Mesothelin/mucin 16 signaling in activated portal fibroblasts regulates cholestatic liver fibrosis

Yukinori Koyama, … , David A. Brenner, Tatiana Kisseleva


Cholestatic liver fibrosis is caused by obstruction of the biliary tract and is associated with early activation of portal fibroblasts (PFs) that express Thy-1, fibulin 2, and the recently identified marker mesothelin (MSLN). Here, we have demonstrated that activated PFs (aPFs) and myofibroblasts play a critical role in the pathogenesis of liver fibrosis induced by bile duct ligation (BDL). Conditional ablation of MSLN+ aPFs in BDL-injured mice attenuated liver fibrosis by approximately 50%. Similar results were observed in MSLN-deficient mice (Msln−/− mice) or mice deficient in the MSLN ligand mucin 16 (Muc16−/− mice). In vitro analysis revealed that MSLN regulates TGF-β1–inducible activation of WT PFs by disrupting the formation of an inhibitory Thy-1–TGFβRI complex. MSLN also facilitated the FGF-mediated proliferation of WT aPFs. Therapeutic administration of anti-MSLN–blocking Abs attenuated BDL-induced fibrosis in WT mice. Liver specimens from patients with cholestatic liver fibrosis had increased numbers of MSLN+aPFs/myofibroblasts, suggesting that MSLN may be a potential target for antifibrotic therapy.

Find the latest version:

http://jci.me/88845/pdf
Mesothelin/mucin 16 signaling in activated portal fibroblasts regulates cholestatic liver fibrosis

Yukinori Koyama,1,2 Ping Wang,1,2 Shuang Liang,1,2 Keiko Iwaisako,1,5 Xiao Liu,1,2 Jun Xu,1,2 Mingjun Zhang,1 Mengxi Sun,1,2 Min Cong,1,2,4 Daniel Karin,1,2 Kojiro Taura,3 Chris Benner,1 Sven Heinz,1 Tapan Bera,6 David A. Brenner,1 and Tatiana Kisseleva2

1Department of Medicine and 2Department of Surgery, UCSD, La Jolla, California, USA. 3Department of Surgery, Graduate School of Medicine, Kyoto University, Kyoto, Japan. 4Beijing Friendship Hospital, Capital Medical University, Beijing, China. 5Department of Target Therapy and Oncology, Graduate School of Medicine, Kyoto University, Kyoto, Japan. 6Laboratory of Molecular Biology, Center for Cancer Research, National Cancer Institute, NIH, Bethesda, Maryland, USA.

Cholestatic liver fibrosis is caused by obstruction of the biliary tract and is associated with early activation of portal fibroblasts (PFs) that express Thy-1, fibulin 2, and the recently identified marker mesothelin (MSLN). Here, we have demonstrated that activated PFs (aPFs) and myofibroblasts play a critical role in the pathogenesis of liver fibrosis induced by bile duct ligation (BDL). Conditional ablation of MSLN- aPFs in BDL-injured mice attenuated liver fibrosis by approximately 50%. Similar results were observed in MSLN-deficient mice (Msln–/– mice) or mice deficient in the MSLN ligand mucin 16 (Muc16–/– mice). In vitro analysis revealed that MSLN regulates TGF-β1-inducible activation of WT PFs by disrupting the formation of an inhibitory Thy-1-TGFβRII complex. MSLN also facilitated the FGF-mediated proliferation of WT aPFs. Therapeutic administration of anti-MSLN–blocking Abs attenuated BDL-induced fibrosis in WT mice. Liver specimens from patients with cholestatic liver fibrosis had increased numbers of MSLN+ aPFs/myofibroblasts, suggesting that MSLN may be a potential target for antifibrotic therapy.

Introduction
Fibrosis is an aberrant wound-healing process, in which chronic injury in any tissue activates myofibroblasts to produce a fibrous scar (1). Myofibroblasts are rare in normal tissues, and their origin depends on the tissue and type of injury. Many types of research suggest that liver-resident hepatic stellate cells (HSCs) and portal fibroblasts (PFs) are major sources of myofibroblasts in fibrotic liver (2). However, the composition of myofibroblasts varies significantly, depending on the etiology of hepatic fibrosis; hepatotoxic liver fibrosis (injury to hepatocytes, such as that caused by chronic viral hepatitis and nonalcoholic steatohepatitis) results from the activation of HSCs (3), while activated PFs (aPFs) are implicated in the pathogenesis of cholestatic liver fibrosis (obstruction to bile flow, such as that caused by primary and secondary biliary cirrhosis and primary sclerosing cholangitis) (4).

PFs are “periductular mesenchymal cells” that maintain the integrity of the portal tract (5, 6). In response to biliary obstruction, portal (myo)fibroblasts proliferate, upregulate expression of collagen type I, α-smooth muscle actin (α-SMA), TGF-β1/2, IL-6, and IL-13, and give rise to myofibroblasts (2, 7–14). Using a flow cytometry–based method of aPF purification (from bile duct ligation [BDL]-injured collagen-null[Il1]–GFP reporter mice, which express GFP under the control of the Colla1 promoter) (4), we demonstrated that myofibroblasts derived from aPFs (referred to herein as aPFs/myofibroblasts) are a major source of collagen type I–producing cells in early cholestatic liver injury, contributing more than 70% of the myofibroblasts by day 5 after BDL. The relative contribution of aPFs/myofibroblasts decreases with progressive injury, as HSCs become activated and contribute to the myofibroblast population (17 and 20 days after BDL). aPFs/myofibroblasts express high levels of collagen type I, serve as a significant source of TGF-β2, secrete IL-13 and IL-6 (4, 6), and facilitate the ductular reaction (cholestasis-induced proliferation of bile ducts) (7). aPFs are activated by taurocholic acid and IL-25, but unlike activated HSCs (aHSCs), are unresponsive to PDGF (4). aPFs can be distinguished from aHSCs by expression of Thy-1, fibulin 2, elastin, and the ecto-ATPase nucleoside triphosphate diphosphohydrolase 2 (NTPD2) and by their lack of aHSC markers (retinoid, desmin, glial fibrillary acidic protein [GFAP], p75NGFR) (15–18). Gene expression profiling of freshly isolated aPFs/myofibroblasts has identified several novel markers of aPFs, including mesothelin (MSLN) and mucin 16 (MUC16) (4).

