

RESEARCH PAPER

A new histone deacetylase inhibitor improves liver fibrosis in BDL rats through suppression of hepatic stellate cells

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BACKGROUND AND PURPOSE

Activation of hepatic stellate cells (HSCs) is a crucial step in the pathogenesis of hepatic fibrosis. Histone deacetylase (HDAC) is an attractive target in liver fibrosis because it plays a key role in gene expression and cell differentiation. We have developed a HDAC inhibitor, N-hydroxy-7-(2-naphthylthio)heptanomide (HNHA), and investigated the anti-fibrotic activity of HNHA in vitro and in vivo.

EXPERIMENTAL APPROACH

We investigated the anti-fibrotic effect of HNHA on mouse and human HSC activation *in vitro* and in the liver of bile duct-ligated (BDL) rats *in vivo* using cell proliferation assays, cell cycle analysis, biochemical assay, immunohistochemistry and Western blots. Liver pathology was assessed with histochemical techniques.

KEY RESULTS

HNHA inhibited proliferation and arrested the cell cycle via p21 induction in HSCs. In addition, HNHA induced apoptosis of HSCs, which was correlated with reduced COX-2 expression, NF-κB activation and cell death signals. HNHA restored liver function and decreased the accumulation of extracellular matrix in the liver via suppression of HSC activation in BDL rats *in vivo*. HNHA administration also increased survival in BDL rats.

CONCLUSIONS AND IMPLICATIONS

HNHA improved liver function, suppressed liver fibrosis and increased survival of BDL rats, accompanied by reduction of cell growth, activation and survival of HSCs. These findings show that HNHA may be a potent anti-fibrosis agent against hepatic fibrosis because of its multi-targeted inhibition of HSC activity *in vivo* and *in vitro*.

Abbreviations

ALT, alanine aminotransferase; AST, aspartate aminotransferase; BDL, bile duct ligation; COL-I, collagen type I; ECM, extracellular matrix; H&E, haematoxylin and eosin; HNHA, N-hydroxy-7-(2-naphthylthio)heptanomide; HSC, hepatic stellate cell; SAHA, suberoylanilide hydroxamic acid; Tbil, total bilirubin; α -SMA, α -smooth muscle actin



Links to online information in IUPHAR/DB and BPS Concise Guide to PHARMACOLOGY

Targets	Ligands
Caspases	Bilirubin
Collagen	Ibuprofen
COX-2, cyclooxygenase 2	Meloxicam
HDAC, histone deacetylase	SAHA (vorinostat)
MMP, matrix metallopeptidase	TGF-β1

This table lists key protein targets and ligands in this article which are hyperlinked to corresponding entries in http://www.guidetopharmacology.org, the common portal for data from the IUPHAR/BPS Guide to PHARMACOLOGY (Pawson et al., 2014) and are permanently archived in the Concise Guide to PHARMACOLOGY 2013/14 (Alexander et al., 2013).

Introduction

Liver fibrosis is an overactive wound healing process in which excessive connective tissue builds up in the liver and results in the distortion of hepatic architecture and liver failure (Branton and Kopp, 1999). Chronic liver injury activates and transforms quiescent hepatic stellate cells (HSCs), which transdifferentiate into activated HSCs, called myofibroblasts, acquiring contractile, proliferative and fibrogenic properties as the primary cellular source of extracellular matrix (ECM) components in the liver (Moreira, 2007). As a consequence, the excessive accumulation of ECM initiates fibrosis by releasing inflammatory mediators such as PDGF, TGF and connective tissue growth factor (Pellicoro et al., 2012). Therefore, suppression of HSC activation and proliferation as well as induction of apoptosis in activated HSCs have been proposed as therapeutic strategies for the treatment and prevention of hepatic fibrosis (Elsharkawy et al., 2005).

Anti-fibrotic drugs are not efficiently taken up by activated HSCs and are often toxic to hepatic cells, producing unwanted side effects, and there are no proven drugs to treat liver fibrosis in patients (Bataller and Brenner, 2005). Recent studies suggest that liver fibrosis can be blocked by induction of histone hyperacetylation by histone deacetylase (HDAC) inhibitors (Kaimori et al., 2010; Liu et al., 2013). We have already described the HDAC inhibitor, N-hydroxy-7-(2naphthylthio)heptanomide (HNHA), as inhibiting nuclear HDAC enzyme activity in human umbilical endothelial cells and breast cancer cells, an effect that was accompanied by histone hyperacetylation and cell cycle arrest, as well as inhibition of ECM regulators such as MMPs (Kim et al., 2007; Park et al., 2011). However, the effects of HNHA on the regulation of fibrogenic responses in HSCs and liver fibrosis have not been studied.

Models of liver injury following bile duct ligation (BDL) in rats have been widely used to dissect the molecular mechanisms underlying acute and chronic liver injury. Here, we evaluated the biological effects of HNHA and other known treatments on the fibrogenic responses, including the proliferation, apoptosis and activation of mouse and human HSCs, and have also evaluated the anti-fibrotic effect of HNHA in liver fibrosis induced by BDL in rats.

