Oxidation Process of Adrenaline in Freshly Isolated Rat Cardiomyocytes: Formation of Adrenochrome, Quinoproteins, and GSH Adduct

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High concentrations of circulating biogenic catecholamines often exist during the course of several cardiovascular disorders. Additionally, coronary dysfunctions are prominent and frequently related to the ischemic and reperfusion phenomenon (I/R) in the heart, which leads to the release of large amounts of catecholamines, namely adrenaline, and to a sustained generation of reactive oxygen species (ROS). This work aimed to study the toxicity of adrenaline either alone or in the presence of a system capable of generating ROS [xanthine with xanthine oxidase (X/XO)], in freshly isolated, calcium tolerant cardiomyocytes from adult rats. Studies were performed for 3 h, and cardiomyocyte viability, ATP level, lipid peroxidation, protein carbonylation content, and glutathione status were evaluated, in addition to the formation of adrenaline’s oxidation products and quinoproteins. Intracellular GSH levels were time-dependently depleted with no GSSG formation when cardiomyocytes were exposed to adrenaline or to adrenaline with X/XO. Meanwhile, a time-dependent increase in the rate of formation of adrenochrome and quinoproteins was observed. Additionally, as a new outcome, 5-(glutathion-S-yl)adrenaline, an adrenaline adduct of glutathione, was identified and quantified. Noteworthy is the fact that the exposure to adrenaline alone promotes a higher rate of formation of quinoproteins and glutathione adduct, while adrenochrome formation is favored where ROS production is stimulated. This study shows that the redox status of the surrounding environment greatly influences adrenaline’s oxidation pathway, which may trigger cellular changes responsible for cardiotoxicity.

Introduction

An estimated one-third of total global deaths result from the various forms of cardiovascular diseases, ischemic heart disease being the main cause (1). Although they were considered diseases of developed countries, since half of all deaths in the United States are related to cardiovascular diseases (2), a new reality is emerging, and it is estimated that by 2010 cardiovascular diseases will be the leading cause of death in developing countries (3).

When a stressful stimulus occurs, adrenaline and noradrenaline are released, throughout the nervous system and adrenal medulla (3). The cardiac sympathetic nerves are preferentially stimulated in severe heart failure, which includes a 50-fold increase in the rate of spillover of noradrenaline, but also a large release of the sympathetic co-transmitters, adrenaline and neuropeptide Y (4). During an ischemic phenomenon, the concentrations of noradrenaline and adrenaline rise progressively in the interstitial myocardial fluid (3, 5). Elevated concentrations of circulating catecholamines are also found in arrhythmias, myocardial necrosis (3, 5), heart failure (4), exercise (6), pheochromocytoma (7), hypoglycemia, hemorrhagic hypotension, circulatory collapse, and distress (8). Additionally, catecholamines have been widely used for decades in cardiovascular therapy (adrenaline has been used for cardiopulmonary resuscitation for more than 100 years) (9, 10).

Ischemic and reperfusion phenomenon (I/R), a coronary dysfunction, is associated with oxidative stress. In fact, in postischemic myocardium, ROS (1) are formed at an accelerated rate (11–13), playing a major role in pathogenesis. Cardiac myocytes, endothelial cells, and infiltrating neutrophils contribute to ROS production (2). High levels of ROS are also found in exercise (14), and in many pathological conditions, such as inflammation (15, 16), neurodegeneration (17), and aging (18).

