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Heat shock inhibits caspase-1 activity while also preventing its inflammasome-mediated activation by anthrax lethal toxin

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Abstract

Anthrax lethal toxin (LT) rapidly kills macrophages from certain mouse strains in a mechanism dependent on the breakdown of unknown protein(s) by the proteasome, formation of the Nalp1b (NLRP1b) inflammasome, and subsequent activation of caspase-1. We report that heat shocking LT-sensitive macrophages rapidly protects them against cytolysis by inhibiting caspase-1 activation without upstream effect on LT endocytosis or cleavage of the toxin's known cytosolic substrates (MEKs). Heat shock protection against LT occurred through a mechanism independent of de novo protein synthesis, HSP90 activity, p38 activation, or proteasome inhibition and was downstream of MEK cleavage and degradation of an unknown substrate by the proteasome. However, the heat shock inhibition of LT-mediated caspase-1 activation was not specific to the Nalp1b (NLRP1b) inflammasome, as heat shock also inhibited Nalp3 (NLRP3) inflammasome-mediated caspase-1 activation in macrophages. We found that heat shock induced pro-caspase-1 association with a large cellular complex which could prevent its activation. Additionally, while heat shocking recombinant caspase-1 did not affect its activity in vitro, lysates from heat shocked cells completely inhibited recombinant active caspase-1 activity. Our results suggest heat shock inhibition of active caspase-1 can occur independently of inflammasome platform, through a titratable factor present within intact, functioning heat shocked cells.

Introduction

Bacillus anthracis lethal toxin (LT) is composed of protective antigen (PA), a receptor binding protein, and lethal factor (LF), a zinc metalloprotease that is responsible for the cleavage of mitogen-activated protein kinase kinases (MEKs) (Leppla, 2006; Duesbery *et al.*, 1998; Pellizzari *et al.*, 1999 Vitale *et al.*, 2002). Within certain strains of murine macrophages, LT causes a rapid and striking cell death, whereas macrophages from other mouse strains are resistant (Friedlander *et al.*, 1993; Singh *et al.*, 1989). While many of the cellular signals between LT treatment and macrophage death remain to be elucidated, key events required for cell death include the degradation of an unknown protein or proteins by the proteasome and activation of caspase-1 downstream of toxin-induced potassium fluxes (Tang and Leppla,

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1999; Boyden and Dietrich, 2006; Wickliffe *et al.*, 2007; Squires *et al.*, 2007). Macrophage sensitivity has been mapped to variations in the gene encoding Nalp1b (also known as NLRP1b), a Nod-like receptor (NLR) protein involved in innate immunity (Boyden and Dietrich, 2006). Nalp family proteins oligomerize in response to numerous bacterial and 'danger' stimuli including cytosolic flagellin (Miao *et al.*, 2006), gout-associated uric acid crystals (Martinon, *et al.*, 2006) and microbial toxins (for review, see Freche *et al.*, 2007; Lamkanfi *et al.*, 2007; Petrilli *et al.*, 2007) to trigger the formation of an inflammasome, a cytosolic, multi-protein signaling complex. The inflammasome platform promotes pro-caspase-1 autocleavage and activation, an event essential for LT-induced cell death (Boyden and Dietrich, 2006). Activated caspase-1 p10 and p20 subunits form a heterotetramer, which cleaves IL-1 β and IL-18 to their mature forms (Dinarello, 1996; Akita *et al.*, 1997), although this process is not required for LT-induced cell death (Wickliffe *et al.*, 2007). IL-1 β and IL-18 are pro-inflammatory cytokines which can trigger the initiation of fever (Dinarello, 1996; Roth *et al.*, 2006). However, little examination has been made of NLR inflammasome systems once inflammation has begun and cells are subjected to conditions of thermal stress.

The classical heat shock response centers around the up-regulation of heat shock proteins (HSPs). These proteins are involved in a multitude of cellular responses, including serving as chaperones for the re-folding of thermally denatured proteins (for review, see Picard, 2002; Rutherford, 2003; Lepock, 2005). HSP transcription is controlled by heat shock factor-1, a regulatory protein that is required for multi-pathogen resistance in *Caenorhabditis elegans* (McMillan *et al.*, 1998; Singh and Aballay, 2006). The heat shock response may regulate several aspects of innate immunity, as HSPs have been implicated in a variety of molecular models of inflammation and bacterial infection (Tanaka *et al.*, 2007; Tang *et al.*, 2007) and heat-stressed mice and rats were shown to be protected from severe endotoxemia (Hotchkiss *et al.*, 1993; Heidemann *et al.*, 2000). Heat shock has also been linked to differential regulation of signals induced by Toll-like receptors, a group of pathogen-sensing proteins related to NLRs (Zhao *et al.*, 2007; Peng *et al.*, 2006). Therefore, it is not unreasonable to suspect that heat shock may have an impact upon NLR signaling as well.

This study investigates the *in vitro* effects of thermal stress on the killing of murine macrophages by LT. We show that heat shock rapidly halts LT-induced cell death without any impact on toxin uptake or MEK cleavage, by a mechanism independent of novel protein synthesis, p38 activation, HSP90 activity, or proteasome inhibition. Rather, heat shock prevents the activation of pro-caspase-1 in LT-treated cells, apparently by the sequestration of pro-caspase-1 in a large, inhibitory complex. Additionally, heat-shocked cell lysates strongly inhibit the active caspase-1 heterotetramer *in vitro*, independent of a specific inflammasome platform. Our results suggest the presence of a titratable, cellular, heat shock-inducible, caspase-1 inhibiting factor. The potential implications of these findings for the regulation of inflammatory responses are discussed.

Results

Heat shock protects against LT-induced cell death in macrophages

RAW 264.7 cells and bone marrow-derived macrophages (BMDMs) from BALB/cJ mice with LT-sensitive macrophages were heat shocked for 2 h prior to LT treatment and were subsequently protected against cell death (Fig. 1 shows data for RAW 264.7 cells). Similar results were obtained whether the cells were incubated at 37°C (data not shown) or at elevated heat shock temperatures during the period of LT treatment. Heme Oxygenase 1 is a marker for thermal shock (Ewing and Maines, 1995) and its up-regulation was used to monitor the induction of a heat shock state (Fig. 1 inset).

Heat shock does not prevent MEK cleavage by LT

LF cleaves members of the mitogen-activated protein kinase kinase (MEK) family after its delivery into the cytosol by PA (Duesbery *et al.*, 1998; Pellizzari *et al.*, 1999). We hypothesized that heat shock could interfere with the binding, endocytosis or translocation of LF into the cytosol. Therefore, we examined MEK cleavage under heat shock conditions to see if active LF was properly delivered to the cytosol. In RAW 264.7 and BALB/cJ BMDM cells heat shocked for 2 h before LT treatment, MEK1, MEK2, and MEK3 were cleaved similarly in heat shocked cells and those incubated at 37°C (Fig. 2 shows results for RAW 264.7 cells, similar results obtained with BALB/cJ BMDMs), indicating that heat shock did not deactivate LF or prevent its proper translocation into the cytosol.