Methods

Isolation and culture of HSCs

The livers of 6-week-old male C57BL/6 mice (Orient Bio Inc., Seong-nam, South Korea) were sequentially digested with pronase (Roche Applied Science, Mannheim, Germany) and type IV collagenase (Sigma-Aldrich, St Louis, MO, USA) by in situ perfusion for primary HSC isolation. Hepatocytes were removed by centrifuging the digested liver at $50 \times g$ for 3 min. The non-parenchymal cells present in the supernatant were laid on top of a gradient of arabinogalactan (Sigma-Aldrich; with densities of 1.035, 1.045, 1.058 and 1.085), followed by centrifugation at 74 000× g for 40 min at 25°C. The fraction containing HSCs was recovered from the interface between the medium (1.045) and the lowest density (1.035). The purity of HSCs was evaluated by examining the characteristic stellate shape of the cells with phase contrast microscopy and by the presence of lipid droplets by autofluorescence using an excitation light of 320 nm. The viability of HSCs assessed by Trypan blue staining always exceeded 95%. Primary mouse HSCs were cultured in DMEM (Life Technologies, Inc., Grand Island, NY, USA) with 10% FBS and 100 units⋅mL⁻¹ penicillin-streptomycin and were maintained at 37°C in an atmosphere of 5% CO₂. We grew human primary HSCs from human resection specimens (Sciencell, San Diego, CA, USA) and used them for up to five passages. For experiments, we plated cells at a density of 3×10^4 cells·cm⁻². When the cultures reached 70-80% confluence, they were trypsinized and passaged.

Cell proliferation assay, cell viability assay and cell cycle phase distribution by flow cytometric DNA analysis

Mouse or human primary HSCs were pre-cultured for 24 h at a density of 2.3×10^5 cells·mL⁻¹ and the growth medium was replaced with an experimental medium containing ibuprofen, meloxicam, suberoylanilide hydroxamic acid (SAHA) or HNHA at final concentrations of 10 μ M in cell proliferation assay and cell cycle analysis, or ranging from 1 to 160 μ M in the cell viability assay. Growth medium containing 0.1% (v/v) saline was used as a vehicle control. These assays were performed as previously described (Park *et al.*, 2011).



Western blot analysis

Western blots were carried out as previously reported by Park *et al.* (2011) using primary antibodies against Bcl-2, Bcl-w, Bcl-xL, HDAC1, HDAC4, Bad, Bax, Bak, COX-2, cyclin D1 (all from Abcam, Cambridge, UK); phospho-IκB-α (Cell Signaling Technology, Danvers, MA, USA); phospho-NF-κB p65, IκB-α (from Santa Cruz Biotechnology, Santa Cruz, CA, USA); p21 and p53 (Abcam); pro-caspase-3 and cleaved-caspase-3, pro-caspase-7 and cleaved-caspase-7, pro-caspase-8 and caspase-8, pro-caspase-9 and cleaved-caspase-9 (all from Santa Cruz Biotechnology), collagen type I (COL-I) (R&D Systems, Minneapolis, MN, USA), α-smooth muscle actin (α-SMA) (Abcam), TGF-β1 (R&D Systems), MMP-2 and MMP-9 (Abcam), and β-actin (Santa Cruz Biotechnology).

RNA extraction and RT-PCR

Total RNA prepared from HSCs was extracted with RNeasy Mini Kit (Qiagen, Valencia, CA, USA), including treatment with DNase I to prevent genomic DNA contamination using RURBO DNA-free Kit (Ambion, Foster City, CA, USA) according to the manufacturer's instructions. RT-PCR was carried out with total RNA (1 μg) using the one-step RT-PCR kit (Qiagen) according to the manufacturer's protocol. The resultant cDNA was used to carry out 40 PCR cycles consisting of 15 s at 95°C, 30 s at 57°C and 60 s at 72°C on an ABI Prism 7000 sequence detection system (Applied Biosystems, Foster City, CA, USA). The reactions were performed using SYBR green 1 (Qiagen), using primers for p21 (forward 5'-tccacagcgatatccagaca-3', reverse 5'-ggacatcaccaggattggac-3') and Gapdh (forward 5'-atcaagaaggtggtgaagcggaa-3', reverse 5'-tggaagagtgggagttgctgttga-3'). Each PCR was set up in triplicate wells in a total volume of 25 μ L. The reaction mixture contained the cDNA equivalent of 20 ng total RNA. The quantitative values of the genes of interest were normalized using Gapdh as the endogenous reference, and fold-increase over control values was calculated using the relative quantification method of $2^{-\Delta\Delta Ct}$.

Serum biochemistry

For testing the liver function, the activities of aspartate transaminase (AST), alanine transaminase (ALT) and the contents of total bilirubin (Tbil) in the serum were determined after 3 weeks of experimental treatment using an automated analyser (RA-XT; Technicon, Tarrytown, NY, USA).

Animals and experimental design

All animal care and experimental procedures complied with local guidelines and were approved by the Animal Experiments Committee of Yonsei University. All studies involving animals are reported in accordance with the ARRIVE guidelines for reporting experiments involving animals (Kilkenny *et al.*, 2010; McGrath *et al.*, 2010). A total of 111 animals (106 rats, 5 mice) were used in the experiments described here.

