Catalase-overexpressing Thymocytes Are Resistant to Glucocorticoid-induced Apoptosis and Exhibit Increased Net Tumor Growth

Margaret E. Tome, Amanda F. Baker, Garth Powis, Claire M. Payne, and Margaret M. Briehl

ABSTRACT

Glucocorticoids are used for the treatment of lymphoid neoplasms, taking advantage of the well-known ability of these compounds to cause apoptosis in lymphoid tissues. Previously, we have shown that dexamethasone, a synthetic glucocorticoid, causes a down-regulation of several antioxidant defense enzymes and proteins, including catalase and thioredoxin, concomitant with the induction of apoptosis in WEHI7.2 mouse thymoma cells. To test whether this down-regulation plays a critical role in the mechanism of steroid-induced apoptosis, WEHI7.2 cells were transfected with rat catalase. Two clones, expressing 1.4-fold and 2.0-fold higher catalase specific activity, respectively, when compared with vector-only transfectants were selected for further study. An increase to 1.4-fold parental cell catalase activity delayed cell loss after dexamethasone treatment, whereas a 2.0-fold parental cell catalase activity prevented dexamethasone-induced cell loss for 48 h after treatment. Dexamethasone treatment of the WEHI7.2 cells stimulated a release of cytochrome c into the cytosol. Catalase-overexpressing cells showed a delay or lack of cytochrome c release from the mitochondria, which correlated temporally with the delay or prevention of cell loss in the culture after dexamethasone treatment. A decreased amount of cell death from WEHI7.2 cells overexpressing catalase was also seen in tumor xenografts in severe combined immunodeficient mice when compared with tumors from vector-only transfected cells. Similarly, thioredoxin-overexpressing WEHI7.2 cells, shown previously to be apoptosis resistant, showed decreased cell death in tumor xenografts. This resulted in larger tumors from cells overexpressing these proteins.

INTRODUCTION

Apoptosis of lymphoid tissues attributable to glucocorticoid treatment is a classic example of apoptosis cited in the early descriptions of apoptotic morphology (1). This process is important physiologically, especially for the sorting of maturing lymphocytes (2). Clinically, this feature of glucocorticoids has been exploited for the treatment of leukemia and lymphoma; unfortunately, as with many chemotherapeutic agents, the development of drug resistance is a problem (3–5). Glucocorticoids exert their action primarily by binding to a cytosolic receptor that translocates to the nucleus and affects transcription (6). Lack of functional receptors desensitizes lymphocytes and thymocytes to glucocorticoid-induced apoptosis both in tissue culture cell lines and animal models (7–11). In some patients, drug resistance can be traced to either a receptor mutation or a decrease in functional glucocorticoid receptor number (3, 5); however, in other individuals resistance cannot be traced to glucocorticoid receptor alterations (12). The events that occur after translocation of the receptor/steroid complex to the nucleus and prior to the commitment of the cell to die are not understood. Elucidation of the mechanism by which glucocorticoids kill lymphocytes may make it possible to target alternative steps in the process for chemotherapeutic intervention, particularly when drug resistance occurs.

The necessity for functional glucocorticoid receptors and RNA synthesis (13, 14) during steroid-induced lymphocyte apoptosis suggests that identifying alterations in mRNA transcripts may provide insight into critical steps required for this process. Previously, we found that treatment of WEHI7.2 mouse thymoma cells with dexamethasone, a synthetic glucocorticoid, causes down-regulation of a number of antioxidant defense enzymes (15, 16). Glutathione peroxidase and catalase, which metabolize cellular H2O2; superoxide dismutase, which detoxifies O2− and thioredoxin, a small protein involved in producing reducing equivalents, are all down-regulated at the mRNA level (16). Concomitant with the antioxidant enzyme down-regulation is an increase in the mRNA for glutathione S-transferase (15), an enzyme thought to be involved in the removal of lipid peroxidation products (17). This suggests a model whereby steroids down-regulate cellular antioxidant defenses, which decreases the ability of the cell to detoxify ROS2 produced via normal metabolism and results in apoptosis.

