# **Forum Review**

# Hsp27 Consolidates Intracellular Redox Homeostasis by Upholding Glutathione in Its Reduced Form and by Decreasing Iron Intracellular Levels

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### **ABSTRACT**

Small stress proteins [small heat shock proteins (sHsps)] are molecular chaperones that modulate the ability of cells to respond to oxidative stress. The current knowledge concerning the protective mechanism generated by the expression of mammalian heat shock protein-27 (Hsp27) that allows cells to increase their resistance to oxidative stress is presented. We describe the effects mediated by Hsp27 expression toward crucial enzymes such as glucose-6-phosphate dehydrogenase and glutathione reductase that uphold glutathione in its reduced form. New data are presented showing that the expression of sHsps correlates with a drastic decrease in the intracellular level of iron, a catalyzer of hydroxyl radical (OH\*) generation. A decreased ability of sHsps expressing cells to concentrate iron will therefore end up in a decreased level of oxidized proteins. In addition, we propose a role of Hsp27 in the presentation of oxidized proteins to the proteasome degradation machinery. We also present an analysis of several Hsp27 mutants that suggests that the C-terminal part of this stress protein is essential for its protective activity against oxidative stress. *Antioxid. Redox Signal.* 7, 414–424.

### INTRODUCTION

Small stress proteins [also denoted small heat shock proteins (sHsps)] share a sequence (the α-crystallin domain) in their C-terminal moiety with mammalian α-crystallin polypeptides. sHsps were first characterized in *Drosophila* as proteins that displayed increased synthesis following heat shock or conditions or agents that alter protein folding (6). Families of sHsps have been characterized in different organisms (22). The human family contains 10 polypeptides, but only a subset of them are true heat shock proteins. In mammals, the most studied sHsps are heat shock protein-27 (Hsp27) (also called HspB1) and αB-crystallin (HspB3). Several reports have described an increased cellular resistance to various stress in response to sHsp overexpression. Protection has been observed in the case of heat shock (47), oxidative stress (2–4, 6, 10, 56), cancer chemotherapy agents (30, 31, 38, 69, 81), and inflammatory mediators such

as tumor necrosis factor- $\alpha$  (TNF $\alpha$ ) (59, 93). Evidence has been obtained that sHsps share an ATP-independent chaperone activity (25, 43, 44) and are able to bind misfolded polypeptides. This creates reservoirs of folding intermediates (24, 49) counteracting the formation of deleterious protein aggregates. Folding intermediates are then renatured by ATPdependent protein chaperones (Hsp70, Hsp40, Hsp90, and cochaperones) and/or degraded by the ubiquitin-proteasome machinery. sHsp expression was also found to counteract different apoptotic processes such as those induced by agents that do not trigger an oxidative stress (9) or that take place during early differentiation (62). For example, because of the existence of an apoptotic signaling pathway linking cytoskeleton damages to mitochondria, the ability of Hsp27 to protect F-actin network integrity interferes with the release of cytochrome c from mitochondria (75). Hsp27 also binds cytochrome c once it is released from the mitochondria and inhibits its ability to promote apoptosome activation (15).

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Hsp27 can also interfere with procaspase 3 activation (70) and can modulate DAXX (18) and Akt (80) signaling mechanisms

The present report deals with recent observations concerning the ability of mammalian sHsps to modulate intracellular redox status and oxidoresistance. New data are presented concerning different Hsp27 mutants and the ability of sHsps to down regulate iron intracellular levels.

## DYNAMIC CHANGES IN THE INTRACELLULAR LOCALIZATION OF sHsps

Mammalian Hsp27 exists in a number of dynamic states depending on the physiological state of the cell. In the absence of nonionic detergents, biochemical fractionation revealed that Hsp27 partitioned approximatively equally between the soluble and insoluble fractions, whereas in the presence of detergent this polypeptide was in the soluble fraction. Indirect immunofluorescence experiments confirmed that, in cells grown at normal temperature, Hsp27 is present in the cytosol and also at the level of detergent-sensitive structures such as membranes (8). When cells are exposed to a drastic heat shock treatment, Hsp27 has the tendancy to concentrate transiently within the nucleus, where it associates with detergent-resistant structures (8). In contrast, oxidative stress abolishes the presence of Hsp27 in detergent-sensitive insoluble fractions (58), hence suggesting that in oxidative conditions Hsp27 is essentially cytosolic.