Male Sprague Dawley rats (Orient Bio Inc.) weighing 250–300 g (7 weeks old) were used. All animals were housed in cages (five mice or two rats per cage) for 7 days. The light cycle was controlled to provide 12 h light and 12 h darkness; temperature was 22°C and humidity was 40–60%. A standard diet of rodent pellets and tap water (membrane-filter purified and autoclaved) were provided *ad libitum*. Rats were anaes-

thetized with an i.m. injection of Zoletil (30 mg·kg⁻¹; Virbac, Carros, France). After a midline abdominal incision, the common bile duct (CBD) was doubly ligated with 4-0 silk sutures and cut with #11 blade. Sham operation was just exposure of the CBD by passing the cotton applicator below the foramen of Winslow. After saline irrigation, abdominal closure was performed with continuous 3-0 silk suture. After the surgical procedure, animals were divided into five groups in efficacy experiments and survival experiments: sham operation group (SHAM, n = 10); BDL (+Veh) group (n = 20), in which saline was i.p. injected beginning at 24 h after BDL; BDL + meloxicam group (n = 20), in which meloxicam was i.m. injected at 1.6 mg·kg⁻¹ beginning at 24 h after BDL; BDL + SAHA group (n = 20); and BDL + HNHA group (n = 20), in which the treatments (30 mg·kg⁻¹) were given i.p. 24 h after BDL. All drugs were administered once every 3 days for 3 weeks, in efficacy experiments or until death, in survival experiments. For in vivo toxicity experiments, meloxicam was i.m. injected (1.6 mg·kg⁻¹) or HNHA (30 mg·kg⁻¹), beginning at 24 h after the sham operation (n = 16).

Tissue preparation

After 3 weeks, all animals were killed, and the blood, spleens and livers were collected and the organ weights were determined. Plasma was separated by centrifugation at $2300 \times g$ for 5 min at room temperature within 15 min of sample collection. Livers were sliced into several parts. Some were immediately snap-frozen in liquid nitrogen for further analysis and kept at -80° C. Other samples of liver were immersed into 10% neutral buffered formalin solution for histopathological examinations.

Haematoxylin and eosin (H&E) staining, Masson trichrome staining and immunohistochemistry

Liver biopsy specimens fixed in 10% buffered formalin solution were embedded with paraffin. Sections (5 μm) were stained with H&E. The degree of fibrosis was assessed with Masson trichrome staining according to the protocol provided by the manufacturer (Sigma-Aldrich). This procedure stains nuclei dark red, cytoplasm and muscle fibres red, and ECM components blue.

Immunohistochemical analysis was conducted as previously reported (Park *et al.*, 2011) using primary antibodies against acetyl histone H3 (1:200; Cell Signaling Technology, Beverly, MA, USA), HDAC-1 (1:200), COL-I (1:200; Cell Signaling Technology), α -SMA (1:20), TGF- β 1 (1:100; Abcam) and COX-2 (1:500; Santa Cruz Biotechnology). The images were quantified by image analysis using Metamorph 6.0 software (Universal Imaging Corp., Downingtown, PA, USA) according to the software manual. Total area of collagen-stained area and its optical density were calculated from three fields of every slide and compared between groups by non-parametric statistical methods.

Data analysis

Data are expressed as means \pm SEM, unless otherwise stated. Statistical analysis was performed with SPSS software, version 11.5 (SPSS Inc., Chicago, IL, USA). Differences between means were assessed by two-tailed Student's t-test or by one-way



anova followed by Bonferroni's multiple-comparison $post\ hoc$ test. P values lower than 0.05 were considered statistically significant.

Materials

HNHA and SAHA (vorinostat) were provided by Dr H-J. Kwon. Meloxicam and PDGF-B were purchased from R&D Systems.

Results

HNHA inhibits the proliferation and activation of HSCs and induces the cell cycle arrest and apoptosis of HSCs

HNHA treatment for 1 week showed greater inhibition of mouse and human primary HSC proliferation, a feature of HSC activation, compared with ibuprofen (a non-selective COX inhibitor), meloxicam (a COX-2 selective inhibitor) and SAHA (another HDAC inhibitor) (Figure 1A and B).

To assess further the effects of HNHA on HSC activation, the expression of COL-I, $\alpha\text{-SMA}$ and TGF- $\beta1$ in PDGF-stimulated HSCs was examined by Western blot analysis. Stimulation of mouse HSCs with PDGF (20 ng·mL $^{-1}$) for 24 h significantly induced the expression of $\alpha\text{-SMA}$, TGF- $\beta1$ and COL-I expression, activation markers of myofibroblasts. HNHA treatment suppressed this increased expression of $\alpha\text{-SMA}$, TGF- $\beta1$ and COL-I in activated HSCs (Figure 1C).

Incubation with HNHA (10 µM) arrested mouse HSCs in the sub- G_0/G_1 and G_0/G_1 phase (about 90%) more effectively than the other compounds used (about 70%; Figure 1D). Consistent with the observed effects on cell proliferation, this concentration of HNHA also reduced the expression of COX-2, the phosphorylation of $I\kappa B-\alpha$ and NF- κB p65, and the degradation of IκB-α in HSCs, unstimulated or stimulated with PDGF, a key molecule in the sustained activation and proliferation of HSC survival (Figure 1E). Furthermore, HNHA induced p21 and p53, a well-known arrestor of the cell cycle, but reduced cyclin D1, a positive regulator of the cell cycle (Figure 1E, F, H). These results were compatible with the concentration-dependent cytotoxicity of HNHA, which induced greater loss of viability at low concentrations (1-10 µM) than the other inhibitors used (Figure 1G and Table 1).

