

Obeticholic acid protects against hepatocyte death and liver fibrosis in a murine model of nonalcoholic steatohepatitis

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Supplementary materials

Supplementary method

Western blotting

Nuclear proteins were extracted using NE-PER Nuclear and Cytoplasmic Extraction Reagents (Thermo Fisher Scientific, San Jose, CA, USA) supplemented with Halt Protease and Phosphatase Inhibitor Cocktail and sodium butyrate. Proteins were separated by SDS-PAGE and immunoblotted with antibodies against MDM2 (ab16895, abcam), phospho-p53 (Ser15) (9284, Cell Signaling) and acetyl-p53 (Lys379) (2570, Cell Signaling), and Lamin A/C (sc-20168, Santa Cruz).

Supplementary Table S1. Primers used in the present study

<i>Genes</i>	Primers	Primers
<i>Abcb11</i>	Forward	GTCTGACTCAGTGATTCTTCGC
	Reverse	GAGCAATGCGCACACACTTC
<i>Apaf1</i>	Forward	CTCTAGATGAAGCCATGTCTG
	Reverse	AAGGATGGAAAGGTCTGTGT
<i>Ccl2</i>	Forward	CCACTCACCTGCTGCTACTCAT
	Reverse	TGGTGATCCTCTTGTAGCTCTCC
<i>Coll1a1</i>	Forward	CCTCAGGGTATTGCTGGACAAC
	Reverse	ACCACTTGATCCAGAAGGACCTT
<i>Col4a1</i>	Forward	TCCATGGCACCCATCTCTG
	Reverse	CTTCACAAACCGCACACCTG
<i>Ctgf</i>	Forward	CTTCTGCGATTTCTGGCTCC
	Reverse	TACACCGACCCACCGAAGA
<i>Cyp7a1</i>	Forward	CTGATCCGTCTACGCATGTTTC
	Reverse	CAGGAATGGTGTTTGCTTGAGA
<i>Cyp8b1</i>	Forward	ACGCTTCCTCTATCGCCTGAA
	Reverse	GTGCCTCAGAGCCAGAGGAT
<i>Emr1</i>	Forward	CTTTGGCTATGGGCTTCCAGT
	Reverse	GCAAGGAGGACAGAGTTTATCGTG
<i>Mdm2</i>	Forward	AAGGAGGAAACGCAGGACAA
	Reverse	TCTTGCCGTGAACAATGCA
<i>Nr0b2</i>	Forward	AAGGGCACGATCCTCTTCAA
	Reverse	GTACCAGGGCTCCAAGACT
<i>Nr1h4</i>	Forward	AGGCCATGTTTCTTCGTTCG
	Reverse	CATGACCGGCAGGAAGTTTC

<i>P21</i>	Forward	AACATCTCAGGGCCGAAAAC
	Reverse	CTGCGCTTGGAGTGATAGAA
<i>Slc51b</i>	Forward	GACAAGCATGTTCCCTCCTGAG
	Reverse	GATGCAGGTCTTCTGGTGTTTC
<i>Srebflc</i>	Forward	AGCTGTCTGGGGTAGCGTCTG
	Reverse	GAGAGTTGGCACCTGGGCTG
<i>Tgfb1</i>	Forward	CCTGAGTGGCTGTCTTTTGACG
	Reverse	AGTGAGCGCTGAATCGAAAGC
<i>Timp1</i>	Forward	CATCACGGGCCGCCTA
	Reverse	AAGCTGCAGGCACTGATGTG
<i>Tnfa</i>	Forward	ACCCTCACACTCAGATCATCTTC
	Reverse	TGGTGGTTTGCTACGACGT
<i>Tnfs10b</i>	Forward	TGCTGCTCAAGTGGCGC
	Reverse	GGCATCCAGCAGATGGTTG
<i>Tp53</i>	Forward	CTCAAAAACTTACCAGGGC
	Reverse	CACCACGCTGTGGCGAAAAGTCTG
<i>Vcam1</i>	Forward	TCACAATTAAGAAGTTTAACACTTGATGTAA
	Reverse	GAGTGCAAGGAGTTCGGGC
<i>36B4</i>	Forward	GGCCCTGCACTCTCGCTTTC
	Reverse	TGCCAGGACGCGCTTGT

Supplementary Table S2. Effect of OCA treatment on total bile acid in MC4R-KO fed WD for 24 weeks.

