ORIGINAL ARTICLE

NKT-associated hedgehog and osteopontin drive fibrogenesis in non-alcoholic fatty liver disease

Wing-Kin Syn,1,2,3 Kolade M Agboola,1 Marzena Swiderska,1 Gregory A Michelotti,1 Evaggelia Liaskou,2 Herbert Pang,4 Guanhua Xie,1 George Philips,1 Isaac S Chan,1 Gamze F Karaca,1 Thiago de Almeida Pereira,1,5 Yuping Chen,1 Zhiyong Mi,6 Paul C Kuo,6 Steve S Choi,1,7 Cynthia D Guy,8 Manal F Abdelmalek,1 Anna Mae Diehl1

ABSTRACT

Objective Immune responses are important in dictating non-alcoholic steatohepatitis (NASH) outcome. We previously reported that upregulation of hedgehog (Hh) and osteopontin (OPN) occurs in NASH, that Hh-regulated accumulation of natural killer T (NKT) cells promotes hepatic stellate cell (HSC) activation, and that cirrhotic livers harbour large numbers of NKT cells.

Design The hypothesis that activated NKT cells drive fibrogenesis during NASH was evaluated by assessing if NKT depletion protects against NASH fibrosis; identifying the NKT-associated fibrogenic factors; and correlating plasma levels of the NKT cell-associated factor OPN with fibrosis severity in mice and humans.

Results When fed methionine-choline-deficient (MCD) diets for 8 weeks, wild type (WT) mice exhibited Hh pathway activation, enhanced OPN expression, and NASH-fibrosis. Jα18−/− and CD1d−/− mice which lack NKT cells had significantly attenuated Hh and OPN expression and dramatically less fibrosis. Liver mononuclear cells (LMNCs) from MCD diet fed WT mice contained activated NKT cells, generated Hh and OPN, and stimulated HSCs to become myofibroblasts; neutralising these factors abrogated the fibrogenic actions of WT LMNCs. LMNCs from NKT-cell-deficient mice were deficient in fibrogenic factors, failing to activate collagen gene expression in HSCs. Human NASH livers with advanced fibrosis contained more OPN and Hh protein than those with early fibrosis. Plasma levels of OPN mirrored hepatic OPN expression and correlated with fibrosis severity.

Conclusion Hepatic NKT cells drive production of OPN and Hh ligands that promote fibrogenesis during NASH. Associated increases in plasma levels of OPN may provide a biomarker of NASH fibrosis.

INTRODUCTION

Non-alcoholic fatty liver disease (NAFLD) is a leading cause of hepatic dysfunction. Compared with individuals with simple hepatic steatosis, individuals with steatosis plus hepatocytic injury, cell death and inflammatory changes (dubbed non-alcoholic steatohepatitis, NASH) are more likely to develop progressive liver fibrosis, cirrhosis and cancer.1,2 Although the mechanisms which drive disease progression remain obscure, recent studies suggest that cells of the innate immune system, such as Kupffer cells, natural killer T (NKT) cells and NK cells may participate.3–6

The liver contains a large number of NKT cells, which provide intravascular immune surveillance.7,8 NKT cells respond to glycolipid antigens presented by major histocompatibility class I-like molecule, CD1d, and when activated, secrete both Th1 helper (Th1) (pro-inflammatory/anti-fibrotic) and Th2 (anti-inflammatory/pro-fibrotic) cytokines.9–11 In the hepatitis B virus-transgenic mouse,12 NKT-derived interleukin 4 (IL4) and IL13 were reported...
to be responsible for the fibrogenic outcomes. Conversely, CD1d knockout (CD1d−/−) mice, which lack NKT cells, were protected from xenobiotic-induced hepatic inflammation, injury and fibrosis. These findings complement human studies showing NKT accumulation in primary biliary cirrhosis and chronic viral hepatitis C.

The role of NKT cells in NAFLD progression is only beginning to emerge. NKT cells are generally depleted in fatty livers, but appear to accumulate in livers with NASH-related fibrosis. Recently, we reported that hedgehog (Hh)-mediated accumulation of NKT cells contributes to development and progression of NASH in mice and humans. Mice with an overly active Hh pathway express higher levels of the NKT cell chemokine CXCL16, vascular cell adhesion molecule 1 (VCAM-1) and IL15, a factor that promotes NKT viability. These mice accumulate more hepatic NKT cells and develop worse NASH fibrosis. Consistent with this, we found that individuals with NASH cirrhosis harboured fourfold more NKT cells than those with healthy livers.