Synthesis of heat shock proteins is not required for protection

A number of inducible HSPs have been implicated in the activation of both innate and adaptive immunity (Srivastava, 2002). We postulated that the rapid up-regulation of a heat shock-induced protein could be involved in macrophage protection from LT. Therefore, we tested the effect of the transcription inhibitor actinomycin D and of the translation inhibitors puromycin and cycloheximide to see if the protection afforded by heat shock was prevented by the inhibition of protein synthesis. The activity of these inhibitors was confirmed in other experiments (data not shown). We observed strong heat shock protection from LT with all the inhibitors tested over both a range of LT concentrations and drug doses (Fig. 3), suggesting that this protection does not involve the synthesis of a novel protein. Additionally, the inhibitors did not have any protective effect on LT-mediated cytolysis at 37°C (Fig. 3), indicating that transcriptional and translational responses to LT play no role in cell death.

p38 inhibition does not reverse heat shock protection

Previous studies have suggested that the activation of p38 MAP kinase is required for protection from LT-induced cell death in C57BL/6J macrophages (Park *et al.*, 2002). Since p38 activation is a general protective stress response in macrophages which has also been shown to occur in response to thermal stress (Aggeli *et al.*, 2002; Maroni *et al.*, 2002), we hypothesized that our heat shock protection could be attributed to the protective effect of activated p38. We found that p38 was indeed phosphorylated in response to our heat shock conditions in both RAW 264.7 cells and BALB/cJ macrophages, and that this activation was lost after 60 min of LT treatment (Fig. 4A). LF cleavage of MEK3 (Fig. 2), the upstream activator of p38, inhibits p38 activation at these later LT treatment times. To test if early p38 activation under heat shock was responsible for protection against LT we inhibited p38 activity using the inhibitor SB203580. Cells pre-treated with this inhibitor were then heat shocked or incubated at 37°C prior to LT treatment. The continued effectiveness of the SB203580 inhibitor under heat shock and 37°C conditions was confirmed by examining the LPS-induced phosphorylation of mitogen-activated protein kinase-activated protein-2 kinase (MAPKAP-2), a direct target of phosphorylated p38. Using an antibody that detects altered migration of this protein in its activated form, we found that MAPKAP-2 was not phosphorylated in cells treated with p38 inhibitor in either 37°C or heat shock conditions (Fig. 4B inset). MAPKAP-2 was not activated at 37°C without LPS treatment (data not shown). Thus, the p38 inhibitor was effective at both 37°C and heat shock conditions, but was unable to reverse the heat shock protection against LT (Fig. 4B). We conclude that p38 activation is not required for heat shock protection. Similar results were obtained with BALB/cJ BMDM cells (data not shown).

HSP90 inhibition does not reverse heat shock protection

HSP90 is a constitutively expressed and highly abundant molecular chaperone for a large number of signal transduction proteins in eukaryotic cells, including those involved in the inhibition of apoptosis (Picard, 2002; Bishop *et al.*, 2007). This protein has also been implicated

in the translocation of some bacterial toxins from cellular endosomes, but does not play a role in the translocation of LT (Ratts *et al.*, 2003; Haug *et al.*, 2003). Additionally, HSP90 inhibition does not prevent macrophage death via the LT-induced Nalp1b (NLRP1b) inflammasome (Haug *et al.*, 2003; data not shown), contrary to the HSP90 requirement that was previously shown in the Nalp3 (NLRP3) inflammasome system (Mayor *et al.*, 2007). We tested whether HSP90 activity was required for heat shock protection from LT using the potent HSP90 inhibitor geldanamycin. The activity of this inhibitor was confirmed in other experiments (data not shown). Heat shocked cells continued to be protected from LT with a range of inhibitory geldanamycin doses (Fig. 5), showing that HSP90 is not involved in heat shock protection against the toxin.

Heat shock prevents LT-induced activation of caspase-1 in cells

Caspase-1 activation by the Nalp1b (NLRP1b) inflammasome is known to be required for LT-induced macrophage cell death (Boyden and Dietrich, 2006). To determine whether heat shock affected events upstream or downstream of caspase-1 activation, we compared the levels of the activated p10 subunit of caspase-1 in LPS-primed BALB/cJ BMDMs that were treated with LT under heat shock or 37°C conditions. In contrast to cells at 37°C, no active p10 caspase-1 was observed in the heat shocked cells following LT treatment (Fig. 6A). IL-1 β cleavage by caspase-1 was also monitored in the same cells and no mature IL-1 β was observed in heat shocked samples (Fig. 6B), confirming the absence of activated caspase-1 under heat shock conditions. Identical caspase-1 cleavage results were also obtained from BALB/cJ BMDMs that were not LPS-primed (data not shown).

Late heat shock protects against LT

We have previously established a timeline of the known cellular events involved in LT-induced cell death in murine macrophages (Wickliffe *et al.*, 2007). We next asked how rapidly and how late heat shock could protect cells from LT. BALB/cJ BMDMs were given LT and then heat shocked immediately (at t=0 min) or moved to heat shock conditions after various durations of LT incubation at 37°C (between t=45 min and t=75 min). After t=110 min, viability was assessed. We determined the amount of activated caspase-1 at the time when cells were moved to heat shock by lysing a parallel set of identically treated samples at every shift time point at 37°C. Additionally, we assessed the levels of continued caspase activation in a third set of similarly treated samples after the move to heat shock.

We observed that no heat shock incubation before LT treatment was necessary for heat shock protection against the toxin. In fact, both BMDMs and RAW 264.7 cells could be heat shocked well after LT addition and be protected from LT-induced death, especially if heat shock was applied before the active p10 subunit was first observed (Fig. 7B shows results for BALB/cJ BMDMs). In parallel, we determined that heat shock applied before the initiation of caspase-1 activation (i.e. at t=0 min or t=45 min) completely prevented subsequent caspase-1 activation (Fig. 7A, compare 1st and 2nd lanes with 9th and 8th lanes). However, heat shock applied after p10 was observed (i.e. at t=65 min or 75 min) resulted in levels of active p10 comparable to that found in cells that were not heat shocked (Fig. 7A, compare 6th and 7th lanes to 5th lane). These results suggest that heat shock prevents caspase-1 activation by the Nalp1b (NLRP1b) inflammasome, but once caspase-1 activation is initiated, heat shock is ineffective. In parallel, heat shock applied as late as 65 min and 75 min does appear to slow down the death event (Fig. 7B), but is unable to provide complete protection once some caspase-1 has been activated. This may be because at late times the cell death-inducing events downstream of caspase-1 have already occurred.

Heat shock does not inhibit proteasome activity

We and others have previously demonstrated that proteasome activity upstream of caspase-1 activation is necessary for LT-induced cell death, and that proteasome inhibition provides protection from LT (Wickliffe *et al.*, 2007; Squires *et al.*, 2007). As proteasome inhibition has sometimes been associated with the induction of a cellular heat shock response (Bush *et al.*, 1997; Kim *et al.*, 1999; Lee and Goldberg, 1998) and, conversely, heat shock has been associated with differential proteasomal processing (Pajonk *et al.*, 2005; Kuckelkorn *et al.*, 2000; Kelly *et al.*, 2007; Luo *et al.*, 2000), we asked whether our observed heat shock protection acted through proteasome inhibition. We first compared the levels of ubiquitinated proteins in LT-treated or untreated macrophages at 37°C, in the presence of the proteasome inhibitor lactacystin, or under conditions of heat shock. A LT-mediated decrease in ubiquitinated proteins in macrophages has been previously reported (Alileche *et al.*, 2006; Salles *et al.*, 2003), although we observed no difference in ubiquitination when cells were treated with LT (Fig. 8A), possibly because previous studies had examined cells at late times, after lysis from LT had already occurred. As expected, there was a large build-up of ubiquitinated proteins in cells treated with lactacystin (Fig. 8A). We also observed a moderate increase in ubiquitinated proteins in the heat shocked cells compared to 37°C cells, suggestive of either inhibition of the proteasome under heat shock or an increase in protein ubiquitination in heat shocked cells, with potentially no change in proteasome function. Both of these phenomena have been previously reported (Pajonk *et al.*, 2005; Parag *et al.*, 1987). To distinguish between these possibilities, we investigated proteasome activity in cells directly using a cell-permeable proteasome substrate activated to luminesce only after cleavage by the proteasome. Lactacystin was used as a positive control for proteasome inhibition in these studies. We found that the proteasome was not inhibited in heat shocked cells, and that proteasomal processing was actually markedly increased under those conditions (Fig. 8B), much in the manner previously shown for the ER unfolded protein stress response to heat shock (Kelly *et al.*, 2007). The apparent increase in cleavage of the proteasome substrate under heat shock conditions may have been due to changes in its uptake and delivery. In both heat shocked and 37°C cells, the lactacystin control treatment resulted in a significant decrease in luminescence as was expected from proteasome inhibition. From these data, we conclude that heat shock does not inhibit proteasome activity, and therefore that heat shock protection from LT is not a result of proteasome inhibition.

