratinocytes from the mouse skin construct model and analyzed for Hsp27 and Hsp70 expression. Interestingly, both Hsp's were found to be selectively expressed in caveolae (Fig. 8, upper panel). Moreover, CEES treatment resulted in increases in Hsp27 and Hsp70 that were also selectively localized in caveolae (Fig. 8, lower panel). Treatment of the keratinocytes filipin (20 μM) or methyl- β -cyclodextrin (MBCD; 5 mM), which disrupt caveolar integrity (Simons and Toomre, 2000), inhibited CEES-induced increases in expression of mRNA and protein for Hsp27 and Hsp70 (Fig. 9, left panel, Fig. 10, and not shown).

Role of MAP kinase signaling in CEES-induced alterations in heat shock proteins

Both sulfur mustard and CEES are known to activate MAP kinase signaling in intact skin and cultured keratinocytes (Dillman *et al.*, 2004; Rebholz *et al.*, 2008; Pal *et al.*, 2009). CEES (100–1000 μM) also activates JNK and p38 MAP kinases in mouse keratinocytes in the skin construct model (Black *et al.*, 2010b). To assess the role of JNK and p38 MAP kinases in regulating CEES-induced alterations in expression of Hsp27 and Hsp70, we used specific inhibitors of these enzymes. Treatment of the mouse keratinocytes with the p38 MAP kinase inhibitor, SB203580, or the JNK inhibitor, SP600125, suppressed CEES-induced expression of Hsp27 and Hsp70 mRNA and protein (Fig. 9, right panel, and Fig. 10). p38 kinase inhibition was more effective than JNK inhibition in suppressing expression of caveolin-1 mRNA and protein.

Discussion

Dermal vesicants such as sulfur mustard or CEES are known to induce oxidative stress (Shakarjian *et al.*, 2010; Laskin *et al.*, 2010). A characteristic response to oxidative stress is the upregulation of Hsp's, an important class of molecular chaperones which function to minimize tissue injury (Helmbrecht *et al.*, 2000). Previous studies have demonstrated that Hsp27 and Hsp70 are rapidly upregulated in human and mouse skin, as well as cultured keratinocytes in response to UVB light, ozone, heavy metals and inflammatory mediators (Maytin, 1995; Trautinger, 2001; Valacchi *et al.*, 2004; Matsuda *et al.*, 2010). We found that CEES also effectively upregulates expression of these Hsp's in a full-thickness human skin equivalent and in a mouse keratinocyte skin construct. In the full-thickness human skin equivalent model, Hsp27 was evident in both basal and suprabasal layers of the epidermis. This is consistent with reports demonstrating that Hsp27 is increased in these cell layers in mouse skin during wound repair (Laplante *et al.*, 1998). Overexpression of Hsp27 has been shown to upregulate cellular actin levels and to stimulate wound contraction during the healing process (Lavoie *et al.*, 1993). These data suggest that increases in Hsp27 are key to triggering tissue repair. Hsp27 is also associated with the epidermal differentiation markers, involucrin, filaggrin, and transglutaminase suggesting that it contributes to the differentiation process (Robitaille *et al.*; Jonak *et al.*, 2002).

Following CEES exposure, Hsp70 is also markedly increased in basal and suprabasal keratinocytes. Studies with mice overexpressing or lacking Hsp70 have demonstrated that Hsp70 is critical for epidermal repair following exposure to UVB light (Kwon *et al.*, 2002; Matsuda *et al.*, 2010). Additionally, *in vivo* delivery of Hsp70 accelerates the rate of full-thickness wound closure in mice (Kovalchin *et al.*, 2006). In clinical studies, decreased Hsp70, as well as Hsp27, have been linked to delayed and impaired healing of chronic wounds commonly associated with diabetes (Atalay *et al.*, 2009). Similarly, in diabetic mice, Hsp70 expression is delayed following wounding, leading to an increased healing time (McMurtry *et al.*, 1999). These data suggest that Hsp70, like Hsp27, is critical for the early stages of wound repair. The mechanisms by which Hsp27 and Hsp70 contribute to this process are not known. As chaperones, these Hsp's may protect critical proteins modified by vesicants. For example, recent studies have shown that both sulfur mustard and CEES alkylate components of the cytoskeleton, resulting in keratin aggregation and actin polymerization (Dillman *et al.*, 2003; Hess and FitzGerald, 2007; Sayer *et al.*, 2009). Heat shock proteins are key to maintaining the structure and organization of cytoskeletal filaments including actin and keratins (Liang and MacRae, 1997; Loffek *et al.*, 2010). Increases in Hsp27 and Hsp70 in keratinocytes following CEES exposure may be important in maintaining cytoskeletal organization, thereby promoting cellular survival and wound repair.

As indicated above, vesicant-induced injury involves oxidative stress, a process that can result in protein and DNA oxidation and the formation of lipid peroxidation products (Pal *et al.*, 2009; Black *et al.*, 2010b). Protein damage resulting from oxidative stress such as structural denaturation and thiol oxidation results in increased expression of Hsp's which target damaged molecules for degradation (Kalmar and Greensmith, 2009). Hsp's also modulate cellular redox state, an activity that may also help to protect against further oxidative injury (Kalmar and Greensmith, 2009). In this regard, both Hsp27 and Hsp70 have been shown to maintain glutathione in its reduced form, and to decrease intracellular free iron concentrations, thus preventing excessive production of reactive oxygen species (Arrigo *et al.*, 2005; Guo *et al.*, 2007). Hsp70 has also been shown to control activation of histone H2AX, a critical signaling molecule in the repair of double strand breaks in DNA induced by oxidative stress (Gabai *et al.*, 2010). This activity is consistent with our findings that Hsp70 localizes in the nucleus following CEES treatment. We have previously shown that CEES induces expression of the activated form of H2AX in the full-thickness human skin equivalent (Black *et al.*, 2010a), and it may be that upregulation of Hsp70, as well as Hsp27, which can also localize in the nucleus, is key to regulating H2AX activation and subsequent DNA repair following exposure to vesicants.