MSLN (19), a membrane-anchored glycosylphosphatidyl inositol–linked (GPI-linked) 71-kDa membrane protein (MSLN precursor), is a mesothelial cell marker that is proteolitically cleaved to generate mature MSLN (40 kDa) and megakaryocyte-potentiating factor (MPF) (30 kDa) (20). Both molecules appear to be biologically active, although their exact function remains unknown. MSLN is expressed in several human malignancies, including mesotheliomas and ovarian cancer. Studies of patients with ovarian cancer have identified the cancer antigen CA125 as an MSLN ligand (20–23), which is widely used as a diagnostic marker (with the exception of pregnancy and liver cirrhosis, which are considered false-positives) (24). Since high expression of MSLN is linked to increased tumor proliferation and invasion, MSLN is a novel
cholestatic fibrosis in mice. Mechanistic studies identified what we believe to be a novel TGF-β1 pathway, in which interaction of MSLN with a MUC16–Thy-1–TGF-βRI complex regulates TGF-β1–mediated activation of aPFs/myofibroblasts. Analysis of patients’ liver biopsies revealed that MSLN+Thy-1+ aPFs contribute to cholestatic fibrosis in patients with secondary biliary fibrosis and biliary atresia (BA), suggesting that MSLN may serve as an attractive target for antifibrotic therapy.

Results
Msln-KO mice are resistant to BDL-induced cholestatic, but not hepatotoxic, liver fibrosis. To test the role of MSLN in the activation of aPFs/myofibroblasts, WT and Msln−/− mice were subjected to BDL and CCl4 injury (Figure 1, A–C). We observed no difference in liver fibrosis in CCl4-injured WT or Msln−/− mice (at 1.5 months, Figure 1B). In concordance with this finding, the aPF markers MSLN and fibulin 2 were not upregulated in the livers of CCl4-injured mice.
control diet–fed WT mice) (Supplemental Figure 1A; supplemental material available online with this article; https://doi.org/10.1172/JCI88845DS1). We observed significant downregulation of Picrosirius red staining (↓1.7-fold decrease) and expression of the fibrogenic genes Col1a1 (↓2-fold), α-SMA (↓2-fold), and Timp1 (↓1.7-fold) in the livers of Msln–/– mice (Supplemental Figure 1A), which was associated with decreased expression of the aPF-specific gene fibulin 2. Our data indicate that MSLN critically contributes to the pathogenesis of cholestatic liver fibrosis of different etiologies.

The number of BDL-activated myofibroblasts was reduced in Msln–/– mice. To further assess the role of MSLN in PF/myofibroblast activation, WT and Msln–/– mice were crossed with reporter collagen α1(I)–GFP mice (in which all activated myofibroblasts upregulate collagen α1(I)–GFP expression) and subjected to sham or BDL treatment (5 and 17 days of BDL), Figure 2, A and B, and Supplemental Figure 1, B–D). Livers were costained with Thy-1 and desmin to visualize aPF- and aHSC-derived myofibroblasts, respectively (Figure 2A), and the area of Thy-1+GFP+ and Thy-1–GFP+ costaining was calculated (Figure 2B). Although expression of collagen α1(I)–GFP was increased in both WT and Msln−/− mice following BDL (5 days of BDL vs. sham-operated mice), the number of activated GFP+ myofibroblasts was reduced by nearly 3-fold in BDL-injured Msln−/− mice (1.6% ± 0.2% vs. 4.8% ± 0.2% in BDL-treated WT mice). Meanwhile, the composition of GFP+ myofibroblasts (considered 100%) was not significantly changed between WT and Msln−/− mice after 5 days of BDL. Thus, WT GFP+ myofibroblasts (Figure 1C), confirming that aPFs/myofibroblasts do not activate in response to hepatotoxic liver injury.

In contrast, the development of BDL-induced cholestatic liver fibrosis was strongly attenuated in Msln−/− mice compared with that seen in WT mice (Figure 1, A–C). We observed significant downregulation of Picrosirius red staining (2-fold) and mRNA expression of the fibrogenic genes Colla1 (2-fold), α-SMA (1.7-fold), and Timp1 (1.7-fold) in the livers of Msln−/− mice 5 days after BDL, which correlated with low levels of serum alanine transaminase (ALT) and aspartate transaminase (ALP) (vs. levels in WT mice). We also observed inhibition of liver fibrosis in Msln−/− mice on day 17 after BDL, as shown by decreased Picrosirius red staining (↓1.8-fold) and downregulation of Colla1 (↓1.8-fold), α-SMA (↓1.4-fold), and Timp1 (↓1.7-fold) mRNA expression, as well as reduced levels of serum ALP (↓2-fold). Our data indicate that MSLN critically contributes to the pathogenesis of cholestatic liver fibrosis of different etiologies.

Msln-KO mice are resistant to diethoxycarbonyl dihydrocollidine–induced cholestatic fibrosis. Similar results were obtained when WT and Msln−/− mice were subjected to another model of cholestatic liver injury induced by 3 weeks of diethoxycarbonyl dihydrocollidine (DDC) feeding. As expected, hepatic expression of Msln mRNA was strongly increased in DDC-fed WT mice (7-fold increase vs. control diet–fed WT mice) (Supplemental Figure 1A; supplemental material available online with this article; https://doi.org/10.1172/JCI88845DS1). We observed significant downregulation of Picrosirius red staining (↓1.7-fold decrease) and expression of the fibrogenic genes Colla1 (↓2-fold), α-SMA (↓2-fold), and Timp1 (↓1.7-fold) in the livers of Msln−/− mice (Supplemental Figure 1A), which was associated with decreased expression of the aPF-specific gene fibulin 2. Our data indicate that MSLN critically contributes to the pathogenesis of cholestatic liver fibrosis of different etiologies.

The number of BDL-activated myofibroblasts was reduced in Msln−/− mice. To further assess the role of MSLN in PF/myofibroblast activation, WT and Msln−/− mice were crossed with reporter collagen α1(I)–GFP mice (in which all activated myofibroblasts upregulate collagen α1(I)–GFP expression) and subjected to sham or BDL treatment (5 and 17 days of BDL), Figure 2, A and B, and Supplemental Figure 1, B–D). Livers were costained with Thy-1 and desmin to visualize aPF- and aHSC-derived myofibroblasts, respectively (Figure 2A), and the area of Thy-1+GFP+ and Thy-1+costaining was calculated (Figure 2B).