Heat shock prevents nigericin-induced caspase-1 activation by the Nalp3 inflammasome

We next sought to determine if heat shock can prevent the activation of caspase-1 by other inflammasome protein complexes. Nigericin has been shown to rapidly induce the Nalp3 (NLRP3) inflammasome in BMDMs, resulting in the activation of caspase-1 and release of IL-1 β (Mariathasan *et al.*, 2006). Because nigericin also requires an LPS co-signal and TLR4 signaling is affected by heat shock conditions (Data not shown; Zhao *et al.*, 2006), BALB/cJ BMDM cells were LPS pre-treated at 37°C prior to heat shock incubation and subsequent nigericin treatment. There was a distinct absence of active caspase-1 and mature IL-1 β in the heat shocked, nigericin-treated cell lysates (Fig. 9). This result could suggest that heat shock can prevent the activation by the Nalp3 (NLRP3) inflammasome and may affect a component necessary for the induction of multiple inflammasome complexes. However, as the precise timing and mechanism of LPS/nigericin activation of the Nalp3 (NLRP3) inflammasome is poorly understood, it is also possible that the lack of activated caspase-1 was due to altered TLR signaling in heat shock conditions, as reflected by heat shock's inhibitory effect on LPS-mediated I κ B- α breakdown (data not shown).

Whole cell lysates from heat shocked cells inhibit caspase-1 activity

We tested the hypothesis that higher temperatures could alter the proteolytic activity of caspase-1. Sucrose lysates of LPS-treated RAW 264.7 cells which contained highly up-regulated, uncleaved IL-1 β were treated for 1 h with recombinant caspase-1 at 37°C or under heat shock. Samples were then analyzed for IL-1 β cleavage. We found that caspase-1 effectively cleaved IL-1 β to its mature form in both heat shock and 37°C conditions (Fig. 10A), even when the enzyme and substrate were pre-heat shocked for extended periods before mixing (data not shown). This implies that a high temperature does not interfere directly with the proteolytic activity of activated caspase-1.

We next asked whether a component in heat shocked, intact cells was necessary for inhibition of caspase-1 activity. In this scenario, heat shock could initiate a signaling event, relocalization, protein modification or other event requiring whole, live cells, which could alter the proteolytic activity of caspase-1. To this end, we heat shocked LPS pre-treated RAW 264.7 cells for 1 h before making sucrose lysates. These lysates were then treated with recombinant active caspase-1 at 37°C for 1 h and examined for IL-1 β cleavage (Fig. 10B). We observed a complete absence of IL-1 β and IL-18 (data not shown) processing within lysates from heat shocked cells, even though we had added recombinant, active caspase-1. To address the hypothesis that the recombinant caspase-1 was degraded in these lysates, we examined the levels of caspase-1 within the same samples (Fig. 10C). We observed that a clear caspase-1 p10 band was present in all samples receiving the recombinant active caspase-1 (Fig. 10C), suggesting that the lack of IL-1 β cleavage was not due to caspase-1 breakdown. No protein synthesis is required for this recombinant caspase-1 inhibition, as identical cleavage patterns were observed when cells were treated with translation inhibitors puromycin and cycloheximide before heat shock (data not shown). As this inhibition was independent of the Nalp1b inflammasome platform, we were not surprised to find identical patterns of inhibition when using heat shock lysates from LT-resistant macrophages (data not shown).

The titratable nature of the heat shock-induced inhibitory activity was shown by combining lysates from 37°C and heat shocked cells in different ratios before incubation with the recombinant caspase-1 (Fig 11A), suggesting the presence of a factor that non-enzymatically binds or modifies caspase-1. We hypothesized that this factor could be a protein or protein complex that binds and inactivates caspase-1. Therefore, we examined recombinant caspase-1 treated with heat shock or 37°C lysates by native gel in order to observe any differential migration of protein complexes. The diffuse band representing recombinant caspase-1 migrated similarly in the 37°C and heat shock samples (Fig. 11B, upper). It is possible that the recombinant caspase-1 heterotetramer in heat shock lysates is inhibited by a modification or binding event that is too small to be detected on these gels.

We also observed that while endogenous pro-caspase-1 was present in both of the lanes with 37°C lysates on a native gel, independent of the addition of recombinant active caspase-1, there was no observable endogenous pro-caspase-1 band in the heat shock lysates (Fig. 11B, upper). However, heat shock samples clearly showed the presence of pro-caspase-1 when subjected to SDS electrophoresis (Fig. 11B, lower). This suggests that endogenous pro-caspase-1 is not broken down, but may be part of a large protein complex in heat shocked cells.

Centrifugation of native lysates showed that although pro-caspase-1 was found in both the pellet and supernatant of lysates made from cells at 37°C, the lysates from heat shocked cells contained pro-caspase-1 almost entirely in the pellet fraction alone, confirming that pro-caspase-1 was part of a larger complex (Fig. 11C). Thus, it appears that heat shock induces the binding of endogenous pro-caspase-1 to other cellular protein/s which may prevent caspase-1 activation by inflammasomes. Additionally, when we tested whether heating 37°C lysates could relocalize pro-caspase-1 to this larger complex we found no difference between heated

and unheated samples (data not shown). This indicated that the sequestration of pro-caspase-1 by heat shock also requires an event within intact cells, as observed for inhibition of active recombinant caspase-1.

Discussion

The general phenomenon of heat shock has been widely studied and documented, but its molecular and cellular effects on immunity are only beginning to be explored. Some studies have shown heat shock conditions to protect from endotoxemia *in vivo* (Hotchkiss *et al.*, 1993; Heidemann *et al.*, 2000) and to affect macrophage and dendritic cell activation (for review, see Ostberg and Repasky, 2006; Srivastava, 2002), TLR signaling (Zhao *et al.*, 2006), cytokine secretion (Peng *et al.*, 2006; Tanaka *et al.*, 2007), and T-cell responses to MHC-presented antigens (Polla *et al.*, 2006; Blachere *et al.*, 1997). To date, there is little published data on the effects of heat shock on NLR signaling, inflammasome formation, or caspase-1 activity.

In this study, we found that heat shock can rapidly prevent anthrax LT-induced macrophage death by inhibiting caspase-1 activation, an event known to be required for cell death from LT (Boyden and Dietrich, 2006; Cordoba-Rodriguez *et al.*, 2004). This inhibition of caspase-1 activation was also observed for the nigericin-induced Nalp3 (NLRP3) inflammasome, and may be due to binding of pro-caspase-1 to a large, inhibitory complex. Independently, heat shock was found to induce a titratable cellular factor, signaling, or modification event that inhibits the active caspase-1 heterotetramer without binding to a large complex. This factor could also be a general cellular condition, such as an alteration of the redox state of the cell. Previous studies (Cruz *et al.*, 2007; Dostert *et al.*, 2008) as well as a recent paper have implicated reactive oxygen species in the regulation of caspase-1 activity (Meissner *et al.*, 2008). Experiments are currently underway to assess the redox state of cells following heat shock as well as possible modifications of caspase-1 in this context. Previous work has provided preliminary evidence that thermal shock can alter the redox state of cells (Jackson *et al.*, 2006; Murapa *et al.*, 2007)

Although the majority of studies of the heat shock response focus on HSP synthesis and activity, we found that *de novo* production of proteins was not important to heat shock protection from LT or for LT-mediated cell death. Other events induced by heat shock and previously shown to induce protective cellular responses, including p38 activation and the recruitment of HSP90, were also not involved in heat shock protection from LT. The breakdown of one or more cellular proteins by the proteasome is a specific requirement for the formation of the LT-induced Nalp1b (NLRP1b) inflammasome (Wickliffe *et al.*, 2007; Squires *et al.*, 2007). Previous studies have suggested that heat shock can either increase (Luo *et al.*, 2000; Meinander *et al.*, 2007; Smith *et al.*, 2005) or decrease (Kuckelkorn *et al.*, 2000; Pajonk *et al.*, 2005) proteasomal degradation in mammalian cells. We found that proteasome activity was increased in macrophages after heat shock and therefore heat shock protection from LT was not a result of general proteasome inhibition.