	WT-SD	MC4R-WD	
	Vehicle	Vehicle	OCA
Serum bile acid ($\mu\text{mol/L}$)	3.69	21.25	2.33
Liver bile acid ($\mu\text{mol/g}$ tissue)	0.16	0.34	0.11
Ileum bile acid ($\mu\text{mol/g}$ tissue)	0.62	2.13	0.73

Serum and tissue bile acid levels were measured using samples pooled from 6 mice for WT-SD, 10 mice for vehicle-treated MC4R-WD, 9 mice for OCA-treated MC4R-WD.

Supplementary Table S3. Effect of OCA on hepatic bile acid composition (%) in MC4R-KO fed WD for 24 weeks.

	WT-SD		MC4R-WD	
	Vehicle	Vehicle	OCA	
TCA	30.4	40.2	10.1	
CA	0.8	0.4	0.0	
TCDCa	2.4	3.2	6.5	
CDCA	0.1	0.0	0.0	
GUDCA	0.0	0.0	0.0	
TUDCA	2.3	3.7	10.7	
UDCA	0.2	0.1	0.8	
TDCA	4.7	0.7	0.0	
DCA	0.0	0.0	0.0	
THCA	2.3	0.7	0.9	
HCA	0.0	0.0	0.0	
TLCA	0.0	0.0	0.0	
LCA	0.0	0.0	0.0	
T- α -MCA	3.0	5.9	7.2	
α -MCA	0.7	0.4	0.8	
T- β -MCA	22.8	32.2	40.0	
β -MCA	12.7	6.1	12.0	
T- ω -MCA	11.1	5.2	7.7	
ω -MCA	3.5	0.6	1.4	

CA, cholic acid; TCA, tauro-CA; CDCA, chenodeoxycholic acid; TCDCa, tauro-CDCA;

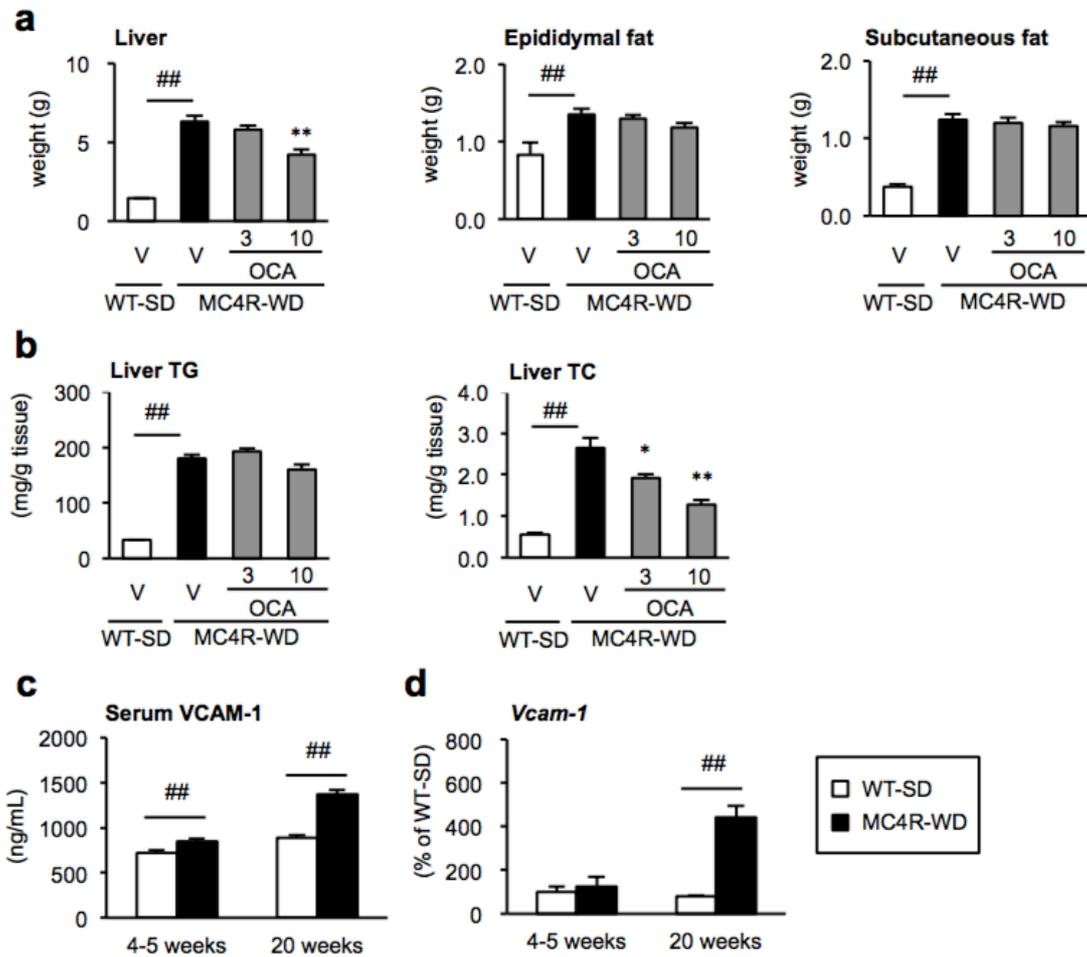
UDCA, ursodeoxycholic acid; GUDCA, glycol-UDCA; TUDCA, tauro-UDCA; DCA, deoxycholic acid; TDCA, tauro-DCA; HCA, hyocholic acid; THCA, tauro-HCA; LCA, lithocholic acid; TLCA, tauro-LCA; $\alpha/\beta/\omega$ -MCA, $\alpha/\beta/\omega$ -muricholic acid; T- $\alpha/\beta/\omega$ -MCA, tauro- $\alpha/\beta/\omega$ -MCA. Hepatic bile acid composition was measured using samples pooled from 6 mice for WT-SD, 10 mice for vehicle-treated MC4R-WD, 9 mice for OCA-treated MC4R-WD.