Caveolae are specialized membrane lipid rafts that control a variety of biochemical signaling molecules and cellular processes including growth and differentiation (Cohen *et al.*, 2004). The present studies demonstrate that Hsp27 and Hsp70 are localized, not only in the nucleus, but also in caveolar fractions of keratinocytes. CEES treatment was found to upregulate caveolar-associated expression of Hsp70, as well as Hsp27, although to a lesser extent. These data are in agreement with previous studies showing Hsp expression in detergent-resistant fractions containing lipid rafts and caveolae in many tissues and cell types (Nedellec *et al.*, 2002; Triantafilou *et al.*, 2002; Broquet *et al.*, 2003; Chen *et al.*, 2005; Lancaster and Febbraio, 2005). These latter studies also showed that cellular stressors such as heat shock, calcium depletion, endotoxin, and glucocorticoids increase Hsp expression in the lipid raft fractions. Caveolae are associated with cytoskeletal and other structural proteins including actin, annexin II, filamin, and dynamin which are thought to be important in regulating caveolar functioning (Stahlhut and van Deurs, 2000; Viola and Gupta, 2007). Hsp27 has been reported to associate with actin in lipid rafts fractions, possibly controlling actin and microtubule depolymerization (Piotrowicz and Levin, 1997). Thus, increases in Hsp27 following CEES-induced keratinocyte stress may be important in minimizing damage to caveolar-associated cytoskeletal proteins critical to caveolae function. Hsp70, on the other hand, appears to be primarily involved in responses to oxidative stress and innate immunity (Macario and Conway de Macario, 2007). In this regard, caveolar localization of Hsp70 has been shown to downregulate NADPH oxidase, thereby reducing reactive oxygen species production (Bocanegra *et al.*, 2010). Hsp70, as well as Hsp27, have been detected in the extracellular environment, and it has been proposed that caveolae function as vehicles by which Hsp's are transported to the cell membrane for externalization (Broquet *et al.*, 2003). It is possible that Hsp70 localized in caveolae functions to prevent oxidative damage to proteins and lipids while extracellular Hsp70 acts as a danger signal to the innate immune system (Vega *et al.*, 2008).

Of interest was our finding that CEES treatment also upregulated keratinocyte expression of caveolin-1, the major structural protein in caveolae (Cohen *et al.*, 2004). Increases in caveolin-1 after vesicant exposure may be important in wound healing and in regulating keratinocyte growth and differentiation (Sando *et al.*, 2003; Zheng and Bollinger Bollag, 2003; Roelandt *et al.*, 2009). Upregulation of caveolin-1 has been shown to be important in protecting against radiation-induced DNA damage by activating repair mechanisms (Zhu *et al.*, 2010). Caveolar localization of Hsp70 may allow it to interact with caveolin-1, thereby facilitating repair of DNA.

An unexpected finding in our studies was that Hsp27 and Hsp70 are not only localized in caveolae, but are also regulated by these structures. Thus, disruption of caveolae by cholesterol depletion markedly decreased CEES-induced Hsp27 and Hsp70 gene and protein expression. These results are in accord with previous reports showing that disruption of caveolae results in decreased heat shock-induced expression of Hsp's (Broquet *et al.*, 2003; Chen *et al.*, 2005). It has also been shown that disruption of caveolae interferes with cellular release of Hsp70, providing additional support for the idea that caveolar internalization of Hsp70 is important in its transport to the cell membrane (Broquet *et al.*, 2003; Lancaster and Febbraio, 2005).

A question remains as to the mechanism by which CEES upregulates Hsp27 and Hsp70. In previous studies we demonstrated that CEES activates JNK and p38 MAP kinase signaling pathways in the mouse skin equivalent model (Black *et al.*, 2010b). Using inhibitors of these MAP kinases, we found that both JNK and p38 MAP kinase regulate expression of Hsp27 and Hsp70. Consistent with these results are studies showing that activation of Hsp27 and Hsp70 are p38 MAP kinase-dependent (Kato *et al.*, 1999; Garmyn *et al.*, 2001; Kim *et al.*, 2005a; Kim *et al.*, 2005b; Dasari *et al.*, 2006; Banerjee Mustafi *et al.*, 2009). Interestingly, activation of the p38 MAP kinase-Hsp27 signaling pathway has been shown to result in the formation of blisters in autoimmune blistering diseases (Berkowitz *et al.*, 2005; Berkowitz *et al.*, 2006; Berkowitz *et al.*, 2008a; Berkowitz *et al.*, 2008b). Using a pemphigus vulgaris mouse model, inhibition of p38 MAP kinase was found to prevent blister formation, as well as cytoskeletal reorganization and loss of cell-cell adhesion (Berkowitz *et al.*, 2005; Berkowitz *et al.*, 2006). It may be that inhibitors of p38 MAP kinase, as well as JNK kinase, can suppress sulfur mustard-induced epidermal damage and subsequent skin injury including blistering. In this regard, an inhibitor of p38 MAP kinase has been shown to downregulate sulfur mustard-induced cytokine release in human epidermal keratinocytes (Dillman *et al.*, 2004).

In summary, the present studies showed that CEES exposure upregulates expression of Hsp27, Hsp70 and caveolin-1 in a full-thickness human skin equivalent and a mouse keratinocyte skin construct model. Moreover, Hsp27 and Hsp70 are localized in caveolar subcellular fractions and expression of these proteins is regulated by both caveolae and JNK and p38 MAP kinase signaling. Further studies are needed to determine if localization of the Hsp's in the caveolae are important in protecting keratinocytes from vesicant-induced cellular damage. It is possible that caveolae-associated protein transport and control of signaling pathways play key roles in modulating cellular responses to vesicant exposure in the skin.

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List of Abbreviations

CEES	2-chloroethyl ethyl sulfide
cav-1	caveolin-1
Hsp27	heat shock protein 27
Hsp70	heat shock protein 70