When we extended our study of heat shock to the better characterized Nalp3 (NLRP3) inflammasome, we observed that caspase-1 and IL-1 β were strongly activated in response to nigericin at 37°C but not in heat shock conditions. This was suggestive of an inhibition of pro-caspase-1 activation within the Nalp3 pathway, similar to that observed for the Nalp1b inflammasome. However, it was uncertain if this inhibition was due to heat shock interference with the LPS co-signal required for Nalp3 inflammasome activation.

Our *in vitro* studies show that recombinant caspase-1 activity is directly inhibited by a titratable factor in heat shocked cell lysates, suggesting that heat shock can also inhibit activated

caspase-1, independent of the NLR inflammasome system of activation. We therefore expect our findings to apply to all inflammasome systems in addition to other caspase-1-related events in murine macrophages. Further studies are needed to see if a similar inhibitory activity also exists in human cells.

The precise mechanism of caspase-1 inhibition under heat shock remains to be determined. The events involved may include protein modification events (such as phosphorylation), altered ion fluxes, the production of reactive oxygen species, or the relocalization or altered conformation of a protein, among other possibilities. We hypothesize that within the intact cell system heat shock mediated events alter pro-caspase-1 binding in a complex which prevents its activation by the inflammasome. Separately, heat shock mediated factors can also inhibit the active caspase-1 heterotetramer independent of any effects on pro-caspase-1. However, an intact cellular structure is required for the cellular modifications or signaling events leading to heat shock inhibition of the caspase-1 heterotetramer, as heating the bulk cell lysate components does not inhibit caspase-1 activity.

Caspase-1 inactivation by heat shock could represent a regulatory pathway for the IL-1 β inflammatory cascade. If a similar inhibitory effect is also shown for the activation of numerous other inflammasome systems that lead to the IL-1 β response (see Lamkanfi *et al.*, 2007 for review), thermal heat shock may represent a general negative feedback mechanism for regulating IL-1 β mediated inflammation, possibly extending to the febrile response induced by this cytokine *in vivo*. Although there have been previous studies that have remarked on the *in vitro* inhibition of IL-1 β production in heat shock conditions (Kappel *et al.*, 1991; Velasco *et al.*, 1990; Boneberg and Hartung, 2003), no study has linked this phenomenon to an inhibition of caspase-1 activity and to date, no correlation between *in vitro* heat shock regulation and *in vivo* events has been found. *In vivo* studies in LT-treated mice have shown a unique and rapid IL-1 β release seen only in mice with LT-sensitive macrophages (Moayeri *et al.*, 2004), which is rapidly shut down before any indications of malaise in animals. We are currently investigating the possibility of a temperature-based shutdown of the IL-1 β response in these animals.

In summary, we have discovered that heat shock inhibits the activation of pro-caspase-1 by the Nalp1b (NLRP1b) and Nalp3 (NLRP3) inflammasomes while also directly inhibiting activated caspase-1 independent of any inflammasome platform. The identification of the cellular events and proteins involved in the shutdown of the caspase-1 pathway will have substantial ramifications in the study of the continually expanding set of inflammatory conditions which are known to act through caspase-1 (for review, see Wilmanski *et al.*, 2008).

Experimental Procedures

Materials

Protective antigen (PA) and lethal factor (LF) were purified in our laboratory as described previously (Park and Leppla, 2000; Ramirez *et al.*, 2002; Varughese *et al.*, 1998). Concentrations of LT given for each experiment correspond to the concentration of each toxin component (i.e. 1 $\mu\text{g mL}^{-1}$ LT has 1 $\mu\text{g mL}^{-1}$ PA and 1 $\mu\text{g mL}^{-1}$ LF). Anti-MEK1 NT (444942) antibody, ultra-pure LPS, and nigericin were purchased from Calbiochem (San Diego, CA). Caspase-1 p10 (sc-514), Mek2 NT (sc-524), Mek 3 NT (sc-959), p38 CT (sc-535-G), and Heme Oxygenase 1 (HOX) (sc-10789) antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Phospho-p38 (4631) antibody was from Cell Signaling Technology (Danvers, MA). Anti-ubiquitin (Z0458) antibody was from Dako (Glostrup, Denmark). Anti-IL-1 β (AF-401-NA) was from R&D Systems (Minneapolis, MN). Anti-goat infrared dye secondary antibody (#605-731-125) was from Rockland (Gilbertsville, PA). Anti-rabbit infrared dye secondary

antibody was from Licor Biosciences (Lincoln, NE) or from Rockland Immunochemicals Inc. (Gilbertsville, PA). The p38 inhibitor SB203580 was purchased from Invitrogen (Carlsbad, CA). Puromycin, cycloheximide and LPS were obtained from Sigma-Aldrich (St. Louis, MO). Geldanamycin was purchased from Stressgen (San Diego, CA). Active recombinant caspase-1 was purchased from MBL International (Woburn, MA). Proteasome-Glo Cell-Based Assay was purchased from Promega (Madison, WI) and used as directed by manufacturer protocol with slight modification. Briefly, after heat shock and lactacystin treatments, all cells were equilibrated to room temperature before a 1:1 addition of the Proteasome-Glo substrate. Cells were shaken and allowed to equilibrate for 10 min before luminescence was measured.

Cell Culture

RAW 264.7 and L929 mouse fibroblast cells were grown in Dulbecco's modified Eagle medium (DMEM) supplemented with 10% fetal bovine serum, 10 mM Hepes, and 50 $\mu\text{g mL}^{-1}$ gentamicin (all obtained from Invitrogen, Carlsbad, CA) at 37°C in 5% CO₂. Bone marrow-derived macrophages (BMDMs) were cultured from BALB/cJ mice (Jackson Laboratories, Bar Harbor, ME) in complete DMEM (as described above) with 30% L929 cell culture supernatant. Bone marrow cells were grown for 7-9 days to allow time for differentiation before use in assays. All experiments involving animals were performed under protocols approved by the Animal Care and Use Committee of the National Institute of Allergy and Infectious Diseases, National Institutes of Health.

Heat shock

Cells were heat shocked in a cell culture incubator at 5% CO₂. Because of incubator variability, heat shock temperatures ranged from 41C-45C for experiments performed on different days.

Cytotoxicity assays

RAW 264.7 and BALB/cJ BMDM cells were grown in 96-well plates to 90% confluence. In assays involving transcription, translation, and HSP90 inhibitors, cells were pre-treated for 1-2.5 h at 37°C with a dose range of each inhibitor as follows: Actinomycin D (0-6250 ng mL⁻¹); Cycloheximide (0-625 ng mL⁻¹); Puromycin (0-3125 ng mL⁻¹); and Geldanamycin (0-10,000 nM). All cells were then treated with 1 $\mu\text{g mL}^{-1}$ LT for 2.5 h and viability was assessed by addition of 0.5 mg mL⁻¹ MTT [3-(4,5-Dimethyl-2-Thiazolyl)-2,5-Diphenyl Tetrazolium Bromide] (USB Corporation, Cleveland, OH). Following 45 min of incubation with MTT dye, cell culture medium was removed and cells were dissolved with 100 μL /well of 0.5% SDS, 25 mM HCl in 90% isopropanol. A microplate reader was used to read the A₅₇₀ of each well, and percent viabilities were calculated relative to untreated controls. In all other assays, cells were treated with a dose range of LT (0-1000 ng mL⁻¹), incubated for 2-3 h (as described in the figure legends) and stained with MTT as described above.