Although expression of collagen α1(I)–GFP was increased in both WT and Msln−/− mice following BDL (5 days of BDL vs. sham-operated mice), the number of activated GFP+ myofibroblasts was reduced by nearly 3-fold in BDL-injured Msln−/− mice (1.6% ± 0.2% vs. 4.8% ± 0.2% in BDL-treated WT mice). Meanwhile, the composition of GFP+ myofibroblasts (considered 100%) was not significantly changed between WT and Msln−/− mice after 5 days of BDL. Thus, WT GFP+ myofibroblasts
were composed of 71% ± 3% aPFs and 28% ± 2% aHSCs, while MSLN-deficient GFP+ myofibroblasts were composed of 61% ± 2% aPFs and 38% ± 4% aHSCs.

Expression of MSLN has been reported in both aPFs/myofibroblasts and hepatic mesothelial cells, identified by expression of the GPM6A marker (34). Costaining with anti–glycoprotein M6A (anti-GPM6A) Ab revealed that mesothelial cells contribute to a small number of collagen α1(I)–GFP–positive (Col+) myofibroblasts residing in the liver capsule of the WT mice subjected to 5 days of BDL injury (Supplemental Figure 2A), while the majority of Thy-1+Col+ myofibroblasts were located in the portal areas and lacked GPM6A expression (Supplemental Figure 2B). Since no other cell types in the liver (including HSCs, cholangiocytes, endothelial cells, hepatocytes, and Kupffer cells) (Supplemental Figure 2, C and D) were reported to coexpress collagen α1(I) and MSLN (and Thy-1), the loss of activated myofibroblasts in BDL-injured Msln−/− mice was attributed to the MSLN deficiency in aPFs/myofibroblasts.

Figure 3. Ablation of Msln attenuates development of cholestatic liver fibrosis in mice. (A) Tamoxifen administration (12 × 5 mg/mouse by oral gavage) irreversibly labeled all MSLN+ aPFs by nuclear LacZ (nLacZ) expression in WT MslnnLacZ mice (n = 14) and caused ablation of aPFs in MslnDTA littersates (C57BL/6, 8-week-old male mice, n = 17 from 3 independent experiments). Livers were stained for LacZ (micrographs show gross liver tissue) and (B) analyzed for LacZ expression (original magnification, ×40). (C) Liver sections were immunostained with DAPI for Thy-1 and α-SMA expression. The number of aPFs in sham-operated MslnnLacZ and MslnDTA mice and the efficiency of aPF/myofibroblast ablation in BDL-operated MslnDTA mice were calculated in comparison with the number of Thy-1+α-SMA+DAPI+ cells in livers of BDL-MslnnLacZ mice (considered as 100%). Representative micrographs are shown (original magnification, ×20). (D) Livers were stained with Picrosirius red (original magnification, ×4). (E) Quantification of Picrosirius red and Thy-1+α-SMA+ areas is shown as a percentage. Expression of fibrogenic, aPF-specific, and inflammatory gene mRNA was analyzed by qPCR and is shown as a fold induction. *P < 0.05 and **P < 0.01, by 2-tailed Student’s t test. (See also Supplemental Figure 3.)
Ablation of MSLN+ aPFs attenuates BDL-induced liver fibrosis. MSLN was minimally expressed in adult mice but was strongly upregulated in response to BDL (Figure 1C). To gain insight into the role of MSLN+ aPFs/myofibroblasts in the pathogenesis of cholestatic liver fibrosis, we generated mice devoid of MSLN+ cells. We assumed that, if MSLN+ aPFs significantly contribute to cholestatic liver fibrosis, then ablation of MSLN+ aPFs would attenuate the development of cholestatic liver fibrosis. To examine this, MslnDTA mice (MslnnLacZ mice, in which the cassette harboring the reporter nuclear [n]LacZ and tamoxifen-inducible ER-Cre was knocked into the Msln gene) were crossed with Rosa26 flox-Stop-flox-DTA mice (Figure 3A). The resulting MSLN-ablated Msln DTA mice and WT MslnnLacZ littermates were treated with tamoxifen and then subjected to 5 days of BDL injury. Upon administration of tamoxifen to MslnnLacZ mice, MSLN+ aPFs/myofibroblasts had increased nLacZ+ expression and were scattered throughout the liver parenchyma of MslnnLacZ mice (Figure 3B and Supplemental Figure 3A). In turn, the disappearance of hepatic nLacZ+ expression in MslnDTA mice was indicative of successful ablation of MSLN+ aPFs/myofibroblasts and correlated with marked downregulation of the aPF-specific markers MSLN and fibulin 2 in the livers of Msln DTA mice (Figure 3E).

Delayed administration of anti-MSLN Abs inhibits cholestatic fibrosis in mice. To test whether anti-MSLN-blocking Abs can avert the development of cholestatic fibrosis, BDL-operated collagen α(I)–GFP mice were treated with 2 consecutive doses of anti-MSLN–blocking Abs or isotype IgG (24 and 48 hours after BDL) (Figure 4A) and sacrificed 5 days after BDL surgery. Delayed administration of Ab no. 1 (Ab1) (200 ng/g or 400 ng/g BW; Abbiotec; 250519) to BDL-treated collagen α(I)–GFP mice inhibited fibrosis and activation of GFP+ myofibroblasts in a dose-dependent manner. Similar results were achieved using delayed administration of Ab no. 1.25.
The Journal of Clinical Investigation

RESEARCH ARTICLE

1259

jci.org   Volume 127   Number 4   April 2017

2 (Ab2) (400 ng/g BW; MBL; D233-3), suggesting that targeting MSLN may be beneficial for patients with cholestatic liver fibrosis. In support of this, several classes of anti-MSLN Abs (26, 35) have been generated and are currently being used in clinical trials.

