Research Article

Isoniazid and Rifampicin Produce Hepatic Fibrosis through an Oxidative Stress-Dependent Mechanism

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Background & Aims. Chronic hepatitis (CH) has emerged as a distinct outcome of drug-induced liver injury (DILI). Combination therapy of Isoniazid (INH) and Rifampicin (RMP) which is widely used for prolonged periods can cause acute hepatotoxicity and has been also incriminated in chronic DILI. We sought evidence of the production of hepatic fibrosis on long-term INH-RMP treatment through experiments in BALB/c mice exposed to INH-RMP.

Methods. A combined dose of INH (50 mg) and RMP (100 mg) per kg body weight per day was administered to mice by oral gavage, 6 days a week, for 4 to 24 weeks for the assessment of liver injury, oxidative stress, and development of hepatic fibrosis, including demonstration of changes in key fibrogenesis linked pathways and mediators.

Results. Progressive increase in markers of hepatic stellate cell (HSC) activation associated with changes in matrix turnover was observed between 12 and 24 weeks of INH-RMP treatment along with the elevation of liver collagen content and significant periportal fibrosis. These were associated with concurrent apoptosis of the hepatocytes, increase in hepatic cytochrome P450 2E1 (CYP2E1), NADPH oxidase (NOX) activity, and development of hepatic oxidative stress.

Conclusions. INH-RMP can activate HSC through generation of NOX-mediated oxidative stress, leading to the development of liver fibrosis.

1. Introduction

Conclusive evidence demonstrating a cause-effect relationship between drug hepatotoxicity and development of liver fibrosis is lacking. However, several large and well-characterized drug-induced liver injury (DILI) registries, based on prolonged follow-up of well-characterized acute DILI subjects, have shown that chronic hepatitis (CH) can occur as a distinct outcome in DILI [1–5]. Persistent liver function derangements can occur in 5.7% to 18.9% of acute DILI subjects [6, 7]. In addition, histological and clinical features of CH have been observed in drug hepatotoxicity cohorts and case studies [8–10]. In most of these cases, the frequency of development of chronic DILI increases as the period of observation of the DILI cohort increases. In the absence of a biomarker for precise DILI definition as well as one indication of evolution to chronicity, the entity chronic DILI remains enigmatic despite its significance. In order to bring clarity on the issue, well-designed experimental studies are needed. In this context, looking for evidence of activation of hepatic stellate cells (HSCs) as the key player in CH and morphological proof for production of liver fibrosis by the drug is important.

Isoniazid (INH) and Rifampicin (RMP) combination therapy is one of the commonest cause to develop acute hepatotoxicity. INH is the primary toxin, and RMP potentiates its toxicity through altered kinetics of metabolites [11, 12]. Recovery from acute hepatitis, clinical or subclinical, generally occurs in clinical settings. Usually, the drugs can be continued thereafter for the originally planned duration of treatment for at least 6 months (often 9–12 months), often in a modified dosage or schedule depending on the presence...
or absence of liver function alterations [11–17]. Overall, INH-RMP combination treatments are associated with overt or indolent and covert hepatocyte functional changes in a significant group of exposed people and hence have the potential to cause activation of HSCs. In view of the prolonged nature of the whole process, this leads to liver fibrosis. Over and above, we have earlier demonstrated, in short term in vivo studies in BALB/c mice, that INH-RMP causes mitochondrial permeability changes and oxidative stress along with hepatic apoptosis [18]. Each of these has the potential to activate HSCs.

In the present study, we are seeking experimental evidence for a relationship between prolonged INH-RMP treatment and development of liver fibrosis. We describe here the findings of an in vivo study in BALB/c mice treated with INH-RMP. We wanted to address three pertinent questions in this study: (1) Can INH-RMP cause hepatic fibrosis on long-term exposure? (2) Is there any evidence for HSC activation along with associated alterations in the matrix proteins to substantiate establishment of a profibrogenic milieu on long-term INH-RMP treatment? (3) Does oxidative stress contribute to HSC activation and fibrosis on INH-RMP exposure, with an eye to get mechanistic insights in the process?