Supplementary Table S4. Serological parameters of MC4R-KO treated with OCA for 4 or 8 weeks after the development of NASH.

Weeks	MC4R-WD				
	-	Vehicle	OCA	Vehicle	OCA
	0	4	4	8	8
BG (<i>ad lib</i> , mg/dL)	149.6 ± 9.4	150.0 ± 7.0	167.9 ± 8.6	152.1 ± 12.4	179.2 ± 9.2
TC (mg/dL)	320.6 ± 11.3 ^{##}	328.0 ± 27.4	263.0 ± 11.0 [*]	336.3 ± 8.8	270.9 ± 19.3 ^{**}
TG (mg/dL)	60.4 ± 7.1	78.8 ± 11.4	56.5 ± 7.2	56.3 ± 4.8	52.7 ± 6.3
NEFA (mEq/L)	0.96 ± 0.74	0.81 ± 0.40	0.66 ± 0.32	0.72 ± 0.62	0.73 ± 0.50
ALT (U/L)	344.6 ± 26.4 ^{##}	499.5 ± 104.5	323.8 ± 52.2	700.0 ± 92.2	323.6 ± 55.0 ^{**}
AST (U/L)	417.0 ± 45.3 ^{##}	426.5 ± 53.6	329.3 ± 62.1	580.0 ± 63.2	300.9 ± 33.6 ^{**}
VCAM-1 (µg/mL)	1.37 ± 0.05 ^{##}	1.43 ± 0.06	1.21 ± 0.03 ^{**}	1.38 ± 0.07	1.19 ± 0.06

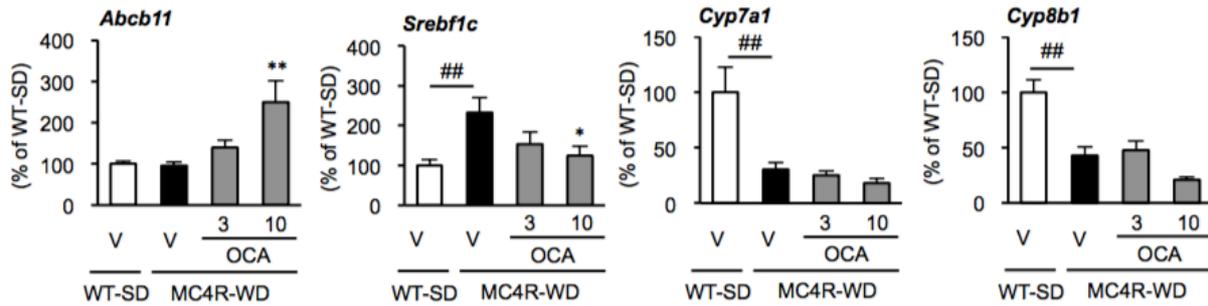
VCAM-1, vascular cell adhesion molecule-1; -, no treatment. Data are expressed as the mean ± SE. n = 5-10. [#]*p* < 0.05, ^{##}*p* < 0.01 MC4R-WD (pre-treatment) group vs. WT-SD group, ^{*}*p* < 0.05, ^{**}*p* < 0.01 vs. MC4R-WD group treated with the vehicle.

