ALTERATIONS OF SURFACE MORPHOLOGY CAUSED BY THE METABOLISM OF MENADIONE IN MAMMALIAN CELLS ARE ASSOCIATED WITH THE OXIDATION OF CRITICAL SULFHYDRYL GROUPS IN CYTOSKELETAL PROTEINS

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Abstract—Incubation of freshly-isolated (rat hepatocytes) or cultured (HeLa, GH₃, and McCoy) mammalian cells with menadione (2-methyl-1,4-naphthoquinone) resulted in the appearance of numerous cell surface protrusions. The perturbation of surface structure was associated with an increase in the amount of cytoskeletal protein and the oxidation of sulfhydryl groups in actin, leading to the formation of high-molecular weight aggregates sensitive to treatment with thiol reductants. Our findings indicate that the oxidation of thiol groups in cytoskeletal proteins may be responsible for menadione-induced cell surface abnormalities in mammalian cells.

The metabolism of menadione (2-methyl-1,4-naphthoquinone) has been widely employed as a model to investigate the mechanisms and consequences of oxidative stress in mammalian cells [1-3]. In hepatocytes, menadione can undergo either one- or twoelectron reduction reactions. One-electron reduction of the quinone results in the formation of the semiquinone free radical which, in the presence of molecular oxygen, is readily reoxidized to the parent quinone with the concomitant production of superoxide anion. At toxic levels of menadione, redox cycling of the quinone results in extensive O_2^- and H_2O_2 formation followed by glutathione oxidation and depletion [2, 4], NAD(P)H oxidation [1], protein thiol modification [2, 3], disruption of intracellular Ca²⁺ homeostasis [1, 3, 5, 6] and, finally, cell death [1, 3].

In addition to the biochemical alterations, morphological changes, characterized by the appearance of numerous surface blebs, are also observed in cells exposed to toxic levels of menadione [7]. We have recently demonstrated that menadione-induced bleb formation in hepatocytes is associated with cytoskeletal abnormalities, of which the oxidation of sulfhydryl groups in actin seems to play a crucial role [8]. We can now report that similar alterations occur during the metabolism of toxic doses of menadione in various cultured cells, suggesting that cytoskeletal thiol oxidation may represent the general mechanism of menadione-induced cell surface abnormalities in mammalian cells.

MATERIALS AND METHODS

Collagenase (grade II) was obtained from

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Boehringer Mannheim (Mannheim, F.R.G.) and menadione was from Sigma (St. Louis, MO). All tissue culture media and sera were from Flow Laboratories (Herts, U.K.). Other reagents were of the highest grade of purity commercially available and were obtained from local suppliers.

Hepatocytes were isolated from male Sprague–Dawley rats by collagenase perfusion of the liver as described in [9]. Incubation conditions were similar to those reported by Mirabelli *et al.* [8]. GH₃ cells were cultured in Ham's F-10 medium supplemented with 15% horse serum and 2.5% fetal calf serum, HeLa cells in Eagle's minimum essential medium supplemented with 10% fetal calf serum, and McCoy cells in RPMI 1640 medium supplemented with 10% fetal calf serum. All cultures were performed in 90 mm plastic Petri dishes. The cultures were washed twice with phosphate-buffered saline (PBS) prior to the addition of the indicated concentrations of menadione in PBS.

Hepatocyte cytoskeleton was prepared as described in [8]. The cytoskeletal fraction from cultured cells was obtained by extracting the cell layer (previously washed twice with PBS) with 5 ml of a solution containing 40 mM KCl, 5 mM EGTA, 5 mM MgCl₂, 50 mM Tris–HCl, pH 7.5, and 1% Triton X-100 for 30 min on ice. The layer remaining adherent to the plastic surface was then scraped with a rubber policeman and centrifuged at 4000 g for 20 min. The pellet obtained was washed twice in the same buffer, dissolved in 8 M urea/1% SDS and used for protein and thiol measurements and for polyacrylamide gel electrophoresis (PAGE).

PAGE was performed on 10-15% gradient slab gels by the procedure of Laemmli [10]. Immunoblotting of actin was performed essentially as described by Mirabelli et al. [8]. Intracellular GSH

was measured by the HPLC technique described by Reed et al. [11]. Protein sulfhydryl groups were assayed using the procedure described in [2], and protein concentration was determined by the method of Lowry et al. [12].

