The right choice of antihypertensives protects primary human hepatocytes from ethanoland recombinant human TGF- β_1 -induced cellular damage

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Correspondence: Andreas Nüssler Eberhard Karls Universität Tübingen, BG Trauma Center, Schnarrenbergstr 95, 72076 Tübingen, Germany Tel +49 7071 606 1065 Fax +49 7071 606 1978 Email andreas.nuessler@googlemail.com **Background:** Patients with alcoholic liver disease (ALD) often suffer from high blood pressure and rely on antihypertensive treatment. Certain antihypertensives may influence progression of chronic liver disease. Therefore, the aim of this study is to investigate the impact of the commonly used antihypertensives amlodipine, captopril, furosemide, metoprolol, propranolol, and spironolactone on alcohol-induced damage toward human hepatocytes (hHeps).

Methods: hHeps were isolated by collagenase perfusion. Reactive oxygen species (ROS) were measured by fluorescence-based assays. Cellular damage was determined by lactate-dehydrogenase (LDH)-leakage. Expression analysis was performed by reverse-transcription polymerase chain reaction and Western blot. Transforming growth factor (TGF)- β signaling was investigated by a Smad3/4-responsive luciferase-reporter assay.

Results: Ethanol and TGF- β_1 rapidly increased ROS in hHeps, causing a release of 40%–60% of total LDH after 72 hours. All antihypertensives dose dependently reduced ethanol-mediated oxidative stress and cellular damage. Similar results were observed for TGF- β_1 -dependent damage, except for furosemide, which had no effect. As a common mechanism, all antihypertensives increased heme-oxygenase-1 (HO-1) expression, and inhibition of HO-1 activity reversed the protective effect of the drugs. Interestingly, Smad3/4 signaling was reduced by all compounds except furosemide, which even enhanced this profibrotic signaling. This effect was mediated by expressional changes of Smad3 and/or Smad4.

Conclusions: Our results suggest that antihypertensives may both positively and negatively influence chronic liver disease progression. Therefore, we propose that in future patients with ALD and high blood pressure, they could benefit from an adjusted antihypertensive therapy with additional antifibrotic effects.

Keywords: primary human hepatocytes, alcoholic liver disease, ethanol, $TGF-\beta_1$, antihypertensives

Introduction

Alcoholic liver disease (ALD) is one of the leading causes of death worldwide. It progresses from steatohepatitis to clinical manifestations, such as fibrosis, cirrhosis, and hepatocellular carcinoma. Ethanol consumption leads to production of reactive oxygen species (ROS), eg, hydroxyl-ethyl free radicals, hydroxyl radicals, and superoxide anions, causing an accumulation of reactive compounds and lipid peroxidation in ALD patients.¹ ROS are produced during ethanol metabolism by both enzymes and, more selectively, through the microsomal ethanol oxidative system with its main

component cytochrome P450 (CYP)2E1.² In a previous study, we showed that ethanol and recombinant human transforming growth factor (rhTGF)- β_1 induce oxidative stress in human hepatocytes (hHeps) with a peak of ROS levels and cellular glutathione depletion after 4 hours. Continuous exposure to both substances results in apoptotic cell death with 40%–60% of total lactate-dehydrogenase (LDH) released into the culture medium after 72 hours.³ In ALD patients, upregulated CYP2E1 expression goes along with iron mobilization4 and reduction of antioxidant enzymes and chemicals, particularly mitochondrial and cytosolic glutathione.3 This causes a disbalance between cell ROS production and cellular enzymatic and nonenzymatic defense mechanisms. Generated ROS then react with and in turn damage complex cellular molecules, eg, lipids, proteins, DNA, mitochondria, and cell membranes.⁵ Such repeated damage causes liver cells to produce excessive extracellular matrix, which eventually prevents organ regeneration. Furthermore, ethanol affects the immune system by altering cytokine production.⁶ This is of particular importance as hHeps exposed to ethanol are more sensitive toward damage from TGF- β_1 , which enhances oxidative stress on hHeps by upregulating nicotinamide adenine dinucleotide phosphate-oxidases (NOX).8 Furthermore, TGF-β signal transduction in this and other liver cell types is critically required for liver disease progression.

