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Abstract

Lipopolysaccharide (LPS) from Gram (-) bacteria induces inflammatory cholestasis by impairing the expression/localization of transporters involved in bile formation (e.g., Bsep, Mrp2). Therapeutic options for this disease are lacking. Ursodeoxycholic acid (UDCA) is the first choice therapy in cholestasis, but its anticholestatic efficacy in this hepatopathy remains to be evaluated. To asses it, male Wistar rats received UDCA for 5 days (25 mg/Kg/day, i.p.) with or without LPS, administered at 8 a.m. of the last 2 days (4 mg/Kg/day, i.p.), plus half of this dose at 8 p.m. of the last day. Then, plasma alkaline phosphatase (ALP), bile flow, basal and taurocholate-stimulated bile acid output, total glutathione output, and total/plasma membrane liver protein expression of Bsep and Mrp2 by confocal microscopy were assessed. mRNA levels of both transporters were assessed by Real-time PCR. Plasma pro- inflammatory cytokines (IL-6 and TNF-α) were measured by ELISA. Our results showed that, UDCA attenuated LPS-induced ALP plasma release and the impairment in the excretion of the Bsep substrate, taurocholate. This was associated with an improved Bsep expression at both mRNA and protein levels, and by an improved localization of Bsep in plasma membrane. UDCA failed to reduce the increase in plasma pro-inflammatory cytokines induced by LPS and Mrp2 expression/function. In conclusion, UDCA protects the hepatocyte against the damaging effect of bile acids accumulated by the LPS-induced secretory failure. This involved an enhanced synthesis of Bsep and an improved membrane stability of the newly synthesized transporters.

Keywords. Ursodeoxycholic acid; hepatocellular transporters; bile acids; bile acid export pump; lipopolysaccharide-induced cholestasis.

Abbreviations: ALP, alkaline phosphatase; Bsep, bile salt export pump; BSP, bromosulfophthalein; DAMP, danger-associated molecular pattern; FXR, farsenoid X receptor; GSH, glutathione; JNK, c-jun-NH2-terminal kinase; LPS, lipopolysaccharide; Mrp2, multidrug resistance-associated protein 2; Ntcp, Na+/taurocholate cotransporting polypeptide; Oatp, organic anion-transporting polypeptide; PAMP, pathogen-associated molecular pattern; PXR, pregnane X receptor; ROI, region of interest; TC, taurocholate; RT-PCR, reverse transcriptase polymerase chain reaction; RXRα, 9-*cis*-retinoic acid receptor-α; UDCA, ursodeoxycholic acid.

1. Introduction

Inflammatory cholestasis is a frequent, often severe clinical condition, characterized by the impairment in bile production induced by pro-inflammatory cytokines [1]. Sepsis-induced cholestasis is prototypical of this kind of cholestasis [2]. It is an early, common complication found in more than 20% of hospitalized, icteric septic patients [3], and a main cause for admission and pediatric mortality in intensive care units [4].

Sepsis-induced cholestasis is causally associated with endotoxin. а lipopolysaccharide (LPS) from the Gram (-) bacterial wall. LPS is a pathogen-associated molecular pattern (PAMP) that is sensed by pattern-recognition receptors in macrophages, mainly Kupffer cells in liver [5]; LPS binding is followed by NLRP3 inflammasome activation, and the further release of pro-inflammatory cytokines [6]. These cytokines induce transcriptional and post-transcriptional impairment in the expression of hepatocellular carriers of bile acids and glutathione (GSH), the main osmotically active solutes driving bile formation [1, 2, 7]. LPS injection to rats induces downregulation of the bile acid basolateral importers Na+/taurocholate cotransporting polypeptide (Ntcp) and organic anion-transporting polypeptide (Oatp) 1, 2, and 4 [8]. In the canalicular pole, the expressions of multidrug resistance-associated protein 2 (Mrp2, aka Abcc2), the bilirubin glucuronide/GSH transporter, and bile salt export pump (Bsep, aka Abcb11), the main bile acid carrier, are also impaired [9], via transcriptional and postranscriptional mechanisms, [10]. Bsep and Mrp2 also suffer endocytic internalization in endotoxemia [11, 12], which may lead to their delivery to degradation compartments [11].

At present, there is no specific treatment for sepsis-induced cholestasis. However, it is imperative to find it, since cholestasis potentiates the development of the final septic

shock [13]. Actually, the increased levels of bile acids occurring in this condition are predictive of sepsis-associated mortality [14], since bile acids act as danger-associated molecular patterns (DAMPs) which, like PAMPs, activate the NLRP3 inflammasome, and synergize with LPS the cytokine release [13].

Ursodeoxycholic acid (UDCA) treatment is the only disease-modifying therapy in primary biliary cholangitis [15, 16], and the "first choice" therapeutic approach for most cholestatic hepatopathies [17]. UDCA may counteract both inflammation and bile acid-induced liver damage via multiple mechanisms, including *i*) protection against bile acid-induced necrotic and apoptotic cell death, *ii*) induction of hepatocellular metabolic changes that reduces bile acid levels and toxicity, and *iii*) anti-inflammatory and immunomodulatory properties [17-19]. Some of these effects seem to involve its role as an activating ligand of the nuclear receptors pregnane X receptor (PXR) [20] and farsenoid X receptor (FXR) [21], two transcription factors that regulate genes involved in synthesis and disposal of bile acids [22, 23]. However, the most relevant UDCA effects *in vivo* are FXR-independent and postranscriptional in nature [24]. For example, UDCA favors reinsertion of canalicular transporters that have been endocytosed after the cholestatic insult [25, 26].

Despite all these alleged protective mechanisms may be of benefit to preserve biliary secretory function in sepsis-induced cholestasis, the anticholestatic efficacy and action mechanisms of UDCA in this hepatopathy have not been evaluated yet. This study sought to address this issue, by using LPS-induced cholestasis in the rat as an experimental model.