## EXPRESSION OF EXOGENOUS sHsps CAN GENERATE A PRO-REDUCED STATUS IN MAMMALIAN CELLS

In normal growing conditions, cells such as murine fibroblasts (i.e., murine L929 or NIH 3T3-ras cells) are devoid of constitutive expression of either Hsp27 or αB-crystallin. These cells are extremely sensitive to oxidative stress and used as standards for TNFα cytotoxicity. Stable transfections of these cells were achieved using expression vectors bearing the coding sequence of different sHsps under the control of a constitutive promoter. In these cell lines, we and others have observed that the expression of either human Hsp27, murine Hsp27, Drosophila Hsp27, or mammalian αB-crystallin upregulated total glutathione level and significantly decreased the basal level of intracellular reactive oxygen species (ROS) (11, 60, 74, 79). However, this redox state modulation depends on the cellular content of sHsps and is therefore difficult to detect in cells that express a high level of sHsps. For example, in L929 fibroblasts, the level of glutathione was not further increased and reached a plateau value when exogenous Hsp27 accumulation reached ~1 ng/μg of total cellular proteins (59). Another example concerns HeLa cells that constitutively express a high level of endogenous Hsp27 (~3 ng/ µg of total proteins). In these cells, a transfection-mediated increase in Hsp27 levels did not significantly modify glutathione and ROS levels. In human colorectal cancer HT-29 cells, the overexpression of Hsp27 down-regulated ROS without altering the reduced glutathione (GSH) level (31). An Hsp27-mediated down-regulation of ROS was also observed in monkey COS-7 cells (W. Firdaus and A.-P. Arrigo, manuscript in preparation).

Some interesting observations have been made concerning the mechanism that generates a pro-reducing state in response to the expression of sHsps. First, it has been reported that the expression of Hsp27 increases the level of extracellular superoxide produced by CCL39 fibroblasts (87). This phenomenon may be related to the protection of NADPH-oxidase activity by Hsp27 and is compatible with the generation of a pro-reduced state inside the cell.

Second, a careful analysis of several key enzymes that regulate intracellular redox state was performed (79). This study showed that in murine L929 cells, the expression of human Hsp27 significantly increases the activity of glucose-6-phosphate dehydrogenase (G6PDH), glutathione reductase, and glutathione transferase (Fig. 1). G6PDH, the first enzyme of the pentose phosphate cycle, has the interesting property of reducing NADP+ to NADPH(H)+ and thus is the key enzyme that provides the reducing power of the cell (Fig. 1). In this respect, we and others have shown that genetically manipulated L929 and HeLa cells expressing high levels of G6PDH display an increased level of GSH and show oxidoresistance (79, 83). Hence, the possibility exists that the modulation of GSH and ROS levels observed in Hsp27-expressing cells results from the increased G6PDH activity detected in these cells. The oxidoresistance generated by high G6PDH activity may then be responsible for the protection observed at the level of cell morphology, cytoskeletal architecture, and mitochondrial membrane potential (74, 77, 79). Further analysis revealed that the cellular content of G6PDH was increased as a consequence of the presence of large aggregates of Hsp27 (61, 78). How the large oligomers of Hsp27 increase G6PDH levels is unknown. It can be excluded that this phenomenon results from a transcriptionally regulated event because similar levels of G6PDH mRNA were detected in normal and Hsp27-expressing L929 cells. Hsp27 may then affect G6PDH half-life or the translability of G6PDH mRNA. These hypotheses should be tested because several reports have suggested an important role of sHsps in the control of protein turnover (71, 72) and mRNA translability (20, 68).

One major problem encountered by in vivo analysis is that Hsp27 knockout mice are not viable. This lethality is linked to the fact that embryonic stem cells that are unable to express Hsp27 during early differentiation undergo apoptosis (62). Recently, a very interesting study from Ivor Benjamin's laboratory provided evidence that the modulation of the intracellular redox state by sHsp was not restricted to cell cultures, but was also observed in animals (95). Using heat shock factor 1 (Hsf1) knockout mice as the model, these authors observed a strong reduction in the constitutive expression of Hsp27 and αB-crystallin in cardiac cells (72% in the case of Hsp27 and 34% in the case of  $\alpha$ B-crystallin). In these *hsf1*<sup>-/-</sup> cardiac cells, the down-regulation of these sHsps correlated with a 40% decrease in the ratio of GSH to oxidized glutathione (GSSG) and with a 43% increase in the level of superoxide ions. These phenomena were also associated with a 34% decreased activity of G6PDH. Taken together, these ob-

