Figure S1. Full-length of western blot images for Fig. 1A. For all experiments, the same membrane was re-probed with p-Smad2, t-Smad2, p-Smad3, t-Smad3, Smad4, p-ERK1/2, t-ERK/1/2, p-p38, t-p38 and GAPDH antibodies. (A-J) Full-length western blot analysis images [(A) p-Smad2, (B) t-Smad2 (C) p-Smad3, (D) t-Smad3, (E) Smad4, (F) p-ERK1/2, (G) t-ERK/1/2, (H) p-p38, (I) t-p38 and (J) GAPDH] were showing. p-, phosphorylated; t-, total; ERK, extracellular signal-regulated kinase; p38, p38 mitogen-activated protein kinase.



Figure S2. Effect of galunisertib on TGF- $\beta$  signaling in HepG2 cells. (A) Levels of p-Smad2 protein were evaluated by western blot analysis (n=4). Galunisertib treatment dose-dependently decreased p-Smad2 expression levels in HepG2 cells. (B-D) Complete western blot images for part A. TGF- $\beta$ , transforming growth factor- $\beta$ ; p-, phosphorylated.



Figure S3. Liver images for all groups. Gal, galunisertib;  $CCl_4$ , carbon tetrachloride; Intact, no  $CCl_4$  or galunisertib treatment; Vehicle,  $CCl_4$ -treated with no galunisertib treatment; Gal 50, livers from mice treated with low-dose galunisertib; Gal 150, treatment with middle-dose galunisertib; Gal 300, treatment with high-dose galunisertib.



Gal 150

Gal 300

Figure S4. Expression levels of genes associated with fibrosis in livers from mice treated with  $CCl_4$  for 8 weeks. (A-C) Reverse transcription-quantitative PCR analysis of (A) *Collal*, (B) *Fn1* and (C) *Acta2* mRNA expression demonstrated increases after 8 weeks of  $CCl_4$  treatment compared with those in the untreated liver group. The mRNA expression levels were similar to those for vehicle-treated mice at all galunisertib dosages. \*P<0.05. Error bars represent the means ± SEM. Gal, galunisertib;  $CCl_4$ , carbon tetrachloride; Intact, no  $CCl_4$  or galunisertib treatment; Vehicle,  $CCl_4$ -treated with no galunisertib treatment; Gal 50, livers from mice treated with low-dose galunisertib; Gal 150, treatment with middle-dose galunisertib; Gal 300, treatment with high-dose galunisertib; n.s., not significant.



Figure S5. Full-length of western blot analysis and gelatin zymography images for MMP protein expression. For the western blot analysis, the same membrane was re-probed with (A) MMP-1, (B) MMP-13 and (C) GAPDH antibodies. (D) Full-length of gelatin zymography image is shown. MMP, matrix metalloproteinase; Gal, galunisertib;  $CCl_4$ , carbon tetrachloride; Intact, no  $CCl_4$  or galunisertib treatment; Vehicle,  $CCl_4$ -treated with no galunisertib treatment; Gal 50, livers from mice treated with low-dose galunisertib; Gal 150, treatment with middle-dose galunisertib; Gal 300, treatment with high-dose galunisertib.



Figure S6. Semi-quantitative analysis of MMP-1 and MMP-13 protein expression and RT-qPCR analysis of *Timp1* gene expression. (A and B) MMP-1 and MMP-13 expression levels in the livers from galunisertib-treated mice were similar to that observed in the vehicle-treated mice using semi-quantitative western blot analysis, in which two membranes were assessed (n=4/group). The samples were derived from the same experiment and the gels/blots were processed in parallel. (C) RT-qPCR analysis indicated that the *Timp1* gene expression level in the livers from galunisertib-treated mice was not significantly upregulated compared with the vehicle-treated mice. \*P<0.05. Error bars represent the means  $\pm$  SEM. MMP, matrix metalloproteinase; Timp1, TIMP metalloproteinase inhibitor 1; RT-qPCR, reverse transcription-quantitative PCR; Gal, galunisertib; CCl<sub>4</sub>, carbon tetrachloride; Intact, no CCl<sub>4</sub> or galunisertib treatment; Vehicle, CCl<sub>4</sub>-treated with no galunisertib treatment; Gal 50, livers from mice treated with low-dose galunisertib; Gal 150, treatment with middle-dose galunisertib; Gal 300, treatment with high-dose galunisertib; n.s., not significant.



Figure S7. Full-length of western blot images for PNCA protein expression. In all experiments, the same membrane was re-probed with (A) PCNA and (B) GAPDH antibodies. PCNA, proliferating cell nuclear antigen; Gal, galunisertib; CCl<sub>4</sub>, carbon tetrachloride; Intact, no CCl<sub>4</sub> or galunisertib treatment; Vehicle, CCl<sub>4</sub>-treated with no galunisertib treatment; Gal 50, livers from mice treated with low-dose galunisertib; Gal 150, treatment with middle-dose galunisertib; Gal 300, treatment with high-dose galunisertib.



Figure S8. Immunohistochemical detection of cleaved caspase-3 in apoptotic hepatocytes in mouse liver. (A) The number of hepatocytes that were positive for cleaved caspase-3-positive (arrowhead) in tissue from galunisertib-treated mice was similar to that for vehicle-treated mice. Scale bar=200  $\mu$ m. (B) For semi-quantitative analyses, hepatocytes that were positive for cleaved caspase-3 were counted using image analysis software. No significant difference was identified in the number of cleaved caspase-3-positive hepatocytes from galunisertib-treated mice compared with that of the vehicle-treated mice. \*P<0.05. Error bars represent the means ± SEM. Gal, galunisertib; CCl<sub>4</sub>, carbon tetrachloride; Vehicle, CCl<sub>4</sub>-treated without galunisertib treatment; Gal 50, low-dose galunisertib treatment; Gal 150, middle-dose galunisertib treatment; Gal 300, high-dose galunisertib treatment; n.s., not significant.