2. Materials and Methods

2.1. Animals and Their Treatment Schedule. Male mice (BALB/c; 7–8 weeks of age) were purchased from National Center for Laboratory Animal Sciences (NCLAS; Hyderabad, India). Treatment of animals and procedures performed were done in accordance with the guidelines stipulated by the animal ethics committee of the Institute of Post Graduate Medical Education & Research (I.P.G.M.E.&R.), Kolkata, India.

Mice (n = 24) were treated with a combined dose of INH (50 mg) and RMP (100 mg) per kg body weight per day by gavage, 6 days a week, for 4 to 24 weeks. The dosage regimen was based on our previous report [18]. Control mice received an equal volume of vehicle by gavage in the same schedule of the INH-RMP-treated mice.

2.2. Animal Sacrifice and Sample Collection. During the period of sacrifice, the blood was obtained by cardiac puncture and the serum samples were stored at -20°C for the measurement of alanine aminotransferase (ALT). The liver was removed, rinsed with phosphate-buffered saline (PBS), and divided into four portions: (a) fixed in 10% buffered formaldehyde (formalin) and embedded in paraffin; (b) homogenized in appropriate buffer(s) and aliquots were frozen at -70°C for biochemical assays; (c) placed in RNA later (from Ambion) solution for RNA expression study; and (d) snap frozen at -70°C for future use.

2.3. Serum Aminotransferases. ALT activity of serum was measured with a commercial kit (DiaSys Diagnostic Systems GmbH, Germany) according to the manufacturer’s instruction.

2.4. Hepatic Biochemical Assays. A 10% liver homogenate was used for the determination of triglyceride (TG) content using a spectrophotometric kit from Sigma Diagnostics (St. Louis, MO, USA), levels of reduced glutathione (GSH), oxidized glutathione (GSSG), thiobarbituric acid reactive substances (TBARs), and protein content [19–22]. Activities of hepatic catalase, glutathione peroxidase (GPx), superoxide dismutase (SOD), cytochrome P450 2E1 (CYP2E1), and NADPH oxidase (NOX) were also determined [23–27]. The collagen content of the liver tissue was measured as described previously [28].

2.5. Histology and TUNEL Assay. Liver tissues embedded in paraffin were cut in sections (5 μm) and stained with hematoxylin and eosin (H&E) and Sirius red for collagen I detection using standard procedures. Terminal deoxynucleotidyl transferase-mediated deoxyuridine triphosphate nick end labeling (TUNEL) assays were performed using the in situ cell death detection kit (Roche, Germany) according to the manufacturer’s instruction. The extent of injury, apoptosis, and fibrosis was evaluated by an investigator, who was blinded to the experimental protocol and graded for steatosis by determining the overall percentage of liver parenchyma containing lipid vacuoles, with 0 = none, 1 = mild (<25%), 2 = mild to moderate (25 to <50%), 3 = marked (50 to <75%), and 4 = severe (>75%) [29]. Inflammation was graded by the presence or absence of inflammatory cells, with 0 = absent, 1 = minimal or focal occasional single clusters of inflammatory cells present in a few microscopic fields, 2 = mild inflammation, 3 = moderate inflammation, and 4 = marked inflammation [30]. The pattern of fibrosis was graded with 0 = none, 1 = portal fibrosis, 2 = perilobular fibrosis or rare septa, 3 = septal fibrosis and architectural distortion but not true cirrhosis, and 4 = cirrhosis, widespread fibrosis, and hepatocyte nodule formation [31].

2.6. Immunostaining. Immunohistochemistry of α-smooth muscle actin (α-SMA) was performed from the paraffin-embedded sections of the liver. Briefly, deparaffinized liver sections were washed in deionized water for 1 minute and in PBS for 5 minutes, followed by permeabilization in 0.1 M citrate buffer and then blocked using PBS with 3% bovine serum albumin (BSA). The liver section was then incubated with Cy3 conjugate α-SMA antibody (C6198; Sigma) at 4°C overnight. After washing, the nuclei were stained with Hoechst (Sigma; 33270) for 5 minutes, washed with PBS, and mounted using Prolong Gold Antifade reagent (Invitrogen; P 36934). Slides were examined by confocal microscopy (Leica, TCS SPE, Germany).