Reducing oxidative stress by upregulating the endogenous antioxidative enzyme heme-oxygenase-1 (HO-1) with cobalt protoporphyrin protected hHeps from ethanol-induced cytotoxicity.9 However, most HO-1 inducers, including cobalt protoporphyrin and hemin, show toxic side effects. Just recently we showed a protective effect of the naturally occurring polyphenol quercetin against ethanol-induced cellular damage in hHeps being dependent on HO-1.10,11 Similar effects were shown for the structurally related calcium channel blockers nifedipine and verapamil,³ both formerly used as standard drugs to regulate blood pressure. Therapeutic targets for constantly controlling high blood pressure have continued to evolve, depending on the patients' disease pathology, age, and other factors, and corresponding drugs have been developed. In addition to the aforementioned calcium channel blockers (calcium antagonists), these include diuretics, angiotensin-converting enzyme (ACE)-inhibitors, aldosterone antagonists, renin inhibitors, as well as adrenergic receptor agonists and antagonists (α - and β -blockers). There are hints that certain antihypertensives influence fibrosis in vivo, as reported for myocardial and renal fibrosis. 12,13 There are reports that amlodipine and captopril may also be effective against lung and liver fibrosis as well as keloid formation. 14-17 However, the available data are contradictory, eg, the ACE-inhibitor captopril may reduce as well as enhance left ventricular fibrosis in rats. 12,13 As patients with ALD often suffer from high blood pressure, they rely on an antihypertensive treatment. Therefore, the aim of this study was to investigate the influence of the commonly used antihypertensives, amlodipine (calcium channel blocker), captopril (ACE-inhibitor), furosemide (diuretic), metoprolol (β_1 -blocker), propranolol (β_1 / β_2 -blocker), and spironolactone (aldosterone antagonist) on alcohol- and TGF- β_1 -induced damage in hHeps.

Materials and methods

rhTGF- β_1 was purchased from Peprotech (London, UK). Chemicals, cell culture medium, and supplements, if not stated differently, were purchased from Sigma (Munich, Germany). Antibodies were purchased from Santa Cruz Biotechnology (Heidelberg, Germany).

Isolation of primary hHeps

Human liver tissue was obtained according to institutional guide-lines ("Klinikum rechts der Isar," Technical University Munich, Munich, Germany) from liver resections of tumor patients with primary or secondary liver tumors. Written informed consent was obtained from all patients. hHeps were isolated by a two-step collagenase perfusion technique followed by Percoll gradient centrifugation.¹⁸ hHeps viability (trypan blue exclusion) was consistently above 85%. hHeps were cultured on collagen-coated culture plates in Williams' Medium E (10% fetal calf serum, 1% nonessential amino acids, 1 mM sodium pyruvate, 32 IE/L insulin (Actrapid®; Novo Nordisk Pharma GmbH, Mainz, Germany), 15 mM HEPES, 0.8 μg/mL hydrocortisone, 100 U/mL penicillin, and 100 mg/mL streptomycin). Prior to all experiments, hHeps were serum starved overnight.

ROS measurement

hHeps, stimulated for 4 hours with 100 mM ethanol and/or 5 ng/mL rhTGF- β_1 in the presence or absence of antihypertensives as indicated, were washed twice with Dulbecco's phosphate-buffered saline. Cells were then incubated with 10 μ M 2′,7′-dichlorfluorescein-diacetate (ROS) for 30 minutes at 37°C in serum-free culture medium. The amount of ROS is in concordance with the fluorescence measured at an excitation/emission of 485/520 nm.³ Basal ROS levels were determined in untreated cells.

Cellular damage

LDH activity in the culture supernatant was determined by the LDH test kit from Fa Hitado (Möhnesee, Germany) according to the manufacturer's protocol. Results are given as percentage of total LDH released. A 100% LDH release was determined by cell lysis with 0.1% Triton X-100. Untreated cells were used to determine the basal cell death over the culture period.

Conventional reverse-transcriptase polymerase chain reaction

Total cellular RNA was isolated with Trifast (Peqlab, Erlangen, Germany) according to the manufacturer's protocol. First-strand cDNA was synthesized from 1 μ g RNA according to the manufacturer's instructions by using the First Strand cDNA Synthesis kit from Fermentas (St Leon-Rot, Germany). Primer data are summarized in Table 1. Products resolved by gel electrophoresis were visualized with ethidiumbromide.

Western blot

hHeps were lysed in ice-cold lysis buffer (50 mM TRIS [Trisaminomethane], 250 mM NaCl, 2% Nonidet P40, 2.5 mM EDTA [Ethylenediaminetetraacetic acid], 0.1% sodium dodecyl sulfate [SDS], 0.5% deoxycholic acid, complete protease inhibitor, 1.0% phosphatase inhibitor, pH 7.2). Protein concentration was determined by micro-Lowry. Thirty micrograms total protein was separated by SDS-polyacrylamide gel electrophoresis and transferred to nitrocellulose membranes (Roth, Karlsruhe, Germany). After overnight incubation with primary antibodies (1:1000 in Tris-buffered saline with Triton-X-100) at 4°C, membranes were incubated with the corresponding horseradish peroxidase-labeled secondary antibodies for 2 hours at room temperature. Chemiluminescent signals were detected with X-ray film.

Statistics

Results are expressed as mean \pm standard error of the mean of at least three independent experiments (N \geq 3) measured

in triplicate or more ($n \ge 3$). Densitometric analysis was performed using ImageJ software (National Institutes of Health, Bethesda, MD, USA). Data sets were compared by one-way analysis of variance followed by Bonferroni's multiple comparison test (GraphPad Prism software; GraphPad Software, Inc, La Jolla, CA, USA). P < 0.05 was taken as the minimum level of significance.