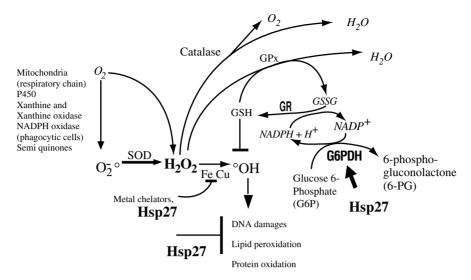


FIG. 1. Illustration of Hsp27 putative mode of action during oxidative stress. Hsp27 generates a pro-reduced state inside the cell by modulating the activity of G6PDH, glutathione reductase (GR), and glutathione transferase. Hsp27 also decreases the intracellular level of iron and therefore interferes with the formation of hydroxyl radical (OH•) via the Fenton reaction. This leads to a decreased level of ROS, increased level of GSH, and the upholding of this redox modulator in its reduced form. GPx, glutathione peroxidase; O<sub>2</sub>•, superoxide ion; SOD, superoxide dismutase.

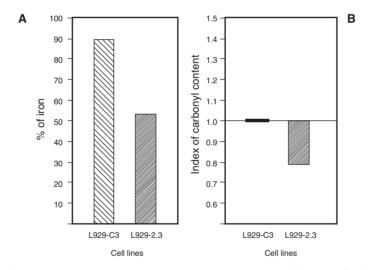


FIG. 2. Hsp27 decreases intracellular iron level and protein carbonyl content. (A) Determination of the intracellular concentration of iron determined by the colorimetric method of Fish (26). Stable transformants of L929 murine fibroblasts expressing (L929–2.3) or not expressing (L929-C2) 0.2 ng of human Hsp27/µg of total cellular proteins (79) have been analyzed [L929–2.3 cells are also indicated as L929-27-3-97 cells in Préville et al. (79)]. Cells were washed and scraped from the dish in phosphatebuffered saline, and the cell pellets were resuspended in 1 ml of water. An equivalent of 2 mg of total protein was used per assay. Buffer A (500  $\mu$ l) (0.6 M HCl, 0.142 M KMnO<sub>4</sub>) was added, and the samples were incubated for 2 h at 60°C. This procedure allowed iron to be released and solubilized. Buffer B (100 ul) was then added (6.5 mM Ferrozine [disodium 3-(2-pyridyl)-5.6-bis(4phenylsulfonate)-1,2,4-triazine]; 3.1 mM neocuproine (2,9-dimethyl-1,10-phenanthroline), 2 M ascorbic acid, and 5 M ammonium acetate]. After 60 min of incubation, absorbance of Ferrozine–Fe(II) complex was read at 562 nm. In addition to Fe(II), only Cu(I) and Co(II) can form a colored complex with Ferrozine. Neocuproine is added to the reaction mixture to chelate Cu(I). This chelator allows a quantification of iron with <5% error in the presence of up to a 2:1 weight ratio of Cu:Fe. Co(II) interferes <5% in the iron assay for quantities of Co(II) up to a 5:1 weight ratio over Fe. Similar analysis was performed using 1 ml of the growth medium as starting material. The ratio between the level of iron present in the cell extract and that observed in the growth medium was calculated. The corresponding percentage is presented. All chemicals were from Sigma-Aldrich (St-Quentin-Fallavier, France). (B) Protein carbonyl content, which is taken as presumptive evidence of an irreversible protein oxidative modification, was measured using 2,4-dinitrophenylhydrazine (51) as already described (79). Index of carbonyl content was calculated as the ratio between the value determined in Hsp27-expressing cells and that measured in control cells.

servations indicate that the constitutive expression of sHsps is directly and functionally linked to the maintenance of redox homeostasis and antioxidative defenses at normal temperature.