2.7. RNA Isolation and Real-Time Quantitative PCR. Total RNA from liver tissue was obtained using the TRIzol Reagent (Invitrogen, Carlsbad, CA). A High-Capacity cDNA Reverse Transcription kit (Applied Biosystems) was used to generate cDNA from extracted RNA. Primers used for quantitative real-time polymerase chain reaction (qRT-PCR) are shown in Table 1. qRT-PCR was carried out on cDNA using StepOnePlus thermocycler (ABI), primer sets, and SYBR® green PCR master mix (Applied Biosystems) according to the
manuactor’s instructions. Data were normalized to the expression of β-actin, a housekeeping gene.

2.8. Caspase Activity. The activity of caspase 3 was determined in liver homogenates by measuring proteolytic cleavage of the specific fluorogenic substrate DEVD-AFC (Asp-Glu-Val-Asp) (AFC: 7-amino-4-trifluoromethyl coumarin, respectively; BioVision). The results are expressed as the percentage of control.

2.9. Cytokine Quantification. Hepatic transforming growth factor β-1 (TGF-β1) levels were evaluated by enzyme-linked immunosorbent assay (ELISA) using quantikine kits of R&D system.

2.10. Western Blotting. For western blotting, tissue lysate was prepared using RIPA buffer (Cell signaling Technology, USA) with protease inhibitors cocktail (Roche Diagnostics, Mannheim, Germany). The lysates were centrifuged at 14,000 × g for 20 min at 4°C. The protein content of the supernatant was determined with the Bradford protein assay (Sigma, USA). Forty micrograms protein was resolved on sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE) after denaturing in sample buffer and transferred onto polyvinylidene fluoride membranes (PVDF; Thermo Fisher Scientific, USA). After blocking with 5% BSA, the blots were probed with the following antibodies: mouse monoclonal α-SMA (1:300; Santa Cruz Biotechnology), mouse monoclonal beta actin (1:1000; Santa Cruz Biotechnology).

### Table 1: PCR primers.

| Primer | Primer sequence |
|--------|----------------|
| NOX1 Forward 5′-CTGACAAGTACTATTACACGAGA-3′ |
| NOX1 Reverse 5′-CATATATGCCACCCGATTAGGGA-3′ |
| NOX2 Forward 5′-CTTCTCAGGGGTCTCCAGTG-3′ |
| NOX2 Reverse 5′-TCTTCAAAACTCTCCGAGT-3′ |
| NOX4 Forward 5′-TAGACTGAGTACAGCTGGA-3′ |
| NOX4 Reverse 5′-TAGGCTCAATGGGAGTCAC-3′ |
| p22phox Forward 5′-TAGGCTCAATGGGAGTCAC-3′ |
| p22phox Reverse 5′-CACCGAGATCTACGATCCGAGT-3′ |
| p47phox Forward 5′-ACAAAGCCTTCGGGAAT-3′ |
| p47phox Reverse 5′-GCCACATGCTTCAATGG-3′ |
| p67phox Forward 5′-GCCACATGCTTCAATGG-3′ |
| PDGF-Rβ Forward 5′-GCCACATGCTTCAATGG-3′ |
| PDGF-Rβ Reverse 5′-GCCACATGCTTCAATGG-3′ |
| α-SMA Forward 5′-ATCAATGCTGACGCAGAT-3′ |
| α-SMA Reverse 5′-ATCAATGCTGACGCAGAT-3′ |
| TGF-β Forward 5′-ATCAATGCTGACGCAGAT-3′ |
| TGF-β Reverse 5′-ATCAATGCTGACGCAGAT-3′ |
| COL1A1 Forward 5′-GAAACCAGGAGTATGCTTG-3′ |
| COL1A1 Reverse 5′-GAAACCAGGAGTATGCTTG-3′ |
| TIMP-1 Forward 5′-GAAACCAGGAGTATGCTTG-3′ |
| TIMP-1 Reverse 5′-GAAACCAGGAGTATGCTTG-3′ |
| MMP2 Forward 5′-GAAACCAGGAGTATGCTTG-3′ |
| MMP2 Reverse 5′-GAAACCAGGAGTATGCTTG-3′ |
| MMP9 Forward 5′-GAAACCAGGAGTATGCTTG-3′ |
| MMP9 Reverse 5′-GAAACCAGGAGTATGCTTG-3′ |
| β-Actin Forward 5′-GAAACCAGGAGTATGCTTG-3′ |
| β-Actin Reverse 5′-GAAACCAGGAGTATGCTTG-3′ |