RESULTS AND DISCUSSION

As shown in Fig. 1, incubation with menadione resulted in the appearance of cell surface abnor-

malities characterized by the protrusion of large (hepatocytes), intermediate (HeLa and McCoy cells), or small (GH₃) blebs. In all instances, the formation of surface blebs preceded cell killing, as measured by the uptake of Trypan blue. The appearance of the morphological changes were, in turn, preceded by biochemical alterations, including glutathione and protein thiol depletion (Table 1). These occurred, although at different rates and to various extents, in all four cell types investigated,

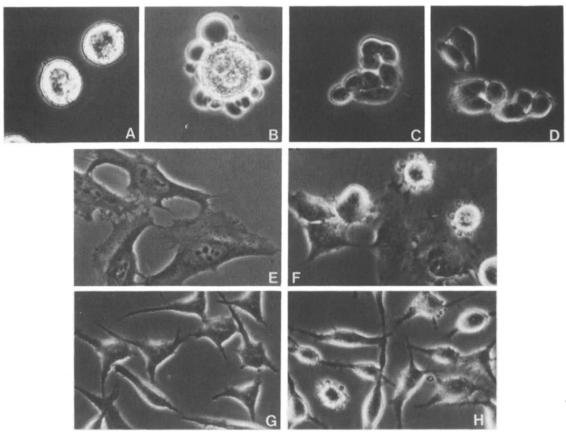


Fig. 1. Alterations in surface morphology following the metabolism of menadione in mammalian cells. Hepatocytes (A, B), GH₃ cells (C, D), HeLa cells (E, F), and McCoy cells (G, H) were incubated for 60 min in the absence (A, C, E, G) or presence (B, D, F, H) of 200 μM menadione. The cells were then harvested and submitted to light microscopy under phase contrast (magnification: 400×).

Table 1. The metabolism of menadione causes glutathione and protein thiol depletion and alterations of cell surface morphology in mammalian cells

	Cell types			
	Hepatocytes	GH ₃	HeLa	МсСоу
Cells exhibiting surface				
protrusions (%)	100	64	75	68
Intracellular GSH				
(% of control)	5	14	12	10
Protein thiols				
(% of control)	52	54	56	49

Hepatocytes, GH₃, HeLa, and McCoy cells were incubated with 200 μ M menadione as described in Methods. After 60 min, samples were taken for evaluation of surface abnormalities and for GSH and protein thiol measurements. The levels of GSH and protein thiols (nmol/mg protein) in untreated hepatocytes, GH₃ cells, HeLa cells, and McCoy cells were 22 \pm 5, 8 \pm 2, 14 \pm 2, 26 \pm 6 and 96 \pm 22, 88 \pm 14, 90 \pm 16, 98 \pm 18, respectively (mean \pm SD of three experiments).

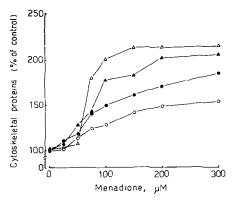


Fig. 2. Menadione-induced increase in the amount of cytoskeleton-associated protein in mammalian cells. Hepatocytes (\bullet), GH₃ cells (\triangle), HeLa cells (\triangle), and McCoy cells (\bigcirc) were incubated for 60 min with the indicated concentrations of menadione. The cytoskeleton was then extracted and the amount of protein measured as described in Methods. The amounts of cytoskeleton-associated protein (μ g/mg of cellular protein) in untreated cells were (mean \pm SD of three experiments): Hepatocytes, 66 \pm 8; GH₃ cells, 78 \pm 12; HeLa cells, 69 \pm 13; and McCoy cells, 52 ± 11 .

suggesting that common pathways may be involved in the metabolism of menadione and in menadione-induced cell damage. This assumption received further support from the finding that GH₃, HeLa, and McCoy cells, like hepatocytes, can catalyze redox cycling of menadione [1, 13, 14], leading to intracellular thiol depletion [2, 4, 15, 16] (F. Mirabelli and G. Bellomo, unpublished observation).

Previous studies with hepatocytes have suggested that plasma membrane bleb formation following toxic cell injury is associated with a perturbation of

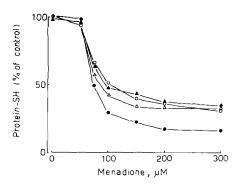


Fig. 3. Menadione-induced loss of cytoskeletal protein thiols in mammalian cells. Hepatocytes (\blacksquare), GH_3 cells (\triangle), HeLa cells (\blacktriangle), and McCoy cells (\bigcirc) were incubated for 60 min with the indicated concentrations of menadione. The cytoskeleton was then extracted and protein thiols measured as described in Methods. The amounts of cytoskeletal protein thiols (nmol/mg protein) in untreated cells were (mean \pm SD of four experiments): Hepatocytes, 55 ± 15 ; GH_3 cells, 44 ± 8 ; HeLa cells, 61 ± 6 ; McCoy cells, 58 ± 12 .

normal cytoskeletal organization [17, 18]. A relationship between the two events is also indicated by the finding that two cytoskeletal toxins, cytochalasin B and phalloidin, cause bleb formation in hepatocytes [19, 20].