In addition to secreting the classical cytokines, IL4 and IL13, NKT cells can also secrete osteopontin (OPN), a cytokine and matricellular protein that exacerbates concavalin A induced hepatitis, and promotes NAFLD progression. Recently, we showed that NKT cells also secrete the fetal morphogen, sonic Hh (Shh). Shh induces the transition of quiescent hepatic stellate cells (HSCs) into activated, collagen-producing myofibroblasts and amplifies the repair-associated inflammatory response. Intriguingly, inhibition of the Hh pathway in vivo leads to the amelioration of fibrosis in mice with cholestatic liver injury.

In this study, we evaluated the hypothesis that fibrosis progression in NASH is NKT cell dependent. We fed mice that were genetically deficient in NKT cells (Jα18−/− and CD1d−/−) methionine-choline-deficient (MCD) diets to induce NASH, and assessed if NKT-cell-mediated fibrogenesis required OPN and Hh. Findings in mice and cultured HSCs were then corroborated by evaluating similar endpoints in patients with NASH.

METHODS

Animal experiments

Mice

Wild type (WT) (Jackson Laboratories, Bar Harbor, Maine, USA), CD1d-deficient (CD1d−/−) and Jα18-deficient (Jα18−/−) mice (RIKEN, Saitama, Japan) of similar background (B6) were fed MCD diet or control chow for 8 weeks. Animal care and procedures were approved by the Duke University Institutional Animal Care and Use Committee as set forth in the Guide for the Care and Use of Laboratory Animals published by the National Institutes of Health.

Analysis of liver architecture, morphometry and immunohistochemistry

Formalin-fixed paraffin embedded (FFPE) livers were analysed. Haematoxylin and eosin (H&E) and picrosirius red staining with quantification by morphometric analysis was performed as previously described. The primary antibodies used were OPN (R&D, Minneapolis, MN, USA, AF1433; 1:40) and α-smooth muscle actin (αSMA) (Abcam, Cambridge, MA, USA, Abcam 5694; 1:800). Polymer-HRP anti-rabbit (Dako, Carpinteria, CA, USA, DAKO0403) and anti-goat (Santa Cruz, California, USA, sc-2768; 1:250) were used as secondary antibodies. Antigens were demonstrated by dianaminobenzidine (DAB) (Dako). Omitting primary antibodies eliminated staining, demonstrating specificity. For OPN and αSMA quantification, 15 randomly selected, 40× fields (excluding the major bile duct in each portal tract from consideration) were analysed with the MetaView software.

ALT determination

Serum alanine aminotransferase (ALT) was measured using kits commercially available from Biotron Diagnostics (66-D; Hemet, California, USA) according to the manufacturer’s instructions.

RNA extraction and mRNA quantification

Total liver RNA extraction and mRNA quantification by real-time PCR (qRT-PCR) were performed as published. Details of primer sequences are given in online supplemental table 1.

Hydroxyproline assay

Hydroxyproline content in whole liver specimens was quantified colorimetrically as previously described.

Effect of primary liver NK cells on activation of primary HSC

Primary liver mononuclear cells (LMNCs) were isolated as previously described, and cultured in complete NKT media, with or without the NKT cell ligand α-galactosylceramide (αGC) (100 ng/ml; San Diego, California, USA; Cat. No 506027) for 24 h. αGC specifically activates invariant NKT (iNKT) cells. αGC-activated LMNC conditioned medium (LMNC-CM) was then added to primary murine HSCs in the presence or absence of the Hh neutralising antibody 5E1 (10 μg/ml; Iowa Hybridoma Bank, Iowa City, Iowa, USA), the Smoothened antagonist (inhibitor of Hh signalling) GDC (GDC-0449; Selleck, Houston, Texas, USA) or vehicle for 48 h and RNA was harvested for qRT-PCR. Experiments were performed in duplicate wells and repeated twice.

