Endotoxin Administration to Humans Inhibits Hepatic Cytochrome P450-mediated Drug Metabolism

Steven I. Shedlofsky,* Beth C. Israel,† Craig J. McClain,* Daniell B. Hill,‡ and Robert A. Blouin†

*Department of Medicine, VA Hospital and Division of Digestive Diseases, University of Kentucky College of Medicine; †University of Kentucky College of Pharmacy; and ‡General Clinical Research Center, University of Kentucky College of Medicine, Lexington, Kentucky, 40511

Abstract

In experimental animals, injection of gram-negative endotoxin (LPS) decreases hepatic cytochrome P450-mediated drug metabolism. To evaluate this phenomenon in a human model of gram-negative sepsis, LPS was administered on two consecutive days to healthy male volunteers during which time a cocktail of antipyrine (AP-250 mg), hexobarbital (HB-500 mg), and theophylline (TH-150 mg) was ingested and the apparent oral clearance of each drug determined. Each subject had a control drug clearance study with saline injections. In the first experiment, six subjects received the drug cocktail 0.5 h after the first dose of LPS. In the second experiment, another six subjects received the drug cocktail 0.5 h after the second dose of LPS. In both experiments, LPS caused the expected physiologic responses of inflammation including fever with increases in serum concentrations of TNFα, IL-1β, IL-6, and acute phase reactants. In the first experiment, only minor decreases in clearances of the probe drugs were observed (7–12%). However, in the second experiment, marked decreases in the clearances of AP (35, 95% CI 18–48%), HB (27, 95% CI 14–34%), and TH (22, 95% CI 12–32%) were seen. The decreases in AP clearance correlated with initial peak values of TNFα (r = 0.82) and IL-6 (r = 0.86). These data show that in humans the inflammatory response to even a very low dose of LPS significantly decreases hepatic cytochrome P450-mediated drug metabolism and this effect evolves over a 24-h period. It is likely that septic patients with much higher exposures to LPS have more profound inhibition of drug metabolism. (J. Clin. Invest. 1994. 94:2209–2214.) Key words: cytokines • inflammation • sepsis • tumor necrosis factor • lipopolysaccharide

Introduction

Gram-negative bacterial sepsis is still an important cause of morbidity and mortality (1, 2). Animal studies of sepsis using endotoxin (LPS) to elicit the inflammatory response have demonstrated that there are profound decreases in hepatic cytochrome P450-mediated drug metabolism (3–8). These decreases may be mediated by inflammatory cytokines such as interleukin-1 (9–11), tumor necrosis factor (11–14), and possibly interleukin-6 (14, 15). Whether decreases in hepatic P450-mediated drug metabolism occur in humans with sepsis is not known, but is likely. Since critically ill septic patients are often given therapeutic drugs that require P450-mediated metabolism, adverse effects of these drugs may be a common but unrecognized problem.

Studying hepatic drug metabolism in septic patients is hampered by the inability to control for most of the variables that affect regulation of the cytochromes P450, including smoking status (16), diet (17), and concurrent medication use that could induce or inhibit various P450 isozymes. Another method to study effects of the inflammatory response on drug metabolism is to administer purified LPS to human volunteers and measure clearances of probe drugs. LPS administration to volunteers has been performed in a number of recent studies describing the physiologic responses of sepsis (18–22). In the current study, male volunteers ingested a “cocktail” of three probe drugs including antipyrine (AP),1 hexobarbital (HB), and theophylline (TH) (23, 24) whose clearances were measured on one occasion after saline injections and on another occasion after two LPS injections. This study demonstrates that even the relatively minor inflammatory response that occurred after this small dose of LPS was enough to decrease clearances of the probe drugs by 22–35%.

Methods

12 male volunteers aged 21–42 yr were enrolled in this study which was approved by the University of Kentucky Institutional Review Board and the General Clinical Research Center. Written consent was obtained at enrollment. All subjects were healthy as confirmed by medical history, physical exam, blood chemistries, urinalysis, and EKG. Subjects were nonsmokers and received no caffeine, theophylline products, nonsteroidal anti-inflammatory drugs, or other medications for at least 3 d before the study. All clearance studies were conducted in the General Clinical Research Center.