NOX = NADPH oxidase; PDGF-Rβ = platelet-derived growth factor receptor β; α-SMA = α-smooth muscle actin; TGF-β = transforming growth factor β; COL1A1 = collagen 1A1; TIMP = tissue inhibitor of matrix metalloproteinase; MMP = matrix metalloproteinase.
Biotechnology), mouse monoclonal collagen 1A1 (1:300; Santa Cruz Biotechnology), mouse monoclonal anti Bax (1:300; Santa Cruz Biotechnology), mouse monoclonal Bcl2 (1:300; Santa Cruz Biotechnology), and mouse monoclonal cytchrome c (1:300; Santa Cruz Biotechnology). The immune complexes were visualized using the enhanced chemiluminescence (ECL) method.

2.1.1. Statistical Analysis. Results are expressed as the mean ± SD. Student’s t-test was used to evaluate statistical differences between groups, and the Mann-Whitney test was used for the comparison of histological findings. A p value less than 0.05 was considered significant.

3. Results

3.1. Oxidative Stress and Liver Injury. Liver injury due to prolonged INH and RMP cotreatment in mice was assessed by measuring serum ALT level as well as histological evaluation of liver specimens. We observed a trend towards increase in serum ALT level at 4, 12, and 24 weeks of INH and RMP cotreatment (52.80 ± 3.91 IU/L at 4 weeks of treatment compared to 26.40 ± 5.41 IU/L in control mice, p < 0.01; 63.75 ± 4.45 IU/L at 12 weeks of treatment vs. 26.60 ± 2.26 IU/L in control mice, p < 0.001; and 35.50 ± 5.50 IU/L at 24 weeks of INH and RMP cotreatment against control 24.66 ± 1.52 IU/L, p < 0.001). We therefore assessed activities of hepatic antioxidant enzymes like SOD and other GSH-related enzymes in mice cotreated with INH and RMP. Following 4 weeks of INH-RMP treatment, activities of GPx and catalase were significantly increased (Table 2). From 12 weeks and onwards of cotreatment of INH and RMP to mice, the activities of hepatic SOD, GPx, and catalase were significantly decreased (Table 2), indicating persistent oxidative stress in the liver.

3.2. Prolonged INH-RMP Treatment Causes Hepatic Fibrosis. Histological changes in the liver due to prolonged INH-RMP treatment included fat infiltration, necrosis, inflammation, and, most importantly, hepatic fibrosis. Steatosis was pronounced all through the 24 weeks of exposure (Figures 1(a) and 1(b)).

Liver TG levels increased also on INH-RMP treatment (Figure 1(c)) in a temporal sequence that paralleled hepatic steatosis. Minimal inflammation and absence of fibrosis were observed at 4 weeks, whereas mild to moderate inflammation and mild portal fibrosis in the liver were evident at 12 weeks (Figures 1(d) and 1(e)). There was a progressive increase in the inflammatory cell infiltration, and the extent of periportal fibrosis was observed in the liver of mice treated with INH-RMP at 24 weeks (Figures 1(d) and 1(e)).

INH-RMP treatment induced collagen 1A1 (COL1A1) mRNA (Figure 2(a)) expressions at 12 weeks, which showed further increase at 24 weeks (Figure 2(a)). Changes in hepatic hydroxyproline content, an amino acid specially contained in collagen, paralleled the induction of COL1A1 mRNA expressions, at different time periods of INH and RMP cotreatment (Figure 2(b)).

3.3. INH-RMP Treatment Is Associated with Stellate Cell Activation and Matrix Remodeling. We examined the number of activated HSCs by immunohistochemistry of α-SMA using a confocal microscope at different time periods of INH-RMP treatment to seek evidence for activation of HSCs (Figure 2(c)).