In separate experiments, LMNC-CM was added to primary murine HSCs with either sham aptamers (OPN-R3-2; 100 nmol/litre) or OPN aptamers (OPN-R3; 100 nmol/litre) (both synthesised by Dharmaco, Lafayette, Colorado, USA) for 48 h, and RNA harvested as above. This concentration of OPN aptamer has been shown to inhibit adhesion, migration and invasion in the MDA-MB-231 breast cancer cell line (which highly expresses OPN and is a standard tool for evaluating OPN actions).

ELISA assays for plasma OPN

Peripheral blood was collected at the time of sacrifice, and plasma obtained and stored at −80°C till analysis. Plasma OPN was measured using the commercially available OPN ELISA kit (R&D, Minneapolis, MN, USA, DY441) in accordance with the manufacturer’s instructions. All samples were run in duplicate, and expressed as pg/ml.

Human subjects

Immunohistochemistry

FFPE liver sections of de-identified subjects with biopsy-proven NASH-related early and advanced fibrosis (n=10/group) from the Departments of Pathology at Duke University were studied in accordance with National Institutes of Health and Institutional guidelines for human subject research.

The primary antibodies used were OPN (R&D, AF1453; 1:40); Indian hedgehog, Ihh (Abcam, Ab59634; 1:1000), Shh (Epitomics, Burlingame, CA, USA, 1843-1; 1:7500) Gli2 (Genway, San Diego, CA, USA, 18-752; 1:4500) CD57 (NeoMarkers, Fremont, CA, USA, MS156). Polymer-HRP anti-mouse (Dako) K4001 was used as secondary antibodies. Antigens were demonstrated by
diaminobenzidine (DAB) (Dako). Omitting primary antibodies eliminated staining, demonstrating specificity.

Plasma OPN measurements
Peripheral blood was taken from patients with early (n=25) or advanced NASH (n=25) at the time of liver biopsies, and plasma stored at −80°C until analysis. Plasma OPN was measured using the commercially available OPN ELISA kit (R&D; DY1433) and in accordance with the manufacturer’s instructions. All samples were run in duplicate, and expressed as pg/ml.

Statistical analysis
Results are expressed as mean ± SEM. Statistical significance was determined using the Student t test. Significance was accepted at the 5% level.

RESULTS
NKT-deficient mice develop less fibrosis than WT mice after MCD diets
Compared with WT mice that were fed normal chow (n=10), MCD-diet-treated mice (n=10) developed significant steatohepatitis and fibrosis after 8 weeks (figure 1A, online supplemental figure 1). The latter was demonstrated by increased Sirius red staining (figure 1B,C), and hepatic hydroxyproline quantification (figure 1D). Collagen deposition was accompanied by the accumulation of α-SMA-immunoreactive cells (figure 2A) and induction of pro-fibrogenic genes, including α-SMA, collagen 1α1, and transforming growth factor β (figure 2C,D, online supplemental figure 2A).

Jα18−/− mice, which lack iNKT cells, developed significantly less hepatic fibrosis (figures 1B−D, 2A−D) and injury (figure 2E) than WT mice after MCD diet. This finding in mice that lack iNKT cells is consistent with our previous report that CD1d−/− mice (which lack all NKT subsets) exhibited reduced fibrogenic responses after diet-induced NASH21 (figures 1C,D, 2B−D, online supplemental figure 2A).

We reported that Hh pathway activation induces the transition of quiescent HSC to collagen-producing myofibroblasts,26 and promotes fibrogenic repair in NASH.29 Consistent with this, Indian Hh (a Hh ligand), and Gli2 (a Hh-regulated transcription factor) were significantly induced during diet-induced NASH (online supplemental figure 2B,C). Conversely, NKT-deficient mice, which developed less fibrosis, exhibited attenuated Hh signalling (online supplemental figure 2B,C).
NKT-deficient mice express less OPN during MCD-diet-induced NASH than WT mice

We recently showed that OPN acts in paracrine and autocrine fashions to promote HSC activation and fibrogenesis. After MCD diet, mice genetically deficient in OPN exhibited reduced fibrosis, while mice that overexpress OPN developed worse fibrosis. Here, we show that livers from MCD-fed WT mice upregulate OPN protein expression by nearly fivefold (figure 5A, C). In contrast, livers of both NKT-cell-deficient strains that developed less fibrosis than WT mice after diet-induced NASH (figures 1B–D, 2A–D, online supplemental figure 2) had significantly reduced levels of OPN (figure 5A–C, online supplemental figure 3). These findings raise the possibility that NKT cell-associated factors modulate OPN expression and resultant hepatic fibrogenesis.