2. Materials and methods

- **2.1 Chemicals.** LPS from *Salmonella typhimurium*, UDCA, bromosulfophthalein (BSP), 3α -hydroxysteroid dehydrogenase, NADPH, β -NAD, and GSH reductase were purchased from Sigma-Aldrich (St. Louis, MO). Sodium taurocholate (TC) was purchased from Santa Cruz Biotechnology (Dallas, Texas, U.S.A.). All the other reagents were of the highest analytical grade available from commercial sources.
- 2.2 Animals and treatments. Animals were obtained from our bioterium and received humane care according to the "Guide for the Care and Use of Laboratory Animals" published by the NIH (publication 25-28, revised in 1996). Animal experiments were approved by the Bioethical Committee for the Care and Use of Laboratory Animals of the Faculty of Biochemical and Pharmaceutical Sciences of the National University of Rosario (Resolution N° 489/2015). Adult (90 days), male Wistar rats weighing 300-350 g were used throughout. Animals were maintained on a standard diet and water ad libitum, under a constant 12 hour-light/12 hour-dark cycle. During the experiments, animal welfare was warranted by monitoring vital signs (heart and respiratory rates) and pain sensitivity (paw-withdrawal latency). All the treatments and the experiments with animals were performed in The Institute of Experimental Physiology (IFISE).

Rats were randomized in 4 experimental groups:

LPS group. The animals received UDCA vehicle (propylene glycol) for 5 days. At 8:00 am of the last two day, the animals were given a single, i.p. injection of LPS (4 mg/Kg/day, i.p., dissolved in saline) and, at 8:00 pm of the last day, a third dose of LPS of 2 mg/Kg was injected.

UDCA+LPS group. The animals received a daily dose of UDCA (25 mg/Kg/day, i.p., dissolved in propylene glycol) for 5 days [27-29]; this dosage is within the therapeutic range used in humans to treat cholestatic diseases [30]. Over the last two days, LPS was given as stated in the previous group.

Control group. The animals received UDCA and LPS vehicles throughout.

Experiments were carried out 12 hours after the last LPS (or LPS vehicle) administration.

- **2.3 Surgical procedures.** Rats were anesthetized with Ketamina and Xylazina (100 mg/kg p.c. and 3,5 mg/kg p.c., respectively) and thus maintained throughout. Body temperature was measured with a rectal probe, and maintained at 37°C with a heating lamp. The femoral vein and the common bile duct were cannulated with PE50 and PE10 polyethylene tubing, respectively (Intramedic, Clay Adams, Parsippany, NJ). At the end of each experiment, animals were sacrificed by exsanguination, and the liver was removed and weighed.
- **2.4. Experimental and analytical procedures.** A sample of basal bile was collected for 10 min to monitor basal bile flow, which was determined gravimetrically, assuming a density of 1 g/mL. Basal bile was assayed for both total bile acids and glutathione, the two main biliary solutes acting as driving force for bile formation [31]. Total bile acids were measured by using the method of the α -hydroxysteroid dehydrogenase [32], whereas total glutathione content was assessed by using the recycling method of Tietze [33], as modified by Griffith [34]. Once finished the bile collection period, animals were sacrificed by exsanguination, and the liver removed. The activity of the cholestasis

enzyme alkaline phosphatase (ALP; EC 3.1.3.1) was assessed in plasma by using a commercial kit (Wiener Lab., Rosario, Argentina). A blood sample from the tail vein was also obtained 2 hs after the second LPS injection using heparin as anticoagulant, to measure plasma levels of the inflammatory cytokines IL-6 and TNF-α, by using commercial kits (Rat IL-6 ELISA Kit from Life Technologies and Rat TNF alpha ELISA Ready-SET-Go!® from eBioscience, respectively).

2.5. Bsep/Mrp2 localization and protein expression. Liver samples were sectioned and frozen in isopentane precooled at -70°C in freezer, and stored at this temperature for further immunofluorescence and confocal microscopy analysis (LSM880, Carl Zeiss LLC, Thornwood, NY, USA). Liver sections were fixed and stained as described [35, 36], followed by overnight incubation with a rabbit anti-rat Bsep (1:200; Kamiya Biomedical Co., Seattle, WA, USA), or with a mouse anti-rat Mrp2 (1:200; [M2III-6], Alexis Biochemicals, San Diego, CA, USA), followed by incubation with a Cy2-conjugated donkey anti-rabbit IgG or a Cy2-conjugated donkey anti-mouse IgG, respectively (1:200, 1 h; Jackson ImmunoResearch Laboratory, West Grove, PA). The tight junctionalassociated protein Occludin was used to delimitate the border of the bile canaliculi. Occludin was labeled with a mouse anti-rat Occludin (when Bsep immunostaining was carried out, 1:200, overnight; Invitrogen) or with a rabbit anti-rat Occludin (when Mrp2 immunostaining was carried out, 1:200, overnight; Zymed, San Francisco, USA), followed by incubation with a Cy3-conjugated donkey anti-mouse or a Cy3-conjugated donkey anti-rabbit (1:200, 1 h, Jackson ImmunoResearch Laboratory, West Grove, PA.), respectively. To ensure comparable staining and image capture performance for the different groups belonging to the same experimental protocol, liver slices were prepared on the same day, mounted on the same glass slide, and subjected to the staining procedure and confocal microscopy analysis simultaneously. Quantification of the degree of Bsep and Mrp2 endocytic internalization was performed on confocal images using ImageJ 1.34m (National Institutes of Health), as described elsewhere [37]. For this purpose, intensity of fluorescence associated with the transporters along an 8-µm line perpendicular to the canaliculus (from -4 µm to 4 µm of the canalicular center). For each section, data from at least 15 different canaliculi were collected, and used for statistical comparison. At least two animals per group were used to assess the localization profile of the transporters, and the number of photographs analyzed per animal was 4-7. Each measurement was normalized to the sum of all intensities of the respective measurement. For quantification of total Bsep- and Mrp2-associated fluorescence intensity as a measure of protein expression, all sections were stained in a serial manner, and the same settings of confocal parameters were applied for each slide. The images were then loaded into ImageJ, the region of interest (ROI) traced, grey level intensities analyzed, and background subtracted systematically [38]. The images were then analyzed with ImageJ/FIJI image processing software, and the total fluorescence intensity of Bsep and Mrp2 were calculated. The results were then referred to control values.