MEK, caspase-1, and IL-1 β cleavage, HOX-1 up-regulation, p38 and MAPKAP-2 phosphorylation, and protein ubiquitination assays

RAW 264.7 and BALB/cJ BMDM cells grown in supplemented DMEM at indicated temperatures were treated with 1 $\mu\text{g mL}^{-1}$ LT for various times. Where indicated, cells were first primed with 500 ng mL⁻¹ LPS prior to toxin addition. At the end of each experiment, cells were washed three times with ice-cold PBS before lysates were prepared with RIPA lysis buffer (1% Nonidet-P40, 0.5% sodium deoxycholate, and 0.1% SDS in PBS) containing EDTA-free Complete protease inhibitor cocktail (Roche Diagnostics, Indianapolis, IN). Lysate protein concentrations were quantified using a BCA protein assay (Pierce, Rockford, IL), and equal amounts of protein were loaded for SDS-PAGE. Western blotting was performed using anti-MEK1 NT (1:7500), anti-MEK2 NT (1:2000), anti-MEK3 NT (1:2000), anti-Heme Oxygenase 1 (1:500), anti-phospho-p38 (1:1000), anti-p38 CT (1:1000), anti-ubiquitin (1:2000), anti-

caspase-1 (1:250), and anti-IL-1 β (1:700). Primary antibodies were detected using IR-dye-conjugated secondary antibodies (Licor Biosciences anti-rabbit, 1:25,000; Rockland Immunochemicals Inc. anti-rabbit, 1:5000; Rockland Immunochemicals Inc. anti-goat, 1:15,000) and the Odyssey Infrared Imaging System (Licor Biosciences, Lincoln, NE).

***In vitro* caspase-1 and IL-1 β assays**

RAW 264.7 cells were grown in supplemented DMEM and treated with ultra-pure LPS (1 μ g mL⁻¹) for 2 h. After heat shock incubations, cells were washed three times with ice-cold PBS before preparing sucrose lysates (250 mM sucrose, 3 mM imidazole). Lysates were passed through a 29G needle 30X prior to protein quantification by BCA assay. Lysates were combined with recombinant caspase-1 (1 unit/40 μ L, MBL) in 3 mM NaCl and incubated for 1 h before SDS or native electrophoresis. Native gel electrophoresis loading dye was 80 mM Tris, pH 7.5, 10% glycerol, and 0.02% bromophenol blue, with or without 5% β -mercaptoethanol. Western blotting was performed using anti-caspase-1 (1:250), and anti-IL-1 β (1:700). Odyssey Infrared Imaging System (Licor Biosciences, Lincoln, NE) and accompanying application software (version 2.0) were used to integrate band intensity for a semi-quantitative measure of protein abundance.

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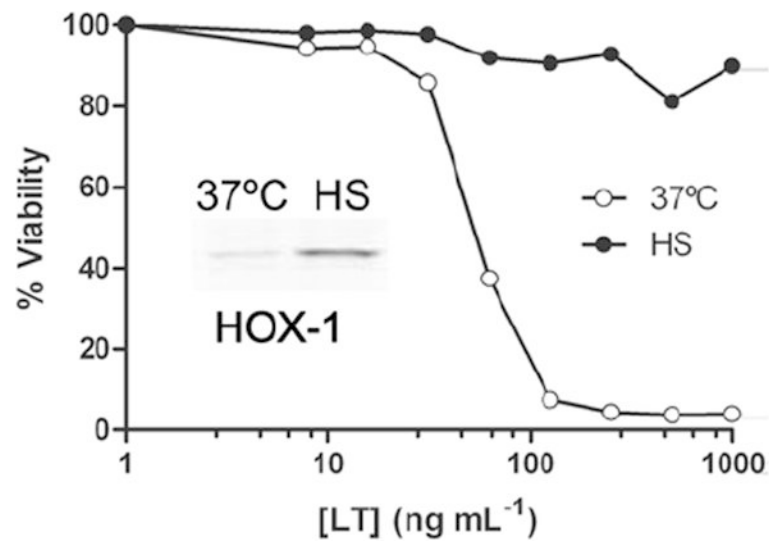


Figure 1. Heat shock protects against LT-induced macrophage death

RAW 264.7 cells were treated with LT after 2 h of heat shock and compared to non-heat shocked toxin-treated controls. Cell viability was assessed after 2.5 h of toxin treatment and percent viability was calculated relative to medium-treated cells. Heat shock induction of Heme Oxygenase-1 in cells receiving similar treatment was used to verify effective heat shock (inset).

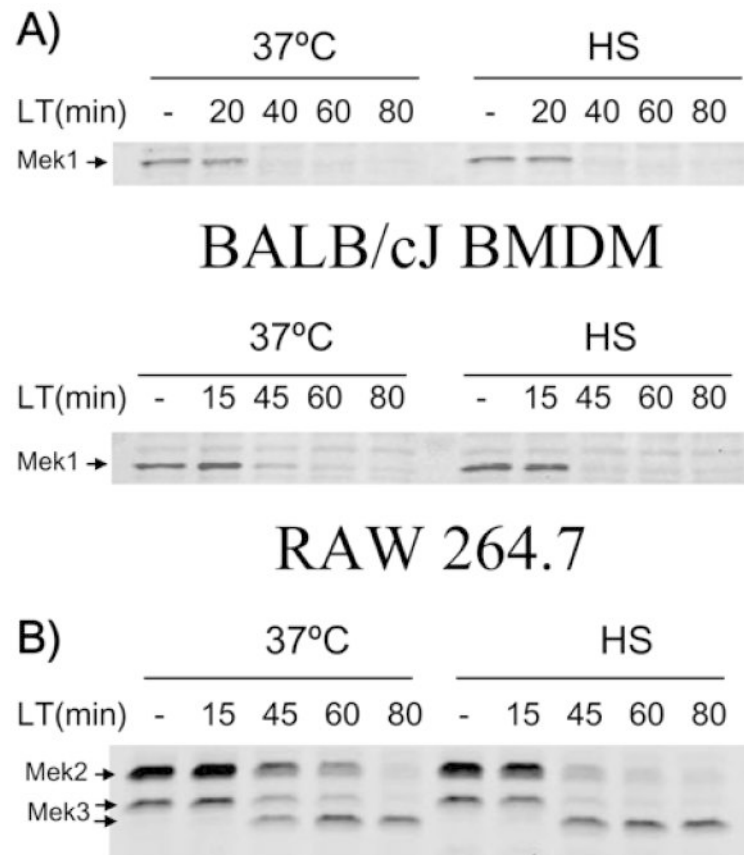


Figure 2. Heat shock does not prevent Mek cleavage by LT

BALB/cJ BMDM and RAW 264.7 cells were heat shocked 2 h prior to LT addition ($1 \mu\text{g mL}^{-1}$ for 0 to 80 min) and compared to toxin-treated cells at 37°C . Cleavage of Mek1 (A), Mek2, and Mek3 (B, RAW 264.7 only) was monitored at various times after toxin treatment in heat shock or 37°C conditions by Western blot.

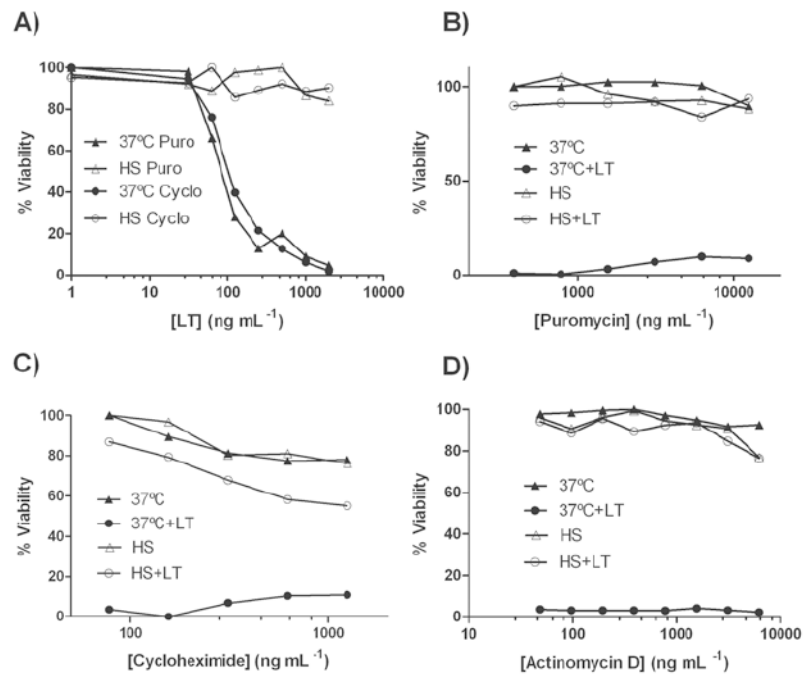


Figure 3. Protein synthesis is not required for protection

RAW 264.7 cells were treated with the translation inhibitors cycloheximide (312 ng mL⁻¹ in (A) or concentration range in (C)) or puromycin (1.56 μg mL⁻¹ in (A) or concentration range in (B)) for 1 h before a 2 h heat shock treatment. LT was added in a concentration range (A) or at 1 μg mL⁻¹ (B, C) and viability was assessed after 2 h. In (D), cells were incubated with the transcription inhibitor Actinomycin D (1 μg mL⁻¹) for 1 h at 37°C prior to continued incubation at this temperature or a 2 h heat shock. LT was added and viability assessed as above.

