Original Research Communication

Effects of Acute γ-Hexachlorocyclohexane Intoxication in Relation to the Redox Regulation of Nuclear Factor-κB, Cytokine Gene Expression, and Liver Injury in the Rat

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ABSTRACT

 γ -Hexachlorocyclohexane-induced hepatotoxicity is associated with oxidative stress. We tested the hypothesis that γ-hexachlorocyclohexane triggers the redox activation of nuclear factor-κB (NF-κB), leading to proinflammatory cytokine expression. Liver NF-kB activation (electrophoretic mobility shift assay), tumor necrosis factor- α (TNF- α) and interleukin- 1α (IL- 1α) mRNA expression (reverse transcription-polymerase chain reaction), and their serum levels (enzyme-linked immunosorbent assay) were measured at different times after γ-hexachlorocyclohexane treatment (50 mg/kg). The relationship between these and hepatic O, uptake, glutathione and protein carbonyl levels, and sinusoidal lactate dehydrogenase (LDH) efflux in liver perfusion studies was determined. γ-Hexachlorocyclohexane increased liver NF-κB DNA binding at 14-22 h after treatment, concomitantly with significant glutathione depletion and an increase in the rate of O, consumption, the content of protein carbonyls, and the sinusoidal LDH efflux. In these conditions, the expression of $TNF-\alpha$ and IL-1α is enhanced, with maximal increases in their respective mRNA content and serum levels of the cytokines being elicited at 18 h after γ -hexachlorocyclohexane treatment. All these changes are suppressed by the administration of α-tocopherol (100 mg/kg) or the Kupffer cell inactivator gadolinium chloride (10 mg/kg) prior to γ-hexachlorocyclohexane. γ-Hexachlorocyclohexane-induced TNF-α levels in serum are suppressed by pretreatment with an antisense oligonucleotide (ASO TJU-2755; daily doses of 10 mg/kg for 2 days) targeting the primary transcript for the cytokine, whereas those of IL-1 α are not modified. It is concluded that γ hexachlorocyclohexane-induced liver oxidative stress triggers the DNA binding activity of NF-κB, with the consequent increase in the expression of NF- κ B-dependent genes for TNF- α and for IL-1 α , factors that may mediate the hepatotoxicity of the insecticide. Antioxid. Redox Signal. 6, 471–480.

INTRODUCTION

Y-HEXACHLOROCYCLOHEXANE (lindane) is an important pesticide extensively used for public health and agricultural purposes in developing countries, which induces toxicity in

several tissues, including the liver. γ -Hexachlorocyclohexaneinduced hepatotoxicity is associated with the production of reactive species, including (a) electrophilic γ -hexachlorocyclohexane-derived metabolites generated during the biotransformation of the insecticide by cytochrome P450 (2, 45), (b)

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reactive O2 intermediates, such as superoxide radical due to cytochrome P450 induction (19, 20), and (c) reactive nitrogen species, such as nitric oxide (9). Enhancement of nonspecific free radical formation, measured by chemiluminescence, and superoxide radical generation is also elicited in hepatic microsomes and mitochondria upon addition of γ-hexachlorocyclohexane in vitro (1). γ-Hexachlorocyclohexane-induced free radical activity leads to depletion of hepatic antioxidant stores, including reduced glutathione (GSH) (4) and lycopene (34) levels, and the inactivation of superoxide dismutase and catalase (19). Thus, disruption of the steady-state balance between generation of prooxidant species and their utilization in favor of the prooxidants is established in the liver (45), with the consequent enhancement in lipid peroxidation (1, 14, 19, 20, 31, 48) and protein oxidation (9) indices. In agreement with these findings, γ -hexachlorocydohexane administration to rats increases the total O2 consumption by the liver, an effect that involves both O₂ equivalents related to oxidative stress (44) and uncoupling of mitochondrial oxidative phosphorylation by lindane-derived chlorophenol metabolites (15). Development of oxidative stress by γ-hexachlorocyclohexane coincides with the onset and progression of liver injury (20). These features are readily reversible depending on the hepatic content of the insecticide and on the extent of microsomal superoxide generation (21), and are triggered in different cell types (1, 13, 28) and tissues (27, 32, 37) of experimental animals, plants (36), and also evidenced in the blood from poisoning human cases (3).