As iron is an important factor in the cellular damage generated by oxidative stress, we have analyzed the intracellular level of this metal in cells that do or do not express sHsps. Iron catalyzes the Haber-Weiss/Fenton reactions that lead to the formation of the highly reactive and toxic hydroxyl radical (OH\*, see Fig. 1) (37, 89). The ability of cells to concentrate iron will therefore facilitate the formation of intracellular OH. High levels of intracellular iron thus stimulate oxidative damage and have been associated with a number of oxidative injury-dependent, age-related conditions and diseases (63, 76, 92). We have observed that the expression of human Hsp27 in murine L929 cells decreased the intracellular level of iron by almost 50% (Fig. 2A). We made a similar observation in genetically modified HeLa cells expressing αB-crystallin, a polypeptide that is normally not constitutively expressed in these cells (S. Virot and A.-P. Arrigo, manuscript in preparation). Moreover, oxyblot analysis (88) revealed a decrease in the level of oxidized proteins in L929 cells expressing Hsp27 (79) (Fig. 2B) and in HeLa cells expressing α-crystallin (S. Virot and A.-P. Arrigo, manuscript in preparation). Hence, at least two sHsps appear to share the ability to decrease intracellular iron levels and to interfere with the level of proteins oxidized as a consequence of OH. activity. This observation is of prime importance because in vivo iron deprivation or chelation acts as a potent antioxidant, preventing oxidative stress in tissues and organs. Iron chelators have also been described to favor successful ageing in general (76). The mechanism responsible for decreasing iron intracellular level is unknown, but could be related to an inhibitory effect of sHsps toward the action of iron-regulatory proteins (IRP-1,2) (17, 54). IRP-1,2, whose action is sensitive to cellular iron concentration, are activated and bind stem-loop untranslated regions (IRE regions) of transferrin and ferritin mRNAs, hence activating and inhibiting their translations, respectively. This concerted mechanism permits a fine tuning of iron homeostasis in the cell. It is likely that IRP-1,2 sense Fe(II) and that Fe(II) oxidation to Fe(III) by hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) treatments of the cells sets up a program for increasing iron uptake (17, 54). Whether sHsps modulate the translation of transferrin and ferritin mRNAs will merit investigation because it could represent the second example of an mRNA translation modulation by sHsp expression that generates a pro-reduced state in cells.

The question remains as to whether the modulation of the intracellular redox status by sHsps results from the biochemical action of these proteins or an adaptation of the cell to their presence. In other words, could it be that sHsps are prooxidant proteins that generate a drastic antioxidant defense of the cell, such as a decrease in the intracellular level of iron? Among the experiments that are in favor of this hypothesis, one can cite those describing a sensitization of the cells to oxidative stress. For example, Hsp27 overexpression sensitizes a squamous carcinoma cell line (91), as well as KMST-6 human immortalized fibroblasts (1) and a sub species of L929 fibrosarcoma cells (53), to oxidative stress. Hence, the cellular context in which Hsp27 is expressed appears impor-

tant. However, care should be taken in analyzing these results because it is difficult to compare studies devoid of Hsp27 expression quantification. For example, in murine L929 fibroblasts, a low level of expression of Hsp27 generates oxidoresistance, whereas a high level of this protein is usually toxic and sensitizes cells to oxidative stress.

# SHSPS PROTECT AGAINST OXIDATIVE STRESS-INDUCED CYTOTOXICITY

Depending on its intensity, oxidative stress can induce two types of cell death process. In the case of cells exposed to moderate levels of ROS, apoptosis is usually observed probably because of the depletion of GSH, which triggers the mitochondrial apoptotic pathway leading to caspase activation (99). In contrast, high levels of ROS drastically oxidize proteins, lipids, and nucleic acids. This oxidation process triggers cell necrosis, which, *in vivo*, can lead to inflammation (42). Apoptosis is inhibited by a high level of ROS, which decreases ATP levels and inhibits the activation of caspases (50, 67, 84).

Several reports have described a protection against the deleterious effects induced by oxidative stress in cells (L929, NIH 3T3-ras, CHO, HeLa, colorectal cancer HT-29) expressing sHsps (human Hsp27, murine Hsp27, rodent Hsp27, Drosophila Hsp27, and mammalian αB-crystallin) (31, 40, 41, 59, 73, 93). This protection correlated with a significant decrease in the intracellular burst of ROS generated by oxidative stress ( $H_2O_2$ , menadione, TNF $\alpha$ ) (58–61, 82). This modulation of ROS production was found to be glutathionedependent because sHsps cannot protect against the oxidative stress generated by drugs that interfere with GSH activity or synthesis, such as buthionine sulfoximine or diethyl maleate (60). Hence, ROS-dependent phenomena, such as lipid peroxidation, protein oxidation, nuclear factor-κB activation, and disruption of F-actin architecture, were decreased by the expression of Hsp27 (60). Further analysis revealed that the pattern of irreversibly oxidized proteins (carbonylation of side chains) in response to H<sub>2</sub>O<sub>2</sub> treatment observed in Hsp27-expressing cells was different from that observed in control cells pretreated with the antioxidant drug glutathione ethyl ester (79). This suggests that Hsp27 may act in different ways: (a) by exerting unspecific protections (upholding of glutathione in its reduced form, decrease of intracellular iron and ROS) and (b) by acting at the level of specific proteins to either avoid their oxidation and/or induce their rapid degradation once they are irreversibly oxidized. Among the different enzymes protected by Hsp27 expression one can cite glutathione S-transferase and glutathione peroxidase that are involved in the ROS detoxification machinery (79).