Although the toxicity of catecholamines is mainly related to the stimulation of adrenoreceptors, there is a growing body of evidence which shows that their oxidation is also responsible...
for cardiotoxicity (13, 19, 20). The oxidation of catecholamines at physiological pH seems to occur very slowly; however, it increases considerably by enzymatic or metal catalysis (19, 21, 22) or in the presence of superoxide anion (O$_2^-$) (23). Therefore, oxidation of adrenaline in vivo is possible in some circumstances, as in I/R. In accordance, products of this oxidation have been described in the heart, skeletal muscle, liver, and blood (20). The oxidative pathway of catecholamines generates various highly reactive intermediaries, like o-quinones, aminochromes, aminolutins, and melanins (19), which can react with external nucleophilic groups, especially SH groups, present in cysteine, glutathione, and proteins (24–26) or OH and NH$_2$ groups also in proteins (27). For instance, adrenochrome is capable of inhibiting the activity of several enzymes (19, 28), thus greatly modifying cellular metabolism (20). The interaction of adrenochrome with SH groups and the induced depletion of oxygen, ascorbate, and glutathione may cause noxious effects toward the cellular function and defenses (28).

In addition, the conjugation of oxidation products of catecholamines with GSH is no longer considered a simple detoxification route. In fact, these adducts have been reported to induce toxicity. Actually, they seem to be involved in the detoxification route. In fact, these adducts have been reported in the presence of ROS.

In the X/XO system was used to mimic an oxidative stress environment. The studies on tolerant calcium cardiomyocytes were performed for a maximum period of 3 h, and cellular viability, oxidation products of adrenaline, glutathione status, quinoproteins, and glutathione adduct were evaluated during the time course of the experiments to determine the mechanistic pathway of adrenaline’s oxidation when alone and compare it with that in the presence of ROS.

Materials and Methods

Animals. Adult male Sprague-Dawley rats (Charles River Laboratories, Barcelona, Spain) weighing 250–350 g were used. The animals were housed in cages with a temperature- and humidity-controlled environment. Food and water were provided ad libitum, and animals were subjected to a 12 h light–dark cycle. Animal experiments were licensed by the Portuguese General Directory of Veterinary Medicine. Housing and experimental treatment of the animals were in accordance with the Guide for the Care and Use of Laboratory Animals from the Institute for Laboratory Animal Research (ILAR 1996). The experiments complied with current Portuguese laws.

Chemicals. All reagents used in this study were of analytical grade. Collagenase type II was obtained from Worthington (Lake-wood, NJ). Collagenase (type IA), bovine serum albumin (fraction V), N-(2-hydroxyethyl)piperazine-N-(2-ethanesulfonic acid) (HEPES), reduced glutathione (GSH), oxidized glutathione (GSSG), glutathione reducase (GR, EC 1.6.4.2), 2-phenylpyridine, reduced β-nicotinamide adenine dinucleotide phosphate (β-NADPH), reduced β-nicotinamide adenine dinucleotide (β-NADH), 5,5-dithiobis(2-nitrobenzoic acid) (DTNB), adenosine triphosphate (ATP), pyruvic acid, phenylmethanesulfonyl (PMSF), nitroblue tetrazolium (NBT), 1-chloro-2,4-dinitrobenzene, 1-octanesulfonic acid, mushroom tyrosinase, γ-glutamyltranspeptidase (γ-GT, EC 2.3.2.2), luciferase, luciferin, and all reagents for enzymatic determinations were obtained from Sigma-Aldrich (St. Louis, MO). Citric acid, methanol (gradient grade), perchloric acid, and all other chemicals were purchased from Merck (Darmstadt, Germany).

5-Glutathion-S-yladrenaline Synthesis and RMN Analysis. The synthesis of the adduct of adrenaline with GSH followed previously published methods (34) for other catecholamine–GSH adducts.