Current evidence indicates that oxidative stress is implicated in the genesis of cell injury through redox regulation of the transcription of genes involved in the pathogenesis of diseases (33, 40), in addition to direct mechanisms determining the oxidative damage to essential biomolecules (45). Nuclear factor-kB (NFκB) is a prototype redox-sensitive transcription factor having a prominent role in the regulation of immune and inflammatory genes, apoptosis, and cell proliferation (18). Activation of NF-κB in Kupffer cells is important in the homeostatic response to acute or mild liver injury or as a pathological event underlying chronic liver disease (40). Therefore, the aim of present study was to test the hypothesis that γ -hexachlorocyclohexane-inducedoxidative stress in the liver activates NF-kB, triggering the expression of NF- κ B-dependent gene for tumor necrosis factor- α (TNF- α) and for interleukin- 1α (IL- 1α) at the Kupffer cell level. For this purpose, liver NF-κB DNA binding, hepatic TNF-α mRNA and IL-1 α mRNA contents, and serum TNF- α and IL-1 α levels were determined in control rats and in animals administered γ -hexachlorocyclohexane at different times after treatment. Separate groups of control and γ -hexachlorocyclohexane-treated ats were pretreated with the antioxidant α-tocopherol or the Kupffer cell inactivator gadolinium chloride (GdCl₃) (17), and results obtained were correlated with changes in biochemical parameters related to oxidative stress and liver injury.

MATERIALS AND METHODS

Animals and treatments

Male Sprague–Dawley rats (Bioterio Central, Instituto de Ciencias Biomédicas, Facultad de Medicina, Universidad de Chile) weighing 200–300 g were housed on a 12-h light/dark

cycle and were provided rat chow (Champion S.A., Santiago, Chile) and water *ad libitum*. Animals received a single intraperitoneal injection of 50 mg of γ -hexachlorocyclohexane/kg or an equivalent volume of vehicle (corn oil; controls), and studies were carried out at the indicated times after treatment. Separate groups of rats were subjected to either (a) 100 mg of α -tocopherol/kg i.p. 17 h prior to γ -hexachlorocyclohexane or (b) 10 mg of $GdCl_3$ /kg i.v. 24 h before γ -hexachlorocyclohexane, and studies were performed at the indicated times after γ -hexachlorocyclohexane treatment. Animals used were cared according to the Guide for the Care and Use of Laboratory Animals by the National Academy of Sciences (NIH publication no. 86-23).

Nuclear extracts and electrophoretic mobility shift assay (EMSA)

Nuclear protein extracts were prepared on ice from liver homogenates as described by Deryckere and Gannon (10). In brief, liver homogenates were centrifuged for 30 s at 2,000 rpm and 4°C, and the supernatant was incubated for 5 min and centrifuged for 5 min at 5,000 rpm and 4°C. The pellet was resuspended in a buffer containing 20 mM HEPES (pH 7.9), 25% glycerol, 420 mM NaCl, 1.2 mM MgCl₂, 0.2 mM EDTA, 0.5 mM dithiothreitol, 0.5 mM phenylmethylsulfonyl fluoride, 2 mM benzamidine, and 5 µg/ml of the protease inhibitors pepstatin, leupeptin, and aprotinin, incubated on ice for 20 min, and centrifuged for 30 s at 13,000 rpm and 4°C. The supernatant was stored at -80° C. Protein concentration was determined according to Bradford (6). The samples were subjected to EMSA to assess the amount of active NF-kB involved in protein-DNA interactions (49), using labeled and unlabeled oligonuclectides containing the consensus sequence 5'-GATCTCAGAGGGGACTTTCCGAG-3' (Genset Corp., La Jolla, CA, U.S.A.). The specificity of protein binding in nuclear extracts was confirmed in competition experiments using 100-fold excess of unlabeled probe. To assess the subunit composition of DNA binding protein, specific antibodies were used for supershift assay (43) (goat and rabbit immunoglobulin G raised against NF-κB p50 and p65, respectively; Santa Cruz Biotechnology Inc., Santa Cruz, CA, U.S.A.). Samples were loaded on nondenaturating 6% polyacrylamide gels and run until the free probe reached the end of the gel; NF-κB bands were detected by autoradiography and quantified by densitometry using Scion Image (Scion Corp., Frederick, MD, U.S.A.).