To investigate pro-apoptotic signalling pathways activated following the incubation with these inhibitors, the expression of pro- (Bad, Bax and Bak) and anti-apoptotic (Bcl-2, Bcl-w and Bcl-xL) members of the Bcl-2 family and the activation of caspase-3, caspase-7, caspase-8 and caspase-9, major effectors in apoptosis, in HSCs as well as the expression of HDACs (HDAC1 and HDAC4) were investigated by Western blotting. HNHA led to increased expression of proapoptotic members of the Bcl-2 family and cleavage of procaspase in HSCs (Figure 1H), which are barely detectable in PDGF-treated HSCs. Moreover, HNHA and SAHA almost abolished PDGF-induced expression of anti-apoptotic members of the Bcl-2 family and decreased HDAC protein levels, key molecules in the signalling pathways of HSC survival, but the COX inhibitors (ibuprofen and meloxicam) were less effective.

HNHA restores liver function in BDL rats

BDL, the most commonly used animal model of obstructive cholestasis, causes obstructive jaundice and cholestasis, which over time leads to histological changes in the liver, including the development of cirrhosis. After 3 weeks of BDL, livers were grossly enlarged and had lost compliance, compared with those from sham-operated rats. BDL caused a significant increase in liver/body weight ratio, serum AST, ALT and Tbil levels, compared with sham-operated rats. In the BDL + HNHA group, the changes in body weights and liver/body weight ratios were significantly less than in the other groups (Table 2). The increased serum activities of AST, ALT and the contents of Tbil were attenuated in the BDL + HNHA group compared with the BDL, BDL + meloxicam or BDL + SAHA groups (Table 3). In addition, there was no evidence of systemic toxicity attributable to the doses of meloxicam, SAHA and HNHA used here. These data suggest that HNHA was the most effective agent for decreasing hepatic damage after BDL in this liver fibrosis model.

HNHA ameliorates histopathological deterioration and ECM accumulation in BDL-liver

Each lobe of the BDL-liver was equally affected grossly by BDL upon examination by light microscopy of liver sections stained with H&E. There was marked architectural distortion of lobular structure and extensive bile duct proliferation infiltrating into hepatocytes with periportal fibrogenesis and occasional fibrogenesis in the parenchymal area, suggesting activation of HSCs. The groups treated with meloxicam, SAHA or HNHA showed an overall reduction of pathological changes compared with the BDL group. The BDL + HNHA group showed a greater suppression of lower bile duct proliferation and periportal fibrosis compared with the meloxicam or SAHA-treated groups (Figure 2A). Because liver fibrosis is characterized by an increased accumulation of ECM, mainly collagens, we assessed the collagen contents based upon Masson's trichrome staining and Western blots for type I collagen. Consistent with the histopathology, HNHA markedly suppressed collagen accumulation in BDL rat livers (Figure 2B) and COL-I expression (Figure 2C). This suggests that HNHA improved the histological changes and reduced ECM accumulation in liver fibrosis.

HNHA suppresses activation of myofibroblasts induced by BDL

Because the appearance of α -SMA-positive myofibroblasts is considered as a key event in the progression of liver fibrosis and TGF- β 1, predominantly expressed by HSCs, is a key mediator for transdifferentiation, we assessed α -SMA or TGF- β 1-positive cells by morphometric quantification to ascertain the accumulation of fibrogenic myofibroblasts. Activated myofibroblasts, as shown by positive staining for α -SMA, were predominant in the peri-ductular zone 1 area and, to a lesser degree, in the peri-sinusoidal area in zone 2 and zone 3. Immunohistochemical staining for TGF- β 1 showed that the positively stained brown-coloured cells were mainly proliferating bile ductules and limiting plate hepatocytes. However, liver samples from rats treated with meloxicam, SAHA or HNHA for 3 weeks after BDL showed less bile duct prolifera-