2.6. Bsep/Mrp2 transport function. Transport activity of Bsep and Mrp2 was assessed by using sodium TC and BSP as model substrates. Based upon its relative affinity for Bsep as compared to other naturally occurring bile acids, TC is considered one of the best substrate to assess Bsep-mediated transport [39]. Regarding BSP, this cholephilic dye is taken up by Oatps and excreted with high affinity via Mrp2 after GSH conjugation

by glutathione-S-transferases [40, 41]; its Mrp2-mediated transfer but not the conjugation step is the rate-limiting one of its plasma-to-bile transport [42].

TC was administered in single, i.v. doses of 8 μ mol/100 g of b.w., and bile samples were then collected in 2-minute intervals over the next 10 minutes; after this time the exogenously administered TC was completely cleared in control animals. Bile salts were measured in bile with the α -hydroxysteroid dehydrogenase method [32], and TC excretion was calculated by assuming that all the excess in biliary excretion rate of bile acids over the basal one after TC injection is fully due to TC excretion. This assumption was confirmed by ancillary experiments in which biliary bile salt composition was analyzed by HPLC (Waters, Mildford, MA), as reported previously [43]; only increments in TC but not in other biliary bile salts in the rat were recorded after TC administration (data not shown).

BSP was administered in single, i.v. doses of 3 mg/100 g of b.w. Bile samples were then collected in 5-minute intervals over the next 60 minutes, a time period sufficient to virtually fully depurate the dye in control animals. BSP in bile was determined spectrophotometry at wavelength of 580 nm, after appropriate dilution with 0.1 N NaOH [44].

2.7 RNA isolation and real time reverse transcriptase polymerase chain reaction (RT-PCR). After treatment, total RNA was isolated from liver tissue using TRIzol® reagent (Invitrogen). Yield of extracted RNA was evaluated spectrophotometrically at 260 nm, through the following conversion: 1 OD = 40 μ g/ μ L of RNA. Purity and integrity of RNA was assessed by the A260/A280 ratio and 1,2 % agarose gel electrophoresis under denaturing conditions (110 V, 40 min), and using SYBER-SAFE as marker. The

gel was then observed in a UV transilluminator. A260/A280 ratios should be between 1.8 and 2.2, whereas the 28S rRNA band should be about twice the intensity of the 18S band, and higher molecular weight bands indicative of DNA contamination should be absent. The first-strand cDNA was synthesized from 1 μg of total RNA using Superscript III Reverse Transcriptase (Invitrogen) and random primers according to the manufacturer's suggested protocol. Primers for Abcb11 were designed using the "Primer-BLAST" program [45], and sequences of primer pairs for Abcc2 were as previously described [46] (Table 1). Real time PCR reactions were performed on a Mx3000P System (Stratagene, La Jolla, CA) with Platinum Taq DNA Polymerase (Invitrogen) and SYBR Green quantification. The thermocycling regime was 2 min at 95 °C for initial denaturation; and then 40 cycles of 95 °C for 15 s, 55 °C for 30 s, and 72 °C for 30 s; and the final step was 1 min at 95 °C, 30 s at 55 °C and 30 s at 95 °C. Results for Abcb11 and Abcc2 mRNA were normalized to the expression of 18S rRNA as the housekeeping gene by using the $2^{-\Delta \Delta Ct}$ method 47]. The primers for 18S rRNA were those previously designed by Ruiz et. al. [46] (Table 1). The specificity of each reaction was verified with a dissociation curve using a temperature range from 55 °C to 95 °C, with continuous fluorescence measurements.

2.8. Statistical analysis. Data are expressed as means \pm SEM. Statistical analysis was carried out by using ANOVA or Student t test, unless another statistical test has been determined to be more appropriate. The variances of the densitometric profiles of Bsep and Mrp2 localization were compared with the Mann-Whitney U test [48]. Values of P<0.05 were considered to be statistically significant.

3. Results

3.1. Blood and biliary parameters of inflammation and cholestasis. As shown in Table 2, LPS significantly increased ALP plasmatic levels, a plasma membrane enzyme that is induced in both hepatocytes and cholangiocytes, and released from the cell surface into blood by detergent bile acids that are retained in cholestasis [49] UDCA pretreatment reduced partially this increase, suggesting protection against intrahepatic bile acid building and/or bile acid-induced plasma membrane damage.

As expected, plasma levels of the pro-inflammatory cytokines TNF- α and IL-6 were both increased by LPS. UDCA pretreatment was unable to significantly reduce these increments; only a tendency was observed for TNF- α , but it did not reach statistical significance (Table 2). This suggests that the beneficial effects of UDCA are not mainly due to an attenuation of the inflammatory response evoked by LPS.

As shown in Table 3, LPS significantly reduced bile flow, and coadministration of UDCA did not attenuate this decrease. UDCA did not significantly increase the basal biliary excretion rate of both bile acids and glutathione when coadministered with LPS. However, when a bolus of TC was administered so that to acutely challenge the Bsepmediated secretory pathway, the decrease in both bile flow and bile acid output induced by LPS was partially counteracted by UDCA coadministration; this suggests that Bseptransport function, and hence bile acid-dependent bile flow, are indeed improved by UDCA, but it is required to force the transport system with higher levels of intracellular bile acids for this recovery to become apparent.