Isolation of hepatic RNA and reverse transcription-polymerase chain reaction (RT-PCR) assay for cytokine mRNA

Total cellular RNA was extracted from liver samples from each rat with TRIzol reagent (GibcoBRL, Rockville, MD, U.S.A.) (8) and quantified by measurement of ultraviolet absorption at 260 nm. For RT-PCR assay of mRNA, first-strand cDNA was synthesized from total RNA using SuperScript RNase H- Reverse Transcriptase (Invitrogen Corp., Rockville, MD, U.S.A.) and random hexamer primers [pd(N)₆] (Promega, Madison, WI, U.S.A.). cDNA was amplified in a PCR reaction using Taq DNA polymerase (Invitrogen Corp.) in the presence of primers specific for rat cytokines. Nucleotide sequences for

sense and antisense primers used were 5'-CACGCTCTTCT-GTCTACTGA-3' and 5'-GGACTCCGTGATGTCTAAGT-3' for TNF- α , and 5'-ATGGCCAAAGTTCCTGACTTGTTT-3' and 5'-TTCATGATGAACTCCTGCTTG-3' for IL-1 α (Invitrogen Corp.). Thermocycling conditions for PCR were defined for each cytokine. RNA concentrations and PCR cycler were titrated to establish standard curves to document linearity and to allow semiquantitative analysis of signal strength. To control the relative amount of total mRNA transcribed in each reverse transcriptase reaction, RNA 18S invariant standards [Classic 18S Internal Standards (489 bp) or Classic II 18S Standards (324 bp); Ambion, The RNA Co., Austin, TX, U.S.A.] were used. PCR products were electrophoresedon 2% agarose (Invitrogen Corp.). The gels were stained with ethidium bromide and analyzed by densitometry using Scion Image (Scion Corp.).

Enzyme-linked immunoabsorbent assay (ELISA) for cytokines

Rats were anesthetized with sodium pentobarbital (50 mg/kg i.p.), and serum from blood obtained by cardiac puncture was separated and stored at -80° C. Levels of serum TNF- α (UltraSensitive CytoscreenTM KRC3013 kit, Biosource International, Camarillo, CA, U.S.A.) and IL-1 α (Endogen Inc.,

Woburn, MA, U.S.A.) were measured by ELISA according to the manufacturers' specifications. Serum samples containing high cytokine levels were repeated after dilution to assure assay results within the standard curve.

Parameters related to oxidative stress and liver cell injury

Livers were perfused via the portal vein in a nonrecirculating system as described previously (44), using Krebs-Henseleit bicarbonate buffer saturated with 95% O₂/5% CO₂ to give pH 7.4, at constant flow rates (3.5-4.0 ml/g of liver/min) and temperature (36-37°C). Livers were allowed to equilibrate for 15 min, and perfusate samples were taken every 5 min for 15 min to measure lactate dehydrogenase (LDH) activity (one unit corresponds to 1 µmol/min at 37°C), with the parallel and continuous polarographic measurement of O2 consumption (44). At the end of perfusion, LDH activity was assessed in the tissue (U/g of liver) (5), concomitantly with the content of total glutathione (GSH) equivalents (38) and that of protein carbonyls as index of protein oxidation (29). Rates of sinusoidal LDH efflux (U/g of liver/min), as a liver cell viability parameter, were calculated from the perfusate activity and the flow rate, integrated between 15 and 30 min of perfusion