MBCD methyl- β -cyclodextrin

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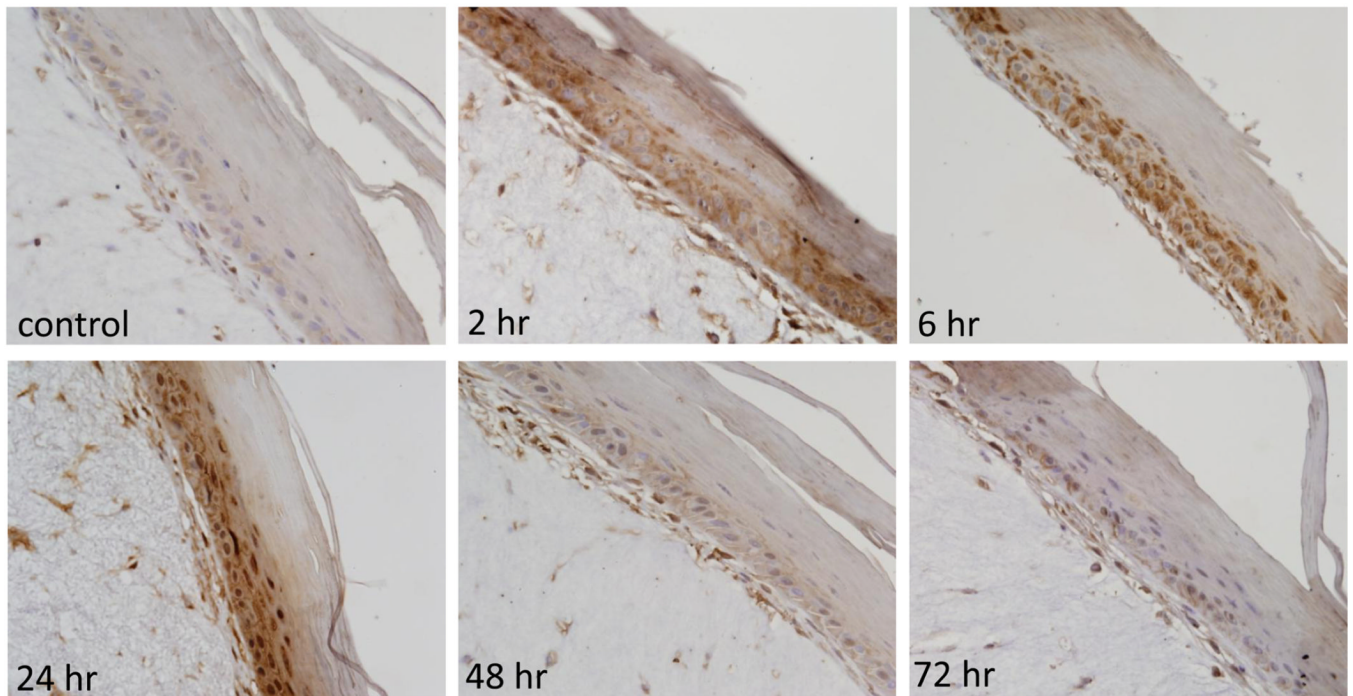


Figure 1. Effects of CEES on Hsp27 expression in a full-thickness human skin equivalent EpiDerm-FT™ was exposed on the air surface with CEES (1000 μ M) or control. Tissues were collected 2–72 hr later and stained with anti-Hsp27 antibody. Binding was visualized using a peroxidase DAB substrate kit. Original magnification, 400 \times .

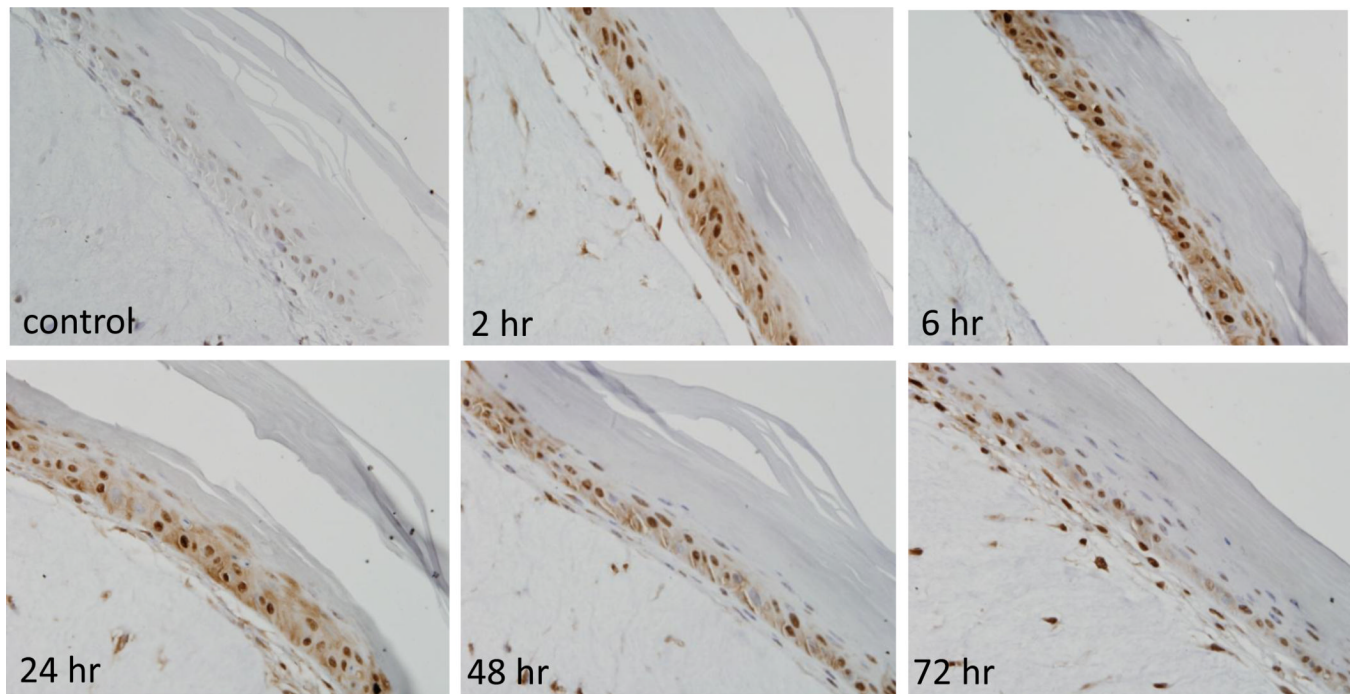


Figure 2. Effects of CEES on Hsp70 expression in a full-thickness human skin equivalent EpiDerm-FT™ was exposed on the air surface with CEES (1000 μ M) or control. Tissues were collected 2–72 hr later and stained with anti-Hsp70 antibody. Binding was visualized using a peroxidase DAB substrate kit. Original magnification, 400 \times .