The number of activated HSCs progressively increased over time and showed a relationship with the duration of INH-RMP treatment in mice (Figure 2(d)). In addition, we observed an increase in α-SMA mRNA expression in the liver tissue, beginning at 12 weeks of INH-RMP treatment (Figure 2(e)).

Proliferation and activation status of HSCs were assessed through expression of candidate molecule platelet-derived growth factor receptor β (PDGF-Rβ). As shown in Figure 3(a), PDGF-Rβ mRNA expression showed incremental increase after 12 and 24 weeks, consistent with the activation of HSCs.

Next, we assessed the tissue inhibitor of matrix metalloproteinase-1 (TIMP-1) expression in the liver tissues of mice and observed the induction of TIMP-1 mRNA after 12 and 24 weeks of INH-RMP treatment (Figure 3(b)). TIMP-1 is synthesized and secreted by activated HSCs in response to fibrogenic cytokines, in particular to transforming growth factor β1 (TGF-β1) [32]. There was a significant increase in hepatic TGF-β protein levels after 24 weeks of INH-RMP treatment compared to control mice (Figure 3(d)), a finding that was also confirmed by mRNA expression for TGF-β1 (Figure 3(c)).

Finally, we also assessed mRNA expression of matrix metalloproteinas 2 and 9 (MMP2 and MMP9), matrix

| Parameters | Cont4 | INH-RMP4 | Cont12 | INH-RMP12 | Cont24 | INH-RMP24 |
|------------|-------|----------|--------|-----------|--------|-----------|
| Cytosolic fraction (n = 8) | | | | | | |
| SOD (Sp activity/min/mg protein) | 125.83 ± 5.48 | 142.18 ± 23.13 | 123.46 ± 7.14 | 72.87 ± 17.02 | 121.04 ± 11.61 | 55.06 ± 10.74 |
| GPx (Sp activity/min/mg protein) | 6.44 ± 0.66 | 7.32 ± 0.34 | 6.24 ± 0.59 | 4.61 ± 0.34 | 6.86 ± 0.62 | 2.56 ± 0.22 |
| Catalase (Sp activity/min/mg protein) | 5.93 ± 0.54 | 6.51 ± 0.31 | 6.06 ± 0.39 | 6.45 ± 0.38 | 6.41 ± 0.66 | 2.67 ± 0.36 |

Mice were cotreated with 50 mg INH and 100 mg RMP/kg body weight by oral gavage for 6 days a week for 4, 12, and 24 weeks, respectively, and were subsequently sacrificed. The results are expressed as the mean ± SD of 8 mice in each group. *p < 0.05 vs. INH-RMP treatment for 4 weeks; †p < 0.05 vs. INH-RMP treatment for 4 and 12 weeks. Abbreviations: Cont4, Cont12, and Cont24 are respective controls of 4, 12, and 24 weeks while INH-RMP4, INH-RMP12, and INH-RMP24 are INH-RMP-treated animals for 4, 12, and 24 weeks, respectively.
remodeling-associated molecules [33] which showed marked upregulation between 12 and 24 weeks of INH-RMP treatment (Figures 3(e) and 3(f)).

3.4. Increased Oxidative Stress Is Related with Hepatic Fibrosis in Long-Term INH-RMP Treatment. During metabolism of INH-RMP, significant stress is being generated within the hepatocytes by the formation of reactive oxygen species (ROS) which is a potential mediator of HSC activation.

We evaluated the oxidative stress markers that revealed a significant decrease in hepatic GSH level (Figure 4(a)) and an increase of lipid peroxidation as evident by thiobarbituric acid reactive substance (TBAR) level (Figure 4(c)). All these data suggested the development of oxidative stress during INH-RMP treatment.

In view of the critical role of CYP2E1 in oxidative stress in INH-mediated hepatotoxicity, we measured hepatic CYP2E1 activity. This showed progressive increase from 4 to 24 weeks of INH-RMP treatment (Figure 4(b)).

Parallel to CYP2E1 activity, NOX activity increased progressively with the duration of INH-RMP treatment (Figure 4(d)). To confirm further that INH-RMP activates NOX in a murine liver, a real-time mRNA expression study of the NOX subunits was performed which revealed a significant increase in the expression of different subunits (Figure 4(e)).