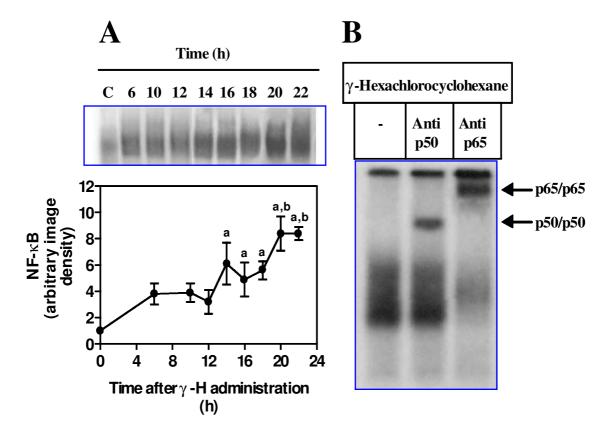


FIG. 1. Effect of acute γ -hexachlorocyclohexane (γ -H) administration on rat liver NF-κB DNA binding activity. (A) Representative autoradiograph of NF-κB DNA binding activity evaluated by EMSA using nuclear extracts from livers of control rats (C, time zero) and γ -H-treated animals (50 mg/kg) at different times after treatment, and the respective densitometric quantification of relative activity of NF-κB. Mean value at time zero was arbitrarily set to unity, and values at other time points were normalized to this. Data are means \pm SEM for three to six different animals. Significance studies: ap < 0.05 versus controls at time zero; bp < 0.05 versus γ -H-treated rats at 6, 10, and 12 h after treatment. (B) Supershift analysis for assessment of the composition of NF-κB induced at 20 h after γ -H treatment.

(U/g of liver), and expressed as percentage of the hepatic LDH activity (fractional LDH efflux).

Chemicals and reagents used were obtained from Sigma Chemical Co. (St. Louis, MO, U.S.A.), except for those stated in the above sections.

Statistical analysis

Values shown are means \pm SEM for the number of separate experiments indicated. The statistical significance of differences in mean values among multiple groups was made by oneway ANOVA and the Student–Newman–Keuls post-hoc test. In all cases, a value of p < 0.05 was considered significant.

RESULTS

 γ -Hexachlorocyclohexane administration increases rat liver NF- κ B DNA binding activity. Suppression by α -tocopherol or $GdCl_3$ pretreatment

NF- κ B DNA binding activity exhibited a progressive enhancement following γ -hexachlorocyclohexane administration, with significant increases starting at 14 h after γ -hexachlorocyclohexane treatment and maximal effects being observed at 20–22 h (Fig. 1A). Supershift analysis using antibodies specific for NF- κ B p50 (anti p50) or p65 (anti p65) disrupted the complexes induced at 20 h after γ -hexachlorocyclohexane treatment, thus confirming the presence of NF- κ B isoforms (Fig. 1B). Rats subjected to α -tocopherol or GdCl₃ exhibited liver NF- κ B DNA binding activity comparable to that in control animals, whereas both pretreatments abolished the eightfold increase in NF- κ B DNA binding activity induced by γ -hexachlorocyclohexane at 20 h after treatment (Fig. 2).

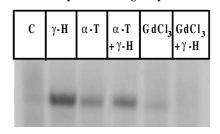
Effects of γ -hexachlorocyclohexane administration on liver O_2 consumption, sinusoidal LDH efflux, and content of GSH and protein carbonyls. Modification by α -tocopherol or $GdCl_3$ pretreatment

The maximal enhancement of liver NF- κ B DNA binding activity observed at 20 h after γ -hexachlorocyclohexanetreatment is accompanied by significant increases in the basal rate of O_2 consumption of the liver (21%; Fig. 3A) and in the sinusoidal fractional efflux of LDH (356%; Fig. 3B), concomitantly with a 24% reduction in the content of hepatic GSH (Fig. 3C) and a 355% elevation in protein carbonyls as index of protein oxidation (Fig. 3D). These γ -hexachlorocyclohexane-inducedchanges are suppressed in γ -hexachlorocyclohexane-treated rats pretreated with the antioxidant α -tocopherol or the Kupffer cell inactivator GdCl₃ (Fig. 3).