Previously, others reported that plasma OPN levels correlated with severity of liver fibrosis in subjects with chronic hepatitis B and C. In this study, plasma from WT mice with fibrosing NASH contained nearly twofold more OPN by ELISA than chow-fed controls (figure 5D). Consistent with reduced hepatic OPN expression, plasma OPN levels were twofold lower in NKT-cell-deficient mice after diet-induced NASH. The aggregate data, therefore, suggest that elevated plasma OPN levels may be a biomarker of advanced NASH fibrosis.

Primary mouse liver NKT-cell-associated OPN promotes myofibroblastic activation of primary HSC

In addition to HSCs and cholangiocytes, OPN is expressed by activated immune cells (figure 3). Therefore, we examined if activated liver NKT cells generate OPN that acts in a paracrine fashion to promote HSC fibrogenesis. LMNCs were isolated from healthy mice and incubated with αGC (which specifically activates iNKT cells) or vehicle overnight. LMNC-conditioned medium (CM) was then collected for OPN ELISA. αGC-activated LMNC-CM contained fourfold more OPN protein than vehicle-treated LMNC-CM, and similar levels of OPN as CM from cholangiocytes, which are known to be a source of OPN in liver (figure 4A). To ascertain if activated LMNCs in vivo also express higher amounts of OPN, we harvested primary LMNCs from chow- or MCD-fed WT mice (n=4/group), and OPN expression was analysed by qRT-PCR. Primary LMNCs from MCD-fed mice harboured 12-fold more OPN mRNA than LMNC from Chow-fed mice (figure 4B).

αGC-activated LMNC-CM was then added to primary cultures of mouse HSC with either OPN aptamers or sham aptamers for 48 h. The addition of OPN aptamers to neutralize OPN activity in LMNC-CM resulted in a significant repression of HSC fibrogenic genes (OPN, collagen) (figure 4C,D).

To ascertain if differences in the fibrogenic outcomes observed in figure 1 could be attributed to fewer NKT cells, and hence, reduced NKT-associated OPN, we isolated primary LMNCs from WT, Jα18–/– and CD1d–/– mice, and incubated them with αGC or vehicle overnight (as above), and LMNC-CM was collected for OPN ELISA or added to primary cultures of HSCs. We found that WT-LMNC mice secreted fivefold more OPN protein (figure 4E) and induced greater collagen expression in HSC than NKT-deficient LMNCs (figure 4F). These findings, in concert, support the hypothesis that activated NKT-cell-associated OPN promotes HSC activation in a paracrine fashion.

NKT-cell-associated Hh promotes myofibroblastic activation of primary HSCs

Previously, we reported that progressive NASH is associated with Hh pathway activation, that activated hepatic NKT cells are capable of secreting Hh ligands, and that Hh ligands could directly induce HSC activation and fibrogenesis. Because OPN expression in HSCs is regulated, at least in part, by Hh signalling, we examined whether NKT-derived Hh ligands could directly promote OPN expression in HSCs and mediate fibrogenic responses. Primary WT LMNCs were incubated with αGC or vehicle (as above). LMNC-CM was then added to primary cultures of mouse HSCs, with or without Hh neutralising antibody, 5E1, or GDC, a pharmacological inhibitor of the Hh pathway. Forty-eight hours later, HSC RNA was harvested for PCR analysis. We confirm by fluorescence activated cell sorting analysis that primary hepatic NKT cells express the Hh ligand (online supplemental figure 4A,B). CM from αGC-LMNCs that contained activated NKT cells upregulated HSC expression of collagen and OPN (figure 5A,B). The addition of 5E1 or GDC (inhibition of Hh signalling) led to a significant abrogation of these fibrogenic effects (figure 5A,B). The aggregate data suggest that in addition to generating OPN (figure 4A,E),
NASH progression in humans is accompanied by increased liver and plasma OPN

NAFLD progression in mice is associated with activation of the Hh pathway, induction of Hh-regulated fibrogenic factors, such as OPN,24 and the accumulation of intrahepatic NKT cells.19 21 We demonstrate that fibrogenesis in mice with diet-induced NASH depends on hepatic NKT cells, and show the mechanism involves NKT-cell-mediated production of Hh ligands and OPN (figures 1–5). NKT cells comprise a much greater proportion of the resident hepatic immune cells in mice than humans.4 35 Therefore, we examined livers from patients with NASH to assess the relationship between hepatic NKT cell accumulation, Hh pathway activation, OPN expression and fibrosis in humans.