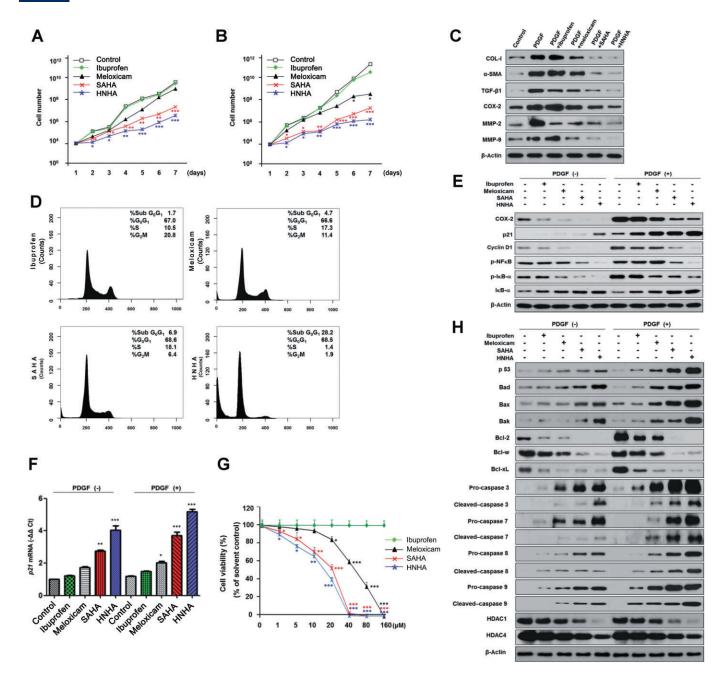


Figure 1

Effects of HNHA on proliferation, apoptosis and activation of HSCs. (A, B) Cell growth assay of mouse (A) or human (B) primary HSCs using cell counting. (C) Western blot analysis of HSC activation markers in mouse HSCs. (D) Cell cycle phase distribution by flow cytometric DNA in mouse HSCs. (E) Western blot analysis of cell proliferation protein levels in activated mouse HSCs. (F) Quantitative RT-PCR analysis of p21 in activated mouse HSCs. Mouse primary HSCs were cultured in complete media with increasing concentrations of drugs for 48 h. (G) Cell viability assay. Mouse primary HSCs were cultured in complete media with increasing concentrations of drugs for 96 h (n = 3). (H) Western blot analysis of apoptosis-related proteins and HDAC levels in mouse HSCs. In the experiments summarized in (C)–(F) and in (H), mouse HSCs were incubated with ibuprofen, meloxicam, SAHA or HNHA at a final concentration of 10 μ M. Data shown are means \pm SD (n = 4 except for G). *P < 0.05, *P < 0.05,

tion and weak or no cytoplasmic immunostaining for α -SMA (Figure 3A) compared with BDL rats. Liver sections from meloxicam, SAHA and HNHA group rats showed weak or no cytoplasmic TGF- β 1 staining in bile duct epithelial cells and limiting plate hepatocytes (Figure 3B). Treatment with HNHA

was more effective in suppressing α -SMA and TGF- β 1 in BDL-livers than treatment with meloxicam or SAHA. We then compared COX-2 expression in liver tissue samples. With vehicle treatment after BDL, COX-2 was expressed mainly in the proliferating bile ductules and, to a lesser degree, in



Table 1 Inhibition of HSC proliferation by drug treatments

	Meloxicam (μM)	SAHA (μM)	HNHA (μM)
Mouse primary HSCs	54 (±1.0)	23 (±1.7)	15 (±1.0)
Human primary HSCs	50 (±1.1)	20 (±1.9)	12 (±1.1)

Each data point represents the mean IC50 value (±SD) from three independent experiments performed in triplicate using sulforhodamine B to assay proliferation.

Table 2 Effects of HNHA on body and organ weight changes in BDL rats

			Organ weight			
Groups		Body weight (g)	Liver (g)	Spleen (g)	Liver/body (%)	Spleen/body (%)
SHAM	Veh	346 ± 7.9	14.00 ± 0.29	1.07 ± 0.07	3.86 ± 0.11	0.30 ± 0.05
	Meloxicam	368 ± 9.0	14.35 ± 0.33	1.12 ± 0.05	3.89 ± 0.10	0.30 ± 0.02
	SAHA	360 ± 4.2	13.56 ± 0.19	1.08 ± 0.02	3.76 ± 0.05	0.29 ± 0.03
	HNHA	353 ± 5.9	13.65 ± 0.17	1.03 ± 0.09	3.86 ± 0.07	0.29 ± 0.05
BDL	Veh	298 ± 4.7	21.13 ± 1.02	2.09 ± 0.12	7.08 ± 0.95	0.70 ± 0.11
	Meloxicam	308 ± 5.5*,##	18.00 ± 0.97*,##	1.80 ± 0.55*,##	5.84 ± 0.45*,##	0.58 ± 0.05*,#
	SAHA	311 ± 6.4*,#	15.74 ± 0.51**,#	1.52 ± 0.43**,#	5.06 ± 0.41**,#	0.49 ± 0.52**,#
HNHA	HNHA	324 ± 5.1**	14.05 ± 2.06***	1.32 ± 0.55***	4.33 ± 0.54**	0.40 ± 0.05**

Meloxicam was injected i.m. at a daily dose of 1.6 mg, and SAHA and HNHA were injected i.p. in rats at a dose of 30 mg·kg⁻¹ once every 3 days for 3 weeks. Data shown are means \pm SEM.