There is some evidence that ROS may be involved in steroid-induced apoptosis of lymphocytes. Exogenous treatment of thymocytes with chemical antioxidants, antioxidant enzymes, or metal chelators protects against dexamethasone-induced apoptosis (16, 18–23). Culture of thymocytes under hypoxic conditions generally affords protection from dexamethasone (16, 22, 24). Furthermore, lipid peroxidation is seen after glucocorticoid treatment in thymocytes (25) and S49.1 lymphocytes (26), suggesting that oxidative damage occurs during steroid-induced apoptosis in these cells. Finally, WEHI7.2 cells overexpressing thioredoxin are resistant to steroid-induced apoptosis (27).

To determine whether the previously observed antioxidant defense enzyme down-regulation is critical during steroid-induced lymphocyte apoptosis, we have isolated two WEHI7.2 catalase transfectants. These cell variants have provided useful tools to test the importance of catalase to the mechanism of steroid-induced apoptosis in vitro and in vivo while helping to elucidate the role of ROS in this process.

MATERIALS AND METHODS

Cell Culture and Transfection. The mouse thymoma-derived WEHI7.2 parental cell line (28) was obtained from Dr. Roger Miesfeld (University of Arizona). Cells were maintained in DMEM-low glucose (Life Technologies, Inc., Grand Island, NY) supplemented with 10% calf serum (Hyclone Laboratories, Logan, UT) at 37°C in a 5% CO2 humidified environment. Cells were maintained in exponential growth at a density between 0.02 and 2 × 106 cells/ml. Stable transfectants of WEHI7.2 cells overexpressing thioredoxin have been characterized previously and were maintained as described (27).

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2 The abbreviation used is: ROS, reactive oxygen species.
Catalase-overexpressing cells were constructed by stably transfecting WEHI.2 cells with a vector containing rat catalase cDNA inserted into the KpnI/Notl sites of the pBK-CMV vector (constructed and kindly donated by Dr. Melissa Williams, University of Texas Health Sciences Center, San Antonio, TX). Cells were transfected, and positive clones were selected as described previously (27). All studies were conducted on clonal lines maintained in 800 μg/ml G418 (Life Technologies, Inc.) and then cultured in medium in the absence of selection drug 1 week prior to each experiment. Immediately before each experiment, live cells were separated from those that underwent spontaneous apoptosis using Ficoll-Plaque Plus reagents and the manufacturer’s suggested protocol (Amersham Pharmacia Biotech, Piscataway, NJ).

Chemicals and Drugs. All chemicals and drugs used in these experiments were obtained from Sigma Chemical Co. (St. Louis, MO) unless otherwise noted.

Glucocorticoid Receptor Number. The number of functional glucocorticoid receptors/cell was determined by using a transient cotransfection of cells with a glucocorticoid response element/chloramphenicol acetyl transferase reporter plasmid and a β-galactosidase expression vector as described previously (27).

Sensitivity to Dexamethasone. Sensitivity to dexamethasone-induced apoptosis in cell culture was measured by treating cells with either 1 μM dexamethasone or an equivalent volume of ethanol vehicle alone (final ethanol concentration, 0.01%). Periodically, an aliquot of cells was removed and resuspended with an equal volume of an Eosin Y solution (500 μg/ml final dye concentration), allowed to stand 2–3 min at room temperature, and counted using a hemocytometer. Cells that excluded dye were scored as viable. To confirm that cell loss in culture after dexamethasone treatment was attributable to apoptosis, an aliquot of cells was prepared for morphological examination as described under microscopy.