3.2 Bsep and Mrp2 transport function. Bsep and Mrp2 transport function was assessed by recording the changes over time in the biliary excretion of TC and BSP, respectively, following a bolus injection of these compounds

As shown in Fig. 1, TC was transported less efficiently by LPS-treated rats, as indicated by the 45% lower value of accumulated biliary excretion of this compound, as compared to controls after 4 min of TC injection. UDCA prevented partially this decrease, reaching a 40% higher value of cumulative TC excretion as compared to LPS group.

As depicted in Fig. 2, BSP was also transported less efficiently by LPS-treated rats, as suggested by the 30% lower value of accumulated biliary excretion of the dye as compared to controls after 60 min of its injection. This impairment was however not counteracted by UDCA, since cumulative BSP excretion in the co-treated rats with UDCA was similar to that of LPS alone (Fig. 2, upper panel).

3.3. Protein expression and localization of Bsep and Mrp2. As shown in Table 4, total Bsep and Mrp2 protein content, as assessed by measuring the total Mrp2- and Bsep-associated green fluorescence in the fields, was significantly impaired by LPS, with reductions of 59% and 31%, respectively.

Figure 3 A shows confocal images of Bsep (green) and the canaliculus border marker, occludin (red). In control livers, transporter-associated fluorescence was confined to the canalicular space. In LPS-treated rats, the carrier relocalized from the canalicular space to the pericanalicular, as indicated by the increased fluorescence at a greater distance from the bile canaliculus, and co-treatment with UDCA counteracted this redistribution. Figure 3 B shows the analysis of the distribution profiles of transporter-associated

fluorescence along a line perpendicular to the canaliculus. LPS-treated livers showed a statistically significant (*P*<0.05) wider profile with increased fluorescence at a greater distance from the canalicular membrane, a finding consistent with endocytic retrieval of these transporters into the intracellular compartment (Table 5) On the other hand, in livers from the LPS+UDCA co-treated group, the distribution of both Bsep and Mrp2 was almost identical to that in control livers (*P*>0.05). Analysis of the distribution profiles of the tight junctional protein occludin as a marker of the canalicular edge demonstrates a conserved width of the canaliculi in all the experimental groups, thus ruling out the influence of changes in this parameter on the Bsep distribution profiles. These data demonstrate that UDCA protects against LPS-induced retrieval of Bsep. Essentially the same results were obtained when the distribution pattern of Mrp2-associated fluorescence was assessed, with re-localization of Mrp2 outside the limits of the canaliculus that is reversed by UDCA co-treatment (Fig. 4).

3.4. Bsep and Mrp2 mRNA expression. RT-PCR analysis shows that both Bsep and Mrp2 mRNA abundance in rat livers from LPS-treated rats was markedly decreased by 85% and 75%, respectively (Table 6). UDCA co-treatment to LPS-treated rats normalized mRNA expressions of both transporters.

4. Discussion

UDCA has been suggested as a rationale alternative to treat sepsis-induced cholestasis due to its multiple anticholestatic mechanisms [50], and because it is often

successful as the first-choice therapy for most cholestatic disorders [17]. However, several authors agree that the available experimental and clinical evidence is still insufficient [50-52]. Therefore, we have provided here for the first time experimental evidence that UDCA bears multiple anticholestatic effects in LPS-induced cholestasis.

First, UDCA helped to preserve hepatocyte integrity against detergent bile acids accumulated in cholestasis, as suggested by the reduction in plasma ALP levels compared to that in LPS-treated rats (see Table 2). ALP is a membrane-bound enzyme that is induced and released into the bloodstream by detergent, endogenous bile acids [53], and represents an early indication of membrane damage leading to hepatocellular lysis [54]. The counteraction of this pathomechanim by UDCA most likely reflects its membrane-protective effects [55] as well as its lack of capability to induce ALP expression [56] and to remove ALP from the plasma membrane in cholestatic livers [57].

In agreement with previous works [7, 58], we show here that LPS represses the gene expression of the main canalicular bile acid transporter Bsep, leading to downregulation of the transporter at a protein level (see Table 4). UDCA upregulated at a transcriptional level these otherwise downregulated bile acid transporter (see Table 6), which resulted in an ameliorated capability of the hepatocyte to excrete an i.v. load of the model Bsep substrate, TC (see Fig. 1).

Improvement of Bsep gene expression by UDCA administration has been reported to occur under other cholestatic conditions, as for example in mice exposed to α -naphtylisothiocyanate [59], but not in normal animals [60] or healthy humans [61]. Therefore, the cholestatic context seems to be important for UDCA to improve transcriptionally Bsep. Transcriptional Bsep downregulation in LPS-induced cholestasis is caused by the cytokines released during the acute-phase response rather than by

accumulated bile acids [7]. Cytokine-mediated repression of FXR, the main nuclear receptor involved in the maintenance of Bsep levels, has been suggested as a chief causal factor [62]. The possibility that UDCA improves Bsep levels by enhancing the remnant Bsep transactivating activity of FXR by acting as a FXR ligand is however unlikely, because it has a very weak (if any) FXR agonist activity [21]. Apart from reducing FXR levels, cytokines impair the DNA binding of FXR to the Bsep IR-1 responsive element [63, 64]. FXR binds to DNA as a heterodimer with the 9-cis-retinoic acid receptor-α (RXRα), and this binding is impaired by cytokines via a c-jun-NH2-terminal kinase (JNK)-dependent RXR phosphorylation [63, 65]. Although RXRα levels are reduced in LPS-induced cholestasis [62, 63], this mechanism may be still operating, since a partial replacement of RXRα by other RXR isoforms not affected by cytokines, such as RXRβ and RXRγ, occurs [62]. In line with this, UDCA can inhibit JNK activity in hepatocytes exposed to different stressor, such as hydrophobic bile acids [66] and lipotoxic fatty acids [67].