Each subject underwent two separate clearance studies: one with saline as a control and the other with LPS treatment. Half the subjects received LPS in the initial clearance study followed by the saline control study at least 2 wk later. For each clearance study, AP (250 mg), HB (500 mg), and TH (150 mg) were given as a “cocktail” (23, 24) after an overnight fast. For the LPS drug clearance study, each subject received two i.v. doses of U.S. FDA Bureau of Biologics endotoxin (LPS) (Lot EC-5, 20 Endotoxin U/kg) on consecutive mornings. Two experiments were conducted with six subjects in each. In the first experiment (EXP 1) LPS was administered 0.5 h before the drug cocktail and again 24 h later. Blood was collected at 0, 10, 20, 30, 45 min and 1, 2, 3, 4, 6, 9, 12, 30, and 48 h after the drug cocktail for plasma drug concentra-

1 Abbreviations used in this paper: APG, α1-acid glycoprotein; AP, antipyrine; C-RP, C-reactive protein; HB, hexobarbital; TH theophylline.
tions and at various time points for plasma cytokine and serum acute phase reactant determinations. In the second experiment (EXP 2), both doses of LPS were given before the drug cocktail, (24.5 ± 0.5 h before). Blood samples were drawn at various time points after the LPS and at the same times as above after the drug cocktail. Blood for complete blood count and clinical chemistry screen was collected prior to both drug clearance studies and 48 h after the first dose of LPS in EXP 1 and 72 h after the first dose of LPS in EXP 2.

Plasma concentrations of the cytokines TNFα, interleukin-1β (IL-1), and interleukin-6 (IL-6) were determined by commercial kits (Quantikine; R&D Systems, Minneapolis, MN) and the acute phase reactants C-reactive protein (CRP) and α1-acid glycoprotein (AGP) were determined in serum by commercial kits (Behring Diagnostic Incorporated, Somerville, NJ). Plasma AP, HB, and TH were analyzed as described previously (25).

AP, HB, and TH plasma concentration versus time data for each clearance study were analyzed by a nonlinear least squares regression analysis program (R-STRIP®, Micromath Scientific Software, Salt Lake City, UT). Inspection of the log plasma concentration vs. time curves for each of the drug probes in both saline and LPS drug clearance studies showed monoexponential slopes for the elimination of each compound. This allowed assumption of a one-compartment model with first order absorption and elimination rate constants. Area under the concentration time curve (AUC) was calculated by the trapezoidal rule from zero to the last time point and then extrapolated to infinity using the following formula: Clast/K where Clast is the last measured concentration time point and K is the terminal elimination rate constant obtained from R-STRIP®. Oral clearance (CLo) of drug was defined as: 

\[ CLo = \frac{Dose}{AUC} \]

Statistical analysis. It was assumed that a 30% change in drug clearance would be clinically significant and a power analysis was performed using the known intra-individual coefficient of variation (6.8–8.2%) from previous clearance studies for antipyrine (26) and determined that an n = 6 was sufficient to detect this change. A statistical package (Pharmacetical Calculation System, Springer-Verlag, NY) was used for data calculations. Drug clearance data between groups (saline control and LPS) were analyzed by the sign test (27). Confidence intervals for the changes in drug clearances were determined at the 95% level. Changes in peak plasma cytokines, peak serum acute phase proteins, bilirubin, albumin, white blood cell count, liver enzymes, temperature, and heart rate were analyzed by paired Student’s t test. The relationship between alterations in drug clearances caused by LPS with changes in plasma cytokines and acute phase proteins were correlated to assess a statistical association using regression analysis and F test. Statistical significance was defined as P < 0.05 for all tests.

### Results

As shown in Table I, endotoxin administration caused physical signs consistent with inflammation, but no obvious evidence of hepatic damage. After each dose of endotoxin, but not saline, subjects from both experiments experienced significant elevations in temperature and pulse, along with clinical symptoms of chills, malaise, and headache. No changes in white blood cell count, platelet counts, liver alanine aminotransferase, alkaline phosphatase, or total bilirubin were observed. There were minor but not significant decreases in subjects’ hematocrits, most likely due to the volume of blood drawn, and there was a minor but statistically significant decrease in albumin concentrations in EXP 2 but not EXP 1.

In EXP 1, (drug cocktail given 0.5 h after the first dose of LPS in six subjects), a number of biochemical indicators documented that an inflammatory response was elicited. As shown in Table II, there were marked increases in concentrations of the cytokines TNF (peaking at 90 min) and IL-6 (peaking at 4 h) after the first dose of LPS. After the second dose of LPS the next morning, the TNF peaks were lower, but the IL-6 responses were not blunted (higher in three subjects and only slightly lower in the others) and IL-6 peaked slightly earlier, just 2 h after the LPS. Data for IL-1β are not shown. Although some IL-1β determinations were made, this cytokine peaked at 9 h in several patients (1.5 pg/ml at baseline rising to 15 pg/ml) after the first dose of LPS, but samples from this time point were not available for three subjects and unavailable for all subjects after the second LPS dose. Both of the acute phase