To a solution of (−)-adrenaline (0.010 g, 5.46 × 10^{-3} mol) in sodium phosphate buffer (20 mL, pH 7.4, 50 mM) at 25 °C was added mushroom tyrosinase (4000 units, 200 units/mL of buffer). The solution became red, indicating the formation of o-quinone. GSH (0.0335 g, 1.09 × 10^{-3} mol) was added, and the red-colored solution changed with time to yellow (19 h). At the terminus of the reaction, 1 mL of 88% formic acid was added, and the solution was carefully concentrated by rotary evaporation without heating. The purification of the product was performed by reverse-phase RP-18 modified silica column chromatography (Merck KGaA) first with water (150 mL) and then 10 × 10 mL of 10% methanol followed by 10 × 10 mL of 20% methanol. Each fraction was checked for the presence of adduct using a UV-vis detector. From the fraction containing adrenaline at 260 and 292 nm were separated and carefully evaporated to dryness. Due to the fact that glutathione was difficult to separate from the adrenalin adduct, further purification by HPLC was necessary using a LiChrospher 100 RP-18 column (Merck KGaA), with two mobile-phase solvents. Solvent A was prepared by adding concentrated trifluoroacetic acid (TFA) to deionized water until the pH reached 2.5. Solvent B was prepared by adding TFA to a 1:1 mixture of acetoniitre (MeCN) and deionized water until the pH reached 2.6. The mobile phase was made of solvents A and B, and the following gradient was used: 100 to 85% solvent A from 0 to 25 min, 85 to 65% solvent A from 25 to 32 min, 65 to 0% solvent A from 32 to 37 min, and 100% solvent A from 37 to 42 min. The compounds eluted within 10 min. The peaks were monitored at 290 nm. 5-(Glutathion-S-yladrenaline (0.006 g) was obtained as an oil in 22.5% yield: 1H NMR (D$_2$O) δ 2.07 (2H, m, Glu-β), 2.41 (2H, m, Glu-γ), 2.67 (3H, s, N-CH$_3$), 3.14 (3H, m, Cys-β), 3.28 (1H, m, Cys-β), 3.75 (1H, m, Gly-α), 3.91 (2H, m, Glu-α), 6.83 (1H, s, ArH$_2$/6), 6.93 (1H, s, ArH$_2$/6); MALDI-TOF (sinapinic acid) m/z 489.5 [M+H]$, 515.5 [M+Na$]^{-2}$.

Calcium Tolerant Cardiomyocytes Isolated from Adult Rat. Calcium tolerant cardiomyocytes were isolated from Langendorff retro perfusion of adult rat heart as previously described (35, 36), with some modifications. The procedure was based on (i) successive treatments with calcium free medium and (ii) digestion with collagenases (collagenase type II and collagenase type IA) in a 200 μM calcium-modified Krebs–Henseleit buffer solution I followed by (iii) gentle mechanical disaggregation. Calcium tolerant cardiomyocytes were obtained by gradual re-introduction of calcium until a final concentration of 1 mM. All steps were performed in modified Krebs–Henseleit buffer containing 102 mM NaCl, 4 mM KCl, 1 mM MgSO$_4$, 10 mM glucose, 5.5 mM NaHCO$_3$, 0.9 mM KH$_2$PO$_4$, and 22 mM HEPES (pH adjusted to 7.2–7.4) saturated with a gaseous stream of carbogen (95% O$_2$ and 5% CO$_2$). At the beginning of the experiments, cell viability was always greater than 60%, evaluated by the lactate dehydrogenase (LDH) leakage assay and by microscopic evaluation of cardiomyocyte morphology. The obtained viability is in accordance with previous reports for calcium tolerant cardiomyocytes (37–39). This viability was obtained after a 5 min preincubation at 37 °C to guarantee the correspondence between values obtained via manual counting and a LDH leakage assay. Incubations were performed in a water bath at 37 °C, using a density of 2.5 × 10$^5$ viable cells/mL in the modified Krebs–Henseleit buffer supplemented with 1 mM CaCl$_2$ (pH 7.4) and saturated with an air stream of carbogen, every hour. After a preincubation for 30 min at 37 °C, the compounds were tested using the following protocol: (i) control cells, with no treatment; (ii) cells incubated with adrenaline (ADR) alone; (iii) cells incubated with ADR and X/XO; and (iv) cells exposed to the X/XO system alone.
The final concentrations were as follows: 0.5 mM ADR (unless otherwise mentioned), 0.1 mM xanthine, and 0.01 unit/mL xanthine oxidase.