3.5. INH-RMP Treatment Increases Apoptosis of Hepatocytes in a Manner Chronologically Relevant to Hepatic Fibrosis. In the current context of long-term exposure to INH-RMP, we studied the incidence of apoptosis in the liver tissue. The number of apoptotic cells showed a gradual increase from 12 to 24 weeks, expressed as an increase in the percentage of TUNEL positive nuclei (Figure 5(a)). Next, we observed a time-dependent decrease in the expression of the specific antiapoptotic protein Bcl-2 (Figure 5(b)), which has been shown to act on the mitochondria and prevent the release of cytochrome c and subsequent caspase activation [34].

As illustrated in Figure 5(b), a progressive increase in translocation of cytochrome c in the cytosol and increased proapoptotic Bax expression at different time points of INH-RMP treatment by western blot were consistent with the findings from the TUNEL assay. To confirm these findings, caspase 3 activity was estimated in the cytosolic fraction...
of the mouse liver and a progressive increase was revealed in INH-RMP-treated mice from 12 to 24 weeks (Figure 5(c)) compared to control mice.

The increased expression of proapoptotic Bax molecules strongly suggests that INH-RMP treatment causes cell death mediated by the mitochondrial apoptotic pathways.

4. Discussion

We show here that INH-RMP treatment in long term in a mouse model can lead to HSC activation and liver fibrosis, acting through liver cell injury mechanisms that involve NOX-dependent oxidative stress and apoptosis of hepatocytes.
Our experiments provide evidence in support of the emerging clinical data for chronic DILI [1–7]. INH caused 2.7% of the chronic DILI in the DILIN data base and is an important component of CH-producing agents in DILI [6, 35]. In this study, we raised two research questions: (a) can INH-RMP produce hepatic fibrosis on long-term exposure, as is commonly used in clinical practice? (b) How can INH-RMP connect with the liver cell injury-repair
mechanisms, including oxidative stress, apoptosis of hepatocytes, and HSC activation pathways in producing liver fibrosis?

Given the exploratory nature of the primary research question, we designed in vivo mouse experiments of long-term INH-RMP exposure to seek relevant morphological
and functional evidence in this respect. This approach provides robust data to suggest existence of a profibrogenic state in the liver on long-term exposure to INH-RMP.

In the in vivo study, we used INH-RMP combinations in order to capture the real life scenario in antitubercular therapy where they are used together for at least 6 months. INH-RMP combination therapy is the most common cause of acute DILI and drug-induced acute liver failure in India [36, 37]. Of the two, INH is the primary hepatotoxic drug in such combinations and RMP modifies the kinetics of toxic metabolite generation through its ability to induce microsomal enzymes. RMP, therefore, primarily plays a role in potentiating INH hepatotoxicity [12, 13, 18]. The doses of INH and RMP used in the present study are about 10 times the human doses on a milligram per kilogram basis; however, they may be equivalent to the human dose but on a body surface area basis [18].

An intriguing aspect of the histology was macrovesicular steatosis along with inflammatory cell infiltration in the early stages of INH-RMP exposure even when fibrosis has already appeared. We have also observed pericellular fibrosis on prolonged therapy—the fibrosis pattern that correlates with steatohepatitis. Histology in INH-RMP hepatotoxicity has been assessed mostly in the setting of acute liver failure and is characterized by varying amounts of necrosis and inflammation. Prominent steatosis has been observed in some human studies of nonliver failure hepatotoxicity [38, 39]. Previously, we have demonstrated that INH-RMP combination causes acute hepatotoxicity through mitochondrial dysfunction, steatosis, and hepatocyte apoptosis [18]. Progressive increase of cell death and inflammation in the liver as observed in the present study are the characteristic features that are associated with chronic liver injury leading to the progression of the development of fibrosis. Cell death is the primary precipitating event that triggers activation of inflammatory and fibrogenic signals. In the current experiments, we observed CYP2E1-dependent and NOX-mediated oxidative stress along with apoptosis increasing linearly over a period of prolonged exposure. Oxidant stress stimulates apoptosis, and we could document an increase in caspase 3, cytoplasmic translocation of cytochrome c, and reciprocal expressions of the proapoptotic Bax and the antiapoptotic BCL2 proteins in the current study, indicating the previously described mitochondrial pathways of apoptosis to be active even during the prolonged therapy periods. Further, we found increased expression of NOX that produces ROS and stimulates HSCs, over the entire duration of experiments, suggesting a nonmitochondrial pathway of oxidative stress generation also to be active. Apart from the conventional phagocytic NOX2, the nonphagocytic NOX4 was progressively expressed in the liver due to INH-RMP treatment.