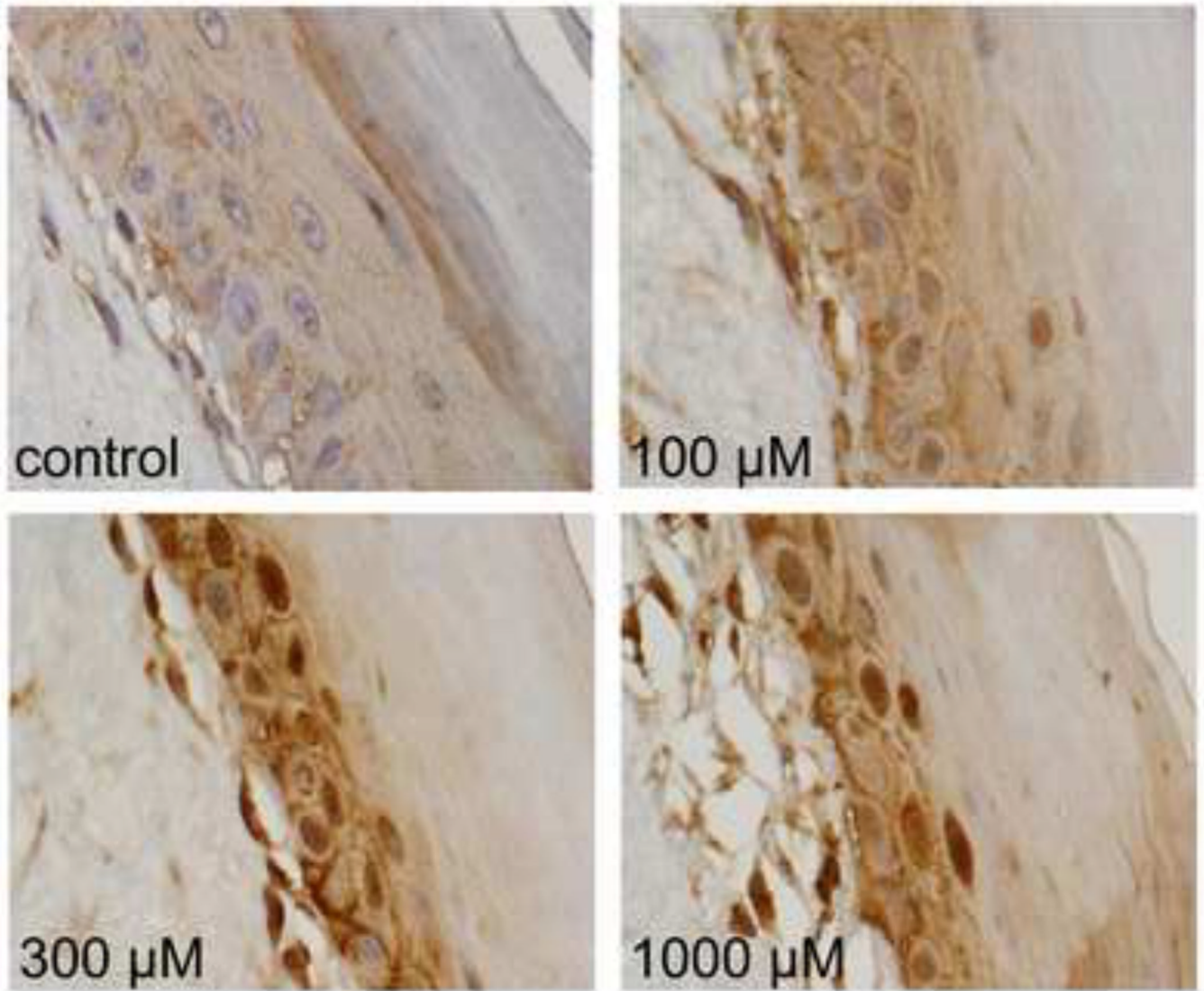


Figure 3. Effects of CEES dose on Hsp27 expression in a full-thickness human skin equivalent EpiDerm-FT™ was exposed to CEES (100–1000 μM) or control. Tissues were collected 24 hr later and stained with antibodies to Hsp27. Binding was visualized using a peroxidase DAB substrate kit. Original magnification, 1000 \times .