Table 3 Serum biochemistry

AST (IU·L ⁻¹)	ALT (IU·L ⁻¹)	Tbil (mg·L ⁻¹)
69.73 ± 7.98	27.17 ± 4.84	1.1 ± 0.2
342.11 ± 81.92	104.84 ± 38.12	81.4 ± 29.7
189.45 ± 47.12*,##	79.48 ± 29.59*,##	68.4 ± 14.5*,##
113.17 ± 19.75**,#	59.08 ± 18.22**,#	50.7 ± 11.1**,#
98.35 ± 20.44**	46.35 ± 17.24***	34.2 ± 9.5***
77.35 ± 10.01	30.27 ± 4.01	1.8 ± 0.4
76.98 ± 9.24	29.59 ± 3.87	1.7 ± 0.4
76.14 ± 9.97	29.87 ± 3.99	1.6 ± 0.3
	69.73 ± 7.98 342.11 ± 81.92 $189.45 \pm 47.12^{*,\#}$ $113.17 \pm 19.75^{**,\#}$ $98.35 \pm 20.44^{**}$ 77.35 ± 10.01 76.98 ± 9.24	$69.73 \pm 7.98 \qquad 27.17 \pm 4.84 \\ 342.11 \pm 81.92 \qquad 104.84 \pm 38.12 \\ 189.45 \pm 47.12^{*,\#} \qquad 79.48 \pm 29.59^{*,\#} \\ 113.17 \pm 19.75^{**,\#} \qquad 59.08 \pm 18.22^{**,\#} \\ 98.35 \pm 20.44^{**} \qquad 46.35 \pm 17.24^{***} \\ 77.35 \pm 10.01 \qquad 30.27 \pm 4.01 \\ 76.98 \pm 9.24 \qquad 29.59 \pm 3.87$

Meloxicam was injected i.m. at a daily dose of 1.6 mg, and SAHA and HNHA were injected i.p. in rats at a dose of 30 mg·kg⁻¹ once every 3 days for 3 weeks. Serum levels of aspartate aminotransferase (AST), alanine aminotransferase (ALT) and total bilirubin (Tbil) of shamoperated (SHAM) and bile duct ligation (BDL) rats were determined. Data shown are means \pm SEM (n=25). *P < 0.05, **P < 0.01, ***P < 0.005 versus BDL; *P < 0.05, **P < 0.01 versus BDL + HNHA.

hepatocytes. In livers from BDL rats receiving meloxicam, SAHA or HNHA, there was decreased hepatic COX-2 expression (Figure 3C) with the lowest expression in the HNHAtreated group. These results were confirmed by Western blot

analysis of total lysates of livers, which showed that produc-

tion of COL-I, α-SMA, TGF-β1, COX-2, MMP-2 and MMP-9 appeared to be most reduced in the BDL + HNHA group compared with other groups (Figure 3D). The acetylation of histone H3 in liver tissue was much stronger in the HNHAtreated group than in the BDL group (Supporting Informa-

^{*}P < 0.05, **P < 0.01, ***P < 0.005 versus BDL; *P < 0.05, **P < 0.01 versus BDL + HNHA.

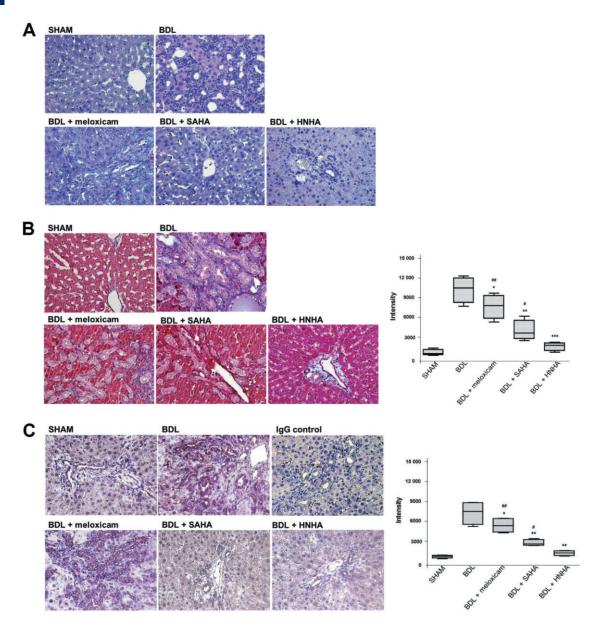


Figure 2

Effects of HNHA on histopathology and ECM accumulation of BDL-livers. (A) The histological changes in the livers of bile duct ligated (BDL) rats as shown by haematoxylin and eosin (H&E) staining. (B, C) Representative staining (left) and densitometric analysis (right) of Masson's trichrome staining (B) and collagen staining (C) of liver tissues of BDL rats. Masson's trichrome staining stains nuclei black, cytoplasm and muscle fibres red, and ECM components blue. Negative control slides (IgG control) were treated with isotype-matched IgG. Quantitative data are shown as box and whisker plots with the median indicated as a horizontal bar, first and third quartiles as the box and the range shown by the whiskers (n = 15). *P < 0.05, *P < 0.05, *P < 0.01, **P < 0.05, *P < 0.05, **P < 0.01, **P < 0.05, **P <

tion Fig. S2), whereas the level of HDAC1 was decreased in the HNHA-treated group compared with the BDL group (Supporting Information Fig. S3).

HNHA increases survival of BDL rats

Meloxicam, SAHA and HNHA were injected i.p. to evaluate their effect on the survival of BDL rat, over 63 days. The survival rate of the HNHA-treated group was higher than that

of the other groups (Figure 4). Administration of the drug treatments to rats without BDL did not affect survival, which was the same as that for sham-operated animals, indicating no systemic toxicity attributable to meloxicam, SAHA and HNHA. These findings were consistent with the results of histopathology or liver function analysis. Taken together, these results showed that HNHA improved liver function and overall survival of rats with BDL.