# Hsp27 ABILITY TO PROTECT AGAINST OXIDATIVE STRESS IS REGULATED BY ITS STRUCTURAL ORGANIZATION

To approach the mechanism of Hsp27 action, the functional consequences of two prominent features of this poly-

peptide, oligomerization and phosphorylation, have been considered.

sHsps are oligomeric proteins that display dynamic changes in their oligomerization profile when cells are exposed to environmental changes (5, 7, 8, 29, 55). For example, in normal growth conditions, human Hsp27 is cytosolic and forms heterodispersed oligomers with native size ranging from 60 to 800 kDa. However, when cells are starved and arrested in G<sub>a</sub>, only the small oligomers of Hsp27 are detected (55). In cells exposed to heat shock, Hsp27 large oligomers transiently accumulate and, depending on the intensity of the stress, can be recovered inside the nucleus (8). This phenomenon is followed by the rapid accumulation of Hsp27 in the form of small oligomers as a consequence of Hsp27 phosphorylation. Of interest, in cells made thermotolerant to the heat stress, no changes in the structural organization and phosphorylation of Hsp27 are observed (8), hence suggesting that these changes are directly correlated with the intracellular damages induced by heat shock. Similarly, the protective activity of Hsp27 in response to oxidative stress (TNF $\alpha$ , H<sub>2</sub>O<sub>2</sub>) correlates with the formation of large structures containing this protein. This phenomenon is followed by the phosphorylation-dependent dissociation of Hsp27 large structures (58, 61, 79). In cells overexpressing the antioxidant enzyme selenoglutathione peroxidase, no changes in Hsp27 structural organization and phospho-isoform composition are observed in response to oxidative stress (57). This suggests that the intracellular burst of ROS observed in response to oxidative stress is responsible for the changes in Hsp27 structural organization and phosphorylation.

In cells exposed to heat shock, Hsp27 large oligomers have been described to act as reservoirs that facilitate the renaturation of misfolded proteins by other chaperones, such as the ATP-dependent chaperones Hsp70, Hsp40, and Hsp90 (24, 49). However, in the case of irreversibly oxidized proteins, selective proteolysis is triggered (33). Hence, the transient accumulation of Hsp27 in the form of large oligomers probably activates the presentation of irreversibly oxidized proteins to the ubiquitin-independent 20S proteasome (4, 10), which is known to have a high affinity for oxidized proteins (33, 34, 85). This phenomenon counteracts the accumulation of proteolysis-resistant large aggregates of oxidized proteins (lipofuscin) that block proteasome activity (13) and are extremely deleterious to the cell (86).

Hsp27 phosphorylation during heat or oxidative stress is regulated by the stress kinase pathway. Murine Hsp27 is phosphorylated at serine residues 15 and 86 (27), whereas human Hsp27 contains three phosphorylated serines residues, 15, 78, and 82 (48). Hsp27 phosphorylation is mediated by numerous stimuli that include heat shock,  $TNF\alpha$ , and other forms of oxidative stress (6) and is catalyzed by mitogen-activated protein kinase-activated protein kinase 2 and 3 (MAP-KAPK2-3) (19, 39, 52, 90) and/or by the inactivation of specific phosphatases (28, 36).

Several studies have demonstrated that phosphorylation generates the formation of small oligomers of Hsp27 (45, 61, 82). For example, inhibition of p38 kinase by the SB203580 inhibitor did not alter the formation of Hsp27 large oligomers in response to oxidative stress, but inhibited their dissociation later (78). Moreover, studies performed with nonphosphory-

latable or phosphorylated mimicry mutants led to the conclusion that the large unphosphorylated oligomers represent the active form of Hsp27 that modulates ROS and glutathione levels and displays *in vitro* chaperone activity (61, 88).