Expression of human MSLN is upregulated in patients with cholestatic liver fibrosis. To gain insight into the role of αPFs in human liver diseases, the myofibroblast composition was assessed in resected livers from patients with cholestatic fibrosis (BA, n = 6; secondary biliary cirrhosis, n = 12; and hepatotoxic HCV, n = 5) using quantitative PCR (qPCR) using immunohistochemical analyses (Figure 4B and Supplemental Figure 4). Expression of MSLN mRNA was upregulated by nearly 3-fold in patients with cholestatic fibrosis (but not in those with nonfibrotic or HCV-fibrotic livers). Immunostaining for MSLN and Thy-1+ expression was detected only in fibrotic lesions of patients with cholestatic fibrosis and overlapped with α-SMA+ (but not desmin+) staining, suggesting that MSLN can serve as a marker of human αPFs (Supplemental Figure 4) and mouse αPFs (4).

Primary Msln−/− αPFs/myofibroblasts have a defect in activation and proliferation. Our findings suggest that genetic and pharmacological ablation of MSLN+ αPFs/myofibroblasts or genetic deletion of Msln in mice attenuates the development of cholestatic liver fibrosis (Figures 1 and 4). To investigate the role of MSLN in αPF/myofibroblast activation, primary αPFs were sort purified from BDL-injured (for 5 days) livers of collagen α1(I)–GFP WT and collagen α1(I)–GFP Msln−/− mice and phenotyped (Supplemental Figure 5). In comparison with WT αPFs, we found that expression of fibrogenic genes was

Figure 5. Primary Msln−/− αPFs have a defect in activation and proliferation. (A) Primary GFP+ vitamin A+ αPFs were sort purified from the livers of BDL-injured (5 days) WT and Msln−/− mice (n = 3/isolation, 6 independent experiments), and mRNA expression of lineage-specific genes (Supplemental Figure 5) and activation markers was analyzed by qPCR. (B) Primary WT and Msln−/− αPFs (105/well) were assessed in a 12-hour scratch assay. Shown are representative bright-field (BF) and fluorescence (Col-GFP) micrographs (original magnification, ×10). The scratch area was calculated as a percentage (average of >3 independent experiments). (C) Ki67 immunostaining of primary WT and Msln−/− αPFs within the scratch area. Representative images (original magnification, ×20) are shown. Graph shows the percentage of Ki67+ αPFs versus total αPFs (100%) for an average of 3 experiments. *P < 0.05, by 2-tailed Student’s t test or ANOVA for comparisons between 2 groups or more than 2 groups, respectively.
IP with anti-MSLN Ab (Figure 6D and Supplemental Figure 6B) revealed that MSLN binds to Thy-1 and MUC16 (but not to TGFβRI or TGFβRII) in WT aPFs. TGFβ-stimulation further increased the formation of an MSLN–Thy-1 complex, while MSLN–MUC16 binding declined. As expected, we found that these complexes were absent in Mshn–/– aPFs.

Finally, IP with anti-MUC16 Ab (Figure 6E and Supplemental Figure 6C) confirmed that MUC16 interacts with MSLN and TGFβRII (but not with Thy-1 or TGFβRI). We found that SMAD7 was also co-coprecipitated as a part of the MUC16-TGFβRII complex and that its interaction with MUC16 was decreased in response to TGF-1. We found that the formation of a MUC16−SMAD7 complex was increased in unstimulated Mshn–/– aPFs, while MUC16−TGFβRII binding was decreased, and detected no MUC16−MUC16 complex. Taken together, we identified what we believe to be a novel TGFβ−M UC16 signaling pathway, in which the activity of the TGFβRII-TGFβRII complex is regulated via interaction with a Thy-1−MSLN-MUC16-SMAD7 complex, and in which MSLN and MUC16 facilitate TGFβ responses in aPFs, while Thy-1 and SMAD7 inhibit TGFβ responses. Our proposed model of MSLN-dependent regulation of TGFβ signaling in aPFs is depicted in Figure 6E.

BDL-induced liver fibrosis is reduced in Muc16+/– mice, but exacerbated in Thy-1−/– mice. To further corroborate the findings on the in vivo role of MSLN, MUC16, and Thy-1 in cholestatic liver injury, we compared the development of BDL liver fibrosis in mice in which individual components of this signaling complex were knocked out. The role of MUC16−MSLN interaction in cholestatic fibrosis was evaluated in vivo in Muc16+/–, Muc16+/– Mshn−/–, and Mshn−/– mice. Liver fibrosis induced by 5 days of BDL was equally attenuated in Muc16+/–, Muc16−/– Mshn−/–, and Mshn−/– mice (Figure 7A and Supplemental Figure 7), as shown by the reduced Picosirius red–stained area and the downregulation of Colla1, α- SMA, and Timp1 mRNA compared with WT mice (Figure 7B), suggesting that MSLN and MUC16 regulate the same signaling pathway. Furthermore, inhibition of BDL-induced fibrosis in Muc16+/– Mshn−/– mice (by 50%) was comparable to that observed in aPF-ablated MshnΔTα mice (Figure 3), indicating that the MSLN/MUC16 pathway might be critical for aPF/myofibroblast activation. Therefore, we studied TGFβ1−mediated activation of PFS in vivo in MshnΔTα mice (generated by crossing MshnLacZ–ER–Cre and Smad4fl/fl mice) and in WT MshnLacZ–ER–Cre mice (Figure 8A). Deletion of Smad4Δ in MshnΔTα aPFs attenuated BDL-induced (5 days) liver fibrosis by approximately 25%, suggesting that MSLN-MUC16 functions in aPFs are not limited to regulation of the TGFβ1 signaling pathway. We also examined the contribution of Thy-1 to PFS activation (Figure 8B) and found that deletion of Thy-1 in aPFs exacerbated cholestatic fibrosis in Thy-1−/– mice by nearly 25% compared with that observed in WT mice, confirming that Thy-1 inhibits fibrosis. Overall, deletion of MSLN/MUC16 signaling inhibits the development of BDL-induced fibrosis by approximately 50%, but only about 25% of this phenotype can be attributed to impaired TGFβ1 signaling (as demonstrated by deletion of Smad4 specifically in MshnΔTα aPFs) and linked to the MSLN−MUC16−Thy-1 complex (since an increase of approximately 25% in liver fibrosis was detected in BDL−injured Thy-1−/– mice). We concluded that MSLN/MUC16 pathway functions are not limited to regulation
of TGF-β1 signaling in aPFs and that the MSLN-MUC16 complex might further facilitate cholestatic fibrosis (by an additional 25%, approximately) through the mediation of other important signaling pathways in aPFs. In the next set of experiments, we investigated the role of the MSLN/MUC16 signaling pathway in aPF proliferation.