Enzyme Activity Assay. Prior to enzyme activity measurements, cells were washed twice at 4°C with PBS, 137 mM NaCl, 2.7 mM KCl, 8.1 mM Na2HPO4, and 1.5 mM KH2PO4, pH 7.4. Cells were resuspended in 10 mM Tris-HCl (pH 7.5), 250 mM sucrose, 1 mM EDTA, 0.5 mM DTT, 0.1 mM phenylmethylsulfon fluoride, and 1% Triton X-100. Samples were incubated for 30 min on ice and centrifuged at 10,000 × g for 15 min at 4°C. Activity was measured in the supernatant fractions as the decrease in H2O2 absorbance at 240 nm as described by Beers and Sizer (29) and calculated using an extinction coefficient of 43.6 M−1 cm−1. Cellular protein was measured using the BCA protein assay kit (Pierce, Rockford, IL) according to the manufacturer’s protocol, and all enzyme activities were normalized to cellular protein.

In Vivo Tumor Growth. Tumor formation in vector-only, thiorodoxin-transfected, and catalase-transfected WEHI.2 cells was compared by injecting 2 × 107 cells in a 0.2-ml Matrigel:0.9% NaCl (50:50) mixture, s.c., into the flanks of female severe combined immunodeficient mice. Catalase and vector-only transfectants were injected into 20 mice, whereas 18 received thiorodoxin-transfected cells. Tumor volume was measured with calipers every 3 days. On the tenth day after tumor cell injection, daily injections of 1 mg/kg body weight dexamethasone in a 10% ethanol:0.9% NaCl vehicle were begun in half the transfectants. Tumor volume was measured with calipers every 3 days. On the tenth day after tumor cell injection, daily injections of 1 mg/kg body weight dexamethasone in a 10% ethanol:0.9% NaCl vehicle were begun in half the mice from each cell variant group. Control mice were injected with vehicle alone. Seventeen days after tumor cell injection, four mice from each cell variant group receiving dexamethasone and four receiving vehicle alone were euthanized, the tumors were excised, and a portion of the tumor was fixed for microscopy. On day 20 after tumor cell injection, the remaining animals were euthanized, and tumors from three mice in each category were weighed.

Microscopy. Portions of the tumors excised on day 17 were fixed in one-half strength Karnovsky’s fixative overnight, postfixed in osmium tetroxide, dehydrated in a graded series of alcohols, and embedded in epoxy resin. One μm-thick sections were prepared and stained with toluidine blue for light microscopy. Cells were scored as apoptotic, mitotic, normal, or para-apoptotic based on morphological features using a ×100 oil immersion objective. For each of six tumors/cell variant, 12 representative fields in nonnecrotic regions were scored. Thus, a total of 72 high power fields were scored for tumors from each cell variant. Ultrathin sections were prepared for electron microscopy, counterstained with 5% uranyl acetate and lead citrate, and examined using a Philips CM transmission electron microscope (FEI, Hillsboro, OR).

Cells from tissue culture were harvested by centrifugation (800 × g for 5 min), washed in PBS, and fixed for 1 h at room temperature in 3% glutaraldehyde in 0.1 M cacodylate, pH 7.2. Subsequently, cells were washed three times in 0.1 M cacodylate (pH 7.2), and the cell pellet was embedded in epoxy resin. One μm-thick sections were prepared as above for light microscopic examination. Apoptotic cells were identified in 1-μm-thick sections (27) using the following characteristics: condensed chromatin, increased nuclear and cytoplasmic stain intensity, cell shrinkage, increased cytoplasmic vacuolization, and apoptotic body formation. Additional features observed by electron microscopy include nuclear segregation, an increase in electron density, and the absence of mitochondrial swelling in cells not showing signs of secondary degeneration (30).

Catalase Antibody Preparation and Immunoblots. Female New Zealand White rabbits were injected i.c. with a 1-ml injection volume containing 363 μg of bovine liver catalase (Amersham Pharmacia Biotech) in water mixed with Freund’s Complete adjuvant. Animals were boosted with a second injection at 3 weeks and a third at 6 weeks using the same catalase emulsified in Freund’s Incomplete adjuvant. Six weeks after the third injection, serum was collected and adjusted to a final concentration of 500 mM NaCl, 50 mM Tris (pH 7.5).