Hence, the intracellular burst of ROS generated early during oxidative stress induces the formation of large oligomers of Hsp27 that bear chaperone and anti-ROS activities. The concomitant stimulation of MAPKAPK2-3 kinase, which phosphorylates Hsp27, then leads to the subsequent formation of small oligomers that are observed when the oligomerization pressure mediated by the ROS burst is decreasing. These small oligomeric forms may either inactivate Hsp27 chaperone activity or induce its recycling through dynamic deaggregation—oligomerization of the protein.

The dynamic changes in Hsp27 oligomerization and phosphorylation may also play a role in the protection of the cytoskeleton because the F-actin microfilament network is one of the earliest targets of oxidative stress (12, 21). In this respect, unphosphorylated Hsp27 small oligomers have been described to inhibit *in vitro* F-actin polymerization (14, 65). *In vivo* Hsp27 also appears to modulate F-actin filament dynamics (35, 66) and protects F-actin against oxidative stress-induced dissociation (40), probably by upholding glutathione in its reduced form (77) and decreasing iron intracellular levels.

### **ANALYSIS OF Hsp27 MUTANTS**

We have obtained several mutants of human Hsp27 (point mutations or deletions) (see Fig. 3). These mutants were transiently expressed in HeLa cells and their protective ability against a 24-h oxidative stress performed with 600  $\mu$ M H<sub>2</sub>O<sub>2</sub> has been analyzed. Figure 4 compares the protective activity

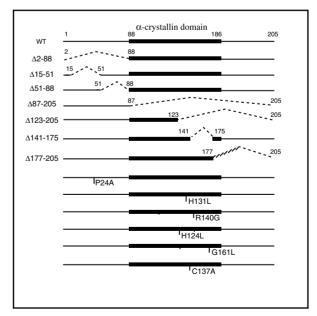
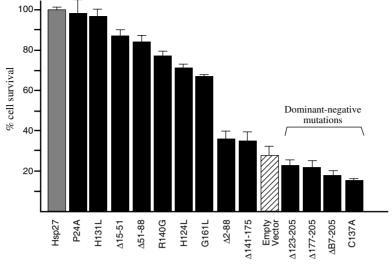


FIG. 3. Schematic illustration of human Hsp27 mutants. Point mutations, as well as deletions, are indicated. Black bars correspond to the  $\alpha$ -crystallin domain shared by sHsps.