Supplementary Figures



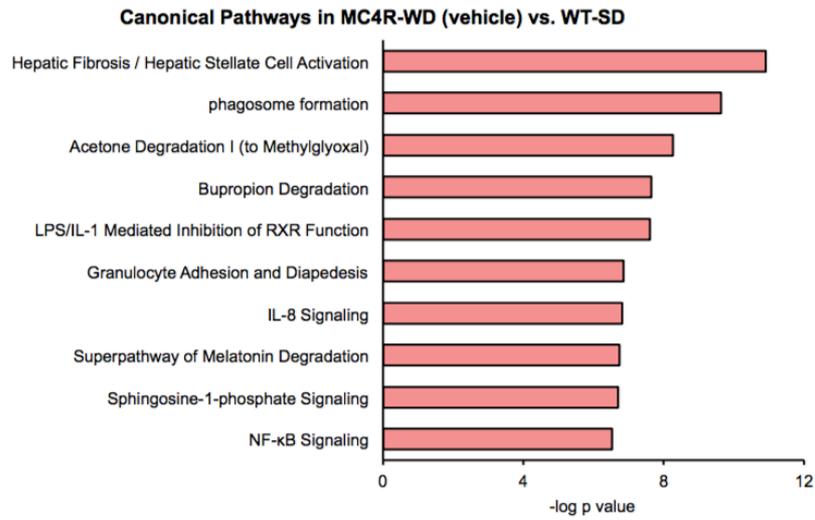
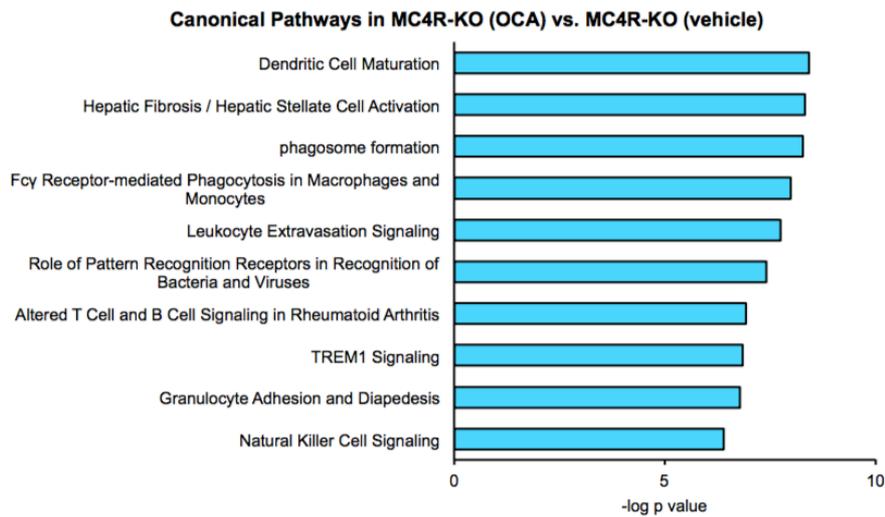
Supplementary Figure S1. Effect of OCA on tissue weight and liver triglyceride and cholesterol levels in MC4R-KO mice.

(a) Body weight and liver, subcutaneous and epididymal adipose tissue weights in MC4R-KO and WT mice. (b) Liver triglyceride (TG) and total cholesterol (TC) levels at 24 weeks. (c) Serum soluble vascular cell adhesion molecule-1 (VCAM-1) concentrations and (d) hepatic mRNA expression levels of VCAM-1 in MC4R-KO mice fed WD for 4-5 weeks or 20 weeks compared to age-matched WT mice fed SD. V, Vehicle. ^{##} $p < 0.01$; ^{*} $p < 0.05$, ^{**} $p < 0.01$ vs. the vehicle-treated MC4R-WD group.