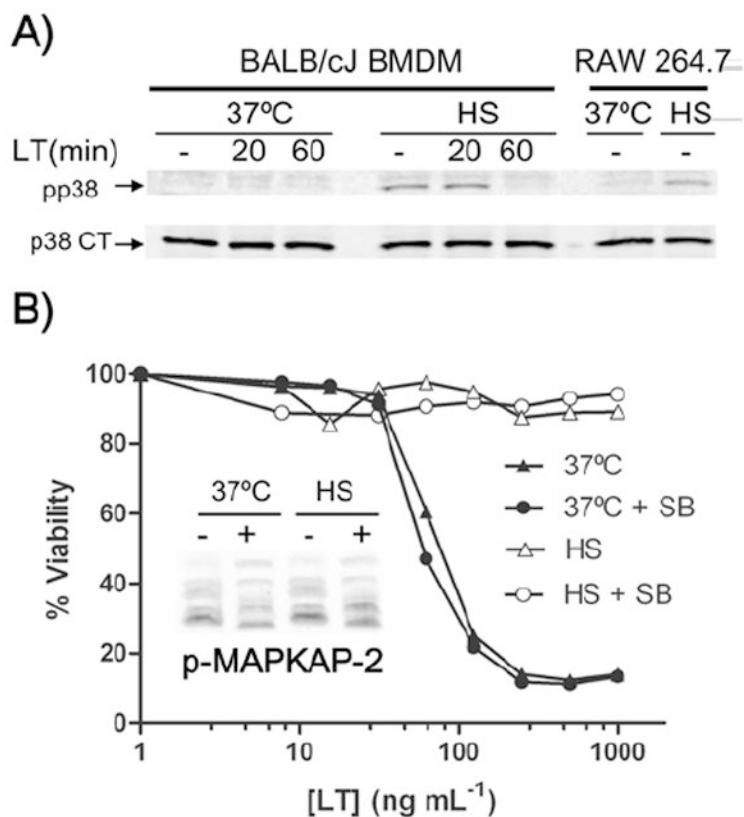


Figure 4. p38 inhibition does not reverse heat shock protection

(A) BALB/cJ BMDM and RAW 264.7 cells were heat shocked (2 h) before LT treatment ($1 \mu\text{g mL}^{-1}$) and compared to non-heat shocked controls. Phosphorylated p38 (upper) and p38 CT (lower) were monitored at various times after toxin treatment. (B) RAW 264.7 cells were incubated for 30 min with SB203580 p38 inhibitor (SB, $26 \mu\text{M}$) or media alone (control) at 37°C before a 1 h heat shock incubation. LT was added and cell viability was assessed after 2 h. Phosphorylation of MAPKAP-2 was monitored in cells stimulated with 500 ng mL^{-1} LPS for 15 min and treated with (+) and without (-) SB203580 as described above to verify effective p38 inhibition (inset).

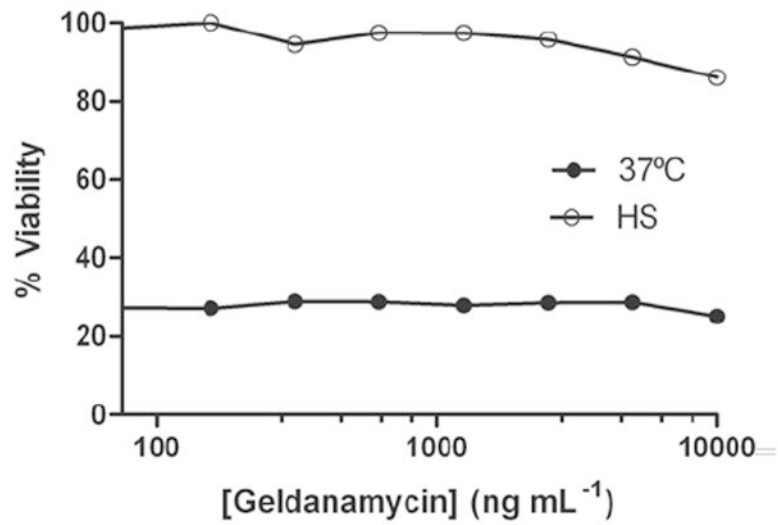


Figure 5. HSP90 inhibition does not reverse heat shock protection

RAW 264.7 cells were incubated with geldanamycin at 37°C for 5 h before being heat shocked or incubated at 37°C for 2 h. LT was added (1 $\mu\text{g mL}^{-1}$) and cell viability was assessed after 2 h.

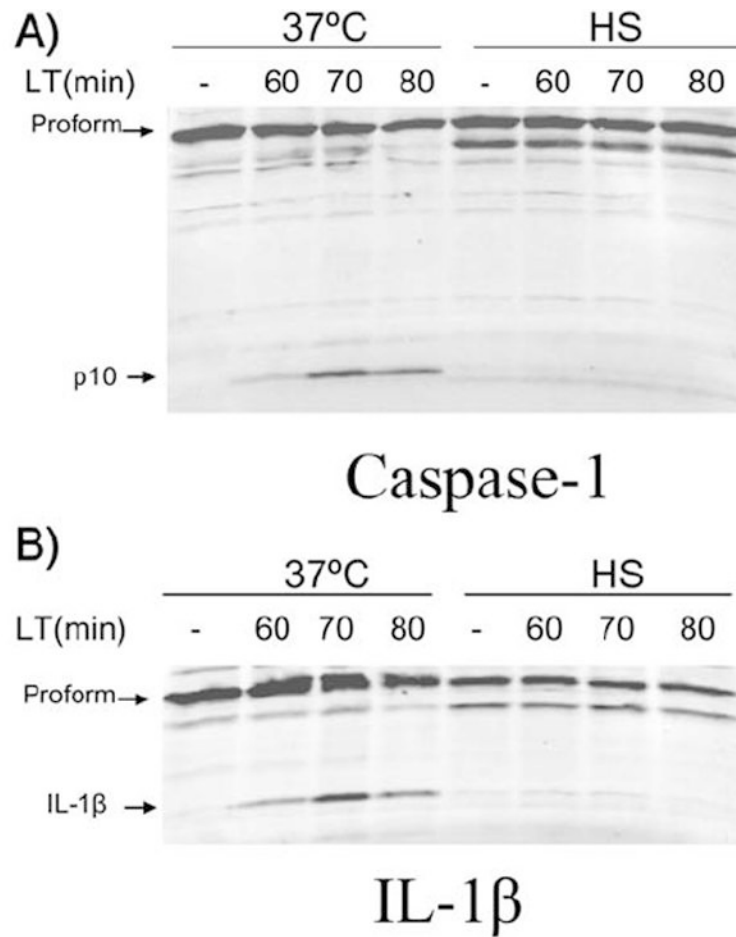


Figure 6. Heat shock prevents caspase-1 activation by LT, but does not inhibit caspase-1 activity
 LPS-primed BALB/cJ BMDM (500 ng mL^{-1}) were heat shocked (2 h) before LT treatment ($1 \text{ } \mu\text{g mL}^{-1}$) and compared to non-heat shock controls remaining at 37°C . Cell lysates were probed for caspase-1 p10 (A) or IL-1 β (B) by Western blot.