Figure S9. RT-qPCR analysis for verification of RT<sup>2</sup> profiler PCR array. RT-qPCR analysis revealed that mRNA expression levels of (A) *Fgf7*, (B) *Egf* and (C) *Hgf* in liver tissue from mice treated with high-dose galunisertib were similar to those from the vehicle-treated group. \*P<0.05. Error bars represent the means  $\pm$  SEM. RT-qPCR, reverse transcription-quantitative PCR; *Fgf7*, fibroblast growth factor-7; *Egf*, epithelial growth factor; *Hgf*, hepatocyte growth factor; n.s., not significant; Gal, galunisertib; CCl<sub>4</sub>, carbon tetrachloride; Vehicle, CCl<sub>4</sub>-treated without galunisertib treatment; Gal 50, low-dose galunisertib treatment; Gal 300, high-dose galunisertib treatment.



Table SI. Primers used for reverse transcription-quantitative PCR.

Target gene	Assay ID	Species
TaqMan COL1a1	Hs00164004_m1	Human
TaqMan <i>MMP1</i>	Hs00899658_m1	Human
TaqMan <i>GAPDH</i>	Hs02758991_g1	Human
TaqMan Acta2	Mm00725412_s1	Mouse
TaqMan Collal	Mm00801666_g1	Mouse
TaqMan Egf	Mm00438696_m1	Mouse
TaqMan Ereg	Mm00514794_m1	Mouse
TaqMan Fgf7	Mm00433291_m1	Mouse
TaqMan Fn1	Mm01256744_m1	Mouse
TaqMan Hgf	Mm01135193_m1	Mouse
TaqMan Igfl	Mm00439560_m1	Mouse
TaqMan Il6	Mm00446190_m1	Mouse
TaqMan Mmp1a	Mm00473485_m1	Mouse
TaqMan Mmp1b	Mm00473493_g1	Mouse
TaqMan Mmp2	Mm00439498_m1	Mouse
TaqMan Mmp9	Mm00442991_m1	Mouse
TaqMan Mmp13	Mm00439491_m1	Mouse
TaqMan Tgfa	Mm00446232_m1	Mouse
TaqMan Timp1	Mm01341361_m1	Mouse
TaqMan Gapdh	Mm99999915_g1	Mouse

Reference number Gene name Fold change NM\_008742 315.4715 Neurotrophin 3 NM 010275 Glial cell line derived neurotrophic factor 12.3961 NM 010556 Interleukin 3 10.1239 NM 008004 Fibroblast growth factor 17 10.0161 NM 007558 Bone morphogenetic protein 8a 9.1085 NM 008109 Growth differentiation factor 5 9.0554 NM\_007445 Anti-Mullerian hormone 6.1883 NM 009971 Colony stimulating factor 3 (granulocyte) 4.9012 NM\_008002 Fibroblast growth factor 10 4.3734 NM 001314054 Interleukin 6 3.5619 NM\_007950 Epiregulin 3.141 Inhibin beta-B NM 008381 2.8123 NM 010834 Myostatin 2.2234 NM 008493 Leptin 2.1067 NM\_008350 Interleukin 11 2.0795 NM 021704 Chemokine (C-X-C motif) ligand 12 1.9672 NM\_008005 Fibroblast growth factor 18 1.8276 NM 008008 Fibroblast growth factor 7 1.7178 NM\_013518 Fibroblast growth factor 9 1.6473 NM 008501 Leukemia inhibitory factor 1.5602 NM 008351 Interleukin 12A 1.5 NM 011313 S100 calcium binding protein A6 (calcyclin) 1.4769 NM\_013611 Nodal 1.4502 NM 010197 Fibroblast growth factor 1 1.4227 NM\_008380 Inhibin beta-A 1.4099 Fibroblast growth factor 22 NM 023304 1.409 NM\_198190 Neurotrophin 5 1.3165 NM 010554 Interleukin 1 alpha 1.2833 NM 008808 Platelet derived growth factor alpha 1.28 NM 008003 Fibroblast growth factor 15 1.2641 NM\_021283 Interleukin 4 1.2424 NM 010113 Epidermal growth factor 1.2218 NM\_007557 Bone morphogenetic protein 7 1.2192 Colony stimulating factor 2 (granulocyte-macrophage) NM 009969 1.1791 NM\_009756 Bone morphogenetic protein 10 1.1775 NM 009263 Secreted phosphoprotein 1 1.1749 NM 010784 Midkine 1.15 NM\_007553 Bone morphogenetic protein 2 1.1405 NM\_009368 Transforming growth factor, beta 3 1.1373 NM 010427 Hepatocyte growth factor 1.1368 NM\_011577 Transforming growth factor, beta 1 1.1349 NM 010200 Fibroblast growth factor 13 1.1323 NM\_013598 Kit ligand 1.1262 NM 008827 Placental growth factor 1.1038 NM 010203 Fibroblast growth factor 5 1.0682 NM 008361 Fibroblast growth factor 2 1.0433 NM 008361 Interleukin 1 beta 1.0368 NM 010094 Left right determination factor 1 1.0131 NM 010216 C-fos induced growth factor 1.0047

Table SII. Expression changes of upregulated growth factors in a liver sample from the high-dose galunisertib-treated group compared with liver tissue from the vehicle-treated group (n=1), as measured by the RT<sup>2</sup> Profiler PCR Array analysis.