In the context of HSC activation, it is important that both oxidative stress and hepatocyte apoptosis are potent mitogens for HSCs [40–42]. We, interestingly, found a steady
time sequence relationship of the events that led to liver fibrosis. In the present study, we have seen the activation of the HSCs that depend largely on oxidative stress. Liver fibrosis results from deposition of type I collagen with simultaneous inhibition of its degradation. In the present study, we observed increased TIMP1 mRNA expression in the liver due to prolonged exposure of INH-RMP treatment in mice. TIMP1 is synthesized and secreted by activated HSCs under the influence of TGF-β1, which is also increased in the liver of mice due to INH-RMP treatment in the present study. Thus, progressive building up of oxidative stress over time was correlated with expression of HSC activation and proliferation markers starting 12 weeks of exposure of INH-RMP. This, along with activation of remodeling matrix (MMP2, MMP9, and TIMP1) and increasing COL1A1 mRNA expressions and collagen content increments, and most importantly, perportal fibrosis evident on histology was at maximum expressions at 24 weeks.

The strength of the present study is in the robustness as well as the novelty of the datasets, with the in vivo designs pursued to address the primary question of HSC activations and fibrogenesis in chronic drug toxicity. Additionally, the ability to demonstrate with precision the relevant pathophysiological changes in a sequential manner beginning with cell injury and finally the pathways that mediate the changes described are all too convincing. We believe this to be the first detailed morphological, functional description of development liver fibrosis, the critical component in CH, in the setting of DILI, and the connotations of the findings are fairly wide.

In conclusion, we have been able to demonstrate that prolonged therapy with INH-RMP can lead to HSC activation and liver fibrosis in a mechanism that is dependent on oxidative stress. Our study provides initial experimental evidence to a simmering body of clinical data suggesting drugs to be important agents in CH.

**Abbreviations**

CH: Chronic hepatitis  
DILI: Drug-induced liver injury  
INH: Isoniazid  
RMP: Rifampicin  
HSCs: Hepatic stellate cells  
CYP2E1: Cytochrome P450 2E1  
NOX: NADPH oxidase  
ALT: Alanine aminotransferase  
PBS: Phosphate-buffered saline  
TG: Triglyceride  
GSH: Reduced glutathione  
GSSG: Oxidized glutathione  
TBARs: Thiobarbituric acid reactive substances  
GPx: Glutathione peroxidase  
SOD: Superoxide dismutase  
H&E: Hematoxylin and eosin  
TUNEL: Terminal deoxynucleotidyl transferase-mediated deoxyuridine triphosphate nick end labeling  
α-SMA: α-Smooth muscle actin  
TGF-β: Transforming growth factor β  
ELISA: Enzyme-linked immunsorbent assay  
COL1A1: Collagen 1A1  
TIMP: Tissue inhibitor of matrix metalloproteinase  
MMP: Matrix metalloproteinase  
PDGF-Rβ: Platelet-derived growth factor receptor β  
ROS: Reactive oxygen species  
SDS-PAGE: Sodium dodecyl sulfate–polyacrylamide gel electrophoresis  
PVDF: Polyvinylidene fluoride  
ECL: Enhanced chemiluminescence  
quRT-PCR: Quantitative real-time polymerase chain reaction  
SD: Standard deviation.

**Data Availability**

The data used to support the findings of this study are included within the article.

**Disclosure**

The present work was published as an abstract in *Journal of Clinical and Experimental Hepatology*. The reference for the abstract is DOI 10.1016/j.jceh.2015.07.160.

**Conflicts of Interest**

The authors declare no conflict of interest regarding the publication of this article.

**Authors’ Contributions**

Ayan Biswas and Suman Santra both equally contributed to this work and are co-first authors.

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