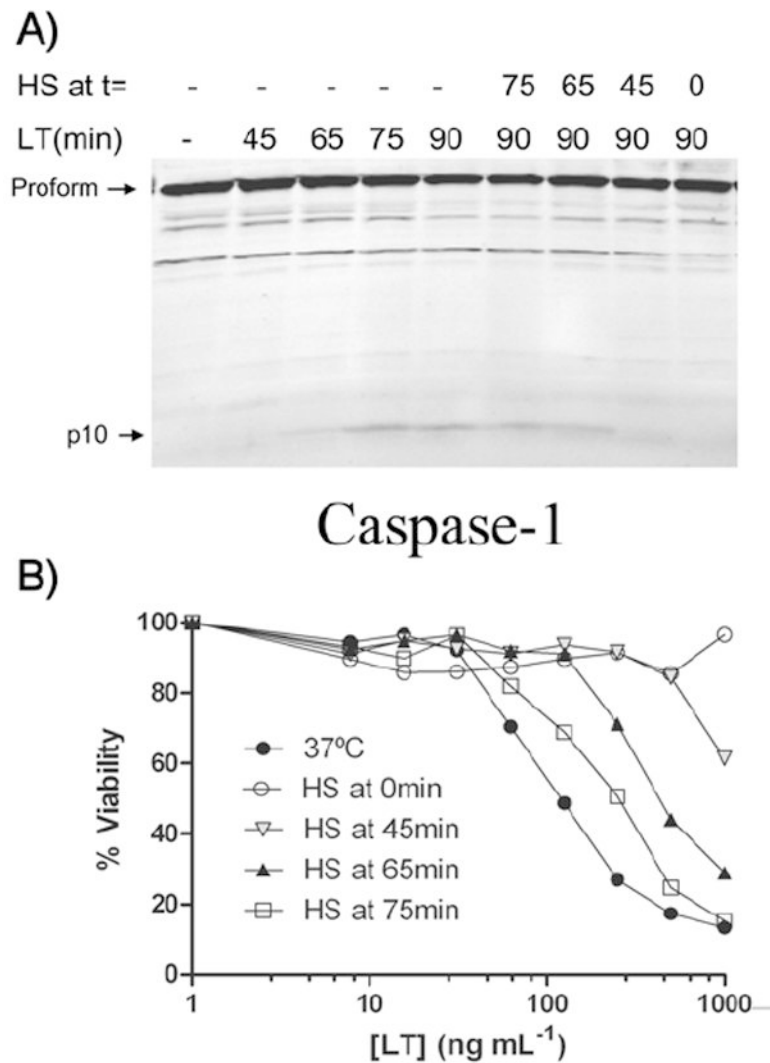


Figure 7. Late heat shock protects against LT α

(A) At t=0 min, LT ($1 \mu\text{g mL}^{-1}$) was added to BALB/cJ BMDMs. Cells were moved to heat shock conditions between 0 and 75 min later, and lysates were made after 0 to 90 min of LT treatment to monitor activated caspase-1 p10. (B) BALB/cJ BMDMs given identical heat shock treatments in parallel to the samples in (A) and a range of LT doses were assessed for viability 110 min after LT treatment.

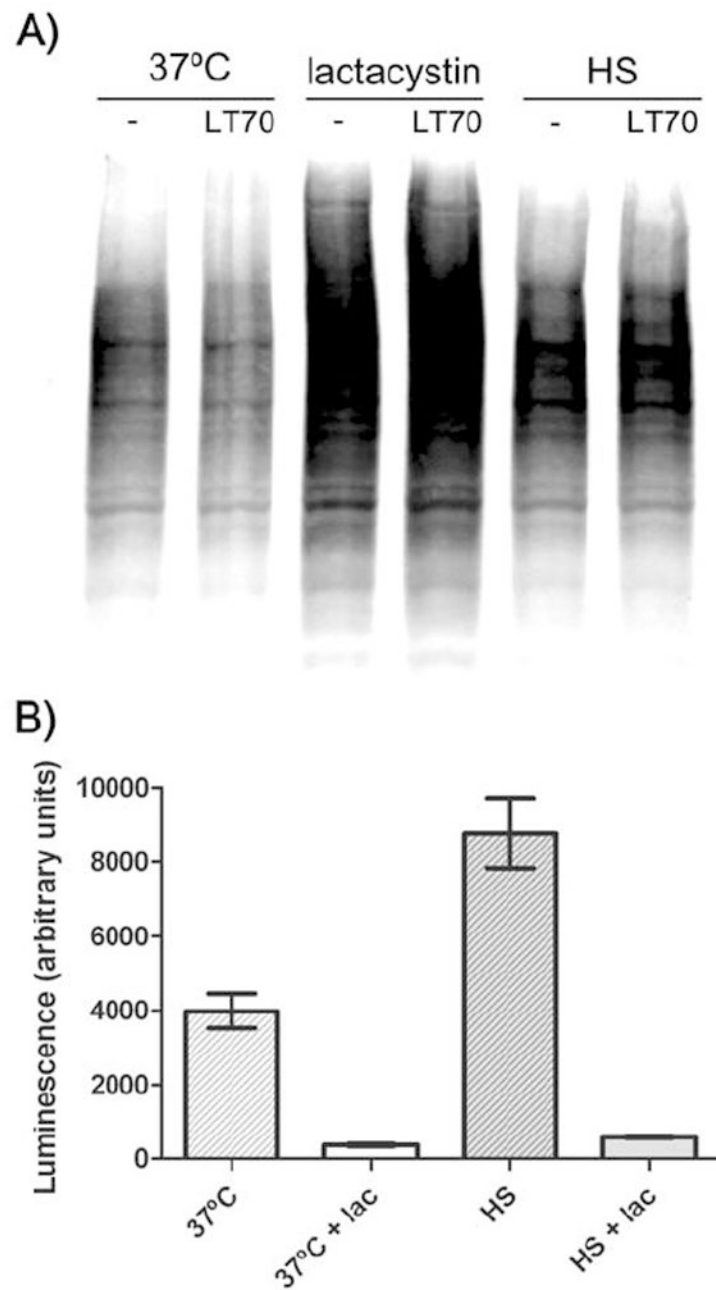


Figure 8. Heat shock does not inhibit proteasome activity

(A) RAW 264.7 cells were heat shocked for 50 min or treated with lactacystin (25 μ M) for 10 min. LT was added (1 μ g mL⁻¹) and after 70 min cells were lysed and protein ubiquitination monitored by Western blot. (B) RAW 264.7 cells were heat shocked or incubated at 37°C for 1.5 h before a 20 min lactacystin treatment (25 μ M). Cells were brought to room temperature and proteasome activity assayed using a cell permeable luminescent substrate as described in Experimental procedures.