Similarly to Bsep, Mrp2 gene transcription was enhanced by UDCA in LPS cholestatic rats (see Table 6); this may reflect, again, the capability of UDCA to counteract cytokine-mediated effects on Mrp2 transcription mechanisms, which is also JNK-dependent in nature [63, 65]. In addition, the well characterized inductive effect that UDCA has *per se* on Mrp2 expression [60] may be another contributing factor. Surprisingly, an enhanced Mrp2 mRNA levels in the co-treated group did not lead to an improved expression of Mrp2 either at the protein level (see Table 4) or at a functional levels, as evaluated by assessing the biliary excretion of its exogenous substrate, BSP (see Fig. 2). Similarly, biliary excretion of GSH, an endogenous Mrp2 substrate [68], was not ameliorated either (see Table 3). Differences in post-transcriptional events (*e.g.*, mRNA processing,

steady-state mRNA stability, translational efficacy), as well as in post-translational events (e.g., transporter sorting and targeting to the canalicular membrane, protein modifications that affect its degradation rate) may be causal factors. Actually, Mrp2 and Bsep have marked differences in their post-transcriptional processing, including intracellular location at different vesicular pools after LPS-induced endocytic internalization [69] and differential intracellular degradation after internalization; Mrp2 and Bsep both suffer ubiquitination, but then Mrp2 traffics from early endosomes into the lysosomal degradation pathway [70], whereas Bsep more likely suffers proteosomal degradation [71] Interestingly, higher rates of lysosomal protein degradation occur in endotoxemic rats [72], which may have selectively enhanced Mrp2 degradation after internalization. The lack of beneficial effect of UDCA on Mrp2 protein expression and function should not be necessarily regarded as a drawback, since it helps to retain both GSH and bilirubin, two endogenous anti-oxidant compounds that have been shown to counteract oxidative stress-induced cholestasis 73, 74], a factor that has been causally linked to LPS-induced cholestasis [75, 76].

The improvement in Bsep activity induced by UDCA in LPS-treated rats was apparent under conditions of increased bile acid load but not under basal conditions, where neither bile flow nor total bile acid excretion was modified (see Table 3). It should be born in mind that basal bile acid excretion depends not only on the intrinsic activity of Bsep but also on the size of the bile acid pool [77] and the extrusion rate via alternative, bile acid basolateral export pumps, such as Mrp3 and Mrp4 [78]; reduction and increments in these two last parameters, respectively, decreases canalicular bile acid output. UDCA may reduce bile acid pool size, since it decreases absorption of endogenous bile acids at the intestinal levels [79] and inhibits Cyp7a1, the rate-limiting

enzyme of bile acid synthesis [80]. As for basolateral bile acid extrusion, LPS increases the expression of Mrp3 [8, 81], as the result of an adaptive response in cholestasis aimed to redirect bile acid flux back to bloodstream for urinary elimination [82]. UDCA can further upregulate Mrp3 in the endotoxemic liver [24], and to upregulate Mrp4 as well [83]. As a corollary, the lack of improvement in bile acid biliary output by UDCA in LPS cholestatic rats under basal conditions should not be regarded as absence of anticholestatic effect of UDCA, but as a consequence of other beneficial effects aimed to limit bile acid accumulation via both reduced bile acid synthesis and enhanced bile acid basolateral efflux [17].

UDCA improved localization of the canalicular transporters Bsep and Mrp2 in their membrane domain by counteracting LPS-induced endocytic internalization (see Figs. 3 and 4, respectively). In line with this, UDCA stabilizes Bsep in the apical membrane of MDCK II cells, by preventing its endocytic internalization [84]. Bsep endocytic internalization occurs via a clathrin-mediated mechanism from clathrin-rich "non-raft" membrane domains under both normal [85] and cholestatic conditions [86], and the same holds true for Mrp2, at least under cholestatic conditions [86]. However, both Bsep and Mrp2 mainly reside in caveolin/cholesterol-rich "raft" domain, where they are fully functional due to its cholesterol dependency [87, 88]. Therefore, they need to be transferred to "non-raft" microdomains to be endocytosed [86]. Since UDCA is a membranotropic agents that was show to expand and stabilize "raft" structures by mimicking cholesterol [89], perhaps it stabilizes Bsep and Mrp2 in these enlarged "raft" microdomains, thus preventing their transfer to "non-raft" structures. UDCA may have also favored the vesicular trafficking of these transporters back to the canalicular membrane after endocytosis. Indeed, UDCA has been shown to stimulate the apical vesicular targeting of both newly-synthesized [90] and pre-existing canalicular transporters localized in subapical endosomes [91]. Actually, anticholestatic effects of UDCA associated with counteraction of canalicular transporter endocytosis have been shown in other experimental models of cholestasis [26, 92].

UDCA attenuated the effects of LPS without reducing expression of TNF- α and IL-6, two critical, sentinel cytokines involved in hepatic acute phase response (see Table 2). In line with our results, UDCA treatment was not instrumental in reducing the hepatic expression of the cytokines TNF- α and IL-1 β in cholestatic rats with bile-duct ligation [93], a cholestatic model largely associated causally with endotoxemia [94]. Lack of effect on cytokine production might, overall, be a rather beneficial UDCA property though, since cytokines are involved in the hepatic regenerative response [95], which, in the setting of liver cell damage, should be conserved to favor healing.