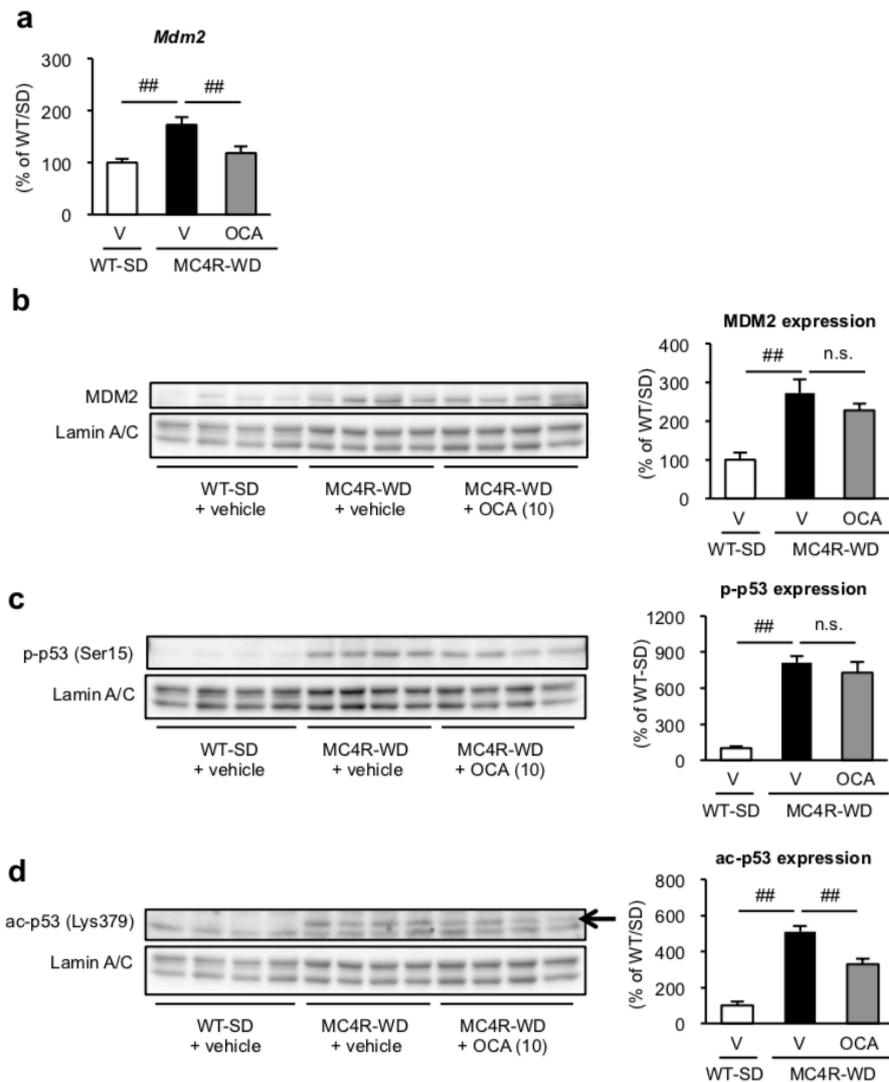
Supplementary Figure S2. Hepatic mRNA expression of FXR-regulated genes after 24-week OCA treatment.

Hepatic mRNA expression levels of ATP-binding cassette subfamily B member 11 (*Abcb11*), sterol regulatory element binding protein 1c (*Srebf1c*), cytochrome P450 7A1 (*Cyp7a1*) and cytochrome P450 8b1 (*Cyp8b1*) after treatment with OCA for 24 weeks. ## $p < 0.01$; * $p < 0.01$ vs. vehicle-treated MC4R-WD group. Vehicle-treated WT-SD group, $n = 7$; vehicle-treated MC4R-WD group, $n = 10$; 3 and 10 mg/kg of OCA-treated MC4R-WD group, $n = 10$ and $n = 9$, respectively.

a**b**

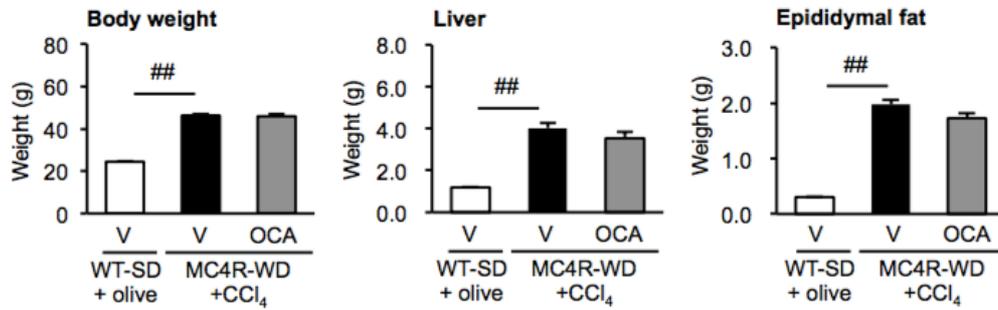
Supplementary Figure S3. Effect of OCA on canonical pathways in NASH.

Ingenuity Pathway Analysis of top 10 canonical pathways for (a) vehicle-treated WT-SD group vs. vehicle-treated MC4R-WD group and (b) vehicle-treated MC4R-WD group vs. OCA-treated MC4R-WD group.