Human livers with advanced NASH (stage 5–4 fibrosis) contained threefold more of the Hh-regulated cytokine, OPN, particularly in the fibrous septa, than those with early NASH (no fibrosis or stage 1–2 fibrosis) (figure 6A–B). Livers from individuals with advanced NASH also contained more cells that expressed the Hh ligands, Shh and Ihh (online supplemental figure 5A–C), and the Hh-regulated transcription factor, Gli2 (online supplemental figure 6A). Previously, we reported that the hepatic content of cells that co-express CD57 and CD3 or CD56 (ie, NKT cells) is increased significantly in patients with NASH-related cirrhosis.21 Here we show that NKT cells that accumulate during fibrosing NASH express Ihh and OPN protein (online supplemental figure 6B–E). Moreover, we found that increased hepatic OPN expression was mirrored by higher plasma OPN levels in NASH patients with advanced fibrosis than those with mild fibrosis (figure 6C). Overall plasma OPN levels in NASH were approximately twofold higher than levels detected in healthy volunteers (mean OPN level, normal: 3000 pg/ml).33 The findings in humans with NASH, therefore, are similar to those in mice with NASH (figure 5) and together suggest that OPN may be a useful biomarker of advanced NASH fibrosis.

**DISCUSSION**

This study shows that hepatic NKT cells are critical drivers of fibrogenesis in NASH. It also demonstrates that the mechanism involves NKT-cell-mediated stimulation of hepatic Hh pathway activation of hepatic NKT cells secreting Hh ligands to promote liver fibrogenesis.
activity and resultant increases in the liver content of OPN, a Hh-regulated, pro-fibrogenic factor. Importantly, plasma levels of OPN reflect the activity of this process in liver, and thus, may prove to be a useful non-invasive marker of advanced NASH.

Previously, we reported that Hh pathway activity correlates with severity of NAFLD-related liver injury, and showed that Hh signalling enhances hepatic accumulation of immune cells, including CD1d-restricted NKT cells. Here, we prove that fibrogenesis in mice with diet-induced NASH directly depends upon NKT cells. Compared with WT mice, two strains of NKT-cell-deficient mice (Jx18−/− and CD1d−/−) exhibited dramatically attenuated fibrosis after 8 weeks of MCD diets. NKT-cell-deficient mice also demonstrated blunted Hh pathway activity (less expression of Hh ligand and the Hh-regulated transcription factor, Gli2), and reduced expression of OPN, a Hh-regulated cytokine which acts in both paracrine and autocrine manners to promote HSC activation and fibrogenesis.

In the present study, LMNCs from WT mice with diet-induced NASH expressed 12-fold higher levels of OPN mRNA than chow-fed controls. CM from WT-mice-derived LMNCs induced NASH expressed 12-fold higher levels of OPN mRNA than CM from similarly treated LMNC populations from NKT-cell-deficient mice. Moreover, neutralising OPN activity with OPN-specific RNA aptamers significantly abrogated the HSC fibrogenic responses to WT LMNC-derived factors, reducing HSC expression of collagen mRNA to basal levels. The aggregate findings, therefore, complement and extend earlier evidence that liver NKT cells are capable of producing OPN by showing that important fibrogenic actions of NKT cells are directly mediated by OPN.

OPN expression is induced by activating Hh signalling in Hh-responsive cells. We previously reported that NKT cells produce and respond to Hh ligands. In the present study we show that adding the pan-Hh-neutralising antibody (5E1) to LMNC-CM completely blocked its ability to upregulate expression of collagen mRNA in HSCs. CM from LMNCs of NKT-cell-deficient mice is also completely unable to induce collagen gene expression in cultured HSCs. Thus, NKT cells appear to be a major source of the Hh ligands that drive fibrogenic responses in HSCs. The latter likely involves both NKT cell-derived OPN and HSC-derived OPN because treating HSCs that were grown in LMNC-CM with a drug that blocks intracellular Hh signalling (GDC) reduced HSC expression of OPN mRNA. Thus, fibrogenesis in NASH seems to result from the combined actions of several autocrine/paracrine signalling loops that re-enforce Hh pathway activation and OPN production in different types of hepatic non-parenchymal cells, particularly NKT cells and HSCs.