Results

Determining median lethal concentration doses of antihypertensives

To determine the median lethal concentration (LC_{50}) of the antihypertensives, LDH release of hHeps, treated with different drug concentrations (dimethyl sulfoxide was used as solvent control) for 72 hours, was determined. LC_{50} values and resulting experimental doses of the drugs are summarized in Table 2.

Antihypertensive drugs reduce ethanol- and rhTGF- β_1 -dependent damage in hHeps

We measured LDH activity in culture supernatants of hHeps treated with 100 mM ethanol and/or 5 ng/mL rhTGF- β_1 in the presence or absence of antihypertensive drugs. After 72 hours both ethanol and rhTGF- β_1 induced cellular damage in hHeps with 50%–55% of total LDH released. Co-incubation with ethanol and TGF- β_1 enhanced the damage (Figure 1A). For ethanol, all drugs investigated significantly reduced the damaging effect in a dose-dependent manner (Figure 1B). Similar results were observed for rhTGF- β_1 , except for furosemide, which had no effect on rhTGF- β_1 -treated hHeps (Figure 1C). Interestingly, when hHeps were stimulated with ethanol and rhTGF- β_1 together, the protective effect of furosemide that was observed with ethanol alone was abolished (Figure 1D).

Table	I Primer	and	pol	ymerase	chain	reaction	conditions
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Gene	GeneBank accession (NM_)	Forward primer 5'-3'	Reverse primer 5'-3'	T _m (°C)	Product length (bp)
Smad2	00100365	CAAACCAGGTCTCT	GAGGCGGAAGT	60	259
		TGATGG	TCTGTTAGG		
Smad3	005902.2	GGAGAAATGGTGC	GAAGGCGAACT	60	258
		GAGAAGG	CACACAGC		
Smad4	005359.3	TGAATCCATATCAC	CAGGCTGACTT	60	294
		TACGAAC	GTGGAAG		
Smad7	005904.1	TTCGGACAACAAG	AAGCCTTGATG	60	201
		AGTCAGC	GAGAAACC		
Alk5	004612.1	TGTTGGTACCCAAG	AACATCGTCGA	60	287
		GAAAGC	GCAATTTCC		
GAPDH	007393	CACCCACACTGTGC	CTCCTGCTTGC	60	607
		CCATC	TGATCCAC		

Table 2 LC_{so} and working concentrations of antihypertensives

Antihypertensive	LC ₅₀ *	Working		
		concentrations		
Amlodipine	I I.73 ± I.06 μM	I/2/4 μM		
Captopril	$>$ 100 μ M $^{\circ}$	25/50/100 μΜ		
Furosemide	36.73 \pm 1.11 μM	5/10/20 μM		
Metoprolol	96.45 \pm 4.71 μM	12.5/25/50 μΜ		
Propanolol	$31.47\pm1.37~\mu\text{M}$	5/10/20 μM		
Spironolactone	$>$ 100 μM°	25/50/100 μΜ		

Notes: *N = 3, n = 4; onot in the range tested. **Abbreviation:** LC_{50} , median lethal concentration.

Antihypertensives reduce ethanoland rhTGF- β_1 -dependent oxidative stress in hHeps

We measured the formation of ROS in hHeps that were treated for 4 hours with 100 mM ethanol and/or 5 ng/mL rhTGF-β, in the presence or absence of antihypertensives. As

expected from previous investigations, ethanol and rhTGF- β_1 induced ROS formation in hHeps. Co-incubation with both substances together further increased ROS formation (Figure 2A). Although all drugs tested reduced ethanol- and rhTGF- β_1 -dependent ROS formation (Figure 2B–D), furosemide and spironolactone had the least effect for rhTGF- β_1 (Figure 2C).

Antihypertensives rescue ethanoland rhTGF- β_1 -depleted glutathione levels in hHeps

We measured glutathione (GSH) levels in hHeps that were treated for 4 hours with 100 mM ethanol and/or 5 ng/mL rhTGF- β_1 in the presence or absence of antihypertensives. As expected from previous investigations, ethanol and rhTGF- β_1 reduced cellular GSH levels in hHeps. Co-incubation with both substances further decreased cellular GSH levels

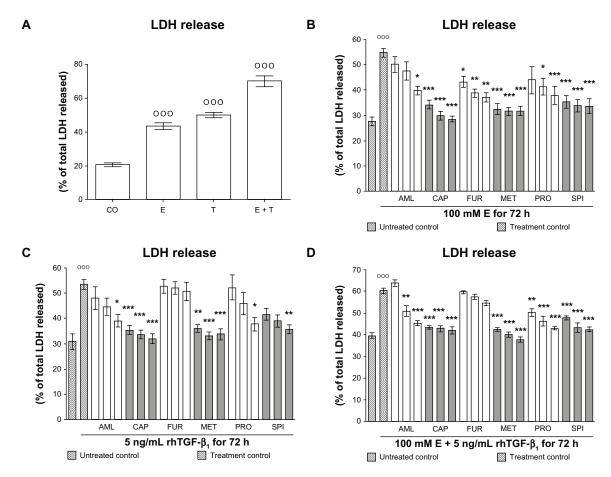


Figure 1 Effect of antihypertensives on E- and rhTGF- β_1 -dependent cellular damage in hHeps. (**A**) LDH release of hHeps (N = 4, n = 4) treated for 72 hours with 100 mM E and/or 5 ng/mL rhTGF- β_1 , LDH release of hHeps (N \geq 3, n = 4) treated with (**B**) 100 mM E, (**C**) 5 ng/mL rhTGF- β_1 , or (**D**) both substances for 72 hours in the presence or absence (treatment control) of AML (1/2/4 μ M), CAP (25/50/100 μ M), FUR (5/10/20 μ M), MET (12.5/25/50 μ M), PRO (5/10/20 μ M), or SPI (12.5/25/50 μ M).