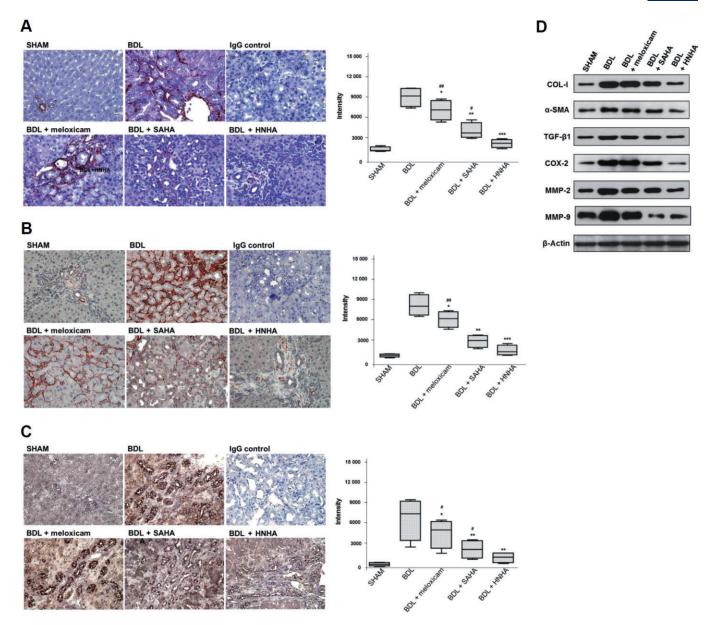


Figure 3

Effects of HNHA on myofibroblast activation in BDL-liver. Representative staining (left) and densitometric analysis (right) of α -SMA (A), TGF- β 1 (B) and COX-2 (C) staining of liver tissues of BDL rats. (D) Representative Western blot of COL-1, α-SMA, TGF-β1, COX-2, MMP-2 and MMP-9 in liver tissues of BDL rats (n = 4). Quantitative data are shown as box and whisker plots with the median indicated as a horizontal bar, first and third quartiles as the box and the range shown by the whiskers (n = 15 except for D). *P < 0.05, **P < 0.01 ***P < 0.005 versus BDL; *P < 0.05, $^{*\#}P < 0.01$ versus BDL + HNHA.

Discussion

The induction of apoptosis in activated HSCs has emerged as a treatment strategy for liver fibrosis (Schuppan and Kim, 2013). HDAC inhibitors can modulate the expression of apoptosis-related genes (Cao et al., 2001) and recent studies have shown that HDAC4 is accumulated during HSC transdifferentiation (Mann et al., 2007; Qin and Han, 2010). For this

reason, the inhibition of HDAC has been considered as a promising suppressor of HSC proliferation and activation in vitro and in vivo (Rombouts et al., 2002; Watanabe et al., 2011). HNHA is a new synthetic HDAC inhibitor with better pharmacological properties than other HDAC inhibitors including SAHA and was more effective in inducing G1/S cell cycle arrest in breast cancer cells, compared with SAHA (Kim et al., 2007). The anti-proliferative effect of HNHA on HSCs

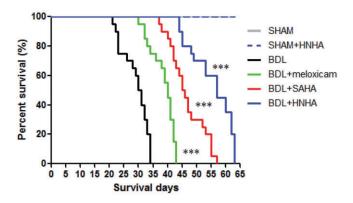


Figure 4

Effects of HNHA on survival in BDL rats. Kaplan–Meier plot for overall survivals. Differences between survival curves for 63 days in groups (SHAM group, n=10; SHAM + HNHA, n=16; BDL group, n=20; BDL + meloxicam group, n=20; BDL + SAHA group, n=20; BDL + HNHA group, n=20) were compared using a log-rank test. ***P<0.005 versus BDL.

appeared also to be mediated through cell cycle arrest. In concordance with the earlier data, we found here that HNHA suppressed cell proliferation and increased, very effectively, cell cycle arrest at the G₀/G₁ phase in mouse and human primary HSC proliferation than ibuprofen, meloxicam or SAHA. These results were related to an increase in the endogenous cyclin-dependent kinase inhibitor, p21, and p53 and accompanied by a decrease in cyclin D1 expression by HNHA, which led to profound inhibition of HSC growth. It is well established that histone hyperacetylation induces apoptosis via p21, whose promoter is amplified from acetylated histone H3- or H4-associated chromatin by HDAC inhibition (Blagosklonny et al., 2002). HNHA also strongly rescued histone acetylation in FM3A and MCF-7 cells (Kim et al., 2007; Park et al., 2011), a function possibly linked to its ability to arrest the cell cycle in HSCs.