In conclusion, UDCA can protect the hepatocyte against the damaging effect of bile acids accumulated by LPS-induced bile acid secretory failure. This beneficial effect involves both transcriptional and posttranscriptional mechanisms, such as improved synthesis and function of the main bile acid canalicular transporter, Bsep, and an enhanced membrane stability of the carrier. Our results may be of relevance not only for sepsis-induced cholestasis, but also for the treatment of other cholestatic liver diseases in which LPS [96] or inflammatory cytokines [97, 98] also play a key role.

Conflict of interest

The authors have no conflict of interest.

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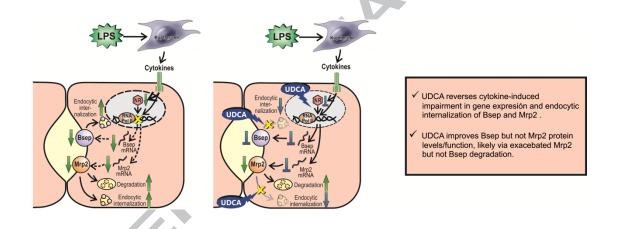
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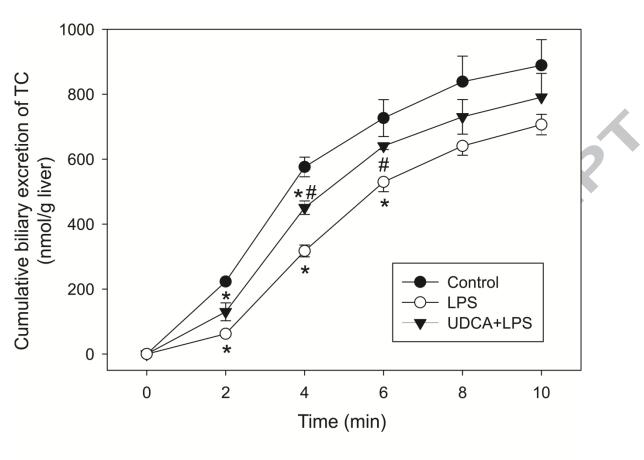
Figure 1. *Upper panel.* Cumulative biliary excretion of TC; bile samples were collected every 2 min for 10 min, after a single, i.v. dose of 8 μmol/100 g of b.w. of the bile salt. *Lower panel.* Area under the curve (AUC) of cumulative biliary excretion of TC graphs. Data are mean value ± SEM for 3-4 individual experiments. **P*<0.05 *vs.* control; **P*<0.05 *vs.* LPS.

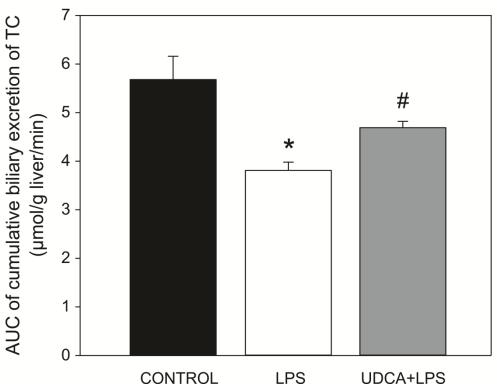
Figure 2. *Upper panel.* Cumulative biliary excretion of BSP; bile samples were collected every 5 min for 60 min, after a single, i.v. dose of 3 mg/100 g of b.w. *Lower panel.* Area under the curve (AUC) of cumulative biliary excretion of BSP graphs. Data are mean value ± SEM for 3-4 individual experiments. **P*<0.05 *vs.* control; **P*<0.05 *vs.* LPS.

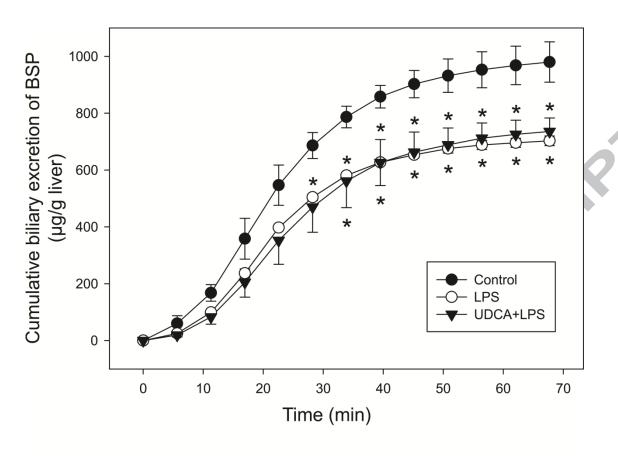
Figure 3. Effect of UDCA co-treatment on LPS-induced endocytic internalization of Bsep. A. Representative confocal images showing co-staining of Bsep (green) with occludin (red; used to visualize the bile canaliculus limits), illustrative of the endocytic internalization of Bsep induced by LPS (arrows), and its protection by UDCA. The lower panels show densitometric analysis of the fluorescence intensity associated with Bsep or occludin along a 8 μ m line perpendicular to the canaliculus (from -4 μ m to +4 μ m from the canalicular centre), corresponding to the confocal images shown in the upper panels. Data are mean values for at least 15 different canaliculi. Data are shown as mean \pm SEM.

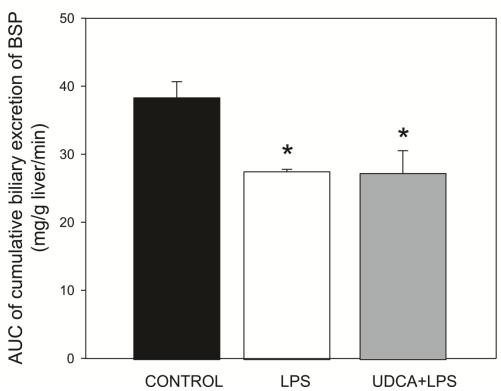
Figure 4. Effect of UDCA co-treatment on LPS-induced endocytic internalization of Mrp2. A. Representative confocal images showing co-staining of Mrp2 (green) with occludin (red; used to visualize the bile canaliculus limits), illustrative of the endocytic internalization of Mrp2 induced by LPS (arrows), and its protection by UDCA. The lower panels show densitometric analysis of the fluorescence intensity associated with Mrp2 or occludin along a 8 μ m line perpendicular to the canaliculus (from -4 μ m to +4 μ m from the canalicular center), corresponding to the confocal images shown in the upper panels. Data are mean values for at least 15 different canaliculi. Data are shown as mean \pm SEM.