Our studies in human subjects support the pathophysiological relevance of excessive OPN production and Hh pathway activation during NASH-related fibrogenesis. Notably, we provide novel evidence that the upregulation of liver OPN production in advanced NASH is mirrored by increased plasma OPN levels. Recent reports suggest that high plasma OPN levels may also be predictive of cirrhosis in patients with chronic hepatitis B and C. In addition, elevated plasma OPN levels are associated with cancer development, recurrence and bad prognosis. Given these observations, it is conceivable that plasma OPN levels might be used to identify patients with NASH who are at high risk of bad outcomes. Further studies, in a larger cohort of individuals with NAFLD, will be necessary to confirm the clinical utility of this novel biomarker.
Supplemental Figure Legends

Figure 1. NKT-deficient mice develop less fibrosing-NASH after methionine-choline deficient (MCD) diet

Wild-type and Jα18/- mice were fed control chow (n=10 per strain) or MCD diet (n=10 per strain) for 8 weeks, and then sacrificed. Representative hematoxylin & eosin and Sirius red stainings from control chow-fed mice.

Figure 2. NKT-deficient mice exhibit reduced fibrogenic response and attenuated Hedgehog signaling after diet-induced NASH.

Mice were fed control chow and MCD diets as described in Figure legend 1. Whole liver tissue was harvested for RNA analysis by QRT-PCR. (A) TGFβ, (B) Indian Hedgehog (Inh), (C) Gli2 mRNA. Results are expressed as fold change relative to WT control chow-fed mice and graphed as mean ± SEM. *P<0.05 vs. WT control chow-fed mice.

Figure 3. Livers from NKT-deficient mice contain less OPN than WT mice after MCD diet.

Mice were fed control chow or MCD diets as described in Figure legend 1. (A) Representative OPN immunostaining after control chow or MCD diets in CD1d deficient mice (final magnification 200X). Whole liver RNA was analyzed by QRTPCR. (B) OPN mRNA. Results are expressed as fold change relative to WT
control chow-fed mice and graphed as mean ± SEM. *P<0.05 vs. WT control chow-fed mice

**Figure 4. Hepatic NKT cell express the Hedgehog ligand, Sonic Hedgehog**

Mouse primary liver mononuclear cells (LMNC) were isolated from whole livers of WT mice (n=3), and used for FACS analysis. (A) CD3/CD1d-tetramer double staining. (B) CD3/CD1d-tetramer double-positive cells express intracellular Sonic Hedgehog ligand.

**Figure 5. NASH progression in humans is associated with Hedgehog pathway activation**

(A-C) Coded liver sections from 10 patients with early and advanced NASH fibrosis were stained for Sirius Red (SR), Indian Hedgehog, and Sonic Hedgehog (both Hh ligands). (A) Representative SR staining, (B) Representative Ihh staining, and (C) Representative Shh staining

**Figure 6. NASH progression in humans is associated with Gli2 overexpression and accumulation of Hedgehog and OPN expressing CD57-positive cells**

(A-C) Coded liver sections from 10 patients with early and advanced NASH fibrosis were stained for Gli2. Liver sections with fibrosing NASH were also double immunostained for CD57 / Inh and CD57 / OPN. (D-E) Liver mononuclear cells from a non-diseased liver were stained for CD3, CD56, OPN, and analyzed
by FACS analysis. (A) Representative Gli2 staining, (B) CD57 (blue) - Inh 
(brown), (C) CD57 (brown) – OPN (blue). (D) CD3-CD56 double positive NKT 
cells, (E) Proportion of CD3-CD56 NKT cells which expressed intracellular OPN
Supplemental Materials and Methods

Animal experiments

**FACS analysis of primary mouse liver mononuclear cells**

Isolated LMNC from WT mice (n=3) were stained with FITC-conjugated CD3 (Santa Cruz; sc18843), APC-conjugated PBS57-loaded-CD1d-tetramers (provided by NIH, Atlanta, GA, USA), and PE-conjugated Shh (R&D, IC4641P), and analyzed using the BD™ LSR II (BD Biosciences) at the flow cytometry core facility at the Duke Human Vaccine Institute Flow Cytometry Core Facility, Duke University Medical Center.