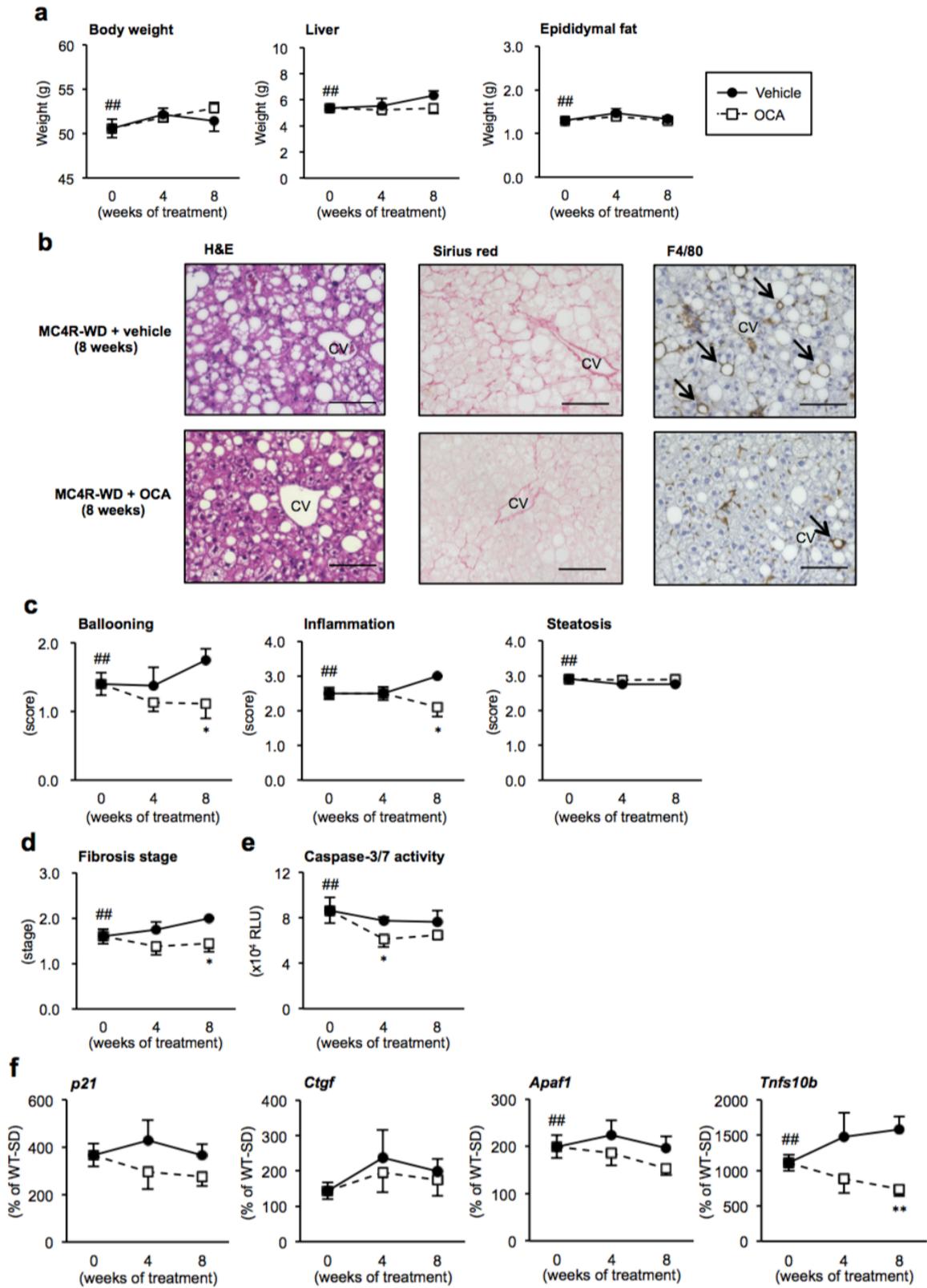
Supplementary Figure S4. Effect of OCA on modification status of p53 in the liver.

(a) Hepatic mRNA expression levels of *Mdm2*. Vehicle-treated WT-SD group, $n = 7$; vehicle-treated MC4R-WD group, $n = 10$; 10 mg/kg OCA-treated MC4R-WD group, $n = 9$. Nuclear protein levels of MDM2 (b), phospho-p53 (Ser15) (c) and acetyl-p53 (Lys379) (d). $n = 4$. $## p < 0.01$, n.s., not significant.