**Sample Treatment.** At incubation times of 0, 1, 2, and 3 h, determinations were performed directly using the cardiomyocyte suspension or after centrifugation at 18g for 2 min, for separation of supernatant and pellet, as previously described (38). The supernatant is termed the incubation medium. The pellet was washed two times with 1 mL of modified Krebs–Henseleit buffer supplemented with 1 mM CaCl₂, centrifuged at 18g for 2 min, and finally treated according to the chemical and biochemical determinations for cardiomyocytes. Washing solutions obtained after centrifugation were rejected.

**Cell Viability Assays. 1. Lactate Dehydrogenase Leakage Assay.** The LDH leakage assay was directly performed in the cardiomyocyte suspensions to evaluate the level of cell injury at time zero (immediately after addition of the compounds) and after incubation for 3 h in all treatments, as previously described (38).

**2. Morphology.** The percentage of rod-shaped cells was determined using a Neubauer chamber, as previously described (40). Cells with a length/width ratio of < 4 were considered rod-shaped cells.

**Measurement of Total GSH (GSH), GSH, and GSSG Levels.** The cardiomyocyte and incubation medium levels of GSH and GSSG were measured by the DTNB–GSSG redutase recycling assay, as previously described (41). Both the incubation medium and cardiomyocytes were acidified to a final concentration of 5% HClO₄ and centrifuged, and the supernatant obtained was used for the measurements.

**Determination of GPx, GR, and GST Intracellular Activities.** For the determination of glutathione peroxidase (GPx), glutathione reductase (GR), and glutathione S-transferase (GST) activities, aliquots of the cell suspension were sonicated for 12 s at intensity 4 in a VibraCell sonicater (Sonics & Materials Inc., Danbury, CT) and then centrifuged at 16000g for 10 min. The GPx, GR, and GST activities were determined in the supernatant that was obtained, as previously described (42). Briefly, GR activity was determined by following NADPH oxidation at 340 nm during the reduction of GSSG to GSH. Selenium-dependent GPx activity was determined by following NADPH oxidation at 340 nm after reduction of GSSG by GR. Finally, GST activity was determined by following the formation of the GST conjugate with 1-chloro-2,4-dinitrobenzene, which was monitored at 340 nm. All measurements were performed in triplicate in a 96-well plate reader.

**Measurement of Cellular and Extracellular ATP Levels.** The incubation medium and cardiomyocytes were acidified to a final concentration of 5% HClO₄ and centrifuged, and the supernatant obtained was used for the measurements. The level of ATP was measured by the bioluminescence test based on the work of De Luca et al. (43).

**Assessment of Protein Carbonylation.** Protein carbonyl groups were quantified in cardiomyocytes, as described by Levine et al. (44), by reaction with 2,4-dinitrophenylhydrazine (DNPH).

**Protein Determination.** The protein levels were determined as previously described by Lowry (45). Protein content for quinoproteins was determined by the method described by Bradford (46).

**Assessment of Lipid Peroxidation.** The extent of lipid peroxidation in suspension cells was measured by the assay for thiobarbituric acid reactive substances (TBARS) at 535 nm, as previously described (42).

**Assessment of Protein-Bound Quinones (Quinoproteins).** For the assessment of protein–bound quinones in cardiomyocytes, the NBT/glycinate colorimetric assay was performed, which is based on the method described by Paz et al. (47), with a few adaptations. The cardiomyocytes were lysed in 200 μL of ice-cold RIPA buffer, supplemented with 5 mM PMSF. The samples were sonicated at intensity 4 in the VibraCell sonicator for 3 s, and the whole protein content was quantified by the Bradford method (46), using BSA as a protein standard. Twenty-five micrograms of the lysates in RIPA buffer was added to 240 μL of a 2 M potassium glycinate (pH 10) solution. To this last solution was added 500 μL of NBT reagent [0.24 mM NBT in 2 M potassium glycinate (pH 10)]. The reaction was performed for 3 h at room temperature in the dark, after which the absorbance was read at 530 nm in a 96-well plate reader.