Samples were prepared as for the enzyme activity assay, and 10 μg of protein/sample were loaded and separated on a 10% SDS-PAGE gel (31). Murine liver catalase (12.5 ng) was used as a positive control. Samples were transferred onto polyvinylidene difluoride membranes (MEN Life Science Products, Boston, MA) using a 25 mA current at 4°C overnight in blotting buffer (32). For immunodetection, membranes were first incubated in 5% BSA fraction V in PBS with 0.5% Tween 20 (PBS-T) at room temperature for 1 h. Next, the membranes were washed three times for 5 min each with PBS-T. Primary antibody was diluted 1:2000 in 1% BSA/PBS-T and incubated with the membranes for 1 h. Membranes were again washed, once for 15 min and then five times for 5 min each with PBS-T. Horseradish peroxidase-linked antirabbit immunoglobulin antibody (Amersham Pharmacia Biotech) was diluted 1:10,000 in 1% BSA/PBS-T and incubated with the membranes for 1 h. Then the membranes were washed once for 15 min and five times for 5 min each with PBS-T. Proteins were detected by chemiluminescence using the manufacturer’s protocol (ECL Western Blotting System; Amersham Pharmacia Biotech) and film (X-OMAT Blue; Eastman Kodak, Rochester, NY).

Measurement of Cytochrome c. Cell cultures were treated with 1 μM dexamethasone or vehicle as described above and harvested by centrifugation (1000 × g for 5 min at 4°C). Cell pellets were washed with PBS, and the cytotoxic (≤100) and mitochondrial fractions were prepared according to Vander Heiden et al. (33), except that cells were resuspended in buffer A without sucrose and allowed to sit for 30 min before the addition of sucrose to make a final concentration of 250 mM. Cells were immediately lysed using a glass tissue homogenizer with a glass pestle (Kontes, Vineland, NJ). Proteins from cytosolic and mitochondrial fractions (30 μg of each sample) were separated on a 15% SDS-PAGE gel (31), and proteins were transferred as described above. Blots were probed for cytochrome c using 1 μg/ml anti-cytochrome c antibody (PharMingen; San Diego, CA) and cytochrome c oxidase with 1 μg/ml anti-cytochrome oxidase subunit I antibody (Molecular Probes, Eugene, OR), using the manufacturer’s suggested protocol accompanying the cytochrome c antibody. Proteins were detected by first incubating with a 1:2000 dilution of horseradish peroxidase-linked antimouse immunoglobulin (Amersham Pharmacia Biotech) and detecting the product by chemiluminescence (Renaissance Western Blot Chemiluminescence Reagent Plus; NEN Life Science Products), as suggested by the manufacturer.

Statistics. The tumor growth rate was compared by first fitting a least square regression line to the cube root of tumor volume by day for each mouse. The line slopes were then compared in a two-way ANOVA to test the difference in tumor growth rate between categories. This analysis was performed by the Statistics Core at the Arizona Cancer Center. The rest of the data were compared using Student’s t test assuming unequal variances. Significance was set at P < 0.05.

RESULTS

To verify the increased expression of catalase in the transfectants, catalase activity was measured, and two clones were selected for further study. Catalase activity in the CAT38 clone was ~1.4-fold that in the vector-only transfectants (Neo2 and Neo3), whereas the CAT2 clone exhibited 2.0-fold greater catalase activity (Fig. 1B).
Cellular catalase protein content corresponded to the activity measured in these cells (Fig. 1A). We were unable to select clones expressing >2-fold the parental cell catalase activity. Increased catalase expression was lost if cells were cultured for prolonged periods in the absence of selection drug (data not shown).