Vectors used for cell transfection

FIG. 4. Protection mediated by wild type and mutants of Hsp27 against H,O, cytotoxicity. Construction of Hsp27 vectors. The cDNA encoding human Hsp27 was inserted into the EcoRI site of pSVK3 vector (Pharmacia, Saint-Quentin-en-Yvelines, France) or pCI-neo vector (Promega, Charbonnieres, France). Empty vector corresponds to either plain pSVK3 or pCI-neo vector. The mutants described in this study were constructed as follows: (A) Mutants constructed in Hsp27-bearing pSVK3 vector using the following PCR primers. H131L (substitution of histidine 131 by a leucine residue): sense 5'-CAGGACGAGCTCGGCTA CATCT-3' and antisense 5'-AGATGTAGCCGAGCTCGTCCTG-3'. H124L (substitution of histine 124 by a leucine residue): sense 5'-ACCGGCAAGCTTGAGGAGCGGC-3' and antisense 5'-GCCGCTCCTCAAGCTTGCCGG-3'. P24A (substitution of proline 24 by an alanine residue): sense 5'-CGACTGGTACGCGCATAGCCG-3' and antisense 5'-CGGCTATGCGCGTA CCAGTCG-3'. G161L (substitution of glycine 161 by a leucine residue): sense 5'-GTCCCCTGAGCTCACACTGACC-3' and antisense 5'-GGTCAGTGTGAGCTCAGGGGAC-3'. DNA amplification was followed by restriction and ligation steps. The Δ2-88 deletion was constructed by introducing two SphI sites in the Hsp27 coding sequence using the following primers: sense 5'-GGGGTCTCGGGCATGCGGCACACTG-3' and antisense 5'-GCGGCGCTCGGGCATGCTGGCTC-3'. Following restriction and ligation steps, the deletion was generated by digestion with SphI. The  $\Delta 15-51$  deletion was generated by digestion with PvuII followed by a ligation step. (B) Mutants generated in Hsp27-bearing pCI-neo vector using the Stratagene Quickchange™ sitedirected mutagenesis kit (Stratagene Europe, Amsterdam Zuidoost, The Netherlands). C137A (substitution of the unique cysteine residue of Hsp27 by an alanine residue), PCR primers: sense 5'-GGCTACATCTCCCGGGCCTTCACGCGGAAATACACG-3' and antisense 5'-CGTCTATTTCCGCGTGAAGGCCCGGGAGATGTAGCC-3'. R140G (substitution of arginine 140 by glycine): sense 5'-CCCGGTGCTTCACGGGGAAATACACGCTGCCC-3' and antisense 5'-GGGCAGCGTGTATTTCCCCGTGAAGCAC CGGG-3'. The Δ51–88 deletion was constructed using the following primers: sense 5'-GGTTAGGCGGCAGCAGCAGCACAC TGCGGACCG-3' and antisense 5'-CGGTCCGCAGTGTGCCGCTGCTGCCGCCTAACC-3'. The Δ141–175 deletion was generated with the following primers: sense 5'-CCGGTGCTTCACGCGGAACGAGATCACCATCC-3' and antisense 5'-GGATGGT GATCTCGTTCCGCGTGAAGCACCGG-3'. For the  $\Delta 87-205$ ,  $\Delta 123-205$ , and  $\Delta 177-205$  deletions, we used the following primers containing an EcoRI site: sense 5'-CGGAATTCATGACCGAGCGCCGCGTCCCC-3' and antisense 5'-GGAATTCTTA CGAGACCCCGCTGC-3', 5'-GGAATTCTTAGCCGGTGATCTCCACC-3' and 5'-GGAATTCTTAGGACTGCGTGGCTAGC-3', respectively. The deleted Hsp27 genes were then cloned in the EcoRI site of pCI-neo plasmid. Transfection experiments: HeLa cells transiently expressing wild type (gray plot) or mutants (black plots) of human Hsp27 were exposed to 600  $\mu$ M H<sub>2</sub>O<sub>2</sub> for 24 h. Control cells transfected with the corresponding empty vector were also analyzed (hatched bar). Cell survival was estimated by crystal violet staining. Percentage of cell survival was calculated by comparing the values obtained for the different mutants with that observed for wild-type Hsp27 (100%). Mutants that induced a sensitization to H<sub>2</sub>O<sub>2</sub> were considered as dominant negative, probably because they interfere with the protective activity of endogenous Hsp27. Standard deviations are indicated (n = 6).

of the different mutants to that of wild-type Hsp27. This figure clearly shows that a hydrophobic replacement at position 131 (H131L mutant) that should alter the major hydrophilic peak seen in the  $\alpha$ -crystallin domain had no effect. The hydrophobic replacement at position 124 (H124L mutant) or 161 (G161L mutant) was more effective. Analysis of the N-terminal part comprised of amino acids (AA) 15–88 revealed that this region is not of prime importance for protecting the cell against  $\mathrm{H_2O_2}$ . In contrast, the C-terminal part of Hsp27 appears essential. Some mutants (in the C-terminal part of the protein, as well as the C137A point mutation) render the cell

hypersensitive to H<sub>2</sub>O<sub>2</sub>. As sHsps of different origins can form oligomeric complexes (96), mosaic structures are probably formed that inactivate endogenous Hsp27 of HeLa cells. These mutants are therefore considered as dominant negative. Concerning the different point mutations that interfered with Hsp27 protective activity, we have already reported that the triple mutation of the phosphorylated serine sites (serines 15, 78, and 82) does not protect against oxidative stress (82). Of interest, mammalian Hsp27 polypeptides contain only one cysteine residue (position 141 in murine and 137 in human) that is highly suceptible to oxidation. It is seen in Fig. 4 that

the substitution of this cysteine by an alanine residue (C137A mutant) renders HeLa cells extremely sensitive to  $\rm H_2O_2$ . The mutation impaired Hsp27 ability to form dimers and induced a dominant-negative effect (C. Diaz-Latoud, E. Buache, E. Javouhey, and A.-P. Arrigo, *Antioxid Redox Signal* 7: 436–445, 2005). The role of the unique cysteine residue of Hsp27 is unclear because the oligomerization of this protein does not appear to rely on disulfite formation (97), but to require phosphorylation-sensitive interactions at the N-terminus (46). However, disulfite formation between two Hsp27 polypeptides and Hsp27 *S*-thiolation have been observed, especially in oxidative stress conditions (23, 97, 98).