**HPLC-DAD/EC Analysis.** Adrenochrome and adrenaline in the incubation medium were quantified by HPLC (Waters model 2690) with a photodiode array detector (DAD), at 279 nm (for catecholamine) and 300 nm (for aminochrome), as previously described (40).

**Statistical Analysis.** Results are given as means ± the standard deviation (SD) from six independent experiments with suspensions of cardiomyocytes proceeding from six different rats. Nonparametric tests were used. Statistical comparisons between groups were performed with a Kruskal–Wallis test (one-way ANOVA on Ranks) followed by the Student–Newman–Keuls post hoc test, once a significant p had been obtained.

When only two treatment groups were compared, the Mann–Whitney Rank Sum test was used. Details of the statistical analysis are described in each figure legend. Significance was accepted at p values of < 0.05.

**Results**

**Changes in the Concentration of Adrenaline and Adrenochrome.** The measurement of adrenaline and adrenochrome levels was performed in cardiomyocytes and incubation medium of cell suspensions incubated with ADR or with ADR and X/XO. In control or X/XO cells, no adrenaline or adrenochrome was detected.

In cardiomyocytes after incubation for 3 h, a significantly higher concentration of adrenaline was found in the ADR group (12 ± 1 nmol of ADR/2.5 × 10⁵ cardiomyocytes) when compared with the group with ADR and X/XO (8 ± 1 nmol of ADR/2.5 × 10⁵ cardiomyocytes).
In the incubation medium, adrenaline was quantified each hour during the 3 h incubation. The concentration of adrenaline decreased rapidly from its initial value of 0.5 mM in both treatments, although more rapidly in suspensions exposed to ADR and X/XO (Figure 1A). The difference between treatments was statistically significant as early as 1 h. After incubation for 3 h, the concentration of adrenaline was 321(52 and 222(80 µM in suspensions with ADR and with ADR and X/XO, respectively.

In the incubation medium, the decrease in adrenaline levels was accompanied by a time-dependent increase in the adrenochrome levels. This result was more evident in the group with ADR and X/XO, where values of adrenochrome reached 155 µM, at 3 h (Figure 1B).

Adrenaline Oxidation Products Bind to Intracellular Proteins. The protein-bound quinone products (quinoproteins) present in cardiomyocytes were evaluated hourly, for all the treatments (Figure 1C). The measurements showed no significant differences between control cells and the X/XO group, at all time points (data not shown). Moreover, in the presence of ADR and ADR with X/XO, quinoprotein levels increased steadily in a time-dependent manner (Figure 1C). Quinoprotein levels in the ADR cells at 2 and 3 h were 2 and 3 times higher, respectively, than in control. In the cells with ADR and X/XO, at 3 h, the quinoprotein content was double control levels.

Alteration in Glutathione Status. In Figure 2, the levels of GSHt (Figure 2A), GSH (Figure 2B), and GSSG (Figure 2C) in cardiomyocytes, during the time course of a 3 h incubation in control, ADR, and ADR and X/XO cells, can be observed. There were no significant differences between the X/XO system as an isolated treatment, when compared with control (data not shown). Results are presented as means ± SD from six different experiments. Statistical comparisons were made using Kruskal–Wallis test, followed by the Student–Newman–Keuls post hoc test: * p < 0.05 and ** p < 0.01 vs control; and # p < 0.05 and ## p < 0.01 for ADR vs ADR and X/XO groups.
differences were registered among the groups when compared to control groups (data not shown), at a maximum incubation of 3 h.

The sampling homogeneity was confirmed by protein levels, which remained fairly constant at all experimental times in each assay (data not shown).