Several cytokines are central to the pathogenesis of fibrosis and the process of HSC activation, particularly PDGF and TGF-β1. PDGF induces the activation of the downstream signalling kinase ERK in activated HSCs, which is associated with myofibroblast differentiation, cell proliferation and stimulation of ECM production, the most important event in tissue fibrogenic progression (Marra et al., 1999; Friedman, 2000; Reif et al., 2003). Simultaneously, inhibition of PDGF signalling pathways have also been shown to induce HSC apoptosis and attenuate liver fibrosis (Borkham-Kamphorst et al., 2004). Furthermore, Bcl-2 enhances the resistance to apoptosis in activated HSCs, which may contribute to the development of fibrosis, while increased HSC apoptosis was associated with reduced anti-apoptotic Bcl-2 expression in the liver (Novo et al., 2006). HNHA showed marked inhibition of HSC activation and Bcl-2 expression and led to activation of caspases in PDGF-stimulated HSCs. Therefore, our results suggested that HNHA not only significantly inhibited the proliferation and activation of HSCs through suppression of NF-κB activation but also had a marked apoptotic effect on HSCs through inhibition of Bcl-2 activation, which is important for fibrogenesis.

Quiescent HSCs do not express COX-2 but activated HSCs do express COX-2, suggesting that the COX-2/PG pathway is involved in hepatic fibrogenesis (Rahman et al., 2000; Cheng et al., 2002). Because the effects of non-selective COX inhibitors such as ibuprofen on the fibrogenic response in HSCs and in the liver still remain controversial (Rahman et al., 2000; Cheng et al., 2002), specific inhibition of COX-2 would appear to be safer. Recent studies reported that selective COX-2 inhibitors, including meloxicam, suppressed proliferation of human HSCs (Furst, 1997). Interestingly, inhibitors of HDAC suppress COX-2 expression by blocking the induction of c-Jun and lead to the inhibition of viability and the induction of apoptosis (Yamaguchi et al., 2005). In our experiment, HNHA mainly acted on portal myofibroblasts in cholestasis, by suppressing COX-2 in bile duct epithelium, resulting in diminished collagen formation in activated myofibroblasts. As shown by histology and image analysis of collagen formation in the present study, HNHA showed powerful repression of bile duct proliferation and periportal fibrosis. Cholestatic liver fibrosis led to hepatocyte apoptosis/ necrosis during acute liver damage, transformation of portal fibroblast or HSCs to activated forms of myofibroblasts producing ECM, proliferation of bile duct and increase in secretion of cytokines such as PDGF and TGF-β (Ramm et al., 2009). In our experiments, TGF-β1 was strongly stained in hepatocyte and HSCs and, to a lesser degree, in cholestatic liver, and this staining was diminished by HNHA administration. Consequently, HNHA suppressed α-SMA expression and collagen formation in cholestasis induced by BDL as well as PDGF and TGF-β, effects confirmed in primary HSCs. In addition to expressing type I collagen, activated HSCs also express MMPs, which are important in the ECM remodelling that accompanies most liver diseases. Our previous study demonstrated that treatment of tumour cell lines with the HDAC inhibitor resulted in modulation of secreted MMP-2 and MMP-9, accompanied by global histone H3 and H4 hyperacetylation (Rodriguez-Salvador et al., 2005). As shown in our present experiments, HNHA inhibited MMP-2 and MMP-9 production in activated HSCs and BDL livers. Because of their role in fibrosis, the inhibition of TGF-β, α-SMA, MMP and COX-2 expression in the present study showed that HNHA could be an effective treatment for fibrotic diseases.

HNHA showed no toxic effect on hepatocytes at the concentrations used in this study (data not shown) and restored the body weight of BDL rats. In addition, a hepatoprotective activity of HNHA was expressed through the reduction in ALT, AST and TBil, which offers an additional therapeutic advantage of HNHA, compared with the other treatments, as it could both improve hepatocyte integrity and reduce stellate cell activation, as two independent but complementary activities. Furthermore, HNHA increased survival in BDL rats. As HNHA itself did not affect liver function and improved survival of BDL rats, this HDAC inhibitor may serve both for the chemoprevention and for the treatment of biliary fibrosis.

In conclusion, we have demonstrated that treatment with HNHA, a new HDAC inhibitor, markedly attenuated the process of liver fibrosis both *in vitro* and *in vivo*, showing a marked decrease of HSC proliferation and increased apoptosis, as well as suppression of profibrogenic factors such as PDGF and TGF-β1. Our data also provided the first *in vivo* evidence that HNHA increased survival in the BDL-induced



liver fibrosis model. Although further studies to determine drug doses from long-term studies are clearly required, our present results encourage the development of HNHA as a possible treatment for hepatic fibrosis.

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Conflict of interest

The authors disclose no conflict of interest.

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Supporting information

Additional Supporting Information may be found in the online version of this article at the publisher's web-site:

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Figure S1 Quantification of Western blots. The box and whisker plots show median, with first and third quartiles (as box) and range (as whiskers). Data are from n = 4 experiments. *P < 0.05, **P < 0.01, ***P < 0.005 versus PDGF, control,

Figure S2 Effects of HNHA on acetylation of histone H3 in BDL-liver. Representative staining (A) and densitometric analysis (B) of acetyl histone H3. The box and whisker plots show median, with first and third quartiles (as box) and range (as whiskers). Data are from n = 5 experiments. *P < 0.05, **P < 0.01, ***P < 0.005 versus BDL.

Figure S3 Effects of HNHA on levels of HDAC1 in BDL-liver. Representative staining (A) and densitometric analysis (B) of HDAC1 levels. The box and whisker plots show median, with first and third quartiles (as box) and range (as whiskers). Data are from n = 5 experiments. **P < 0.01, ***P < 0.005 versus BDL.