**Msln**<sup>−/−</sup> aPFs exhibit defective FGF-induced proliferation. FGF was identified as an important factor that drives the proliferation of aPFs. Primary cultured WT aPFs had a growth advantage over **Msln**<sup>−/−</sup> aPFs (Figure 5, B and C). In an attempt to further characterize MSLN-MUC16 functions, we examined the proliferation and responses to FGF using immortalized WT and **Msln**<sup>−/−</sup> aPFs and found that cyclin D mRNA expression was suppressed by 4.5-fold in FGF-stimulated **Msln**<sup>−/−</sup> aPFs and correlated with the downregulation of **Mmp9** mRNA (Figure 9A). When we examined the components of the FGF signaling pathway, we found that expression of AKT was strongly reduced (by 5-fold) in FGF-stimulated **Msln**<sup>−/−</sup> aPFs (Figure 9C). A novel mechanism for terminating RTK signaling and reducing receptor abundance has been described and showed that, in AKT-impaired cells, RTK expression and trafficking are defective, resulting in prolonged signaling and increased activation of ERK1/2 (45). In accordance with previously published reports, we detected an 11-fold increase in ERK1/2 phosphorylation in FGF-stimulated **Msln**<sup>−/−</sup> aPFs (Figure 9B) that was associated with reduced degradation of FGF receptor 1 (FGFR1) following FGF signaling (Figure 9C). Next, we performed IPs with anti-FGFR1 Ab and anti-MSLN Ab to determine the relationship between MSLN and FGFR1, but did not identify MSLN-MUC16 as binding partners of the FGFR1 complex (Figure 9D), suggesting that MSLN directly regulates AKT rather than the FGF signaling pathway.

**Activation of JAK2-STAT3 and ERK1/2 regulates the proliferation of aPFs.** We further investigated the molecular mechanism of FGF-mediated aPF proliferation (Figure 10 and Supplemental Figure 8). As expected, we found that pretreatment with an inhibitor of FGFR1 completely blocked (100%, vs. DMSO) cyclin D mRNA expression in FGF-induced WT aPFs (Supplemental Figure 8). Cyclin D expression (Supplemental Figure 8) and STAT3 phosphorylation (Figure 10A) were also suppressed by inhibitors of AKT and JAK2, but most effectively (>80%) by AZD1480, an inhibitor of JAK2 and ERK (Figure 10B). In agreement with these results, the proliferation and migration of WT and **Msln**<sup>−/−</sup> aPFs to the scratch area were completely blocked by AZD1480 (Figure 10C) and associated with complete inhibition of ERK1/2 phosphorylation (Figure 10B). Our proposed model of MSLN-MUC16–dependent regulation of FGF signaling in aPFs is depicted in Figure 10D.

**Reconstitution of MSLN rescues the phenotype in **Msln**<sup>−/−</sup> cells.** To confirm these findings, we transfected **Msln**<sup>−/−</sup> aPFs with a pCMV-MSLN construct (Figure 11A and Supplemental Figure 9A). We observed that reexpression of MSLN restored basal and FGF-dependent proliferation in **Msln**<sup>KO+Msln</sup> aPFs, as shown by the
The Journal of Clinical Investigation

RESEARCH ARTICLE

1263

Effect on cyclin D (Figure 12, A and B) or ERK1/2 (Supplemental Figure 10), suggesting that the regulatory function of Thy-1 is restricted to TGF-β1 signaling.

Overexpression of AKT restores Msln–/– cell proliferation. Furthermore, to test whether restoration of AKT phosphorylation can restate the proliferation of Msln–/– aPFs, we infected Msln–/– aPFs with constitutively activated (CA) AKT (AAV-CA-AKT) or with a control adenoviral construct expressing GFP (AAV-GFP). We found that expression of cyclin D and cyclin E (Figure 13A) and FGF-induced degradation of FGFR1 were restored in Msln KO-AAV-CA-AKT aPFs to the levels observed in WT aPFs (Figure 13B). The changes in JAK2 and its phosphorylation observed in MSLN-deficient aPFs were restored to normal levels in Msln KO-AAV-CA-AKT aPFs (Figure 13B and upregulation of cyclin D mRNA (Figure 11B) and the increased numbers of cells populating the scratch area (Figure 11C and Supplemental Figure 9B), both of which could be also blocked by AZD1480 (Supplemental Figure 9C). We found that reexpression of MSLN also restored Colla1 and Muc16 in Msln KO/Msln KO aPFs, but had no effect on the expression levels of Thy-1, which remained elevated, as observed in Msln–/– aPFs (Figure 11A).

Inhibition of Thy-1 restores TGF-β1–induced responses in Msln–/– cells. To test whether suppression of Thy-1 can restore the responsiveness of Msln–/– aPFs to TGF-β1, we transfected Msln–/– aPFs with Thy-1 siRNA or control siRNA (Figure 12A). Inhibition of Thy-1 restored phosphorylation of SMAD2 (Figure 9D) and the expression of Colla1 in TGF-β1–stimulated Msln KO+siThy-1 aPFs, but had no effect on cyclin D (Figure 12, A and B) or ERK1/2 (Supplemental Figure 10), suggesting that the regulatory function of Thy-1 is restricted to TGF-β1 signaling.

Overexpression of AKT restores Msln–/– cell proliferation. Furthermore, to test whether restoration of AKT phosphorylation can restate the proliferation of Msln–/– aPFs, we infected Msln–/– aPFs with constitutively activated (CA) AKT (AAV-CA-AKT) or with a control adenoviral construct expressing GFP (AAV-GFP). We found that expression of cyclin D and cyclin E (Figure 13A) and FGF-induced degradation of FGFR1 were restored in Msln KO-AAV-CA-AKT aPFs to the levels observed in WT aPFs (Figure 13B). The changes in JAK2 and its phosphorylation observed in MSLN-deficient aPFs were restored to normal levels in Msln KO-AAV-CA-AKT aPFs (Figure 13B and

Figure 8. Liver fibrosis is exacerbated in BDL-injured Thy-1–/– mice. (A) Liver fibrosis was compared in 8-week-old male BDL-injured Msln KO+Smad4 mice and WT Msln+/+ littermates (n = 9/group). (B) Liver fibrosis was compared in 8-week-old male BDL-operated (for 5 days) WT and Thy-1–/– littermates (C57BL/6, n = 8/group). (A and B) Micrographs show Picrosirius red staining (original magnification, ×4). Data indicate the percentage of positively stained area. Expression of fibrogenic, inflammatory, and aPF-specific genes was quantified by qPCR, and the data indicate the fold induction versus sham-operated WT mice. **P < 0.01 and *P < 0.05, by 2-tailed Student’s t test.
lioma (26), and both MSLN and CA125 became targets for anti-cancer therapy. Here, we demonstrate that expression of MSLN is not restricted to tumorigenesis, but is strongly induced in aPFs in response to cholestatic liver injury (Figure 1 and Supplemental Figure 1A). In support of this, Msln-KO mice developed a defect in the activation of cancer-associated myofibroblasts that resulted in reduced tumorigenesis (25–29).