Notes: DMSO was used as solvent control. Results are given as percentage of total LDH released into the culture supernatant (cells dissolved with 0.1% Triton X-100 equals 100%). Basal toxicity (untreated control) was approximately 30%. $^{\circ\circ}P < 0.001$ as compared to untreated cells; $^*P < 0.05$; $^{***}P < 0.01$; $^{***}P < 0.001$ as compared to E- or rhTGF- β_1 -treated cells. Data are presented as mean \pm SEM.

Abbreviations: AML, amlodipine; CAP, captopril; DMSO, dimethyl sulfoxide; E, ethanol; FUR, furosemide; hHeps, human hepatocytes; LDH, lactate dehydrogenase; MET, metoprolol; PRO, propranolol; rhTGF- β 1, recombinant human transforming growth factor beta 1; SPI, spironolactone.

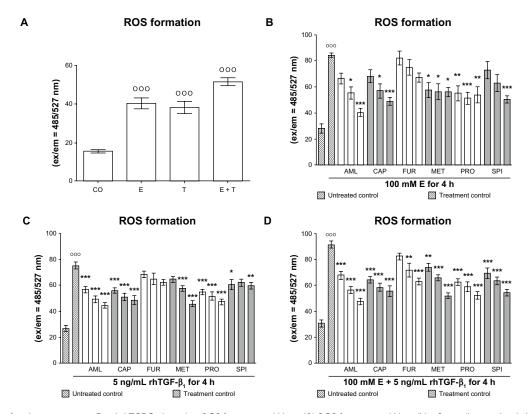


Figure 2 Effect of antihypertensives on E and rhTGF- $β_1$ -dependent ROS formation in hHeps. (A) ROS formation in hHeps (N = 3; n = 4) treated with 100 mM E and/or 5 ng/mL rhTGF- $β_1$ for 4 hours. ROS measurement in hHeps (N = 3, n = 4) treated with (B) 100 mM E, (C) 5 ng/mL rhTGF- $β_1$, or (D) both substances for 4 hours in the presence or absence (treatment control) of AML (1/2/4 μM), CAP (25/50/100 μM), FUR (5/10/20 μM), MET (12.5/25/50 μM), PRO (5/10/20 μM), or SPI (12.5/25/50 μM). Data are presented as mean ± SEM.

Notes: $^{\circ\circ}P < 0.001$ as compared to untreated cells. DMSO was used as solvent control. Results are given as fluorescent intensities (ex/em = 485/527 nm). Basal ROS formation (untreated control) was approximately 25; $^*P < 0.05$; $^{**P} < 0.01$; $^{***P} < 0.001$ as compared to E- or rhTGF- β_1 -treated cells.

Abbreviations: AML, amlodipine; CAP, captopril; CO, control (untreated cells); E, ethanol; ex/em, excitation/emission; FUR, furosemide; hHeps, human hepatocytes; MET, metoprolol; PRO, propranolol; rhTGF-β₁, recombinant human transforming growth factor beta 1; ROS, reactive oxygen species; SPI, spironolactone; T, treated with rhTGF-β1.

(Figure 3A). Almost all drugs tested increased GSH levels reduced by ethanol and rhTGF- β_1 treatment dose dependently (Figure 3B–D). However, furosemide had the least effect for for both substances (Figure 3C).

Antihypertensives induce HO-I expression in hHeps

We determined HO-1 expression levels in hHeps treated with the antihypertensives for 24 hours. For controls, we treated the cells with 100 mM ethanol and/or 5 ng/mL rhTGF- β_1 . Densitometric analysis revealed that all antihypertensives induced the expression of HO-1 in hHeps and showed strong effects for captopril, furosemide, and metoprolol (Figure 4A).

Blocking HO-I activity abolished the protective effect of antihypertensives on ethanol- and rhTGF- β_1 -dependent hepatocyte damage

We measured LDH-release of hHeps treated with 100 mM ethanol and/or 5 ng/mL rhTGF- β_1 in the presence or absence of antihypertensives. In order to block HO-1 activity,

cells were co-incubated with 10 μ M zinc-protoporphyrin (ZnPP)-9. ZnPP9 completely abolished the beneficial effect of antihypertensives on ethanol-dependent hHeps damage (Figure 3B). For rhTGF- β_1 -dependent damage where furosemide was not effective, ZnPP9 did not completely reverse the protective effects of the drugs (Figure 4C and D).