γ -Hexachlorocyclohexane administration promotes liver TNF- α and IL-1 α expression

 γ -Hexachlorocyclohexane-inducæl liver NF-κB DNA binding activity (Fig. 1A) was paralleled by increases in hepatic TNF- α expression, evidenced by a significant enhancement

Experimental groups



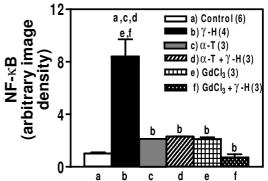


FIG. 2. Representative autoradiograph and densitometric quantification of relative activity of NF- κ B in control and γ -hexachlorocyclohexane (γ -H)-treated animals (20 h) subjected to α -tocopherol (α -T) or GdCl₃ pretreatments. Data are means \pm SEM for the number of rats indicated in parentheses, with significance assessment (p < 0.05) shown by the letters identifying each group.

in the levels of TNF- α mRNA in the 14–18-h time interval after γ-hexachlorocyclohexane treatment, returning toward control values at later times (Fig. 4A). At 18 h after γ-hexachlorocyclohexane administration, the increase in hepatic TNF- α mRNA levels was abolished by pretreatment with either α-tocopherol or GdCl₃ (Fig. 4B). γ-Hexachlorocyclohexane administration also stimulated the hepatic expression of IL-1 α , a response that is observed in the 14–22-h time interval, with a maximal effect being found at 18 h (Fig. 4C). At this latter experimental time, enhancement in hepatic IL-1α mRNA levels by γ -hexachlorocyclohexane was suppressed by both α tocopherol and GdCl₃ pretreatments (Fig. 4D). In these conditions, γ -hexachlorocydohexane elicited significant increases in the serum levels of TNF- α (Fig. 5A) and IL-1 α (Fig. 6A) in the 14-20-h time interval, with maximal effects at 18 h after treatment being abolished by both α-tocopherol and GdCl₃ pretreatments (Fig. 5B and 6B). In addition, pretreatment with the antisense oligonucleotide targeting the primary transcript of TNF-α ASO TJU-2755 (daily doses of 10 mg/kg i.v. for two consecutive days, 24 h prior to γ -hexachlorocyclohexane) (42) suppressed γ -hexachlorocyclohexane-inducedserum TNF- α enhancement at 18 h after treatment (Fig. 5B), without significantly affecting the increase elicited in IL-1 α levels (Fig. 6B).

DISCUSSION

Data reported in this study demonstrate that the administration of a single dose of γ -hexachlorocyclohexane to rats

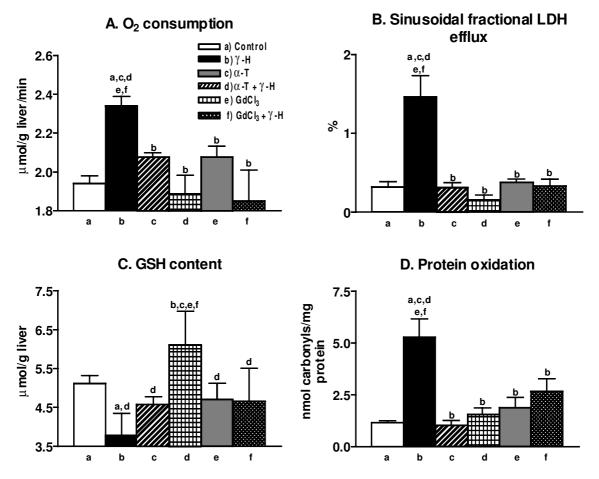


FIG. 3. Effect of pretreatment with α -tocopherol (α -T) and GdCl₃ on γ -hexachlorocyclohexane (γ -H)-induced changes in hepatic (A) O₂ consumption rate, (B) sinusoidal fractional efflux of LDH, (C) total reduced GSH content, and (D) levels of protein carbonyls at 20 h after γ -H treatment in the rat. Data are means \pm SEM for three to 11 different animals, with significance assessment (p < 0.05) shown by the letters identifying each group.

