

**IGF-I induces senescence of hepatic stellate cells and limits fibrosis in a p53-
dependent manner.**

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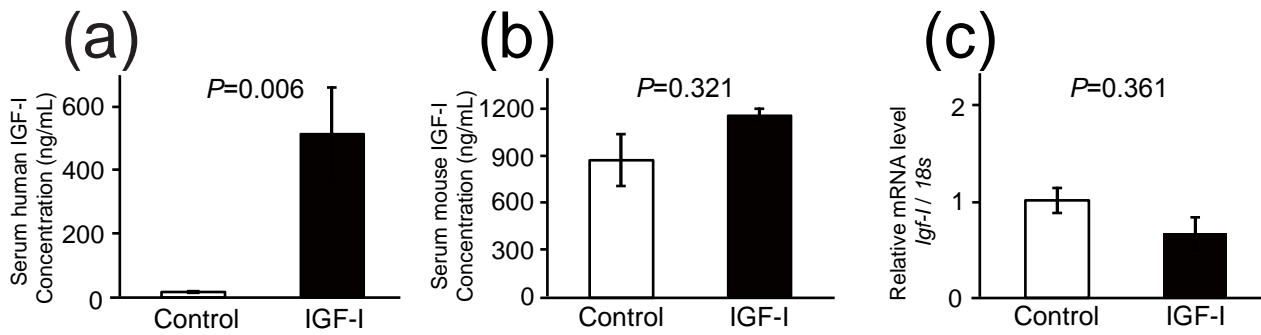
Supplementary Table 1

RT-PCR Primers for Analysis

Gene	Organism	Direction	Sequence
<i>Cd68</i>	mouse	Forward	GACACTTCGGGCCATGTT
		Reverse	GAGGAGGACCAGGCCAAT
<i>F4/80</i>	mouse	Forward	GGAGGACTTCTCCAAGCCTATT
		Reverse	AGGCCTCTCAGACTTCTGCTT
<i>Tnfa</i>	mouse	Forward	CCGATGGGTTGTACCTTGTC
		Reverse	CGGACTCCGCAAAGTCTAAG
<i>Tgfb</i>	mouse	Forward	TTGCCCTCTACAACCAACACAA
		Reverse	GGCTTGCGACCCACGTAGTA
<i>Il-1β</i>	mouse	Forward	TTGACGGACCCAAAAGAT
		Reverse	GAAGCTGGATGCTCTCATCTG
<i>Il-6</i>	mouse	Forward	GCTACCAAAGTGGATATAATCAGGA
		Reverse	CCAGGTAGCTATGGTACTCCAGAA
<i>Procollagen1a1</i>	mouse	Forward	GAGCGGAGAGTACTGGATCG
		Reverse	GTTCCGGCTGATGTACCAGT
<i>Collagen3a1</i>	mouse	Forward	TCCCCTGGAATCTGTGAATC
		Reverse	TGAGTCGAATTGGGGAGAAT
<i>Collagen4a1</i>	mouse	Forward	TTAAAGGACTCCAGGGACCAC
		Reverse	CCCCTGAGCCTGTCCACAC
<i>Mmp3</i>	mouse	Forward	TTGTTCTTTGATGCAGTCAGC
		Reverse	GATTTGCGCCAAAAGTGC
<i>Mmp9</i>	mouse	Forward	ACGACATAGACGGCATCCA
		Reverse	GCTGTGGTTCAGTTGTGGTG
<i>Timp1</i>	mouse	Forward	ATTCAAGGCTGTGGGAAATG
		Reverse	CTCAGAGTACGCCAGGGAAC
<i>Cox3</i>	mouse	Forward	GCAGGATTCTTCTGAGCGTTCT
		Reverse	GTCAGCAGCCTCCTAGATCATGT
<i>Cyt c</i>	mouse	Forward	GCAAGCATAAGACTGGACCAA
		Reverse	TTGTTGGCATCTGTGTAAGAGAATC
<i>Nrf2</i>	mouse	Forward	CCAGAAGCCACACTGACAGA
		Reverse	GGAGAGGATGCTGCTGAAAG
<i>Sco2</i>	mouse	Forward	ATCGCACAGCCCTAAGTCTC
		Reverse	CAGTAGCATCGTGGACCTGA
<i>αsma</i>	mouse	Forward	AAACAGGAATACGACGAAG
		Reverse	CAGGAATGATTTGGAAAGGA
<i>Vimentin</i>	mouse	Forward	CCAACCTTTTCTTCCCTGAA
		Reverse	TGAGTGGGTGTCAACCAGAG
<i>Fibronectin</i>	mouse	Forward	CGGAGAGAGTGCCCTACTA
		Reverse	CGATATTGGTGAATCGCAGA
<i>p21</i>	mouse	Forward	AACATCTCAGGGCCGAAA
		Reverse	TGCGCTTGGAGTGATAGAAA
<i>18s</i>	mouse	Forward	TCAAGAACGAAAGTCGGAGG
		Reverse	GGACATCTAAGGGCATCACA
<i>Gapdh</i>	mouse	Forward	AGCTTGTCAACCGGAAG
		Reverse	TTTGATGTTAGTGGGTCTCG
<i>αsma</i>	rat	Forward	TCCCAAGCTGTGTCTCTG
		Reverse	GTGCCACGTTATGATGATGC
<i>Vimentin</i>	rat	Forward	CTCACCTCGGAAGTGGATG
		Reverse	TCTTTCCATTTACGCATCT
<i>Igf-1</i>	mouse	Forward	GCTTGCTCACCTTCACCAGC
		Reverse	AATGTACTIONCTTCTGAGTCT

