ASK1 Inhibition Reduces Cell Death and Hepatic Fibrosis in an Nlrp3 Mutant Liver Injury Model

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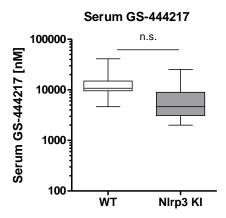
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Supplementary Table. Primer sequences used for quantitative PCR.

Gene	Primer	Sequence 5'-3'
Мро	FW	TTA AAA ACC TGG ATC GGA ACC AA
	RV	GCA TTA GCT TCA GAT TTA CGG GT
Collal	FW	GCT CCT CTT AGG GGC CAC T
	RV	CCA CGT CTC ACC ATT GGG G
Col1a2	FW	AAG GGT GCT ACT GGA CTC CC
	RV	TTG TTA CCG GAT TCT CCT TTG G
Lum	FW	CTC TTG CCT TGG CAT TAG TCG
	RV	GGG GGC AGT TAC ATT CTG GTG
Emp1	FW	TTG GTG CTA CTG GCT GGT CT
	RV	CAT TGC CGT AGG ACA GGG AG
Acta2	FW	GTC CCA GAC ATC AGG GAG TAA
	RV	TCG GAT ACT TCA GCG TCA GGA
Timp1	FW	CTT GGT TCC CTG GCG TAC TC
	RV	ACC TGA TCC GTC CAC AAA CAG
Hprt	FW	AGT CCC AGC GTC GTG ATT AG
	RV	TTT CCA AAT CCT CGG CAT AAT GA

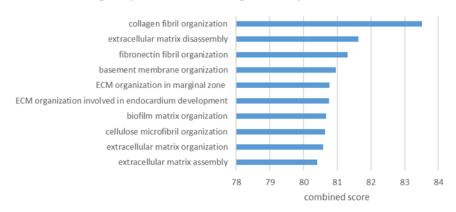
Supplementary Figure S1



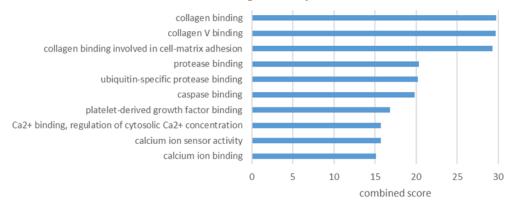
Supplementary Figure S1. Serum concentration of GS-444217. Serum concentration of GS-444217 [nM] were measured in fasted WT and *Nlrp3* KI mice by Mass spectrometry. Data show the median with Min to Max. (n.s not significant, unpaired two-tailed t-test).

Supplementary Figure S2

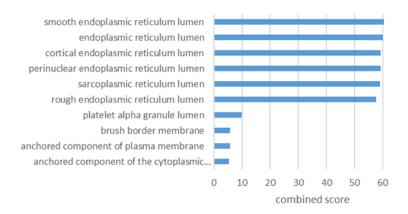
GO Biological processes downregulated by ASK1i



GO Molecular function downregulated by ASK1i



GO Cellular compartment downregulated by ASK1i



Supplementary Figure S1. Enrichment analysis of Gene Ontology (GO) Biological processes, Molecular function and Cell compartment. Top 10 GO's ranked by combined

score associated with down-modulated signatures using EnrichR analysis tool. Benjamini-Hochberg adjusted p-value < 0.05 was used as significance threshold.