enhances the DNA binding activity of hepatic NF-kB, an effect that is related to the enhancement in the oxidative stress status of the liver. This view is in agreement with previous timecourse studies showing that hepatic formation of thiobarbituric acid reactants, NADPH-supported chemiluminescence, and superoxide radical production/superoxide dismutase activity ratios are significantly increased at 6 h after γ-hexachlorocyclohexane administration and maximally enhanced at 24 h (20), which correlate with the maximal NF-KB activation found at 20-22 h (Fig. 1A). Furthermore, the increases in hepatic respiration and protein oxidation, GSH depletion, and activation of NF-κB observed at 20 h after γ-hexachlorocydohexane treatment are abolished by α -tocopherol pretreatment, thus suggesting a cause-effect relationship between oxidative stress and activation of the transcription factor, although the role of mechanisms that are not dependent on the cellular redox status cannot be excluded. γ-Hexachlorocyclohexane-induced DNA binding activity of hepatic NF-κB occurs primarily at the Kupffer cell level, considering its suppression by GdCl, pretreatment. This rare earth metal eliminates large Kupffer cells (17), with the consequent reduction in the phagocytic (7, 46) and respiratory burst (46) activities of liver macrophages. It is of interest that an interrelationship between the oxidative stress status and Kupffer cell activity appears to exist, evidenced by the inhibition of γ -hexachlorocyclohexane-induced liver protein oxidation, lipid peroxidation (35), and GSH depletion by GdCl₂. The suppression of γ-hexachlorocyclohexane-indiced O_2 consumption by GdCl₃ and α -tocopherol suggests that significant mitochondrial uncoupling is not achieved in these conditions, as reported in isolated rat liver mitochondria (15), and indicates O2 equivalents used at the Kupffer cell level that may involve O2 utilization for production of reactive oxygen species involved in NF-kB activation in these cells. In this respect, O2 utilization by liver macrophage NADPH oxidase may play a role in γ-hexachlorocyclohexane-induced GdCl₃sensitive respiration, providing a sufficient prooxidant status to induce NF-kB activation (40, 47). The role of NADPH oxidase in NF-kB activation underlies hydrogen peroxide generation in the respiratory burst of macrophages (22) and is supported by studies with the NADPH oxidase inhibitor diphenyleneiodonium sulfate (23) or by deletion of p47^{phox}, the regulatory component of NADPH oxidase (40), in an animal model of alcoholic liver injury. In addition to liver macrophage NADPH oxidase, Kupffer-cell cytochrome P4502E1 could also play a