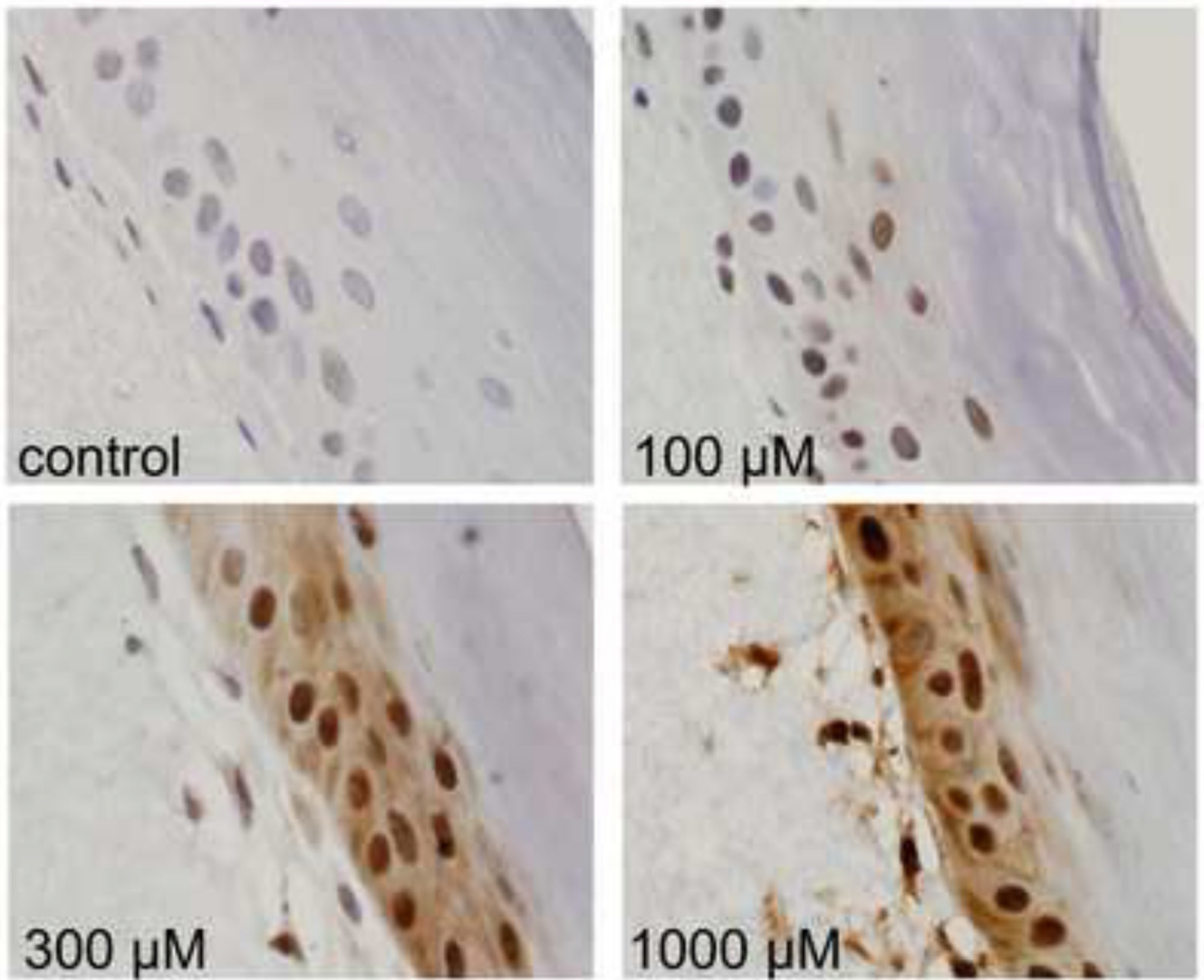


Figure 4. Effects of CEES on Hsp70 expression in a full-thickness human skin equivalent EpiDerm-FT™ was exposed to CEES (100–1000 μM) or control. Tissues were collected 24 hr later and stained with antibodies to Hsp70. Binding was visualized using a peroxidase DAB substrate kit. Original magnification, 1000 \times .

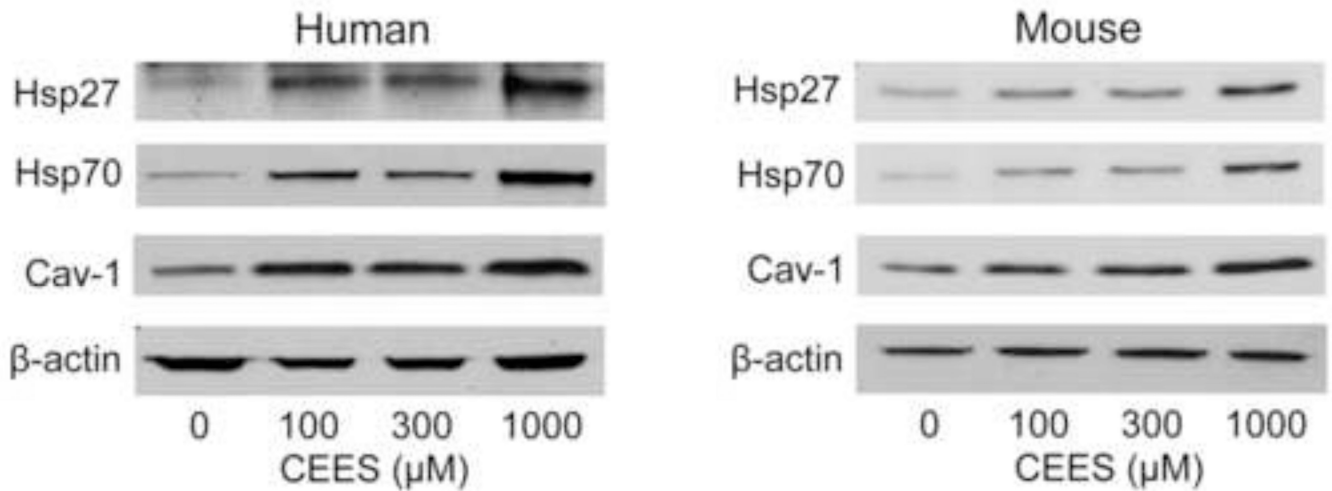


Figure 5. CEES upregulates Hsp27, Hsp70 and caveolin-1 protein expression in a full-thickness human skin equivalent and in a mouse keratinocyte skin construct

EpiDerm-FTTM or mouse keratinocyte skin constructs were exposed to CEES (100–1000 μM) or control. After 24 hr, epidermal sheets from the skin equivalents and keratinocytes from the skin construct were collected and analyzed for Hsp27, Hsp70 and caveolin-1 (cav-1) protein expression by Western blotting. β-actin was used as a control for equal protein loading.

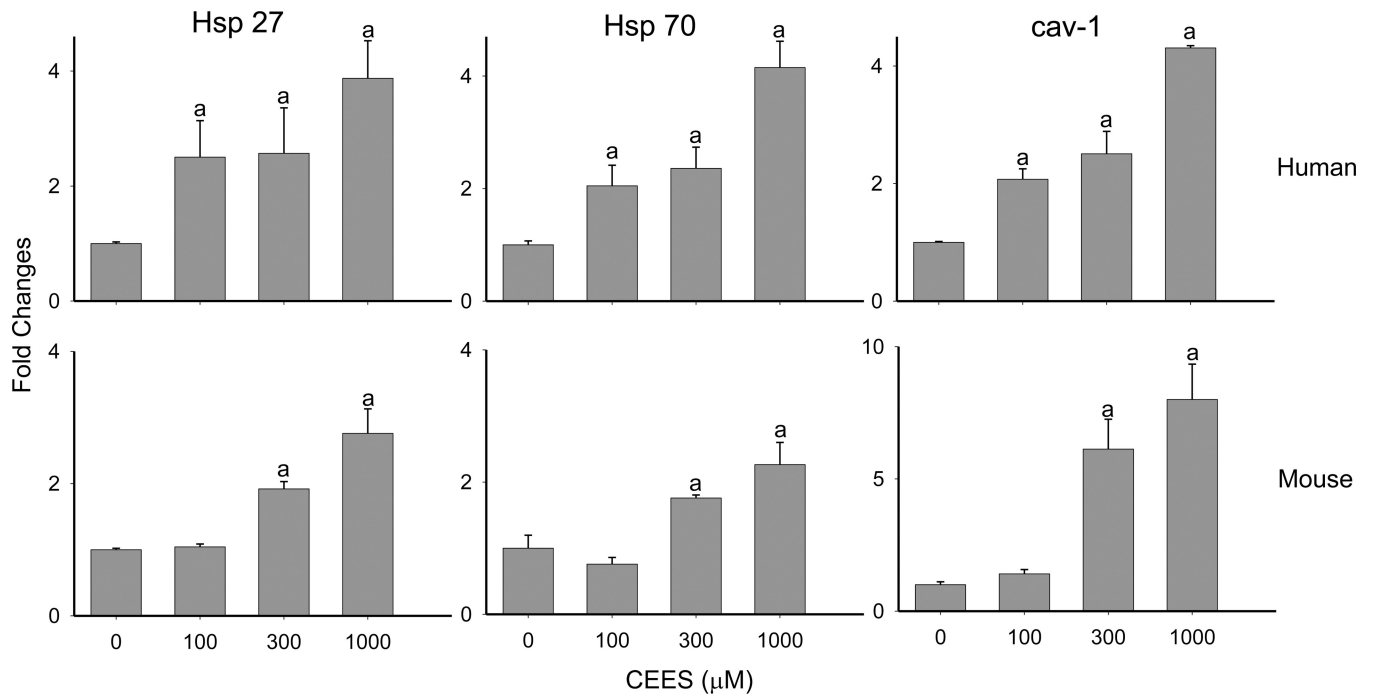


Figure 6. CEES upregulates Hsp27, Hsp70 and caveolin-1 mRNA expression in a full-thickness human skin equivalent and in a mouse keratinocyte skin construct

