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Supplemental information

Therapeutic silencing of SMOC2

prevents kidney function loss in mouse

model of chronic kidney disease

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SUPPLEMENTARY TABLES

ANTIBODY	DILUTION	IDENTIFIER	CAT. NO
SMOC2	1:250	Santa Cruz Biotechnology Inc.	sc-67396
α-SMA	1:500	Sigma-Aldrich	A2547
COLLAGEN 1	1:200	Abcam	ab600408
FIBRONECTIN	1:200	Abcam	ab23750
F4/80	1:200	Abcam	ab6640
PDGFRβ	1:200	Abcam	ab32570

Supplemental table 1 (Table S1), related to STAR Methods, section of *In vivo* immunofluorescence.

List of primary antibodies for immunostaining.

ANTIBODY	DILUTION	IDENTIFIER	CAT. NO
P-P42/44 (T202/Y204)	1:1000	Cell Signaling	4377S
P-P38 (T180/Y182)	1:1000	Cell Signaling	9211S
P-JNK	1:1000	Cell Signaling	9251S
P-AKT (S473)	1:1000	Cell Signaling	9271S
P-MTOR	1:1000	Cell Signaling	2971S
P-SMAD 2 (S465/467)/3 (S423/425)	1:1000	Cell Signaling	13820S
SMOC2	1:1000	Santa Cruz Biotechnology Inc.	sc-67396
BCN1 (E-8)	1:1000	Santa Cruz Biotechnology Inc.	sc-48341
COLLAGEN 1	1:1000	Abcam	ab600408
FIBRONECTIN	1:1000	Abcam	ab23750
GAPDH	1:1000	Abcam	ab181602
α-SMA	1:5000	Sigma-Aldrich	A2547

Supplemental table 2 (Table S2), related to STAR Methods, section of Immunoblotting. List of primary antibodies for Western Blot.

INHIBITOR	INHIBITOR OF	IDENTIFIER	CAT. NO
ATN-161 (AC-PHSCN-NH2)	Integrin α 5	Selleckchem	S8454
CHLOROQUINE DIPHOSPHATE SALT	Lysosome	Sigma	C 6628-50G
GW788388	ALK5/TGFβRII	Selleckchem	S2750
LY2109761	TGFβRII	Selleckchem	S2704
LY294002	Akt	GIBCO	PHZ1144
3-МА	Autophagy	Millipore Sigma	189490-100MG
MHY1485	mTOR Activator	Millipore Sigma	5005540001
SB203580	P38	Selleckchem	S1076
SB273005	Integrin	Selleckchem	S7540
SB431542	ALK5	Selleckchem	S1067
SP600125	JNK	Selleckchem	S1460
U0126	P42/44	EMD Millipore	19-147

Supplemental table 3 (Table S3), related to STAR Methods, section of *In vitro* inhibitional experiments. List of inhibitors.

Collagen 1 Forward Reverse	5'- GAG CGG AGA GTA CTG GAT CG-3' 5'- GTT CGG GCT GAT GTA CCA GT-3'
Fibronectin Forward Reverse	5'- CCA CCC CCA TAA GGC ATA GG-3' 5'- GTA GGG GTC AAA GCA CGA GTC ATC-3'
<i>α-SMA</i> Forward Reverse	5'- ATC ATG CGT CTG GAC TTG G-3' 5'- AAT AGC CA GCT Cag TCA GG-3'
<i>PDGFRβ</i> Forward Reverse	5'- CAC CTT CTC CAG TGT GCT GA-3' 5'- GGA GTC CAT AGG GAG GAA GC-3'
<i>IL-1β</i> Forward Reverse	5'- GCA CTA CAG GCT CCG AGA TGA AC-3' 5'- TTG TCG TTG CTT GGT TCT CCT TGT-3'
IL-6 Forward Reverse	5'- GAG GAT ACC ACT CCC AAC AGA CC-3' 5'- AAG TGC ATA ATC GTT GTT CAT ACA-3'
<i>TNFα</i> Forward Reverse	5'- TAG CCA GGA GGG AGA ACA GA-3' 5'- TTT TCT GGA GGG AGA TGT GG-3'
<i>TGFβ1</i> Forward Reverse	5'- GAA GGA CCT GGG TTG GAA GTG G-3' 5'- CGT AGT AGA CGA TGG GCA GTG G-3'
GAPDH Forward Reverse	5'- ATC TTG GGC TAC ACT GAG GA-3' 5'- CAG GAA ATG AGC TTG ACA AAG-3'

Supplemental table 4 (Table S4), related to STAR Methods, section of Quantitative PCR. List of primer sequences.