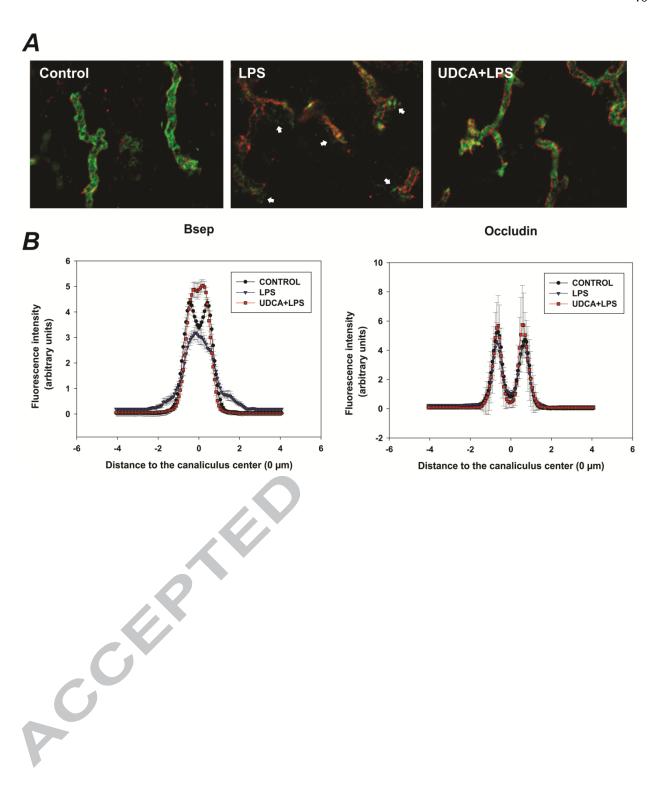


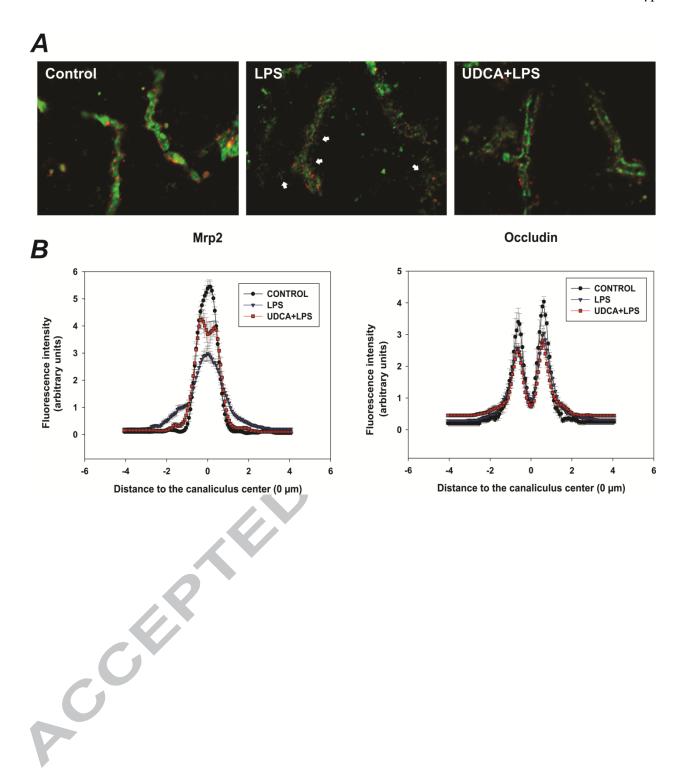


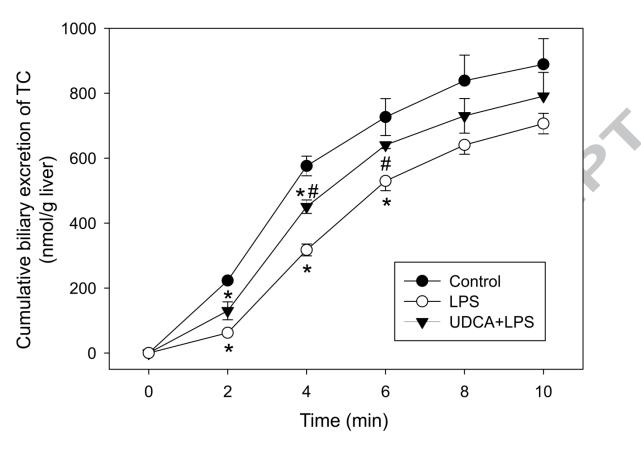


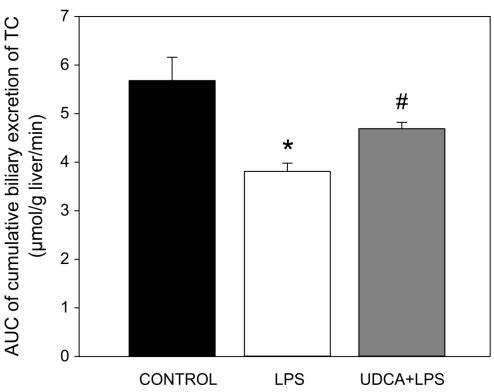


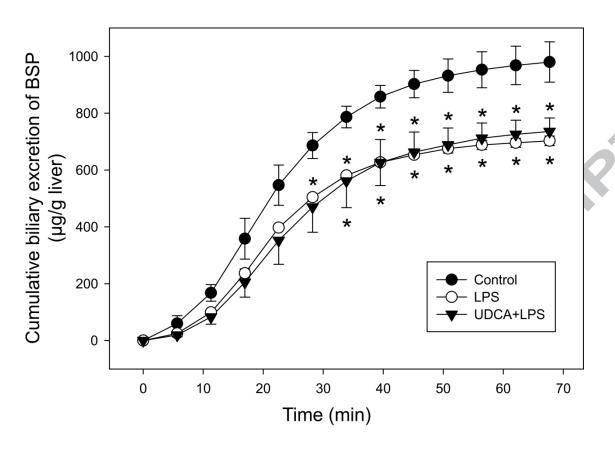


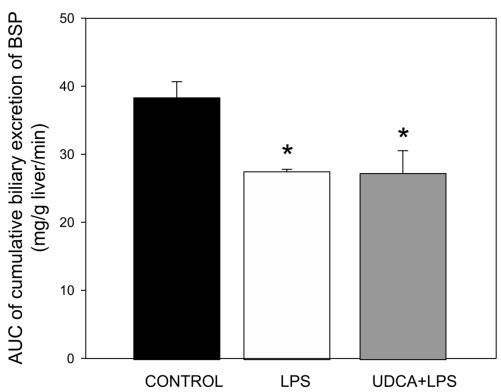












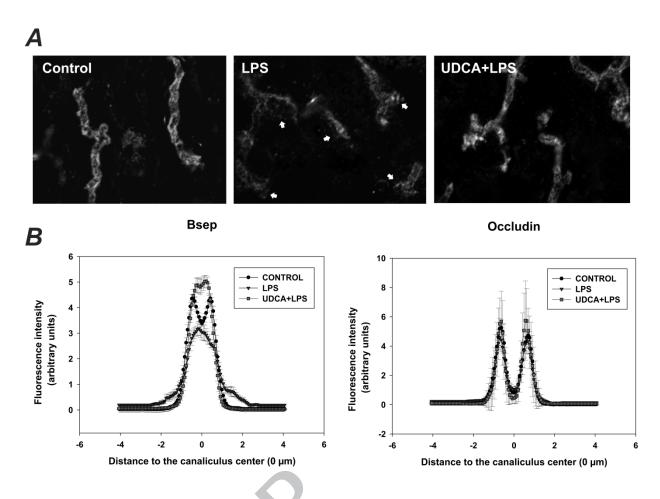


Table 1. Sequence of primers for RT-PCR analysis of rat Abcb11 and Abcc2 mRNAs, and 18s rRNA.

	Primer	Primer sequence (5' \rightarrow 3')
Ababdd (Baan)	Forward	CCGAAGGCTCAGGGTATTGG
Abcb11 (Bsep)	Reverse ATCAGGTGACAT	ATCAGGTGACATGGTGGCAG
A h a a 2 (M/m 2)	Forward	ACCTTCCACGTAGTGATCCT
Abcc2 (Mrp2)	Reverse	ACCTGCTAAGATGGACGGTC
	Forward	GTAACCCGTTGAACCCCATT
18s rRNA	Reverse	CCATCCAATCGGTAGTAGCG

Table 2. Plasma levels of ALP and the pro-inflammatory cytokines, TNF-α and IL-6.

	Control	LPS	UDCA+LPS
ALP (U/I)	207±23	404±21*	316±24*#
TNF-α (pg/ml)	49±6	133±33*	100±22*
IL6 (pg/mL)	N.D.	3604 ±603*	3312 ±558*

^{*}P<0.05 vs. control; *P<0.05 vs. LPS; N.D. = No detectable. n = 6-22.

Table 3. Bile flow and biliary excretion of total bile acids and total GSH.

	Control	LPS	UDCA+LPS	