Addition of antioxidants to the culture medium protects thymocytes from glucocorticoid-induced apoptosis (16), suggesting that ROS are involved in this cell death. To test whether the increased cellular catalase protects cells from steroid-induced apoptosis, the viable cell number over time after the addition of 1 µM dexamethasone in catalase transfectant cell cultures was compared with that in cultures with vector-only transfected cells (Fig. 2). The CAT38 cells showed a delayed cell loss when compared with the vector-only transfected cells. After a 48-h treatment with dexamethasone, a significantly greater number of viable cells remained in the CAT38 cultures than in either of the two neo-transfectant cultures. CAT2 cells, which have the higher catalase activity, had a decreased cell loss when compared with that in the CAT38 cell cultures. CAT2 cell cultures contained significantly more viable cells than either CAT38 or neo-transfectant cultures 48 h after dexamethasone treatment. In the absence of dexamethasone, cells grew exponentially with similar growth rates (data not shown). Morphological assessment of cells pelleted from these cultures verified that cell loss was attributable to apoptosis (data not shown). On the basis of [3H]dexamethasone binding assays, WEHI7.2 cells contain ~33,000 receptors/cell (34). Our receptor measurements indicated equivalent numbers of functional receptors in the vector-only and catalase-transfected cells (data not shown). Increased catalase appears to correlate well with a decrease in the rate of dexamethasone-induced apoptosis.

Data from caspase knockout mice suggest that glucocorticoids induce apoptosis in thymocytes by stimulating the release of cytochrome c from the mitochondria into the cytosol (35–37). In this apoptotic pathway, release of cytochrome c is considered the committed step and causes activation of the caspases that complete the apoptotic process (38). In WEHI7.2 cells, dexamethasone stimulated the release of cytochrome c into the cytosol (Fig. 3). Twenty-four h after the addition of dexamethasone, cytochrome c is easily detectable in the cytosol in the parental, WEHI7.2, and vector-only transfected Neo3 cells. In the CAT38 cells, which showed a delay in apoptosis, the cytochrome c release is delayed so that only a trace is detectable in the cytosol 24 h after dexamethasone treatment; however, after 32 h in drug, cytosolic cytochrome c is readily detectable. CAT2 cells, with the highest catalase activity and the least cell loss, showed a negligible loss of cytochrome c into the cytosol up to 40 h after dexamethasone treatment. The pattern of cytosolic cytochrome c seen in Fig. 3 was not attributable to contamination of the cytosolic preparation with mitochondria because the cytochrome oxidase subunit I (a mitochondrial specific protein) content of all cytosolic preparations was minimal and similar (data not shown).

Appendix B

Fig. 1. Catalase protein and catalase-specific activity in catalase-transfected (CAT) WEHI7.2 cells compared with that in vector-only transfected (Neo) cells. A, representative immunoblot showing 20 µg of total cellular protein probed with anti-catalase primary antibody. Murine liver catalase is shown as a positive control (lane STD). B, catalase activity measurements corresponding to the clones shown in A. Values represent the mean of triplicate measurements in an experiment that has been replicated; bars, SE. *p , significantly different from vector-only (Neo2 and Neo3) controls. **p , significantly different from Neo2, Neo3, and CAT38.

Fig. 2. Comparison of dexamethasone-induced apoptosis in catalase-transfected and vector-only-transfected WEHI7.2 cells. The viable cell number over time after the addition of 1 µM dexamethasone in cultures containing catalase-transfected clones, CAT2 (■) and CAT38 ( ), or vector-only transfectants, Neo2 (○) and Neo3 ( ), are shown. Values represent the average of triplicate cultures in a representative experiment that has been replicated; bars, SE.

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fected cells showed a significantly increased tumor growth rate when compared with that from the vector only cells (Neo3; Fig. 4C). Tumor xenografts from cells with increased catalase also appeared to grow slightly faster than the Neo3 cell tumors, although this difference was not significant. Unexpectedly, no significant effect of dexamethasone on tumor weight or tumor growth rate was seen. This lack of dexamethasone response may be because the dose of dexamethasone was inadequate or the timing of the doses made it difficult to measure an effect.