### Table I. Clinical Characteristics and Laboratory Findings

<table>
<thead>
<tr>
<th>Experiment-1 (n = 6)</th>
<th>Experiment-2 (n = 6)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Baseline</td>
</tr>
<tr>
<td>Temperature (°F)</td>
<td>96.8±0.3</td>
</tr>
<tr>
<td>Pulse (bpm)</td>
<td>63±1</td>
</tr>
<tr>
<td>Hematocrit (%)</td>
<td>43.9±2.7</td>
</tr>
<tr>
<td>WBC (x10³/µl)</td>
<td>5.2±0.5</td>
</tr>
<tr>
<td>Platelets (x10³/µl)</td>
<td>228±49</td>
</tr>
<tr>
<td>Albumin (g/dl)</td>
<td>3.7±0.2</td>
</tr>
<tr>
<td>ALT (IU/l)</td>
<td>25±11</td>
</tr>
<tr>
<td>Alk. Phos. (IU/l)</td>
<td>71±19</td>
</tr>
<tr>
<td>Total Bili (mg/dl)</td>
<td>0.7±0.5</td>
</tr>
</tbody>
</table>

In experiment-1, the drug cocktail was given 0.5 h after the first dose of LPS and in experiment-2, the drug cocktail was given 0.5 h after the second dose of LPS. Peak temperatures and pulses were recorded 2.5 h after the first dose of LPS administration in each experiment and compared with baseline values just before injection. Similar responses were seen after the second dose of endotoxin. For blood values reported, samples were taken 48 h after the first dose of LPS in EXP 1 and 72 h after the first dose of LPS in EXP 2 and compared with baseline values obtained at enrollment in the study. Data are means±SD and * P < 0.05 for LPS compared with baseline.

### Table II. Serum Inflammatory Responses and Changes in Drug Clearance after LPS in Experiment-1

<table>
<thead>
<tr>
<th>Serum markers</th>
<th>Saline control</th>
<th>LPS Day 1</th>
<th>LPS Day 2</th>
</tr>
</thead>
<tbody>
<tr>
<td>TNF (pg/ml)</td>
<td>4±1</td>
<td>53±34*</td>
<td>19±11*</td>
</tr>
<tr>
<td>IL-6 (pg/ml)</td>
<td>3±1</td>
<td>31±36</td>
<td>52±58</td>
</tr>
<tr>
<td>CRP (mg/dl)</td>
<td>0±0</td>
<td>2.7±0.5*</td>
<td>3.3±0.5*</td>
</tr>
<tr>
<td>AGP (mg/dl)</td>
<td>71±21</td>
<td>89±15*</td>
<td>100±15*</td>
</tr>
<tr>
<td>Drug clearance data</td>
<td>Saline</td>
<td>LPS</td>
<td>Percent decrease</td>
</tr>
<tr>
<td>Antipyrine (lh)</td>
<td>2.33±0.43</td>
<td>2.06±0.40</td>
<td>12%</td>
</tr>
<tr>
<td>Hexobarbital (lh)</td>
<td>19.5±5.8</td>
<td>17.9±4.2</td>
<td>8%</td>
</tr>
<tr>
<td>Theophylline (lh)</td>
<td>3.30±0.33</td>
<td>3.08±0.60</td>
<td>7%</td>
</tr>
</tbody>
</table>

TNF peaked 1.5 h after LPS on each day, IL-6 peaked 4 h after LPS after the first LPS dose and 2 h after the second dose. There were no peaks with saline. CRP and AGP were measured 24 h after saline and at 24 and 48 h after the first dose of LPS. In this experiment (EXP 1) the drug cocktail was administered 0.5 h after the first dose of endotoxin or saline. Blood was collected over 48 h and clearance values were determined. Data are means ± SD for six subjects. * P < 0.05 for LPS vs. saline.
reactants C-RP and AGP demonstrated consistent increases after each dose of LPS.

Despite eliciting an inflammatory response, drug clearance values for all three probe drugs showed only modest decreases of 7–12%, none of which were statistically significant by the sign test (Table II). However, studies in experimental animals (7) have shown that maximal depression of P450 occurs 12–24 h after an LPS injection. We therefore predicted that decreases in drug clearances would be greater in EXP 2 in which both doses of LPS were administered before the drug “cocktail” (24.5 and 0.5 h before).