EpiDerm-FTTM or mouse keratinocyte skin constructs were exposed to CEES (100–1000 μM) or control. After 24 hr, epidermal sheets from the skin equivalents and keratinocytes from the skin construct were collected and analyzed for Hsp27, Hsp70 and caveolin-1 (cav-1) mRNA expression by real-time PCR. Data are presented as fold change in gene expression relative to control cells. ^aSignificantly ($p < 0.05$) different from control.

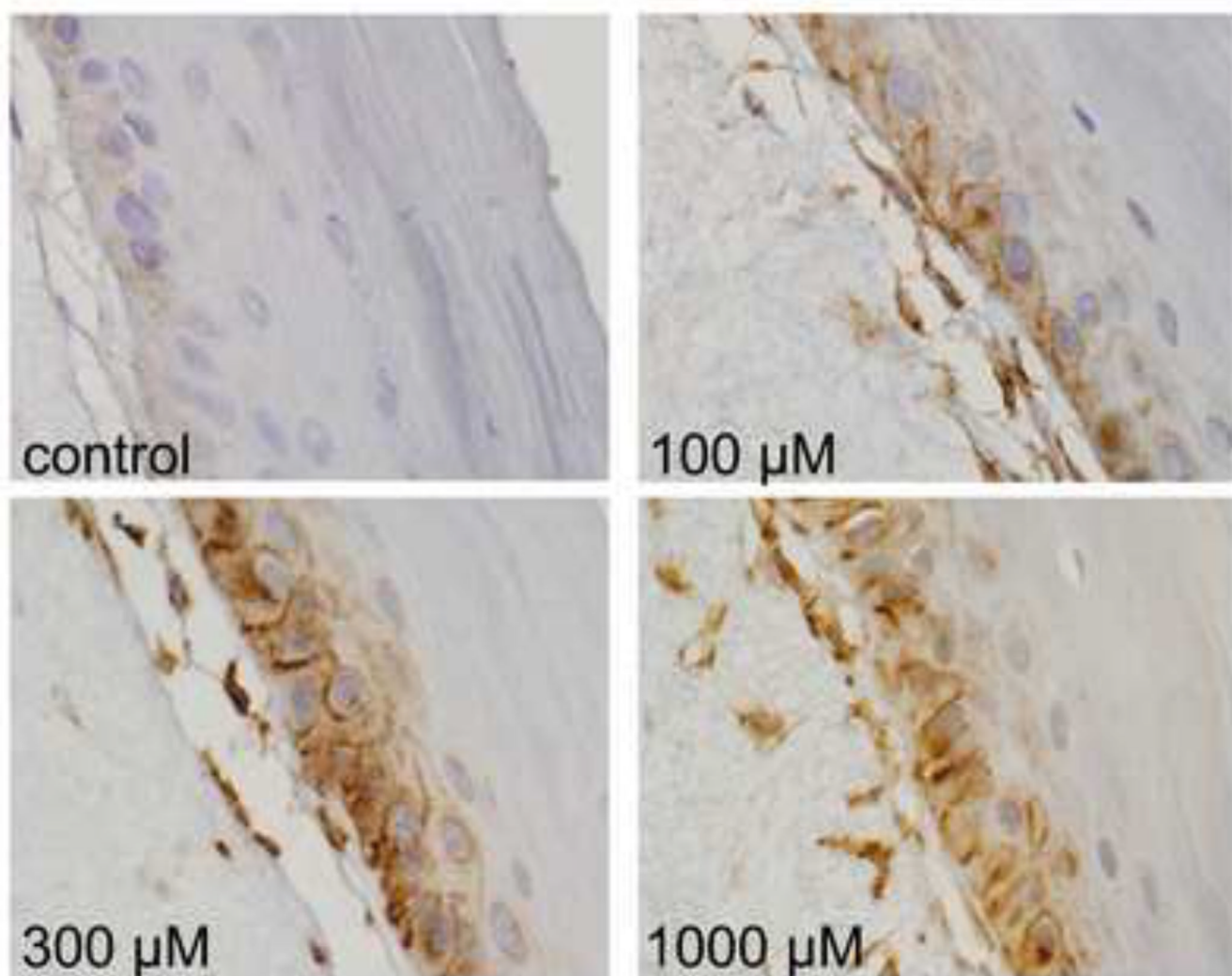


Figure 7. Effects of CEES on caveolin-1 expression in a full-thickness human skin equivalent EpiDerm-FT™ was exposed to CEES (100–1000 μM) or control. Tissues were collected 24 hr later and stained with antibodies to caveolin-1 (cav-1). Protein expression was visualized using a peroxidase DAB substrate kit. Original magnification, 1000 \times .