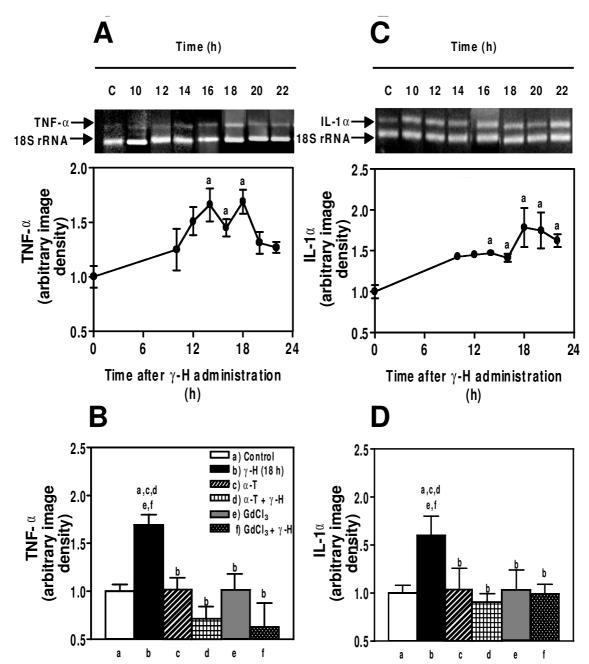


FIG. 4. Effects of acute γ-hexachlorocyclohexane (γ-H) administration on the liver expression of TNF- α and IL-1 α . (A) Representative agarose gel electrophoresis of the RT-PCR products for TNF- α mRNA (580 bp) and 18S rRNA (489 bp) after ethidium bromide staining in total hepatic RNA samples from control rats (C, time zero) and γ-H-treated animals at different times after treatment, and the respective densitometric quantification of RT-PCR products of the mRNA of TNF- α expressed as the TNF- α /18S rRNA ratios to compare lane-lane equivalency in total RNA content. Mean value at time zero was arbitrarily set to unity, and values at other time points were normalized to this. (B) Influence of α -tocopherol (α -T) and GdCl₃ pretreatment on γ-H-induced TNF- α mRNA levels at 18 h after treatment. (C) Agarose gel electrophoresis of the RT-PCR products of IL-1 α mRNA (408 bp) and 18S rRNA (324 bp) in total hepatic RNA samples from controls (C, time zero) and γ-H-treated rats at different times after treatment, and the respective densitometric quantification of the mRNA of IL-1 α expressed as the IL-1 α /18S rRNA ratios. (D) Effects of α -T and GdCl₃ pretreatment on γ-H-induced IL-1 α mRNA levels at 18 h after treatment. Data are means ± SEM for three different rats. Significance studies; ap < 0.05 compared with control values at time zero in (A) and (C); significance in (B) and (D) is shown by the letters identifying each experimental group (p < 0.05).

role in NF- κ B activation by γ -hexachlorocyclohexane, considering that the pesticide enhances its expression and activity in rat liver (12). Cytochrome P4502E1 is characterized by

a substantial prooxidant activity, which generates superoxide radical and hydrogen peroxide, and by its induction by chronic ethanol consumption in rat Kupffer cells (25). However, the

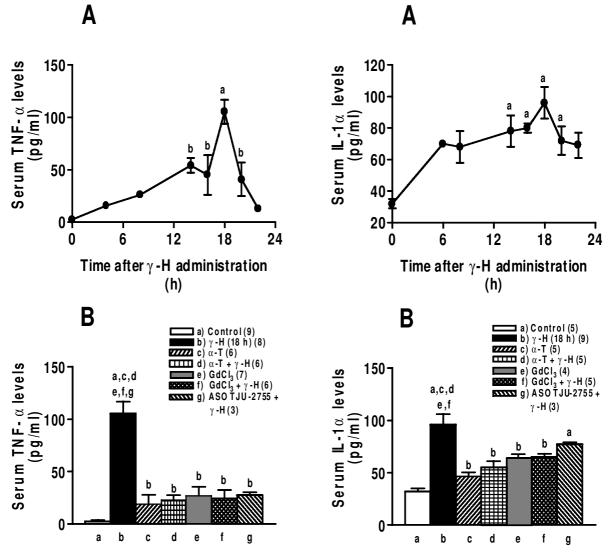


FIG. 5. Effect of acute γ -hexachlorocyclohexane (γ -H) administration on the serum levels of TNF- α at different times after treatment (A) and the influence of pretreatment with α -tocopherol (α -T), GdCl₃, and the antisense oligonucleotide targeting the primary transcript of TNF- α (ASO TJU-2755) assessed at 18 h after γ -hexachlorocyclohexane treatment (B). The serum levels of TNF- α were determined by ELISA. Data are means \pm SEM for three to nine different animals. Significance studies in (A): ap < 0.05 versus controls (time zero) and γ -H-treated rats at 4, 8, 14, 16, 20, and 22 h after treatment, and bp < 0.05 versus controls and γ -H-treated rats at 18 h. Significance in (B) is shown by the letters identifying each group (p < 0.05).

role of cytochrome P4502E1 and that of aldehydic products arising from γ -hexachlorocyclohexane-induced lipid peroxidation and/or reactive γ -hexachlorocyclohexane-derivedmetabolites in NF- κB activation in Kupffer cells remain to be established.