**Adrenaline Conjugated with Glutathione: Formation of 5-(Glutathion-S-yl)adrenaline.** The depletion of GSH and GSSG formation may be explained by the conjugation between GSH and oxidation products of adrenaline. Thus, cardiomyocytes and incubation medium samples were collected and analyzed by HPLC-EC for GSH–adrenaline adduct detection, by a previously described method (48).

The 5-(glutathion-S-yl)adrenaline adduct was found in cardiomyocytes and in the incubation medium in ADR group and in ADR and X/XO group. The chromatographic peak was confirmed as being 5-(glutathion-S-yl)adrenaline after co-elution with a 5-(glutathion-S-yl)adrenaline standard. The sample was also injected with and without addition of the 5-(glutathion-S-yl)adrenaline standard, showing that the peaks have the same retention time (Figure 3A). Additionally, the chromatographic peak of this adduct completely disappeared when samples (cardiomyocytes and incubation medium) were treated with γ-GT, which can be observed in Figure 3B. Treatment with γ-GT showed no interfering peaks (data not shown).

The levels of 5-(glutathion-S-yl)adrenaline were evaluated, and a higher level of formation of the GSH adduct was observed in cells incubated for 3 h with ADR [378 ± 48 pmol of 5-(glutathion-S-yl)adrenaline/2.5 × 10^5 cardiomyocytes] when compared with ADR and X/XO [115 ± 32 pmol of 5-(glutathion-S-yl)adrenaline/2.5 × 10^5 cardiomyocytes] (Figure 4), which represents an increase of more than 3-fold in the first group compared to the latter.
heart, and an extensive uptake of adrenaline from the plasma occurs in some pathologic conditions (8). Concentrations are expected to be higher in the heart tissue than the ones reported in the plasma, where the rate of clearance is higher.

In particular, it has been shown that, during heart ischemia, the release of catecholamines becomes non-exocytotoxic and is thought to involve the uptake of carrier-mediated efflux in reverse of its normal transport direction (5), largely increasing the concentration of those biogenic amines in the interstitial space, where they can reach values of 4.7 nM (5). Furthermore, sympathetic neurons can take up adrenaline from circulation and release it upon stimulation in the heart (5). When the enzymes responsible for their catabolism (monoamine oxidase and catechol-o-methyltransferase) are unable to cope efficiently, the catecholamine levels in the heart rise and the catecholamine can undergo oxidation, which is catalyzed by trace metals and several enzymes (19, 21, 22). The oxidation pathway of adrenaline involves multiple steps (Figure 5). Adrenaline may be converted to an unstable o-semiquinone that, after deprotonation and loss of a second electron, gives rise to the corresponding o-quinone. At physiological pH, partial deprotonation of the amine group of the side chain of adrenaline leads to an irreversible 1,4-intramolecular cyclization, a reaction that occurs through nucleophilic attack of the nitrogen atom at the 6 position of the quinone ring, to give leucoadrenochrome, which is then further oxidized to adrenochrome (19, 51). In vivo, this oxidation pathway may be more complex, since other factors, such as metal ions or other nucleophilic groups, can be involved (24).

Oxidative stress is a condition in which pro-oxidant metabolites exert toxic effects due to their enhanced production and/or an exhaustion in cellular protection mechanisms (52). There is a growing body of evidence which shows that prolonged ischemia reduces the defense mechanisms within the heart against free radicals (53). Meanwhile, during reperfusion, ROS can be formed above the neutralizing capability of the cells, playing a major role in the pathogenesis of post-ischemic reperfusion injury (11).

Since ROS are formed in I/R phenomena in vivo, an \( \mathrm{O}_2^{\cdot-} \)-generating system can be used to mimic an oxidative stress condition (24, 54). In our work, X/XO was applied to freshly isolated cardiomyocytes, since it is a system suitable for generating ROS. This system was used to mimic the in vivo...
The work of Miyazaki et al. (25) using tyrosinase null mice strengthens our hypothesis. Tyrosinase present in wild mice induces rapid oxidation of dopamine to form stable melanin, with dopamine quinone as an intermediate. In contrast, in tyrosinase null mice, the values for quinoproteins are higher, since, in these animals, quinones are more stable (25).