Alterations in tumor weight appeared to be primarily attributable to alterations in spontaneous cell death (Fig. 5A). Total cell death was significantly greater in Neo3 tumors when compared with tumors from thioredoxin- or catalase-transfected cells (Fig. 5B). Mitosis was similar in sections of tumors from all three cell variants (data not shown). Most of the cell death observed in the sections from the Neo3 tumors was apoptosis (Fig. 6A, top). In tumors from the thioredoxin- or catalase-transfected cells, an increased amount of cell death, morphologically distinct from apoptosis, was observed (Fig. 6A, bottom, and B). The classic ultrastructural features of apoptosis, including margination of chromatin, nucleolar exclusion, and cell shrinkage as shown in Fig. 7A, were absent from a large portion of the dead cells from thioredoxin or catalase transfectant tumor xenografts. Instead, these dead cells exhibited a type of cell death termed para-apoptosis (39, 40). The para-apoptotic cells had very extensive processes that squeezed between adjacent cells (Fig. 7B) and displayed a network of cytosolic fibers (Fig. 7C), neither of which is a characteristic of apoptotic cells. In contrast to the electron lucency of necrotic cells, both cytoplasm and nucleus of the para-apoptotic cells were very electron dense. The mitochondria were also distended and showed a

Fig. 4. Growth in tumor xenografts from cells transfected with catalase, thioredoxin, or vector only. A, comparison of representative tumors from vector-only (Neo3, NEO), catalase (CAT38, CAT), or thioredoxin (THX6, THX) transfected WEHI7.2 cells 17 days after cell injection. B, average weight of tumor xenografts 20 days after cell injection (n = 6); bars, SE. *, significantly different from Neo3 values. C, mean tumor volume (days 0–17, n = 8; day 20, n = 4) in tumor xenografts from Neo3, CAT38, or THX6 cells; bars, SE. The daily injections, 1 mg/kg dexamethasone (D) or vehicle only (E), were started on day 10.
loss of cristae, features not observed in apoptosis. These para-apoptotic cells appeared very similar to those described as “type B dark cells” (41, 42).

DISCUSSION

Catalase overexpression protects thymocytes from glucocorticoid-induced apoptosis. The degree of protection is proportional to the level of catalase overexpression such that the higher the catalase activity the greater the protection. The normal cellular function of catalase is to detoxify H$_2$O$_2$, resulting in the formation of H$_2$O and O$_2$ (43). Although catalase is a peroxisomal enzyme in most cells, H$_2$O$_2$ is thought to be readily diffusible; therefore, overexpression of cata-

![Fig. 6. Comparison of the cell death morphology in the tumor xenografts. A, light micrographs of representative areas in tumor xenografts from the Neo3 and THX6 cell variants (high power light microscope field). The dark staining cells with extended processes are the para-apoptotic cells in the THX6 micrograph. B, average number of apoptotic (■) and para-apoptotic (□) cells/high power light microscope field (n = 72 fields/cell variant); bars, SE. Abbreviations are as in Fig. 4. *, significantly different from Neo3 values.]

Fig. 5. Comparison of cell death frequency in the tumor xenografts. A, light micrographs (low magnification) showing representative areas in the tumor xenografts from each cell variant. B, average number of dead cells/high power light microscope field (n = 72 fields/cell variant); bars, SE. Abbreviations are as in Fig. 4. *, significantly different from Neo3 values.