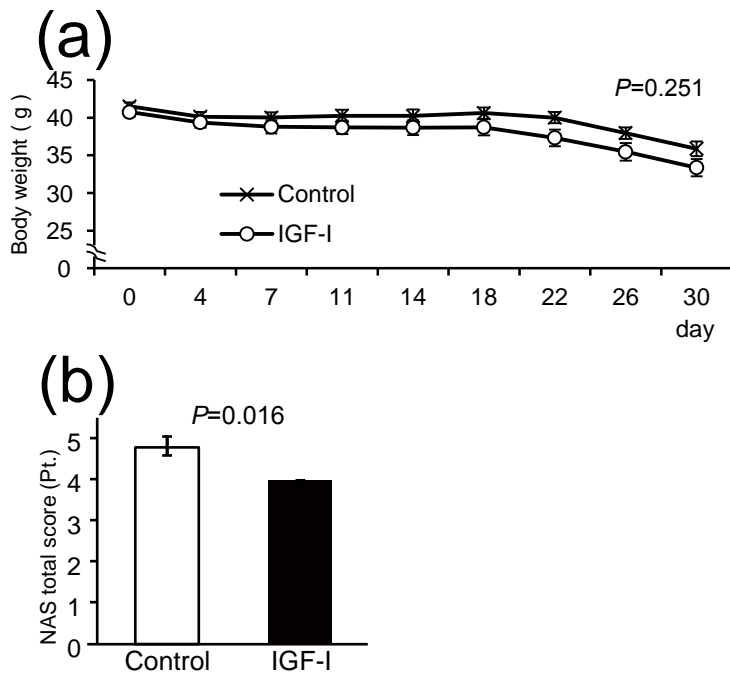
Supplementary Table 1 Primer sequence for realtime PCR analysis