Antihypertensives interfere with rhTGF- β_1 -induced Smad3/4 signaling

Primary hHeps were infected with Ad5-CAGA₉-MLP-Luc adenoviral particles (Smad3/4-reporter construct) and stimulated for 24 hours with 5 ng/mL rhTGF- β_1 in the presence or absence of antihypertensives. Luciferase activity was measured in cell lysates. Interestingly, all drugs except furosemide (tendency for induction) were able to inhibit Smad3/4 signaling induced by rhTGF- β_1 (Figure 5A). To confirm that the interference was not at the receptor level, we repeated the experiment with hHeps co-infected with Ad5-CAGA₉-MLP-Luc adenoviral particles and viral particles encoding for a constitutive active TGF- β receptor type I

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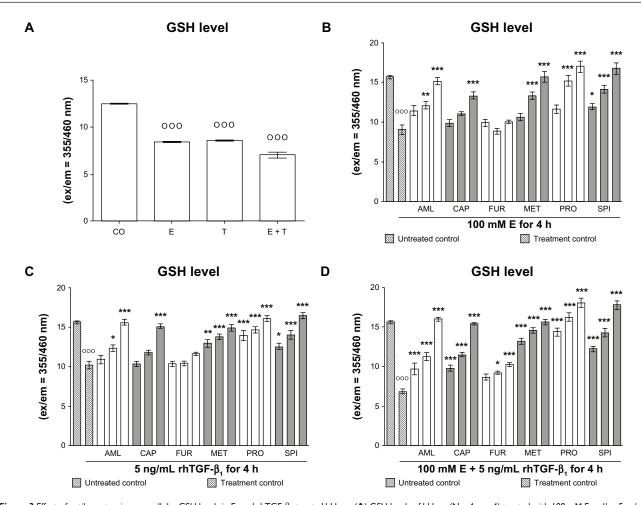


Figure 3 Effect of antihypertensives on cellular GSH levels in E- and rhTGF- β_1 -treated hHeps. (**A**) GSH levels of hHeps (N = 4; n = 4) treated with 100 mM E and/or 5 ng/mL rhTGF- β_1 for 4 hours. GSH levels of hHeps (N = 4, n = 4) treated with (**B**) 100 mM E, (**C**) 5 ng/mL rhTGF- β_1 , or (**D**) both substances for 4 hours in the presence or absence (treatment control) of AML (1/2/4 μM), CAP (25/50/100 μM), FUR (5/10/20 μM), MET (12.5/25/50 μM), PRO (5/10/20 μM), or SPI (12.5/25/50 μM). Data are presented as mean ± SEM.

Notes: DMSO was used as solvent control. Results are given as fluorescent intensities (ex/em = 355/460 nm). Basal GSH levels (untreated control) were approximately 16. $^{\circ\circ}P < 0.001$ as compared to untreated cells; $^*P < 0.05$; $^{***}P < 0.01$; $^{***}P < 0.001$ as compared to E- or rhTGF- β_1 -treated cells.

Abbreviations: AML, amlodipine; CAP, captopril; CO, control (untreated cells); DMSO, dimethyl sulfoxide; E, ethanol; ex/em, excitation/emission; FUR, furosemide; GHS, glutathione; hHeps, human hepatocytes; MET, metoprolol; PRO, propranolol; rhTGF-β₁, recombinant human transforming growth factor beta 1; SPI, spironolactone; T, treated with rhTGF-β1.

(Alk5-receptor) (Ad5-caAlk5), which does not require ligand binding for activating Smad3/4 signaling. For amlodipine, captopril, propranolol, and spironolactone, the inhibiting effect in Smad3/4 signaling remained, whereas in this assay, metoprolol no longer inhibited Smad3/4 signaling (Figure 5B).

Antihypertensives alter expression levels of Smad3 and Smad4 in hHeps

We measured expression levels of Smad2, -3, -4, -7, and Alk5 in hHeps that were treated for 24 hours with antihypertensives. We used β -actin as the house-keeping gene (Figure 5C). While Smad2, Smad7, and Alk5 expression levels were not significantly altered, expression levels of Smad3 were reduced by all antihypertensives (most notably for captopril and metoprolol), except for furosemide, which

in contrast showed a significant induction of *Smad3* and *Smad4*.

Discussion

Although the liver is probably the organ with the best regenerative capacity in the human body, repeated damage, eg, from alcohol abuse, causes liver cells to produce excessive extracellular matrix, which eventually prevents regeneration. As patients suffering from ALD show increased levels of reactive products and lipid peroxidation, it is widely accepted that oxidative stress plays an important role in the pathogenesis of ethanol-induced cellular injury.