**Mouse primary hepatic stellate cell isolation**

Hepatic stellate cells (HSC) were isolated from C57BL/6 mice as previously described (1). Briefly, after in situ perfusion of the liver with pronase (Roche, Indianapolis, IN) followed by collagenase (Roche), cell suspensions were layered on a discontinuous density gradient of Larcoll (Sigma-Aldrich, St. Louis, MO) and Histodenz (Sigma-Aldrich). The resulting upper layer consisted of >95% HSC. Viability of all cells was verified by phase contrast microscopy and the ability to exclude propidium iodide. The viability of all cells was >95%. Isolated HSC were seeded at a density of 5 x 10^3 cells/mm² in DMEM supplemented with 10% fetal bovine serum, streptomycin and penicillin. Day 3 HSC was used in all experiments.
Human tissues:

**Double Immunolabeling of CD57 (+) cells**

CD57 and Indian Hedgehog (Inh), or CD57 and Osteopontin (OPN) double immunostaining were demonstrated using DAB (DAKO) and Ferangi Blue™ (Biocare Med, Cat no FB8125, Concord, CA).

**Analysis of NKT cells subsets in human liver**

Non-diseased liver tissues removed during resection for colorectal hepatic metastases or from split-liver grafts, were used to isolate liver-derived lymphocytes as described (1). Primary hepatic lymphocytes were stained with Pacific Blue-conjugated anti-CD3 (Biolegend; Cat: 300-417), FITC-conjugated anti-CD56 (BD; cat no 34058; Oxford, UK), and PE-conjugated anti-OPN (R&D; cat no IC14331P). CD3-CD56 double positive cells are recognized as NKT cells¹.

Liver samples were obtained with informed consent of donors and in accordance with local ethical approval 04/Q2708/41 and REC 2003/242 from the South Birmingham Research Ethics Committee, UK.

References

Supplemental Fig 1

WT

Jα18-/-

H&E

SR
Supplemental Fig 3

A

CD1d-/- con

CD1d-/- MCD

OPN

B

Osteopontin mRNA Fold change

WT con  WT MCD  Ja18 con  Ja18 MCD  CD1d con  CD1d MCD
Supplemental Figure 6

A

Early NASH

Advanced NASH

Gli2

B

CD57 (blue) Inh (brown)

C

CD57 (brown) OPN (blue)

D

E

CD3-CD56

8.24%

64.9%

CD3

CD56

Count

OPN
## Supplemental Table 1: Primers used for Real time QRT-PCR

Mouse primers:

<table>
<thead>
<tr>
<th>GENE</th>
<th>Sense</th>
<th>Antisense</th>
<th>Sequence</th>
</tr>
</thead>
<tbody>
<tr>
<td>Collagen I α1</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Sense</td>
<td></td>
<td>GAGCGGAGAGTACTGGATCG</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>GCTTCTTTTCTTTGGGGTTTC</td>
</tr>
<tr>
<td>α sma</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Sense</td>
<td></td>
<td>AAACAGGAATACGACGAAG</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>CAGGAATGATTTGGAAAGGA</td>
</tr>
<tr>
<td>Tgf β</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Sense</td>
<td></td>
<td>TTGCCCTCTACAACCAACACAA</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>GGCTTGCGACCCCACGTA</td>
</tr>
<tr>
<td>OPN</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Sense</td>
<td></td>
<td>CTC CAT CGT CAT CAT CAT CG</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>TGC ACC CAG ATC CTA TAG CC</td>
</tr>
<tr>
<td>Gli2</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Sense</td>
<td></td>
<td>ACCATGCCTACCCAAACTCAG</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>CTGCTCCTGTGTGTCAGTCCAA</td>
</tr>
<tr>
<td>Inh</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Sense</td>
<td></td>
<td>ACAGATGGAATGCGTGTA</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>CCGAACCTTCATCTTGGGT</td>
</tr>
<tr>
<td>s9</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Sense</td>
<td></td>
<td>GACTCCGGAACAAACGTCAGG</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>CTTCATCTTGCCCTCGTCCA</td>
</tr>
</tbody>
</table>