Bile flow (µl/min per g liver)				
Basal	3.4 ± 0.1	2.7 ± 0.1*	2.6 ± 0.1*	
TC-stimulated [†]	5.3 ± 0.2	4.1 ± 0.2*	4.7 ± 0.1#	
Bile acid output (nmol/min per g liver)				
Basal	128 ± 16	94 ± 13	86 ± 13	
TC-stimulated [†]	226 ± 16	152 ± 7*	216 ± 5#	
Total GSH output (nmol/min per g liver)	3.7 ± 0.7	1.3 ± 0.2*	0.8 ± 0.1*	

[†]TC-stimulated values correspond to the maximum value obtained after a TC i.v. injection of 8 μ mol/100 g of b.w. *P<0.05 vs. control; *P<0.05 vs. LPS. n= 3-26.

THANAILEXELScence intensity (% of mean control values)	Control	LPS	UDCA+LPS S	Table 4.
Mrp2	10ρ _θ <u>7</u> 7	25 <u>‡</u> 1 <u>Q</u> *2*	190 ₃ 449 ₂ *	Mrp2 andBsep
Bsep	100 ₆₀ 29 ₅	15 ₆₉ 3 <u>*</u> 3*	79 ₆₃ 3 [#] 7*#	expression
				by

measuring total fluorescence intensity.

*P<0.05 vs. control; #P<0.05 vs. LPS

Table 5. Variance ranges of the fluorescence intensity profiles.

	Control	LPS	UDCA+LPS
Bsep	0.25 - 1.00	0.66 - 2.29*	0.12 - 1.24#
Mrp2	0.12 - 1.56	0.78 - 2.83*	0.25 - 2.15#

*P<0.05 vs. control; *P<0.05 vs. LPS; n= 15-22.

Table 6. Mrp2 and Bsep mRNA expression by RT-PCR.

*P<0.05 vs. control; #P<0.05 vs. LPS; n = 3-4.