In ALD patients, metabolism of consumed alcohol in the liver rapidly increases ROS production.² Similarly, in our in vitro experiments, ethanol stimulation rapidly and

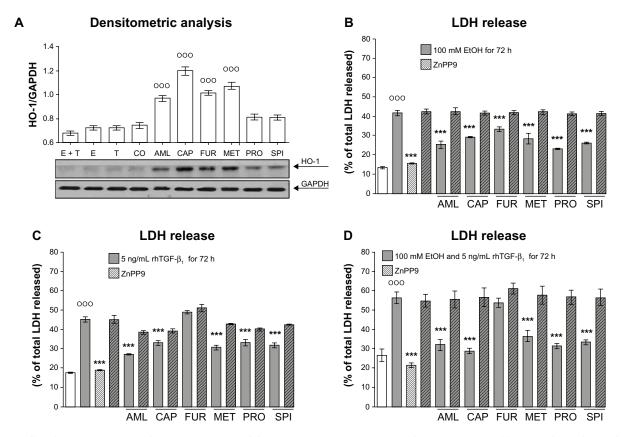


Figure 4 Effect of antihypertensives on HO-1 expression in hHeps. (**A**) Representative Western blot picture of hHeps treated with 100 mM E and/or 5 ng/mL rhTGF- β_1 as well as 4 μM AML, 100 μM CAP, 20 μM FUR, 50 μM MET, 20 μM PRO, or 50 μM SPI for 24 hours. The membrane was probed for HO-1. GAPDH was used as loading control. LDH release of hHeps (N = 3, n = 4) treated for 72 hours with (**B**) 100 mM E and/or (**C** and **D**) 5 ng/mL rhTGF- β_1 in the presence or absence of antihypertensives and 10 μM of the HO-1 inhibitor ZnPP9.

Notes: $^{\circ\circ}P < 0.001$ as compared to untreated cells; $^*P < 0.05$; $^{***}P < 0.01$; $^{***}P < 0.001$ as compared to E- or rhTGF- β_1 -treated cells. Data are presented as mean \pm SEM.

Abbreviations: AML, amlodipine; CAP, captopril; CO, control (untreated cells); E and EtOH, ethanol; FUR, furosemide; hHeps, human hepatocytes; HO-I, heme-oxigenase-I; LDH, lactate dehydrogenase; MET, metoprolol; PRO, propranolol; rhTGF- β_1 , recombinant human transforming growth factor beta I; SPI, spironolactone; T, treated with rhTGF- β_1 ; ZnPP9, Zinc (II) Protoporphyrin 9.

dose-dependently induced ROS formation, which was even further enhanced by co-incubation with rhTGF- β_1 . These results resemble our earlier observations in primary mouse hepatocytes.7 Increased formation of ROS in combination with reduced production of antioxidant enzymes and chemicals, particularly cellular glutathione, 19 causes damage to hepatocytes. In our experiments approximately 40%-60% of the hHeps were damaged after 72 hours exposure to 100 mM ethanol, with damage mainly due to apoptosis and not necrosis.3 In ALD patients, ethanol affects the immune system by altering cytokine production.⁶ Infiltrating macrophages in liver secrete profibrotic cytokines, eg, TGF- β_1 , whose signal transduction in the different liver cell types is critically required for chronic liver disease progression. TGF- β , activates the resident hepatic stellate cells to produce excessive extracellular matrix and even more TGF-β,, which in turn further activates yet quiescent surrounding hepatic stellate cells. rhTGF-β, stimulates ROS production in rat hepatocytes by upregulating NOX.8

In our in vitro experiments, exposure to $rhTGF-\beta_1$, similar to ethanol, dose dependently induced ROS formation. Interestingly, co-incubation of hHeps with ethanol and $rhTGF-\beta_1$, in contrast to murine hepatocytes in which pretreatment of the cells with ethanol sensitized them toward damage from $rhTGF-\beta_1$, did not significantly increase the sensitivity of the cells.⁷ However, this might be due to the higher basal toxicity levels in the hHeps.

We showed earlier that the calcium channel blockers nifedipine and verapamil protected hHeps from ethanoland rhTGF- β_1 -dependent damage by upregulating the antioxidative enzyme HO-1.³ Today these calcium channel blockers are mainly used for the treatment of hypertensive crises. For a more general control of high blood pressure, a great variety of drugs are available, eg, novel calcium channel blockers, diuretics, ACE-inhibitors, aldosterone antagonists, renin inhibitors, angiotensin II receptor antagonists, adrenergic receptor agonists, or α - and β -blockers. Therefore, we tested the effect of the

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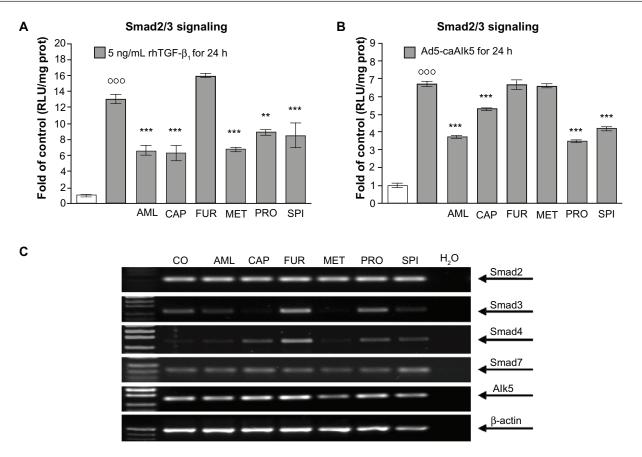