Despite extensive studies of human cancer cells, the functional properties of the MSLN/MUC16 signaling pathway remain poorly understood. MUC16 was identified as an MSLN ligand, and MSLN/MUC16 signaling was linked to AKT and ERK1/2 activation and was implicated in the metastatic growth and dissemination of cancer cells (46–49). Like other GPI-anchored proteins, MSLN requires MUC16 and/or utilizes other receptors for intracellular signaling. Our studies of immortalized aPFs show that Msln–/– aPFs exhibit impaired proliferation and responses to FGF and that these defects are associated with defective AKT phosphorylation (Figure 9C). However, co-IP experiments did not detect MSLN-MUC16 as a part of the FGF /FGFR1 signaling complex. A novel mechanism of AKT activation has been identified that implicates AKT in the regulation of RTK trafficking and degradation (45). Accordingly, cells deficient in AKT activation fail to properly express EGFR and PDGFR, but as a compensatory mechanism, strongly upregulate ERK1/2 phosphorylation. We observed a similar phenotype in Msln–/– aPFs (Figure 9B). The loss of AKT activation was associated with decreased phosphorylation of AKT (Supplemental Figure 11A), while activation of ERK1/2 remained unchanged (Supplemental Figure 11B). Furthermore, we found that TGF-β1-mediated phosphorylation of SMAD2 was not rescued in MslnKO-AAV-CA-AKT aPFs (Supplemental Figure 11C), suggesting that activation of AKT does not affect TGF-β1 signaling pathway in aPFs.

Discussion

We demonstrate here that aPFs mediate hepatic fibrosis at the onset of cholestatic injury and that ablation of MSLN+ aPFs in mice decreases BDL-induced (5 days) liver fibrosis by approximately 50%. This study links MSLN/MUC16 signaling to the pathogenesis of cholestatic fibrosis and shows that deletion of MSLN and/or MUC16 in mice reduces cholestatic liver fibrosis by nearly 50%. We determined that MSLN facilitates both TGF-β1–induced activation of aPFs and FGF-induced proliferation of aPFs. We detected a similar upregulation of MSLN+ aPFs in patients with biliary fibrosis of different etiologies. Our data demonstrate that therapeutic administration of MSLN-blocking Abs attenuates BDL-induced liver fibrosis in mice, suggesting that MSLN could serve as a target for antifibrotic therapy.

MSLN (19) is highly expressed during embryonic development (17, 32) but is downregulated in adulthood (20). However, Msln–/– and Muc16–/– mice do not have developmental abnormalities (31, 33). In turn, human MSLN and CA125 (the human counterpart of MUC16) is highly upregulated in several malignancies such as ovarian cancer, hepatocellular carcinoma, and mesothelioma (26), and both MSLN and CA125 became targets for anti-cancer therapy. Here, we demonstrate that expression of MSLN is not restricted to tumorigenesis, but is strongly induced in aPFs in response to cholestatic liver injury (Figure 1 and Supplemental Figure 1A). In support of this, Msln-KO mice developed a defect in the activation of cancer-associated myofibroblasts that resulted in reduced tumorigenesis (25–29).

Despite extensive studies of human cancer cells, the functional properties of the MSLN/MUC16 signaling pathway remain poorly understood. MUC16 was identified as an MSLN ligand, and MSLN/MUC16 signaling was linked to AKT and ERK1/2 activation and was implicated in the metastatic growth and dissemination of cancer cells (46–49). Like other GPI-anchored proteins, MSLN requires MUC16 and/or utilizes other receptors for intracellular signaling. Our studies of immortalized aPFs show that Msln–/– aPFs exhibit impaired proliferation and responses to FGF and that these defects are associated with defective AKT phosphorylation (Figure 9C). However, co-IP experiments did not detect MSLN-MUC16 as a part of the FGF/FGFR1 signaling complex. A novel mechanism of AKT activation has been identified that implicates AKT in the regulation of RTK trafficking and degradation (45). Accordingly, cells deficient in AKT activation fail to properly express EGFR and PDGFR, but as a compensatory mechanism, strongly upregulate ERK1/2 phosphorylation. We observed a similar phenotype in Msln–/– aPFs (Figure 9B). The loss of AKT activation was asso-
associated with an inability of Msln−/− aPFs to mount FGF-mediated responses. Because of reduced phosphorylation of JAK2 (despite upregulation of JAK2 expression), we observed that FGF-dependent activation of STAT3 was also impaired in Msln−/− aPFs. In turn, we observed that basal and FGF-induced ERK1/2 activation was highly increased in Msln−/− aPFs and was driving the residual proliferative capability of Msln−/− aPFs (via phosphorylation of STAT3). Although the exact mechanism by which MSLN-MUC16 activates AKT remains unknown, our data indicate that the MSLN/MUC16/AKT pathway is a key regulator of cholestasis-induced proliferation of aPFs/myofibroblasts, but does not affect the fibrogenic properties of aPFs.
Msln−/− aPFs exhibit defective TGF-β1 signaling manifested by reduced expression of profibrogenic genes and TGF-β1 target genes (Figure 6A). Here, we demonstrate the involvement of MSLN in the TGF-β1 signaling pathway and propose a mechanism in which the MSLN-MUC16 complex interacts with the TGFβRI receptor and facilitates TGF-β1 signaling in WT aPFs via disruption of TGFβRI binding to its inhibitory receptor Thy-1 (Figure 6, C and D). This enables TGFβRI-TGFβRII interaction, phosphorylation and nuclear translocation of SMAD2/3/4, and transcription of target genes. In turn, deletion of Msln in Msln−/− aPFs resulted in overexpression of Thy-1 (>150-fold vs. WT levels), which binds to TGFβRI and prevents TGF-β1 signaling. Consistently, inhibition of Thy-1 partially restored TGF-β1 signaling in Msln−/− aPFs, but had no effect on ERK1/2 activation.