A comparison of the results described in Fig. 4 with those obtained when cells are exposed to etoposide (an apoptotic agent inhibiting topoisomerase II) reveals different domains of Hsp27 that protect against these agents (Table 1). For example, the region comprised of AA 51–88 is essential for the binding to cytochrome c and for the protection against apoptosis (15). In contrast, the  $\Delta 51$ –88 deletion only weakly alters Hsp27 protective activity against  $H_2O_2$  induced cell death. The C-terminal part of the protein (AA 141–205) also shows differential protective activity against etoposide or  $H_2O_2$ . Hence, different domains appear to be involved in the protection against  $H_2O_2$  or etoposide. As Hsp27 appears to protect against oxidative stress through its chaperone activity (82), several other activities of Hsp27 may be required to control the mitochondrial apoptotic pathway.

Table 1. Protection Mediated by Wild-Type and Mutant Hsp27 Against H<sub>2</sub>O<sub>2</sub> and Etoposide Cytotoxicity

Mutations in human Hsp27	% cell death	
	$H_2O_2$	Etoposide
Wild type	$21.3 \pm 4.0$	$37 \pm 4.9$
P24A	$22.2 \pm 4.9$	ND
H131L	$22.3 \pm 3.5$	$33 \pm 4.6$
C137A	$88.3 \pm 4.1$	$92 \pm 4.3$
R140G	$39.0 \pm 6.9$	ND
H124L	$43.5 \pm 5.0$	ND
G161L	$47.2 \pm 3.5$	ND
$\Delta 1-88$	$68.6 \pm 2.0$	$97.4 \pm 4.3$
$\Delta 15 - 50$	$31.7 \pm 1.6$	$40 \pm 2.4$
$\Delta 51 - 88$	$35.1 \pm 2.6$	$110 \pm 4.1$
$\Delta 141 - 175$	$78.2 \pm 4.1$	$49 \pm 3.3$
$\Delta 123-205$	$82.3 \pm 1.0$	$87 \pm 6.5$
$\Delta 177 - 205$	$82.7 \pm 7.0$	$34 \pm 3.1$
$\Delta 87 - 205$	$85.8 \pm 4.8$	$92 \pm 4.3$
Empty vector	$100.0\pm0.0$	$100.0 \pm 0.0$

HeLa cells transiently expressing wild type or mutants of human Hsp27 were exposed for 24 h to  $600 \, \mu M \, \text{H}_2 \text{O}_2$  or  $500 \, \mu M$  etoposide. Control cells transfected with the corresponding empty vector were also analyzed. Cell death was estimated by crystal violet staining. Percentage of cell death was obtained from the ratio between the values obtained for the wild type and the different mutants and the value determined when transfection was performed with the empty vector (100%). Standard deviations are indicated (n=6). ND, not determined;  $\Delta$ , deletion.

### CONCLUSIONS AND PERSPECTIVES

Hsp27 is a negative regulator of programmed cell death with a broad spectrum of action. In addition to its antiapoptotic effect, Hsp27 also appears to play key roles in pathological situations where its expression can either aggravate the pathological state (*i.e.*, cancer cells with impaired apoptosis) (16, 32) or be beneficial to protect against oxidative stress, inflammation, or acute cell death (as, for example, in the case of neurodegenerescence and asthma pathologies where high levels of Hsp27 have been detected) (64, 94). We can therefore be confident that future studies will bring us a better understanding of the molecular mechanisms induced by this chaperone protein that modulate key enzymes of the ROS–glutathione pathway and intracellular iron levels.

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### **ABBREVIATIONS**

AA, amino acids; G6PDH, glucose-6-phosphate dehydrogenase; GSH, reduced glutathione; GSSG, oxidized glutathione;  $H_2O_2$ , hydrogen peroxide; Hsf1, heat shock factor 1; Hsp27, heat shock protein-27; IRP, iron regulatory protein; MAPKAPK2-3, mitogen-activated protein kinase-activated protein kinase 2 and 3; OH $^{\star}$ , hydroxyl radical; ROS, reactive oxygen species; sHsp, small stress protein or small shock protein; TNF $\alpha$ , tumor necrosis factor- $\alpha$ .

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