Fig. 1 illustrates the changes in plasma cytokines (TNF, IL-1 and IL-6) and serum acute phase proteins (C-RP, AGP) observed after the LPS or saline injections in EXP 2. Cytokines and acute phase proteins were markedly elevated after endotoxin administration when compared to control values (after saline). Concentrations of TNF (Fig. 1 a) peaked within 90 min after each LPS injection, but the response to the second LPS dose was much less than the first. IL-1β (Fig. 1 b) peaked 4–6 h after the first dose, did not return to baseline, and then increased even further with the second LPS dose before returning to baseline at 48 h. IL-6 (Fig. 1 c) peaked sharply 4 h after the first LPS dose, and had a much lower response after the second LPS dose. The C-RP and AGP responses again showed a steady increase after both doses of LPS. Values at 72 h (data not shown) were down slightly from 48 h, but were well above baseline and saline levels.

In EXP 2 much larger and significant changes (P < 0.05) in drug clearances were apparent as shown in Fig. 2. There was a 35% decrease in AP clearance, (95% CI, 18 to 48%), a 27% decrease in HB clearance, (95% CI, 14 to 34%), and a 22% decrease in TH clearance (95% CI, 12 to 32%). Correlations between the changes in drug clearances and LPS-induced cytokine and acute phase protein concentrations were examined. Although no statistically significant correlations were seen between any of the cytokine and acute phase protein responses and the decreases in HB and TH clearances, significant (P < 0.05) associations were found for decreased AP clearance and the peak TNF (r = .82) and peak IL-6 (r = .86) concentrations seen after the first dose of LPS (Fig. 3). These were cytokine responses that had been elicited before the drug cocktail was administered. Correlations were not significant between IL-1β or acute phase protein responses and decreased AP clearances.

Discussion

Because of the well known inhibition of hepatic cytochrome P450-mediated drug metabolism during inflammation in animal studies (3–8), it is very likely that similar inhibition occurs in humans. However, few studies evaluating regulation of P450 metabolism in critically ill septic patients have been performed despite the fact that these patients receive numerous drugs that require P450 for their clearances. Drug toxicities due to decreased clearance may be common but unrecognized phenomena that contribute to patient morbidity and mortality. Although Sangrador et al. reported in ICU patients a 34% decrease in TH clearance in one study (28) and a 50% decrease in caffeine clearance in another study (29), inflammatory responses were not assessed.

There are many difficulties in studying P450 metabolism in critically ill patients, including an inability to control for such.

Figure 1. Plasma cytokine and serum acute phase reactant concentrations after LPS injections at 0 and 24 h (■—■) or after saline injections at same times (●—●). The drug cocktail was given at 24.5 h. (a) Plasma tumor necrosis factor; (b) plasma interleukin-1β; (c) plasma interleukin-6; (d) serum α1-acid glycoprotein; and (e) serum C-reactive protein. Each point is mean±SEM of six subjects.
Figure 2. Oral drug clearances of antipyrine (250 mg), hexobarbital (500 mg), and theophylline (150 mg) administered as a “cocktail” in six male subjects given on one occasion saline IV 0.5 h before the drugs and on another occasion LPS (20 endotoxin units IV) 24.5 and 0.5 h before the drugs (EXP 2). The numbers identify data for individual subjects for each drug clearance comparison. The bar is the arithmetic mean for clearance data.

Factors as smoking, diet, concurrent drug use, and hemodynamic alterations (e.g., shock) that might alter hepatic blood flow and function. Administration of endotoxin (LPS) to human volunteers offers a more controlled and reproducible model to study the effects of the inflammatory response on hepatic P450-mediated drug metabolism. Because volunteers are clinically stable, the clearances of drug probes can be analyzed over relatively prolonged periods of time (24–48 h), and several probes can be administered simultaneously using the cocktail approach. In the current study, each of the three drug probes used (23, 24) were chosen because their oral clearances are dependent on activities of one or more hepatic microsomal cytochromes P450 and not on hepatic blood flow (30). It is also unlikely that extrahepatic sites such as the lung or intestine play an important role in their clearances (24). Furthermore, because the intrapatient variation for measuring oral clearances is low (26), using each subject as his own control allowed the use of a relatively small number of subjects to achieve significant results.

The current study demonstrates unequivocally that LPS administration to male human volunteers depresses hepatic cytochrome P450-mediated drug metabolism. Decreased clearances of all three drugs used in this study were seen, (AP [35%], HB [27%], and TH [22%]). Despite the very low LPS dose used (4 ng/Kg or 20 “endotoxin” U/Kg), these decreases are comparable to findings in most rodent studies (3–8) where doses ranging from 30 mg to 5 μg/Kg caused 35–50% decreases in total cytochromes P450 concentrations and various drug metabolizing activities in hepatic microsomes. Humans are known to be much more sensitive to LPS than rodents (31). However, because septic patients are probably exposed to much higher concentrations of LPS and suffer multisystem organ failure from these doses, it is likely that the decreases in P450-mediated metabolism reported in the volunteers are an underestimation of what occurs clinically.