Figure 5 Effect of the antihypertensives on TGF- β signaling. Adenoviral Smad3/4-reporter assay (Ad5-CAGA $_9$ -MLP-Luc) of hHeps (N = 4, n = 6) stimulated for 24 hours with/without antihypertensives and (**A**) 5 ng/mL rhTGF- β_1 or (**B**) co-infected with constitutive active Alk5. (**C**) Representative RT-PCR picture for Smad2, Smad3, Smad4, Smad7, and Alk5 of hHeps treated with 4 μM AML, 100 μM CAP, 20 μM FUR, 50 μM MET, 20 μM PRO, or 50 μM SPI for 24 hours.

Notes: $^{\infty\circ}P < 0.001$ as compared to untreated cells; $^*P < 0.05$; $^{**P} < 0.01$; $^{***P} < 0.001$ as compared to rhTGF- β_1 -treated cells. Data are presented as mean ± SEM.

Abbreviations: AML, amlodipine; CAP, captopril; CO, control (untreated cells); FUR, furosemide; hHeps, human hepatocytes; MET, metoprolol; PRO, propranolol; prot, protein; rhTGF- β_1 , recombinant human transforming growth factor beta 1; RLU, relative light unit; RT-PCR, reverse transcription polymerase chain reaction; SPI,

commonly used antihypertensives amlodipine, captopril, furosemide, metoprolol, propranolol, and spironolactone on ethanol- and rhTGF-β₁-treated hHeps. All substances reduced ethanol-induced ROS accumulation and the resulting cellular damage in a dose-dependent manner. The best protection was provided by captopril, metoprolol, and spironolactone, while a significant effect was only observed for amlodipine and propranolol at very high doses close to the LC₅₀. Similarly, captopril, metoprolol, and spironolactone protected hHeps best from rhTGF-β,induced cellular damage, while the effect of amlodipine and propranolol was strongly dose dependent. Results from the first reports on the in vivo effect of amlodipine and propranolol on myocardial and/or renal fibrosis varied from antifibrotic to noneffective, 20-25 which might be a consequence of the respective applied doses. As a possible protective mechanism of amlodipine, a reduction in ROS by upregulating Cu/Zn superoxide-dismutase was reported;26 captopril was also reported to reduce oxidative stress in vivo and thus to reduce myocardiac fibrosis. 12,27

In the present experiments, all antihypertensives dose dependently reduced ethanol- or rhTGF-\(\beta\),-induced ROS formation. The protective effect observed when (poly)phenolic drugs were added to hHeps together with ethanol and/ or rhTGF- β , can be explained partly by the fact that (poly) phenolic substances due to their structure have strong free radical scavenging properties. As in the example of nifedipine and verapamil, they also are able to induce expression of the antioxidative enzyme HO-1.3 HO-1 is of particular interest as it is upregulated during oxidative stress and helps to protect the liver against damage from several chemical compounds, such as acetaminophen, carbon tetrachloride, and heavy metals.²⁸ Thus, induction of HO-1 by cobalt protoporphyrin or the less toxic natural polyphenols quercetin and diallyldisulfide protect hHeps from ethanol-induced cytotoxicity. 9-11 Captopril, furosemide and metoprolol were the strongest inducers of the antioxidative enzyme HO-1. Inhibition of

spironolactone.

HO-1 activity by ZnPP9 partly reversed the hepatocyte damage inhibitory effect of the drugs, identifying HO-1 as a key downstream player of antihypertensives in mediating tissue protection. However, in contrast to ethanol, where furosemide significantly reduced cellular damage, rhTGF- β_1 -dependent cellular damage was not affected by this drug. This suggests that besides antioxidative action, another mechanism is involved in the protective effect of the drugs.

In primary mouse hepatocytes, we found that furosemide reduces expression of alcohol-dehydrogenase 1 (data not shown), which has been described to negatively influence hepatocyte survival to these stimuli in mouse hepatocytes.⁷ TGF-β,-dependent Smad3/4 signaling is crucial for progression of both renal and hepatic fibrosis. Besides furosemide, all drugs tested significantly inhibited rhTGFβ₁-dependent Smad3/4 signaling. The best inhibition of profibrotic Smad3/4 signaling was observed with amlodipine, captopril, and metoprolol, followed closely by spironolactone and propranolol. Furosemide by trend even enhanced Smad3/4 signaling. These results are supported by several in vitro and in vivo findings that showed reduced TGF- β_1 expression and activity upon treatment with these antihypertensives, mostly in renal cells. 21,29-32 For example, amlodipine reduced adriamycin-induced toxicity in rat mesangial cells by affecting the expression of TGF- β_1 , Smad2, and Smad4.31 Furthermore, captopril effectively decreased plasma levels of TGF- β_1 , proteinuria, and uric acid in hypertensive kidney transplant patients.³² In addition, captopril limited TGF-β, over-expression in acute normotensive anti-Thy1 glumerulonephritis.³⁰ In patients with chronic kidney disease, spironolactone effectively reduced renal fibrosis and urinary TGF- β_1 excretion, emphasizing the link between both parameters;²⁹ similarly, it may prevent renal dysfunction and fibrosis induced by tacrolimus.³² Interestingly, metoprolol does not display antifibrotic effects in transgenic mice with TGF-β₁-induced fibrosis.³³