In these conditions, livers from animals subjected to γ -hexachlorocyclohexane administration with increased NF- κB DNA binding activity concomitantly exhibit mRNA expres-

FIG. 6. Effect of acute γ -hexachlorocyclohexane (γ -H) administration on the serum levels of IL-1 α at different times after treatment (A) and the influence of pretreatment with α -tocopherol (α -T), GdCl₃, and the antisense oligonucleotide targeting the primary transcript of TNF- α (ASO TJU-2755) assessed at 18 h after γ -H treatment (B). The serum levels of IL-1 α were determined by ELISA. Data are means \pm SEM for three to nine different animals. Significance studies in (A): ap < 0.05 versus controls at time zero. Significance in (B) is shown by the letters identifying each group (p < 0.05).

sion of the NF-kB responsive gene for TNF- α . In addition, the expression of the IL-1 α gene is also up-regulated by γ -hexachlorocyclohexane, a process that is triggered by several signals, including TNF- α (11). Thus, induction of the expression of IL-1 α in the liver of γ -hexachlorocyclohexane-treated rats might also result from the up-regulation of the TNF- α gene through the redox activation of NF-kB, leading to an enhancement in the serum levels of both cytokines sharing similar biologic activities. As the increases in IL-1 α induced by γ -hexachlorocyclohexane are not significantly reduced by an antisense oligonucleotide that fully blocked the γ -hexachloro-

cyclohexane-induced TNF- α enhancement, these data suggest that the increases in both cytokines are mediated by a third common factor, likely the activation of the NF- κ B system, that is suppressed by pretreatment with α -tocopherol and GdCl₂.

Although Kupffer cells play a central role in the homeostatic response to liver injury with release of TNF- α and IL-1 α , these mediators can be key effector molecules for induction of hepatotoxicity when their expression is sustained or enhanced, or the target cells become sensitized (40). Morphological evidence shows that hepatic cell injury following acute γ-hexachlorocyclohexane intoxication is fully developed at 24 h after treatment (20) and regresses to normal from the third day and thereafter (21). Liver injury observed at 20 h after γ -hexachlorocyclohexane administration, assessed by the substantial elevation in the sinusoidal release of LDH during perfusion studies, coincides with the higher levels of TNF- α and IL-1 α in serum, which are sustained from 14 h to 20 h. Moreover, γ-hexachlorocyclohexane-indwed liver LDH release is suppressed by α -tocopherol and GdCl₃ pretreatments, thus supporting the interrelationship between the redox activation of hepatic NF- κ B, the up-regulation of TNF- α and IL-1 α expression, and hepatotoxicity. This view is in agreement with data showing that the free radical-mediated redox activation of transcription factors (NF-κB and/or activator protein-1) transactivating promoters of inflammatory genes plays a central role in the hepatotoxicity of carbon tetrachloride (7, 16, 26), ethanol (23, 40), or acetaminophen (24). γ-Hexachlorocyclohexane-induced liver oxidative stress and hepatocellular injury are readily reversible (21), a phenomenon that may involve resetting of gene expression at the Kupffer cell level to favor cell survival and reparative responses.

In summary, data presented support the contention that acute γ -hexachlorocyclohexane administration triggers the DNA binding activity of hepatic NF- κ B by a redox mechanism exerted at the Kupffer cell level, with the consequent up-regulation of TNF- α and IL-1 α expression and related hepatotoxicity. Liver oxidative stress-dependent NF- κ B activation occurs in the 14–22-h time period after *in vivo* γ -hexachlorocyclohexane treatment, although NF- κ B is an acute transcription factor (41) that is activated as early as 10 min by the respiratory burst of carbon-stimulated Kupffer cells in liver perfusion studies (30) or within 30 min in isolated Kupffer cells exposed to lipopolysaccharide (39).

ACKNOWLEDGMENTS

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ABBREVIATIONS

ELISA, enzyme-linked immunoabsorbent assay; EMSA, electrophoretic mobility shift assay; $GdCl_3$, gadolinium chloride; GSH, reduced glutathione; IL- 1α , interleukin- 1α ; LDH, lactate dehydrogenase; NF- κ B, nuclear factor- κ B; RT-PCR, reverse transcription-polymerase chain reaction; TNF- α , tumor necrosis factor- α .

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