Like MSLN, Thy-1 is a GPI-anchored protein (50) that is expressed in fibroblasts, T cells, and neurons and is considered a specific marker for these cell types. Thy-1 was implicated in the inhibition of TGF-β1 responses in tissue fibroblasts via interaction with ανβ5 integrins (38–42). While Thy-1 functions are not well characterized, deletion of Thy-1 in mice was shown to exacerbate bleomycin-induced lung fibrosis (36). Our data suggest that Thy-1 blocks TGFβRI and that formation of the MSLN-Thy-1 complex enables TGF-β1 signaling. In accordance with this, downregulation of Thy-1 expression restored TGF-β1 signaling in Msln−/− mice, and deletion of Thy-1 in mice aggravated the development of cholestatic liver fibrosis. We identified TGFβRI as a binding partner of Thy-1 and MUC16 as a binding partner of MSLN. Since knocking out MSLN and MUC16 receptors together or separately yielded similar phenotypes in BDL-injured mice (Figure 7), we propose that the MSLN-MUC16 complex regulates the same signaling pathway in tissue fibroblasts and, therefore, can serve as a unique target for antifibrotic therapy in patients with cholestatic fibrosis.

Overall, our proposed model (summarized in Figure 6F and Figure 10D) suggests that the MSLN/MUC16 signaling pathway facilitates cholestatic fibrosis in mice via regulation of TGF-β1-mediated activation and FGF-induced proliferation of aPFs. In response to TGF-β1 signaling, MSLN disrupts the Thy-1-TGFβRI inhibitory complex and enables binding of TGF-β1 to its cognate receptor, phosphorylation and nuclear translocation of SMAD2/3, and transcription of the TGF-β1 target genes PAI1 and activin. Furthermore, our data suggest that the MSLN-MUC16 complex regulates FGF-dependent proliferation of aPFs, although indirectly, via activation of AKT, which regulates the trafficking and degradation of FGFR1 (45).
In support of these findings, a role for MSLN and CA125 (human analog of murine MUC16) in the regulation of AKT phosphorylation has been suggested in human cancer cells and was associated with increased proliferation and metastatic dissemination of cancer cells (49). CA125 release has been documented in patients with various malignancies and now serves as a standard marker in the diagnosis of ovarian cancer in patients (51). The role of MSLN and MUC16 in patients with liver fibrosis has not been evaluated. However, elevated serum levels of MSLN and CA125 were observed clinically in patients with liver fibrosis, and these levels were considered nonsignificant (52). Here we provide, for the first time to our knowledge, evidence that expression of MSLN is upregulated in the livers of patients with cholestatic fibrosis (Figure 4B) and that MSLN might play a role in the pathogenesis of BA and primary sclerosing cholangitis and possibly serve as a target for antifibrotic therapy. Further investigations are required to evaluate the role of MSLN and MUC16 in patients with cholestatic fibrosis.

Although the origin of aPFs during development has been identified in embryonic mesenchyme (32), which is also rich in MSLN expression (31, 53), the source of proliferating aPFs in the adult injured murine liver remains unclear. In response to cholestatic liver injury, adult aPFs may result from endogenous mesenchymal progenitors (33) or may derive from hepatic mesothelial progenitors. Remarkably, expression of the same fibulin 2 is induced in aPFs throughout BDL injury (Figure 1). And in agreement with previous findings (34), we observed that mesothelial cells, identified by expression of the GPM6A marker, also contributed to a small population of Col1+ myofibroblasts residing in the liver capsule of WT mice that had undergone 5 days of BDL injury (Supplemental Figure 2), while the majority of myofibroblasts were composed of Thy-1+ aPFs/myofibroblasts, located in the portal areas, and desmin+Col1+ aHSCs/myofibroblasts, located in the acinus. Concordant with our previous findings, aPFs/myofibroblasts contributed to 60% to 70% of hepatic myofibroblasts after BDL-induced (5 days) liver fibrosis (4).

**Methods**

**Patient specimens.** Archived liver specimens were obtained from 16 patients with liver cirrhosis (HCV, n = 5 [META VIR score of F2]; obstructive jaundice, n = 5 [F2]; BA, n = 6) and from 3 patients who were undergoing liver resection for reasons unrelated to liver fibrosis (normal liver) at the Kyoto University Hospital (2010 to 2013).

**Mice and liver injury.** Eight-week-old collagen α1(1)–GFP mice (62), Msln–/– mice (31), Mslnα1(1)–GFP Cre mice (53), Thy-1+ mice (50), Rosa26 CreERT2 FRT Cre mice (The Jackson Laboratory; 009669), Smadfl/fl mice (The Jackson Laboratory; 017462), and WT littermates on a C57BL/6 background were maintained under specific pathogen–free conditions at UCSD and had ad libitum access to normal chow and water. Liver injury was induced in mice by ligation of the common bile duct (for 5 days and 17 days), or by oral gavage with CCl4 carbon tetrachloride CCl4 (1:4 dilution in corn oil, 60 μl × 14 injections) (4). An additional model of cholestatic liver fibrosis was induced in WT and Msln–/– mice fed a diethyoxycarbonyl dihydrocolchicine (DDC) diet (AD5001; Custom Animal Diets) or a control diet (AD3012; Custom Animal Diets) for 3 weeks.

**Serum ALT and ALP measurements.** Serum ALT and ALP measurements were analyzed using Infinity ALT (Thermo Fisher Scientific), a QuantiChrom ALP Assay Kit (BioAssay Systems), and a VALIDATE calibration verification kit (Maine Standards Company LLC).