To more closely investigate the Smad3/4-inhibitory mechanism of the drugs, we co-infected the cells with an adenovirus that expresses a constitutively active Alk5-receptor, which induces Smad3/4 signaling independent of ligand binding to the receptors. In this setup, metoprolol could no longer inhibit Smad3/4 signaling, suggesting an interference of the drug with ligand/receptor binding. Interestingly, Alk5 was reduced by metoprolol at the mRNA level, but none of the other drugs that were investigated. Smad2 and Smad7 expression levels were not significantly affected by any of the antihypertensives. However, Smad3 and Smad4 expression levels were strongly reduced, especially in hHeps

treated with amlodipine, captopril, metoprolol, propranolol, and spironolactone. *Smad3* and *Smad4* expression levels were notably increased with furosemide, which may be an explanation for the enhanced signaling observed.

Our data suggest that high blood pressure treatment with even low doses of captopril, metoprolol, or spironolactone may additionally have beneficial effects on liver fibrosis in patients with ALD. Although amlodipine and propranolol also reduced ethanol-and rhTGF-β,-dependent cellular damage, the effect was strongly dose dependent and present only at higher concentrations, exacerbating the choice of the therapeutic dose. This is supported by animal studies effectively using propranolol to delay progression of sclerosing cholangitis in multi-drug resistance protein 2 (MDR2)-knockout mice.¹⁶ Furthermore, captopril but not amlodipine was shown to reduce fibrotic markers in experimental nonalcoholic steatohepatitis (NASH)¹⁷ or after bile-duct ligation.¹⁵ However, captopril is also reported to directly interfere with matrix metalloproteinase activity, thus inhibiting the degradation of extracellular matrix and further favoring fibrotic tissue changes. 13 On the other hand, captopril may induce upregulation of metallopeptidase with thrombospondin type 1 (ADAMTS-1), which is known to accelerate the degradation of type I collagen,34 which would compensate for the above. Interestingly, furosemide protected hHeps only against ethanol-induced damage but not against rhTGF-β₁-induced damage. We could show that despite an induction of HO-1, furosemide enhances profibrotic TGF-β₁induced Smad3/4 signaling, suggesting that this drug might even favor liver fibrosis in vivo. Special care has to be taken when different antihypertensives are used in combination. Eplerenone, for example, is reported to potentiate the effect of amlodipine against cardiac inflammation and fibrosis in salt-sensitive hypertensive rats.³⁵ Similarly, in a rat model of genetic hypertrophic cardiomyopathy, a combination therapy of low-dose spironolactone and captopril more effectively reduced fibrosis markers when compared with the single drugs.36

In summary, our data suggest that antihypertensives might influence progression of liver fibrosis by modulating HO-1 activity in liver cells. Increased HO-1 activity protects hHeps from ROS-dependent damage by increasing cellular GSH. Under these conditions GSH favors formation of nontoxic products from ROS. Although all drugs tested induced HO-1 expression in hHeps, protecting them from excessive ROS production, the effect of amlodipine and propranolol was only observed at high concentrations close to the ${\rm LC}_{50}$. Therefore, drugs, eg, captopril, metoprolol, or spironolactone, already

protecting hHeps from ethanol- and $rhTGF-\beta_1$ -induced damage at low concentrations, might be more suitable to reduce liver fibrosis in vivo. Furthermore, furosemide even induced profibrogenic $TGF-\beta_1$ signaling, suggesting that this drug might facilitate liver fibrosis in vivo. Therefore, the choice of antihypertensive used for reducing high blood pressure might be crucial in patients with ALD as they might, in the worst case, further promote liver fibrosis, while in the best case even reduce or prevent its progression. Thus, our in vitro data suggest antihypertensive therapy could be adjusted in order to support the antifibrotic therapy, which needs to be further explored in vivo.

Acknowledgments

This work was partially supported by the Federal Ministry of Research (BMBF - 0315208E; BMBF - 01GN0984).

Author contributions

Sabrina Ehnert, Teresa Lukoschek, and Andreas K Nussler designed the study. Sabrina Ehnert, Teresa Lukoschek, Anastasia Bachmann, and Juan José Martínez Sánchez carried out the research. Sabrina Ehnert, Terersa Lukoschek, and Sebastian Mueller analyzed the data. Natascha C Nussler, Stefan Pscherer, and Andreas K Nussler provided the material for the study. Sabrina Ehnert, Teresa Lukoschek, Anastasia Bachmann, and Andreas K Nussler wrote the paper. Anastasia Bachmann, Georg Damm, Stefan Pscherer, Steven Dooley, and Sebastian Mueller critically reviewed the paper.

Disclosure

The authors report no conflicts of interest in this work.

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