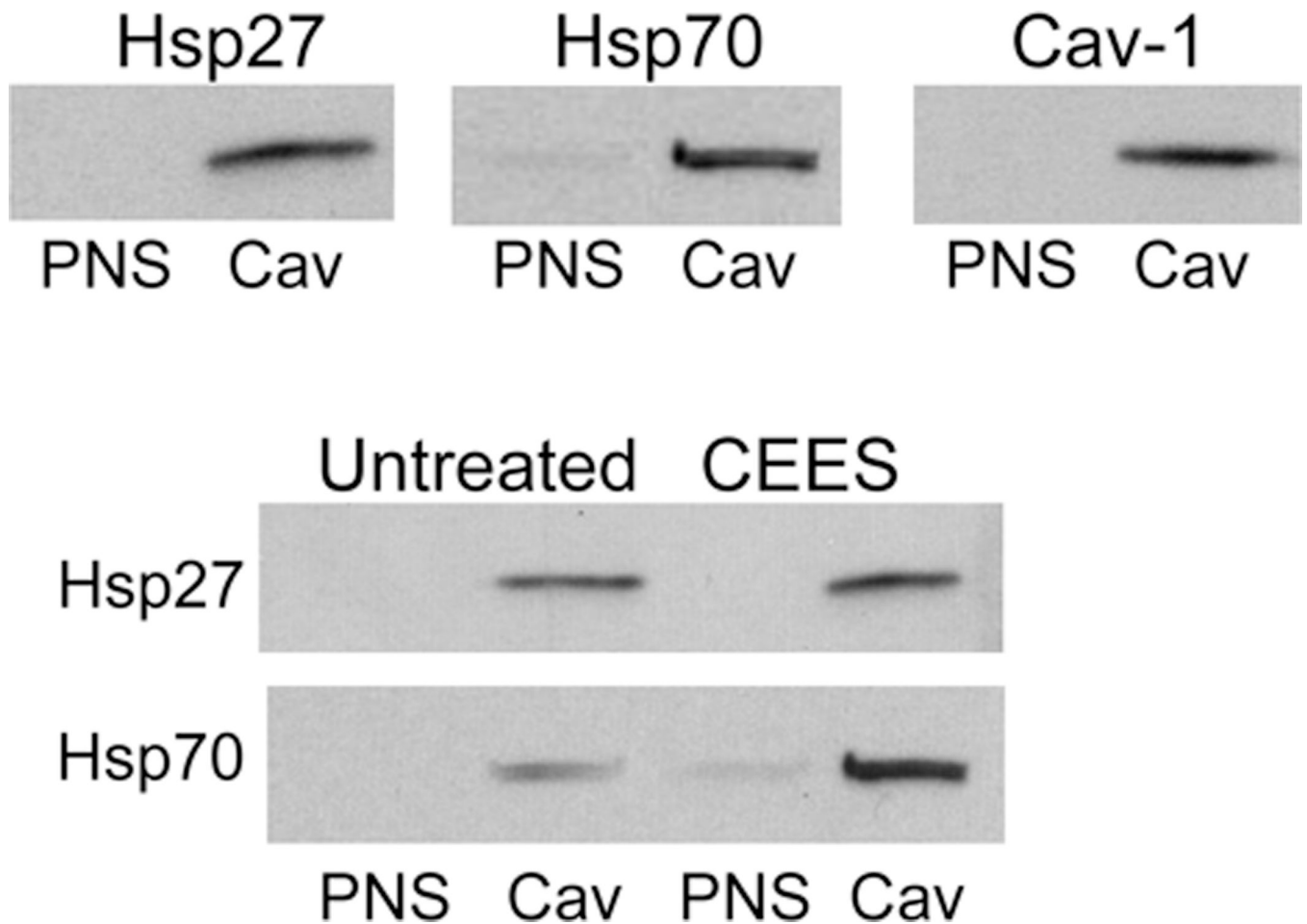


Figure 8. Localization of Hsp27 and Hsp70 in caveolae

Mouse keratinocyte skin constructs were exposed to CEES (300 μ M) or control. After 24 hr, caveolar (Cav) and post-nuclear supernatant (PNS) subcellular fractions were isolated from the cells using sucrose density gradient centrifugation as described in the Materials and Methods. *Upper panel:* Cav and PNS fractions were assayed for Hsp27 and Hsp70 by Western blotting. The relative purity of the caveolar fractions was determined by Western blot analysis using caveolin-1 antibodies. *Lower panel:* Effect of CEES on Hsp27 and Hsp70 expression in cav and PNS subcellular fractions isolated from control and CEES-treated mouse keratinocytes.