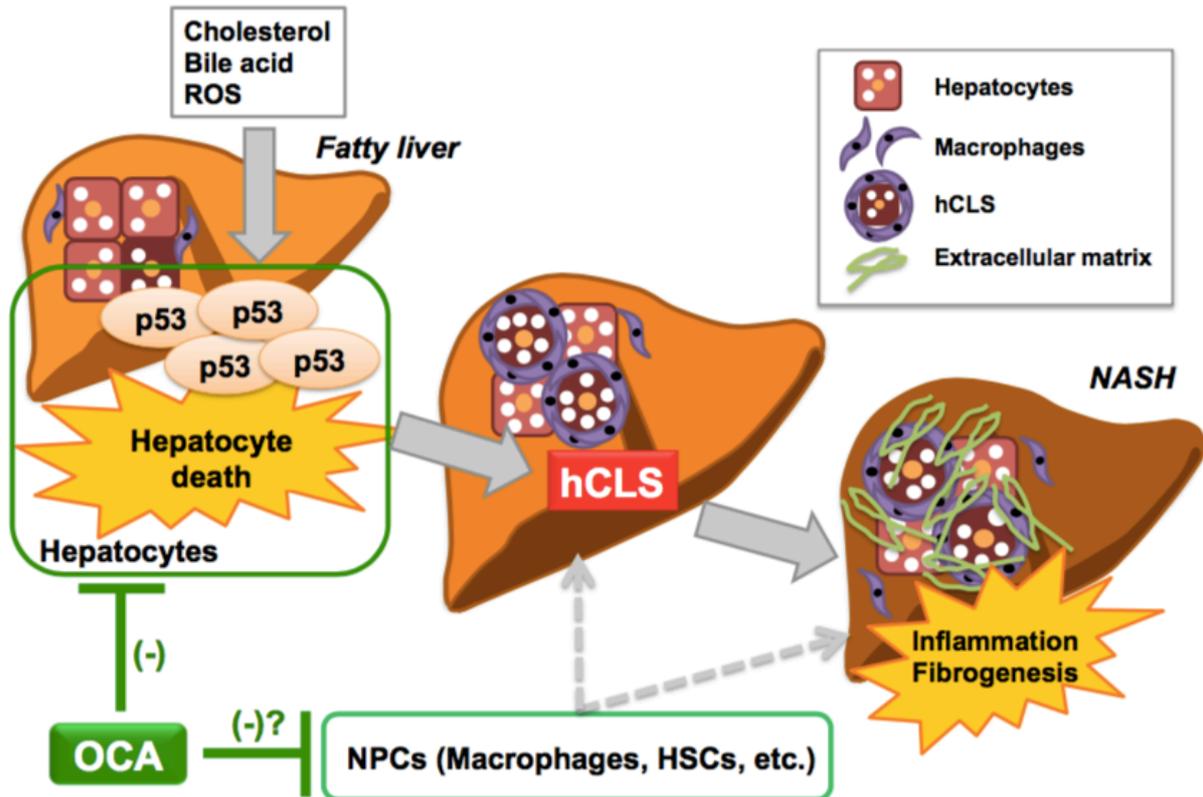
Supplementary Figure S5. Effect of OCA on body and tissue weights in the inducible NASH model.

Body, liver and epididymal fat weights 7 days after CCl₄ injection. V, Vehicle. #*p* < 0.05, ##*p* < 0.01. Vehicle-treated WT-SD group with olive oil injection, *n* = 10; vehicle-treated MC4R-WD group with CCl₄ injection, *n* = 10; OCA (10 mg/kg)-treated MC4R-WD group with CCl₄ injection, *n* = 10.