**Histology.** Formalin-fixed, paraffin-embedded human or mouse livers were stained with Picrosirius red (analyzed by a pathologist) or with Masson’s tris-green.
and LMP cell line (murine pancreatic MSLN + and MUC16 + cancer, blasts, gift of James S. Hagood, UCSD, La Jolla, California, USA). AKT was overexpressed in GEN, transfected using a FlexiTube GeneSolution (GS21838). CA Msln –/– used for IP experiments. Reconstitution of MSLN was achieved in aPFs by transfection of pCMV-Tag3A-N-Myc-MSLN. Thy-1 knockout mice were stained with anti-mouse α-SMA (Abcam; ab5694); anti-mouse desmin (Thermo Fisher Scientific; RB-9014-P0); and anti-mouse F4/80 (eBioscience: 14-4801-82) Abs, followed by staining with DAB (Vector Laboratories) or with secondary Alexa Fluor 594–conjugated Abs. LacZ staining was performed in freshly isolated mouse livers (Sigma-Aldrich; GALS1KT). Images were taken using an Olympus microscope, and positive areas were calculated using ImageJ software (NIH) (see Supplemental Methods).

Mouse livers were immunostained with anti-human α-SMA (Dako; M0851); anti-human desmin (Thermo Fisher Scientific; PA5-16705); anti-human MSLN (Sigma-Aldrich; HPA017172); and anti-human Thy-1 (Abcam; ab92574) Abs. Mouse livers were stained with anti-mouse α-SMA (Abcam; ab5694); anti-mouse desmin (Thermo Fisher Scientific; RB-9014-P0); and anti-mouse F4/80 (eBioscience: 14-4801-82) Abs, followed by staining with DAB (Vector Laboratories) or with secondary Alexa Fluor 594–conjugated Abs. LacZ staining was performed in freshly isolated mouse livers (Sigma-Aldrich; GALS1KT). Images were taken using an Olympus microscope, and positive areas were calculated using ImageJ software (NIH) (see Supplemental Methods).

Cell culture. The RFL-6 CD90.2 cell line (rat lung Thy-1+ fibroblasts, gift of James S. Hagood, UCSD, La Jolla, California, USA) and LMP cell line (murine pancreatic MSLN+ and MUC16+ cancer, blasts, gift of James S. Hagood, UCSD, La Jolla, California, USA) were used for IP experiments. Reconstitution of MSLN in Msln–/– aPFs was performed by transfection of pCMV-Tag3A-N-Myc-MSLN. Thy-1 knockdown was achieved in Msln–/– aPFs with Thy-1 siRNA (Qiagen). CA AKT was overexpressed in Msln–/– aPFs infected with adenovirus AA-CA-AKT expressing an amino-terminal myristylation signal (AdmyrAkt) (63), compared with control AAV-GFP (see Supplemental Methods).

qPCR. Total RNA was isolated from livers or isolated aPFs using RNeasy columns (QIAGEN) (4). Expression levels of selected genes were calculated after normalization to the standard housekeeping gene HPRT (Invitrogen, Thermo Fisher Scientific) using the ΔΔCt method. The data represent relative mRNA levels compared with control levels (mean ± SEM, P < 0.05).

IP and Western blot analysis. IPs from the precleared whole-cell lysates were performed at 4°C overnight using 2 μg anti-MSLN Ab (IBL; catalog 28127), anti-TGFβRI Ab (Abcam; catalog ab31013), or anti-MUC16 Ab (Abicode; catalog R2334-3) and analyzed by Western blotting (see Supplemental Methods).

Statistics. All data represent the mean ± SEM. Comparisons of 2 groups were analyzed using an unpaired, 2-tailed Student’s t test. Comparisons of 3 or more groups were analyzed using ANOVA. ANOVA with a Dunnett’s test was used for comparing multiple groups of mice or treatments with controls. ANOVA with a Bonferroni’s test was used when making multiple pair-wise comparisons between different groups. A P value of less than 0.05 was considered statistically significant. Analyses were performed using GraphPad Prism (Prism 5 version 5.03; GraphPad Software).

Study approval. All mice were maintained under specific pathogen-free conditions at UCSD according to UCSD IACUC protocol S07088. For human subjects, written informed consent was obtained from each patient in accordance with the ethics guidelines for epidemiological research in Japan (E2077 protocol, approved by the Ethics Committee of Kyoto University Graduate School of Medicine).

Author contributions YK performed in vivo experiments, analyzed the data, and wrote the manuscript. PW, SL, and DK performed in vitro experiments. KI performed in vivo experiments and conducted human studies. XL, JX, MS, and MC performed experiments. KT and TB provided expertise in liver fibrosis and mesothelin. CB and SH assisted with data analysis. DAB provided support and critical revision of the manuscript. TK provided support, designed the study, and wrote the manuscript.

Acknowledgments We thank Gianfranco D. Alpini (Baylor Scott & White Healthcare, Temple, Texas, USA) for providing mRNA from isolated cholangiocytes. We thank James S. Hagood (UCSD, La Jolla, California, USA) and Yuval Rinkevitch (Institute of Lung Biology and Disease [ILBD] Helmholtz Zentrum München, Munich, Germany) for providing Thy-1–/– and MslnER–/– mice. We are grateful to Koji Taniguchi (Keio University, Tokyo, Japan) for his technical support on confocal microscopy. We thank Karin Biggle (UCSD, La Jolla, California, USA) for her dedicated management of our laboratory. This research was supported by the NIH (AA022614, DK099205, DK101737, DK111866, AA011999, ES010337, AA021856, and GM041804); the Japanese Ministry of Health, Labour and Welfare; and the American Liver Foundation. This research was supported in part by the Intramural Research Program of the NIH, National Cancer Institute, Center for Cancer Research.

Address correspondence to: Tatiana Kisseleva, 9500 Gilman Drive, #0063, La Jolla, California 92093, USA. Phone: 858.822.5339; E-mail: tkisseleva@ucsd.edu.

Figure 13. MSLN/MUC16/AKT signaling regulates FGF-dependent proliferation of aPFs. (A and B) Msln–/– aPFs were infected with CA AKT (MslnRD-AD-AKT aPFs). (A) Expression of cyclin D and cyclin E was detected by qPCR. (B) Phosphorylation of JAK2 was restored in MslnRD-AD-AKT aPFs and was normalized to JAK2 protein expression levels (see also Supplemental Figure 11). Data shown is representative of more than 3 independent experiments.


61. Kinkevich Y, Mori T, Saho D, Xu PX, Bermingham JR, Weissman IL. Identification and prospective isolation of a mesothelial...