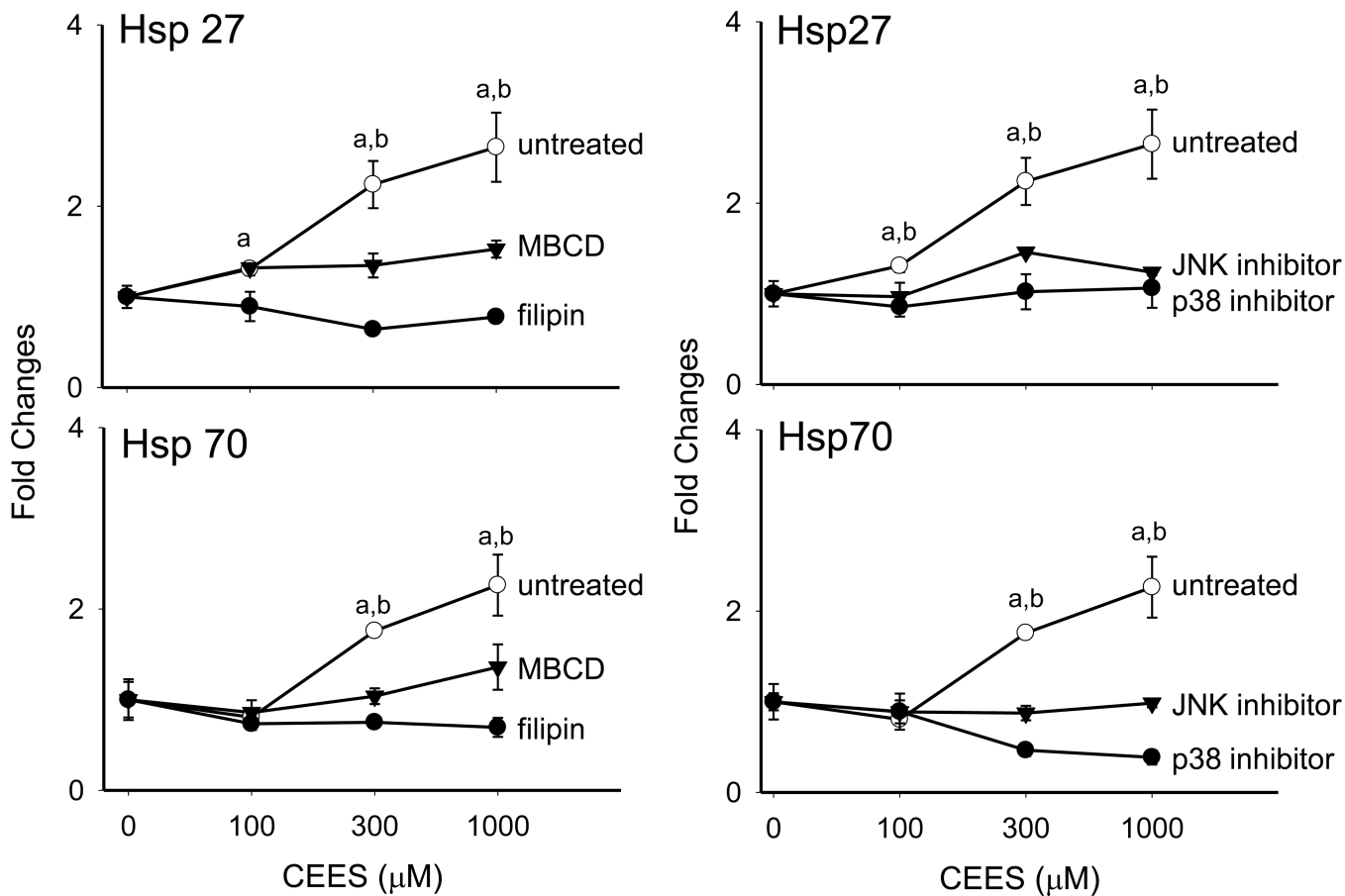


Figure 9. Effects of caveolae and MAP kinase inhibitors on CEES-induced expression of Hsp27 and Hsp70

Mouse keratinocyte skin constructs were preincubated with the caveolae inhibitors, filipin (10 µM), and MBCD (5 mM), or control for 30 min, or JNK (SP600125, 20 µM) and p38 (SB203580, 10 µM) MAP kinase inhibitors or control for 3 hr, and then exposed to CEES (0, 100, 300 or 1000 µM). After 24 hr, mRNA was isolated from the cells and analyzed by real-time PCR. Data are presented as fold change in gene expression relative to untreated cells. *Left panel:* Effects of caveolae inhibition on Hsp27 and Hsp70 expression. ^aSignificantly ($p < 0.05$) different from control (filipin); ^bSignificantly different ($p < 0.05$) from control (MBCD). *Right panel:* Effects of MAP kinase inhibition on Hsp27 and Hsp70 expression. ^aSignificantly ($p < 0.05$) different from control (p38 inhibitor); ^bSignificantly different ($p < 0.05$) from control (JNK inhibitor).

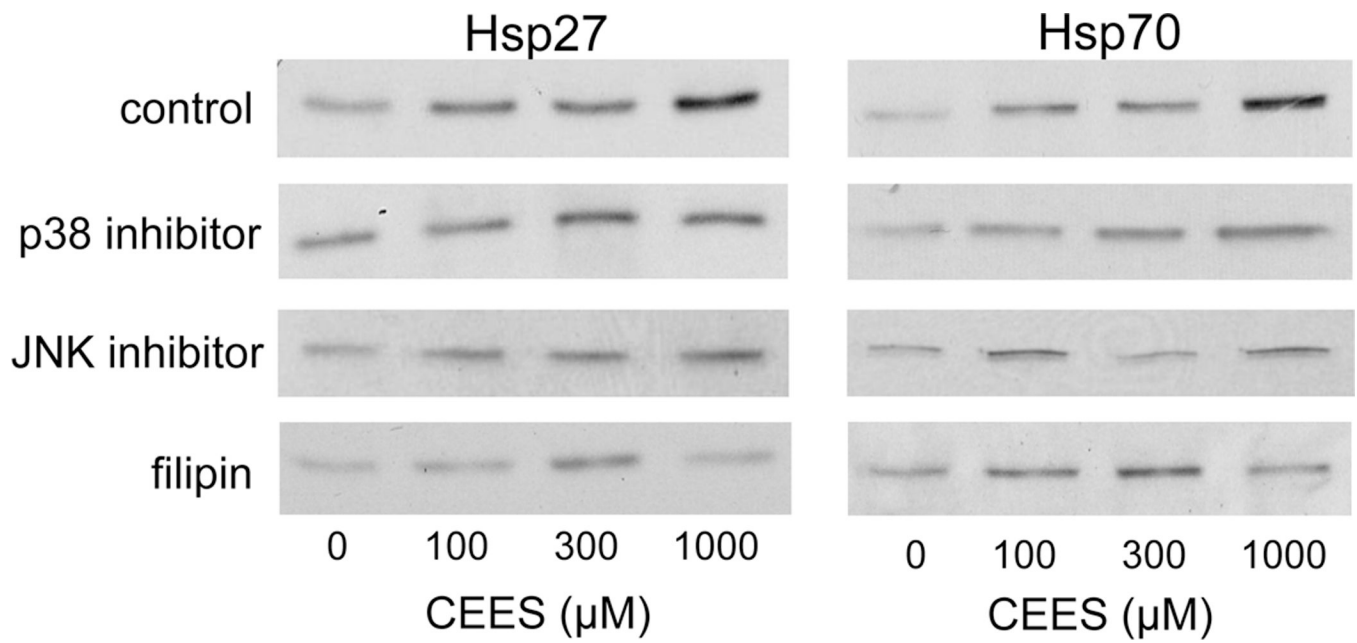


Figure 10. Effects of inhibitors on CEES-induced protein expression of Hsp27 and Hsp70
Mouse keratinocyte skin constructs were preincubated with the caveolae inhibitors, filipin (10 μ M), and MBCD (5 mM), or control for 30 min or JNK (SP600125, 20 μ M), and p38 (SB203580, 10 μ M) MAP kinase inhibitors or control for 3 hr then exposed to CEES (0, 100, 300 or 1000 μ M). After 24 hr total cellular lysates were prepared and Hsp27 and Hsp70 protein expression analyzed by Western blotting.

Table 1

Real-time PCR primer sequences.

Gene	Forward (5'→3')	Reverse (5'→3')
Human		
Caveolin-1	AGACGAGCTGAGCGAGAAGC	TCGATCTCCTTGGTGTGCG
GAPDH	TGGGCTACACTGAGCACCAG	GGGTGTCGCTGTTGAAGTCA
Hsp27	AAGCTAGCCACGCAGTCAA	CGACTCGAAGGTGACTGGGA
Hsp70	ATGTCGGTGGTGGGCATAGA	CACAGCGACGTAGCAGCTCT
Mouse		
Caveolin-1	ACCGTGCATCAAGAGCTTCC	TAGACGCGGCTGATGCACT
GAPDH	TGAAGCAGGCATCTGAGGG	CGAAGGTGGAAGAGTGGGAG
Hsp27	AAGGAAGGCGTGGTGGAGAT	TTCGTCCTGCCTTTCTTCGT
Hsp70	CAGCGAGGCTGACAAGAAGAA	GGAGATGACCTCCTGGCACT