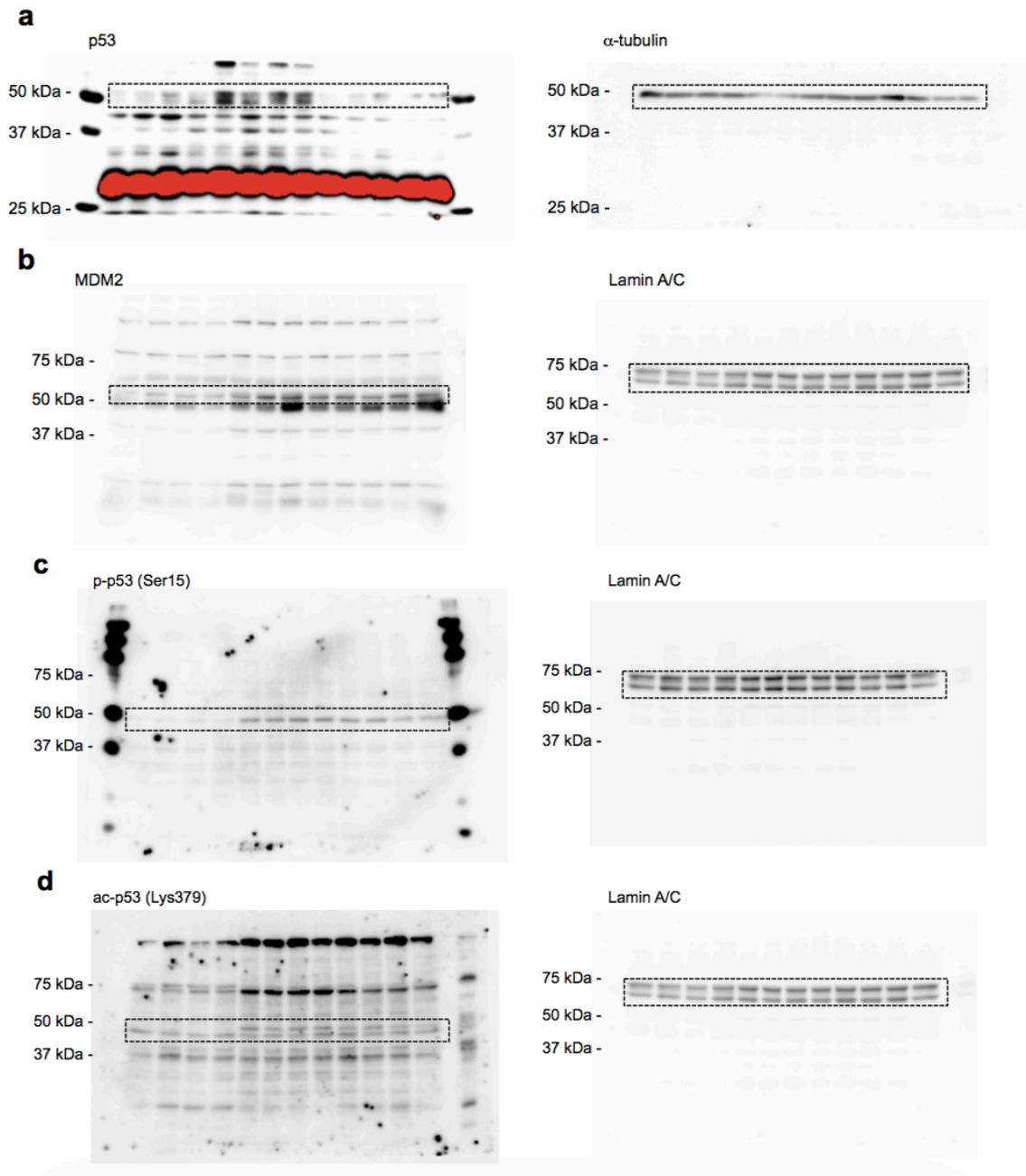
Supplementary Figure S6. Effect of OCA on the progression of NASH in MC4R-KO mice.

(a) Time course of body, liver and epididymal fat weights. (b) Representative images of Hematoxylin and eosin, Sirius red staining, and F4/80 immunostaining of the livers from WD-fed MC4R-KO mice treated with OCA for 8 weeks. Arrows, hCLS. Scale bars, 100 μ m. CV, central veins. Time course of ballooning, inflammation and steatosis scores (c), and fibrosis stage in the liver (d). Caspase-3/7 activity in the liver (e). (f) Time course of hepatic mRNA expression levels of *p21*, *Ctgf*, *Apaf1* and *Tnfs10b* after 4 or 8 weeks treatment with OCA. $^{##}p < 0.01$ MC4R-WD (pre-treatment) vs. WT-SD. $^{*}p < 0.05$, $^{**}p < 0.01$ OCA-treated MC4R-WD group vs. vehicle-treated MC4R-WD group at each point. Open square, vehicle-treated MC4R-WD group (pretreatment, $n = 10$; 4w and 8w, $n = 8$); closed circle, OCA-treated MC4R-WD group (4w, $n = 8$; 8w $n = 9$). WT-SD, $n = 7$.



Supplementary Figure S7. Graphical summary

hCLS, hepatic crown-like structure; HSC, hepatic stellate cell; NASH, non-alcoholic steatohepatitis; NPC, non-parenchymal cell; OCA, obeticholic acid; ROS, reactive oxygen species.



Supplementary Figure S8. Uncropped Western blots for Fig. 3c and Supplementary Fig. S4.

Boxes indicate cropped areas.