The current study also assessed the timing of the inflammatory effect on depression of cytochrome P450-mediated drug metabolism. Drug clearances were not significantly depressed when the probe drugs were given just 0.5 h after the initial LPS exposure (EXP 1), but were significantly depressed when the probe drugs were administered 24 h later (EXP 2). Therefore, it is apparent that depression of drug metabolism in humans is not an immediate effect of the inflammatory response, but one
which develops over a 24-h period. Based on data from animal experiments, maximal depression of cytochrome P450 concentrations and activities occurs from 18 to 24 h after a single dose of LPS with a gradual return to baseline values by 48–72 h (7).

In the current study, LPS administration elicited the expected cytokine and acute phase responses confirming previous reports (18–22) of a safe and well characterized inflammatory response with this dose of LPS. Clinical symptoms of malaise, chills, and headache occurred ~1 or 2 h after LPS injection followed by significant \((P < 0.05)\) increases in temperature and heart rate lasting three to four hours. In EXP 1, the administration of the drug cocktail containing the anti-inflammatory drug antipyrine (250 mg) 0.5 h after the LPS did not mitigate these physiologic responses.

Based on animal studies (8–15), it is likely that depression of cytochrome P450-mediated drug metabolism after LPS is mediated by inflammatory cytokines, since administration of cytokines in these studies reproduced the depressions seen with LPS (8–15). Supporting this contention is our finding that the decreases in AP clearance in EXP 2 correlated with peak TNF and IL-6 responses after the first dose of LPS (Fig. 3, a and b).

The current study administered a second LPS dose 24 h after the first in order to "maximize" the inflammatory response during the drug clearance studies. In EXP 1, (Table II) there did not appear to be any significant blunting of the response to the second LPS dose, a phenomenon referred to as "endotoxin tolerance" (31). In EXP 2, in which the drug cocktail was given 0.5 h after the second LPS dose, there did appear to be marked blunting of the TNF and IL-6 responses (Figs. 1, a and c). However, clinical symptoms were not blunted and because the drug cocktail could have affected the cytokine responses, this protocol doesn’t allow one to address the issue of endotoxin tolerance. Tolerance to the effects of LPS in septic patients was recently addressed by Munoz et al. (32) and Callegari et al. (33), and found to be very complex. Granowitz et al. (34) also reported that a single dose of LPS to male volunteers made their peripheral monocytes less responsive to subsequent stimulation in vitro. How LPS tolerance might relate to regulation of hepatic drug metabolism in humans is unknown, but animal studies by Williams (35), demonstrated that LPS tolerance also extends to effects on hepatic cytochromes P450.

Several animal studies have also addressed the issue of LPS-induced inflammation on specific P450 isoenzymes by assessing enzyme activities, specific P450 proteins and mRNA concentrations (6–8). Stanley et al. (6) observed differential inhibition of P450 isozyme expression in rats with low doses of LPS, whereas high doses of LPS depressed all isoenzymes studied. Morgan (8) recently reported decreased P450 2C6, 2C7, 2C11, 2C12, and 2E1 mRNA but decreased proteins for only P450 2C11, 2C12, and 2E1 after LPS administration. Since P450 isoenzymes vary between species, how these studies apply to humans is not clear. The three probes used in the current study are rather "broad" substrates and are not believed to represent the activities of specific P450s, although the clearances of TH and HB provide assessments primarily of P450 1A2 and members of the P450 2C family, respectively (36–41). Whether LPS-induced inflammation differentially affects different P450 isoenzymes in humans will have to await studies using more specific probes.

In conclusion, this study shows that LPS-induced inflammation in humans is associated with significant depression of hepatic cytochrome P450-mediated drug metabolism. The depression in drug metabolism takes several hours to develop after initiation of the inflammatory response and the extent of the depression correlates with the intensity of the inflammatory response as assessed by elevations of the cytokines TNFα and IL-6. Further use of this experimental model (injecting human volunteers with LPS) can yield information as to the significance of depressed hepatic drug metabolism in critically ill septic patients as well as which drug metabolizing enzymes are most affected.

Acknowledgments

The authors wish to thank the CRC staff, including Patti Nickl, Connie Mitchell, Amy Hughes, and Susan Berryman, and acknowledge the technical assistance of Drs. Alice Swim and Rajna Tosheva-Tousanova and statistical review by Dr. Richard Kryscio.

This work was supported by the Department of Veterans Affairs and by a National Institutes of Health General Clinical Research Center grant M01RR02602-07.

References