Supplementary Figure 1



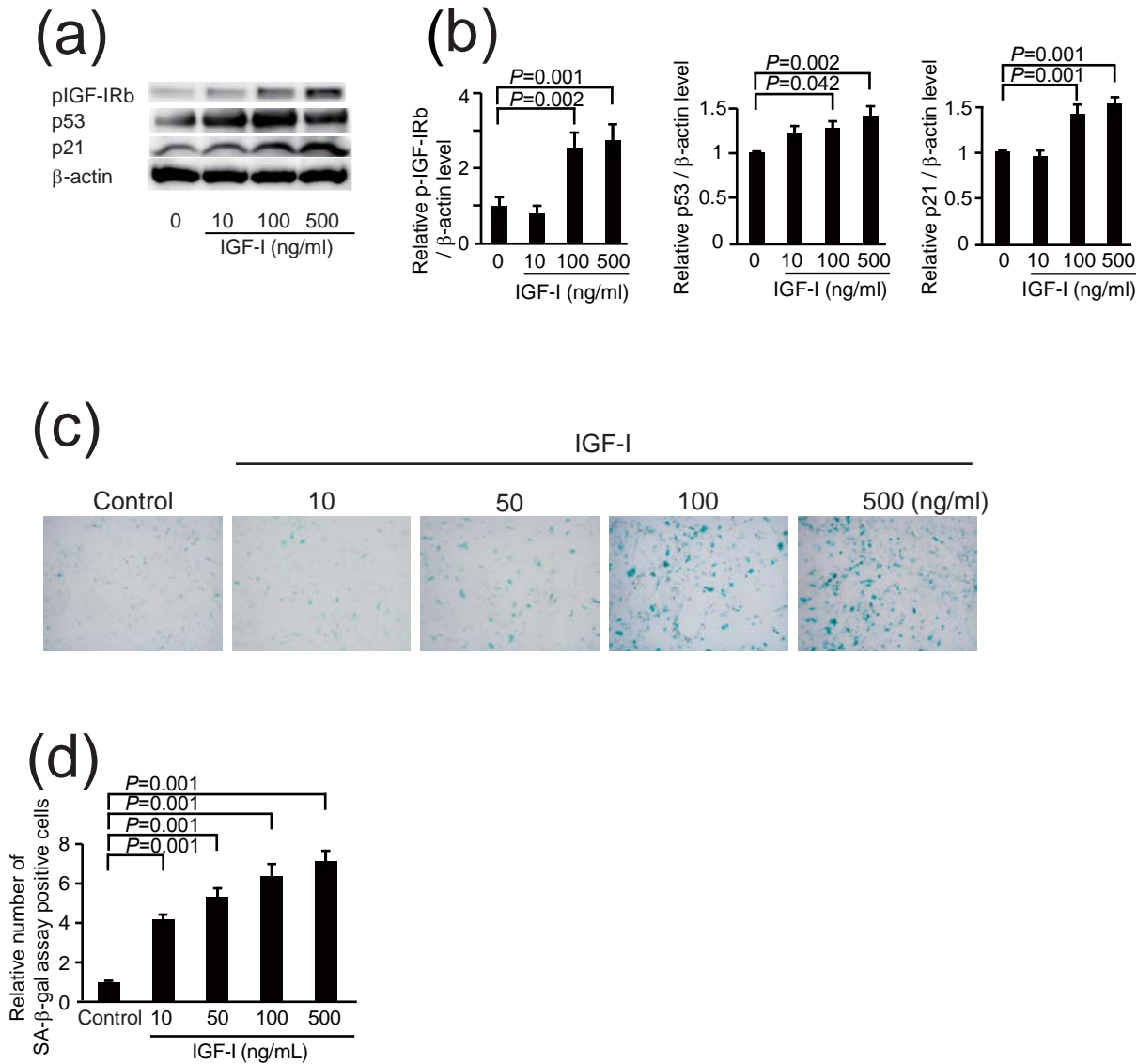
Supplementary Figure 1 (a) Serum human IGF-I concentrations measured using human IGF-I-specific ELISA kit in NASH model mice. (b) Serum mouse IGF-I concentrations measured using mouse IGF-I-specific ELISA kit. (c) Quantitative realtime PCR analysis revealed no changes in mRNA expression of endogenous *Igf-1* in the liver. Data were compared by Student's *t* test.

Supplementary Figure 2



Supplementary Figure 2 (a) Changes in the body weight during IGF-I treatment in the NASH model mice. Data were compared by MANOVA test. (b) Total NAFLD activity score (NAS) was evaluated by a blinded pathologists. NAS was assessed on a scale of 0-8, with higher scores indicating more severe disease; the components of this measure include steatosis (assessed on a scale of 0-3), lobular inflammation (assessed on a scale of 0-3), and hepatocellular ballooning (assessed on a scale of 0-3). SData were compared by tudent's *t* test, $p < 0.05$.

Supplementary Figure 3



Supplementary Figure 3 IGF-I induced cellular senescence in human hepatic stellate cell line, LX2 cells.

(a) IGF-I increased expressions of p53 and p21 proteins in a concentration-dependent manner. Cells were incubated with IGF-I for 72 h. (b) Quantitative analysis demonstrated that IGF-I significantly increased p53 and p21 protein expression and phosphorylation of the IGF-I receptor in a concentration-dependent manner. Densitometric analyses were performed using data from 5 independent experiments. Each value was normalized to that of β -actin. (c) IGF-I induced SA- β -gal activation in LX2 cells. HSCs were incubated with IGF-I at the indicated concentrations for 5 days. (SA- β -gal staining, 200 \times). (d) Quantitative analysis demonstrated that IGF-I significantly increased SA- β -gal-positive cells in a concentration-dependent manner. Data were expressed as the mean \pm SEM of 20 random fields ($n = 10$). Data were compared by Tukey's honestly significant difference test.