SUPPLEMENTARY FIGURES AND LEGENDS



Supplemental Figure 1 (Fig S1), related to Figure 1. Mouse chronic kidney disease (CKD) model. Immunofluorescent staining of different markers (20x magnification and scale bar, 50um) in cortex and medulla of kidney in mouse CKD model compare with sham as indicated (A) collagen 1; (B) α -SMA; (C) PDGFR β ; (D) F4/80. n=6/group unless otherwise stated.



Supplemental Figure 2 (Fig S2), related to Figure 3. Genes expression in whole CKD kidney tissue of SMOC2 KO compared with WT.

(A-D) Quantitative PCR for transcripts of matrix proteins, fibrotic factors and inflammatory factors.

n=6/group unless otherwise stated. Data are represented as mean +/- SEM.



Supplemental Figure 3 (Fig S3), related to Figure 4. SMOC2 siRNA delivery and distribution in mouse kidney.

(A,B) C57BI6 mice were administrated with PBS or fluorescent SMOC2 siRNA (30 mg/200ul) via retroorbital vein injection. Kidneys were harvested at 30 min, 2 h, 8 h, 24 h, 48 h and 72 h. (A) Fluorescent SMOC2 siRNA deposition in the kidney and (B) SMOC2 siRNA distribution in proximal tubules (Megalin+) and identified in endothelial cells (CD31+) and glomerular cells (PDGFR β +) SMOC2, a therapeutic target for CKD (C,D) Mice subjected to experimental CKD then received either SMOC2 siRNA (30 mg/200ul) or PBS at day 21 post nephrectomy surgery twice/week until day 43. SMOC2 depletion was confirmed at protein (C) and mRNA (D) level.

Immunofluorescence staining with 40x magnification and scale bar, 20um. n=6/group unless otherwise stated. Data are represented as mean +/- SEM.



Supplemental Figure 4 (Fig S4), related to Figure 4. Genes expression in whole CKD kidney tissue

of SMOC2 siRNA treatment compared with vehicle treatment.

(A-D) Quantitative PCR for transcripts of matrix proteins, fibrotic factor and inflammatory factors.

n=6/group unless otherwise stated. Data are represented as mean +/- SEM.



Supplemental Figure 5 (Fig S5), related to Figure 5. SMOC2 induced genes expression.

NIH/3T3 cells were incubated with SMOC2 recombinant protein for 24 hours with 1, 10, 20, 50, 100 ng/ml. Quantitative PCR for transcripts of matrix proteins, fibrotic factors and inflammatory factors (A-F). Quantification are representative of three independent experiments. *P<0.05; **P<0.01; ***P<0.001. Data are represented as mean +/- SEM.



Supplemental Figure 6 (Fig S6), related to Figure 5. Western Blots analysis of lysosomeautophagosome signaling pathways involved in SMOC2-induced kidney fibrosis.

(A-I) Ratio of target protein expression to Actin. *P<0.05; **P<0.01; ***P<0.001. Data are represented as mean +/- SEM.



Supplemental Figure 7 (Fig S7), related to Fig 6. SMOC2 induced phosphorylation of MAPKs and Akt.

NIH/3T3 cells were treated with either SMOC2 (100 ng/ml) at 5 min, 15 min, 30 min, 1 h, 4 h and 24 h, or TGFβ1 (10 ng/ml). Cell lysates were performed by Western Blotting analysis and showed phosphorylation of MAPKs (p-P42/44, p-P38, p-JNK) and Akt (p-Akt), and control of Actin expression showed same as in Fig 5A. Blots are representative of three independent experiments.



Supplemental Figure 8 (Fig S8), related to Figure 6. SMOC2 trigged cell differentiation.

NIH/3T3 cells were co-incubated with SMOC2 (100 ng/ml, 24 h) and lysosome inhibitors of Chloroquine (10 uM, 24 h), Bafilomycin (10 uM, 24 h) and Integrin α 5 inhibitor of ATN-161 (10 uM, 24 h), cell differentiation was represented by F-Actin (Green) staining. Immunofluorescence staining with 40x magnification and scale bar, 20um. Images are representative of three independent experiments.