To investigate the effects of menadione exposure on cytoskeletal organization in the various cell types, the cytoskeleton was prepared by the conventional Triton X-100 extraction method [21–23]. As illustrated in Fig. 2, menadione caused a dose-dependent increase in Triton X-100 non-extractable protein, suggestive of an enhanced association of non-cytoskeletal proteins with the cytoskeleton. This effect

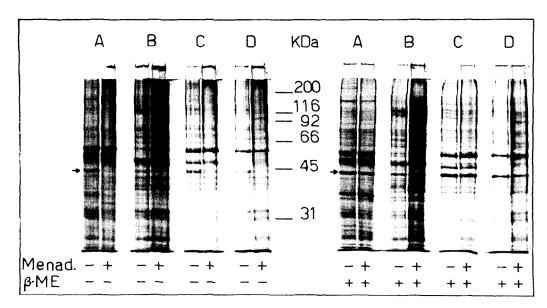


Fig. 4. Alterations in cytoskeletal polypeptide composition induced by the metabolism of menadione in mammalian cells. Hepatocytes (A), GH_3 cells (B), HeLa cells (C), and McCoy cells (D) were incubated for 60 min in the absence or presence of 200 μ M menadione. The cytoskeletal fraction was then extracted and analyzed by PAGE (80–100 μ g protein) under non-reducing (left panel) or reducing (right panel) conditions. Numbers between the two panels refer to the molecular mass (kDa) of protein standards. Arrows indicate a protein band corresponding to actin. β -ME: β -mercaptoethanol; Menad.: menadione.

was most pronounced in GH₃ and McCoy cells but was also apparent in hepatocytes and HeLa cells. Concomitant with the increase in cytoskeleton-associated protein, there was a marked decrease in free sulfhydryl groups in this fraction (Fig. 3). Both the increase in cytoskeleton-associated protein and the decrease in thiol groups occurred well before cell killing.

When the polypeptide composition of the cytoskeleton from control and menadione-treated cells was analyzed by means of PAGE, several major changes appeared, depending on the cell type (Fig. 4, left panel). However, two of these alterations were consistently present in all four cell types, namely, a decrease or disappearance, of a protein migrating like actin (44 kDa) and the concomitant appearance of high-molecular weight aggregates which did not enter the gel. Pretreatment of the same samples with the thiol reductant β -mercaptoethanol prevented both the disappearance of the 44 kDa polypeptide and the formation of protein aggregates (Fig. 4, right panel), suggesting that the latter contained actin molecules cross-linked by disulfide bonds. The identification of the 44 kDa polypeptide was achieved by immunoblotting with monoclonal anti-actin antibodies. As shown in Fig. 5, the cytoskeletal fraction extracted from menadione-treated cells did not react with anti-actin antibodies when analyzed under non-reducing conditions. However, prior

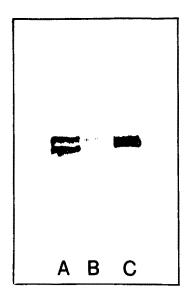


Fig. 5. Immunological identification of actin in the cytoskeletal fraction isolated from control and menadionetreated hepatocytes. Hepatocytes (106 cells/ml) were incubated without (lane A) or with (lanes B and C) 200 μM menadione for 60 min. The cytoskeleton was then extracted and analyzed by PAGE under non-reducing (lanes A and B) or reducing (lane C) conditions. Proteins were then transferred to a nitrocellulose sheet and actin identified using monoclonal anti-actin antibodies, as described in Methods. The presence of two different bands of 44 and 42 kDa molecular mass, reacting with the anti-actin antibodies, in lane A (control cells, non-reduced sample) was due to spontaneous, partial autooxidation of actin. Reduction of the sample with β -mercaptoethanol led to the recovery of a single band migrating in the 44 kDa region (not shown).

reduction of the cytoskeletal fraction from menadione-treated cells with β -mercaptoethanol, led to the recovery of a protein reacting with the anti-actin antibodies.

The data reported in this study demonstrate that the metabolism of menadione in cultured mammalian cells can induce morphological and cytoskeletal changes comparable to those observed in isolated hepatocytes [8]. Our findings suggest that a common mechanism may be responsible for the morphological and cytoskeletal alterations seen in mammalian cells exposed to oxidative stress. A direct relationship between the cytoskeletal alterations and surface blebbing in menadione-treated cells is also strongly indicated by the protective effects of thiol reductants against both surface blebbing and cytoskeletal abnormalities [8]. Thus, it appears that the oxidation of thiol groups in actin is the predominant mechanism underlying surface blebbing in mammalian cells exposed to oxidative stress. Although other mechanisms, e.g. ATP depletion [25] and the activation of Ca²⁺-dependent enzymes acting on the cytoskeleton [26, 27], may contribute to cytoskeletal alterations during toxic cell injury, they probably play a less important role in oxidative cell damage.

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