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Lase, even in the peroxisomes, should result in decreased intracellular H$_2$O$_2$ (43–45). Our data suggest that H$_2$O$_2$ plays a critical role in glucocorticoid-induced thymocyte apoptosis. The protection provided by the modest increases in catalase activity seen here suggests that even small changes in ROS may be critical. Although ROS measurements using whole-cell populations have suggested that little, if any, increase in ROS occurs (25, 26, 46), separation of the apoptotic populations by flow cytometry (23) or Percoll gradients (20) have shown that small increases in ROS occur in the apoptotic cells after dexamethasone treatment. Indications that ROS might play a critical role come from studies that show that treatment of thymocytes in culture with antioxidant enzymes or chemical antioxidants is protective (16, 18–23). Hypoxic conditions, where lower ROS production is expected, are also generally protective (16, 22, 24). Treatment with divalent metal ion chelators protects thymocytes in culture, suggesting a role for Fenton chemistry in the mechanism of steroid-induced killing (18). A decrease in reduced glutathione (15, 20, 21, 25) and the appearance of lipid peroxidation products (25, 26) are also observed after steroid treatment, suggesting that cells are undergoing oxidative stress during glucocorticoid-induced apoptosis. Dexamethasone induces apoptosis in the WEHI7.2 cells by stimulating the release of cytochrome c from the mitochondria into the cytosol. In this apoptotic pathway, some as yet unidentified series of events (the signal) impinges on the mitochondria to stimulate the release of cytochrome c into the cytosol. Release of cytochrome c, considered the committed step, then triggers the activation of apical caspases. Apical caspases activate effector caspases, which in turn degrade cellular substrates in what is considered the execution phase (38). Cytochrome c release is also thought to trigger the hyperproduction of ROS common to the later stages of apoptosis induced by many agents (21, 47, 48). Overexpression of catalase delays or prevents the release of cytochrome c into the cytosol, suggesting that ROS, in particular H$_2$O$_2$, acts in the signaling phase of glucocorticoid-induced lymphocyte apoptosis. This protection occurs before the committed step and is therefore separable from late-stage ROS hyperproduction. Other studies that have attempted to determine the sequence of apoptotic events have suggested that ROS increases may be an early event in steroid-induced apoptosis. The small increases in ROS seen in two studies (23, 25) and the lipid peroxidation seen by Bustamante et al. (25) appeared shortly after steroid treatment. Work by Macho et al. (21) and Zamzami et al. (49) has also shown that addition of catalase or N-acetylcysteine, an antioxidant and glutathione precursor, to lymphocyte cultures delays the irreversible loss of mitochondrial transmembrane potential and the hyperproduction of ROS. To act as biological signals, molecules need only brief appearances at low concentrations to be effective. This may explain how modest increases in ROS play a critical role during steroid-mediated apoptosis. A number of the signaling pathways known to be redox sensitive or affected by ROS directly (50, 51) are involved in apoptosis. Our data suggest that a closer look at redox-sensitive signaling pathways may help determine the mechanism of steroid-induced apoptosis.

Tumor cell resistance to the chemotherapeutic effects of glucocorticoids may be connected to increased resistance to ROS. Loss of functional glucocorticoid receptors is a cause of resistance in some patients (3, 5); however, resistance in other individuals cannot be traced to glucocorticoid receptor alterations (12). Overexpression of catalase (as shown here) or thioredoxin (27), both part of the cellular antioxidant defense, protects thymocytes from glucocorticoid-induced apoptosis in vitro without alterations in functional receptors. Similarly, WEHI7.2 cells selected for resistance to H$_2$O$_2$ show a bolstered

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Fig. 7. Comparison of ultrastructural features in representative apoptotic and para-apoptotic cells. A, electron micrograph showing a typical apoptotic cell from a Neo3 cell tumor xenograft. Electron micrographs of cell processes (B) and cell bodies (C) of para-apoptotic cells in a THX6 tumor xenograft. ×3000.

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modified Hodgkin and Reed-Sternberg cells of Hodgkin's disease. Hodgkin and Reed-Sternberg cells that exhibit this morphology do not show the DNA fragmentation characteristic of apoptosis or alterations in p53 or Bcl-2 (75, 77). The increased incidence of this type of cell death in tumor xenographs from antioxidant enzyme-overexpressing cells suggests that cellular redox status may play a role in para-apoptosis.

These data lend support to a critical role for antioxidant enzymes in tumorigenesis. The cell variants developed here will provide useful tools to investigate antioxidant enzyme status as a predictor of chemotherapeutic response or target for intervention.

ACKNOWLEDGMENTS

We thank Dr. Melissa Williams for the donation of the catalase transfection vector; Gillian Payne and Jeanne Morreale for technical expertise and help to complete the mouse tumor study; Peggy Krasovich and Gina Zhang for expert preparation of the microscopy sections; Huiyen Cui for statistical analysis; and Dr. David Askew for technical advice.

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*Cancer Res* 2001;61:2766-2773.

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