In our work, levels of quinoproteins in ADR cells are superior to those treated with ADR and X/XO (Figure 1C). These results suggest that the stability of the quinone intermediate is higher in the absence of ROS, allowing the quinone to further react with cellular groups, while in the presence of ROS, adrenaline is rapidly converted into more oxidized species, namely, adrenochrome and its metabolites. Quinones, by themselves, are reported to induce cytotoxicity, immunotoxicity, and carcino-genesis in vivo (61, 62). The mechanisms of toxicity are diverse, depending on their chemical structure and the cellular environment in which they are formed. In addition, alklylation of cellular nucleophiles (GSH, proteins, and DNA) by these species may occur to a significant extent, forming covalent adducts that can significantly compromise cellular integrity and function (25, 38, 61). Thus, the reactivity of adrenaline toward glutathione was evaluated. We observed a time-dependent decline in the level of GSH in cells exposed to ADR, which was potentiated in cells treated with ADR and X/XO (Figure 2). In both groups, the decrease in GSH levels was not accompanied by changes in the activities of the enzymes involved in GSH metabolism or in the GSSG levels, leading to the hypothesis of the occurrence of GSH conjugation with electrophilic compounds. The depletion of intracellular GSH as a result of the catecholamine oxidation process has been previously described (28, 36, 40), and it increases the cellular vulnerability to further oxidative injury. This fact can be of relevance during cardiopathologic events in humans, since GSH is the major nonproteic antioxidant present in the cells and reacts with several electrophilic compounds.

Although adducts of GSH and dopamine or catechol metabolites of MDMA have already been described (24, 41), until now no GSH–adrénaline adducts had been reported in cells. The rate of cyclization of adrenaline o-quinone was probably considered to be too fast for that to occur (27). We report, for the first time, the detection of 5-(glutathion- S-yl)adrénaline in cardiomycocytes, even for adrenaline concentrations as low as 25 µM.

The o-quinone formed during the adrenaline oxidation process (51), in the presence of GSH, may conjugate to form glutathio- nyl adducts (63) (Figure 5). The catechol thioether formed by the addition of the sulfur atom to the quinone ring (62) may be much more redox active than an unsubstituted quinone (34, 62, 64). This capacity to further oxidize may improve their ability to undergo a redox cycle (62), which contributes to the cytotoxicity of these adducts (24, 29, 61, 64). GSH conjugates were found to covalently bind with DNA, raising the question of their ability to cause carcinogenicity (62, 64). They were also reported as nephrotoxic (30, 62) and neurotoxic (24, 31, 62) compounds.

The measurement of these compounds (adduct of glutathione and quinoproteins) is an index of oxidation of the parent catechol, but also a direct measurement of nucleophilic modification. The selective increase in the levels of the GSH–adrénaline adduct and quinoproteins in the groups exposed to ADR suggest that these reactions are favored where quinone is more stable (19, 25). This was the case for the levels of the GSH adduct in the cells, but also in the incubation medium. The high values of 5-(glutathion- S-yl)adrénaline in the incuba-
tinction medium of both groups exposed to adrenaline may suggest a possible mechanism of GSH conjugate efflux in cardiomyocytes (65).

In conclusion, our study has provided several lines of evidence suggesting that an increased rate of adrenaline oxidation is associated with cardiotoxicity and that the adrenaline oxidation pathway depends greatly on the surrounding medium. The catecholamine oxidation products are able to form covalent bonds with the cellular nucleophilic groups (belonging to either GSH or macromolecules). Adrenaline-induced modifications to the protein structure may result in alteration of the cellular rate of cell death were not observed in this work. The findings of our work may prove to be important in clarifying adrenaline-induced toxicity but also in highlighting its oxidation pathway in a cellular medium and the influence of the microenvironment where it occurs.

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