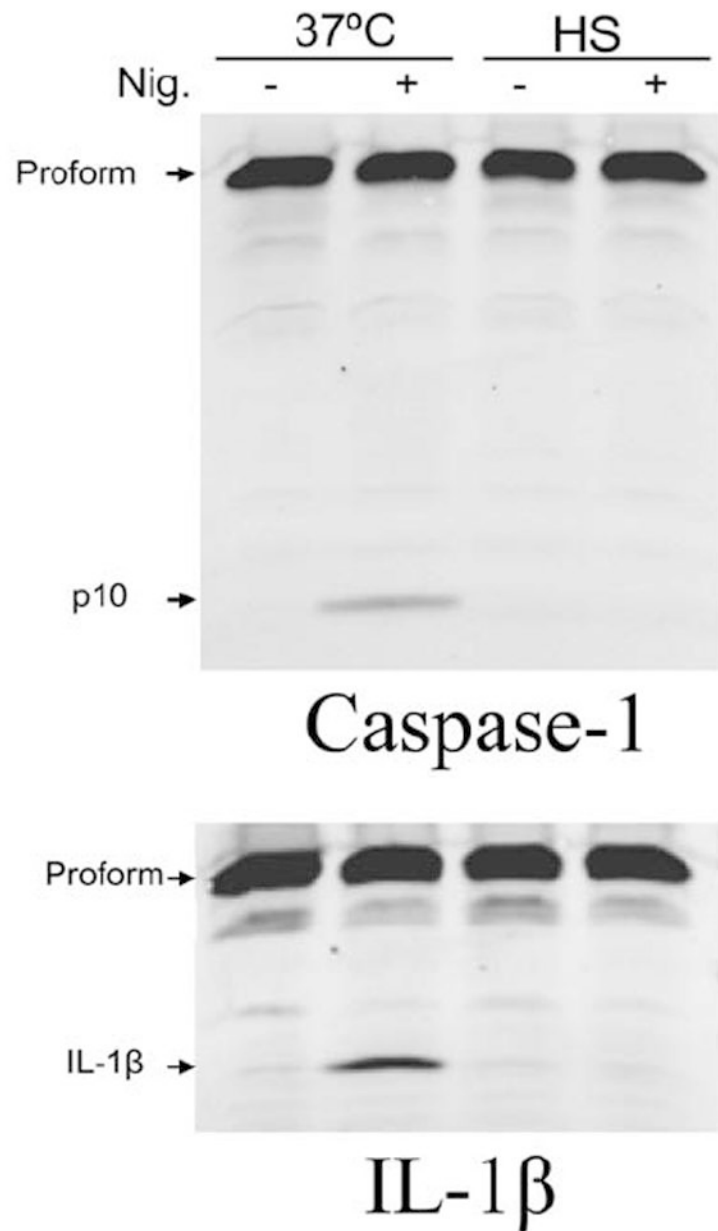


Figure 9. Heat shock prevents caspase-1 activation by nigericin
 BALB/cJ BMDM cells were treated with LPS (500 ng mL^{-1}) for 2 h at 37°C prior to a 1 h incubation in heat shock or 37°C conditions. Heat shocked cells were moved back to 37°C for an additional 30 min, prior to treatment with nigericin ($30 \mu\text{M}$, 20 min) and caspase-1 p10 and IL-1 β were monitored by Western blot.

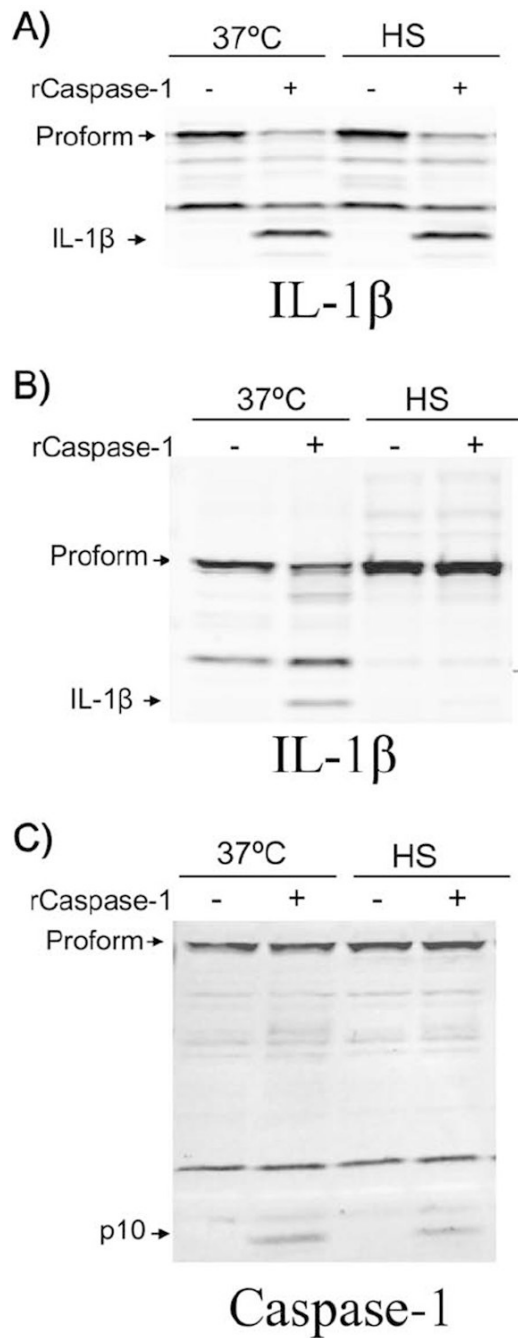


Figure 10. Whole cell lysates from heat shocked cells inhibit caspase-1 activity

(A) Sucrose lysates from LPS-treated ($1 \mu\text{g mL}^{-1}$ for 2 h) RAW 264.7 cells were mixed with recombinant caspase-1 (1 unit/ $40 \mu\text{L}$) or H_2O , and incubated at 37°C or heat shocked for 1h. IL-1 β cleavage was monitored by Western blot. (B) RAW 264.7 cells were LPS treated as above before a 1 h heat shock or 37°C incubation. Sucrose lysates were made and mixed with recombinant caspase-1 or H_2O as above at 37°C and examined for IL-1 β cleavage (B) or caspase-1 (C) by SDS electrophoresis Western blotting.

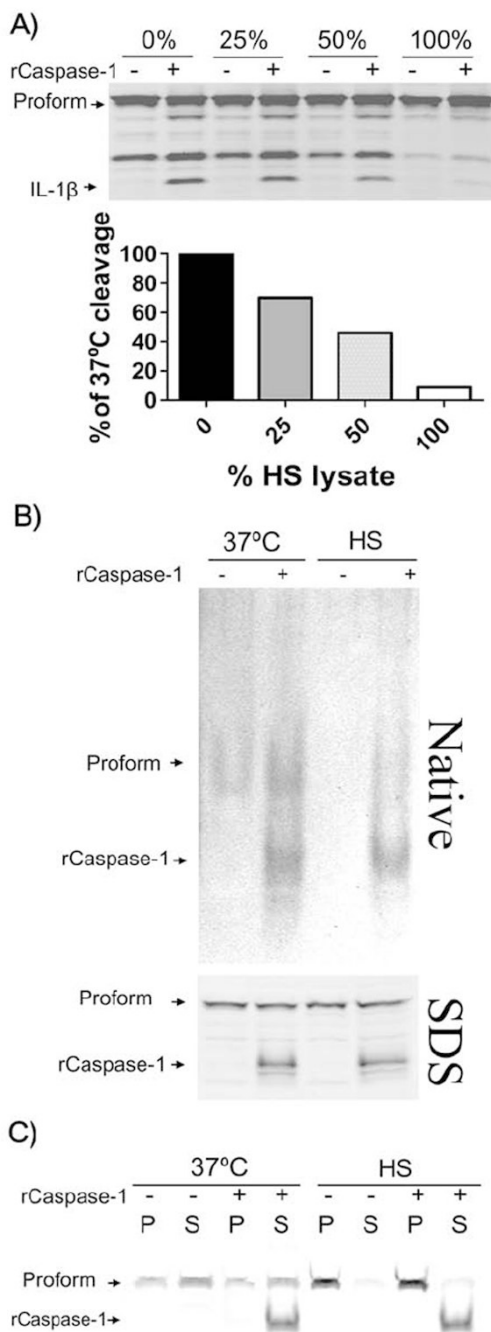


Figure 11. Caspase-1 inhibition by lysates of heat shocked cells is titratable

(A) RAW 264.7 cells were LPS-treated ($1 \mu\text{g mL}^{-1}$ for 2 h) and subsequently heat shocked for 1 h. Sucrose lysates of 37°C and heat shock cells were combined in various ratios (% heat shock lysate, vol/vol) before the addition of recombinant caspase-1 (1 unit/40 μL). After a 1 h incubation at 37°C, samples were examined for IL-1 β cleavage. Semi-quantitative analysis of IL-1 β cleavage band intensity is shown below. (B) RAW 264.7 cells were LPS treated as above before a 1 h heat shock or 37°C incubation. Sucrose lysates of the cells were mixed with recombinant caspase-1 or H₂O at 37°C and examined for caspase-1 by native gel electrophoresis (upper) and SDS-PAGE (lower) by Western blotting. (C) RAW 264.7 cell lysates prepared as above were mixed with native gel loading dye with β -mercaptoethanol and

centrifuged at $16,000 \times G$ for 10 min. Pellets (P) and supernatants (S) were examined for caspase-1 by SDS-PAGE and Western blotting.