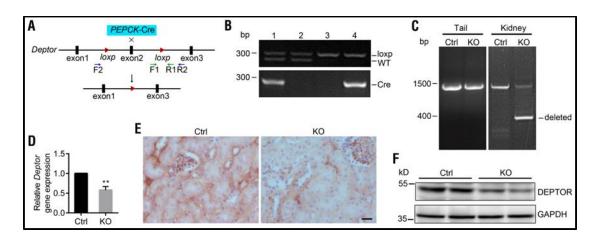
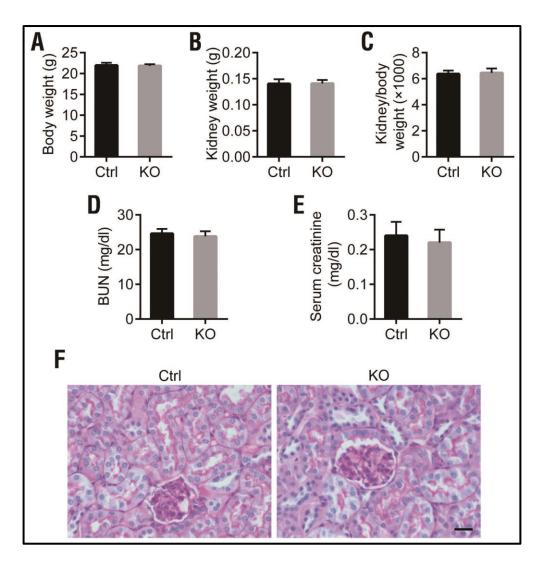
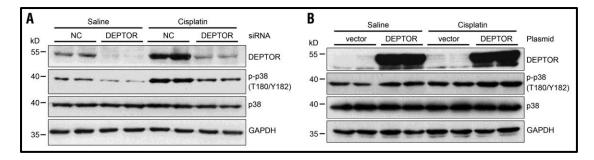
Supplemental Materials



Supplemental Figure 1. Generation of mice with *DEPTOR* specifically deleted in renal proximal tubules. (A) Strategy for the production of *DEPTOR* cKO mice. *DEPTOR* flox mice were crossed with *PEPCK*-Cre mice to produce *DEPTOR* KO mice, in which the second exon of *DEPTOR* gene was deleted. F1/F2: forward primers for PCR identification, R1/R2: reverse primers for PCR identification. (B) Genotyping the F2 offspring by PCR with primers F1/R1 after mating the transgenic mice. M: DNA marker. (C) Identification of gene deletion in renal genomic DNA by PCR with primers F2/R2. Tail genomic DNA was used as the endogenous control. (D) qPCR analysis showed DEPTOR mRNA levels in the renal cortex of KO mice were obviously reduced compared with control mice. Data are expressed as mean values \pm SEM (n = 5). **P < 0.01. (E) IHC staining showed DEPTOR expression was significantly reduced in renal proximal tubules of *DEPTOR* KO mice compared with control mice. Scale bar = 20 µm. (F) Western blotting analysis showed DEPTOR expression in the renal cortex of KO mice was decreased.



Supplemental Figure 2. Normal renal functions and histology in *DEPTOR* KO mice under physiological conditions. No significant difference was observed in bodyweight (A), kidney weight (B), kidney/body weight ratio (C), BUN (D), or serum creatinine (E) between *DEPTOR* KO and control (Ctrl) mice. BUN and creatinine were measured with the IDEXX Catalyst Chemistry Analyzer. (F) Kidney histology of paired mice detected with PAS staining. Mice were 8-12 weeks old. Scale bar = 20 μ m. Bars indicate mean values \pm SEM. n = 8.



Supplemental Figure 3. DEPTOR regulates p38 MAPK activity in HEK293 cells.

(A) Immunoblotting showed that DEPTOR knockdown by siRNA inhibited p38 MAPK activity in HEK293 cells, with or without cisplatin treatment. In the cisplatin group, cells were incubated with 20 μ M cisplatin for 24 h, the same below. (B) Immunoblotting showed that DEPTOR overexpression enhanced p38 MAPK activity in HEK293 cells.

Supplemental Table 1. Primers used for experiments

Gene	Forward primers(5'-3')	Reverse primers(5'-3')	
DEPTOR -	GGAAGTGAAGACCTGTGAAGA	CAGTCTAGAAAAGCAGATAACC	
F1/R1	TAAGAG	CAAGTA	
DEPTOR -	CGTCGAGAAGTTCCTATTCCGA	GTGTGCCATTGAAATTTGTTACC	
F2/R2	AG	AT	
PEPCK-Cre	GCATAACCAGTGAAACAGCATT	GGACATGTTCAGGGATCGCCAG	
	GCTG	GCG	
DEPTOR-	AGCAGAGAGAGCTGGAACGC	CAGAGGCCTCCTTATGTTCA	
qPCR			
Mouse Kim1	ATGCCCATCTTCTGCTTGTCA	CCTTGTAGTTGTGGGTCTTCT	
Mouse NGAL	TGGCCCTGAGTGTCATGTG	CTCTTGTAGCTCATAGATGGTGC	
Mouse TNFa	CCTGTAGCCCACGTCGTAG	GGGAGTAGACAAGGTACAACCC	
Mouse Gapdh	TGTGTCCGTCGTGGATCTGA	TTGCTGTTGAAGTCGCAGGA	
Human TNFα	CCTCTCTAATCAGCCCTCTG	GAGGACCTGGGAGTAGATGAG	
Human Gapdh	ACAACTTTGGTATCGTGGAAGG	GCCATCACGCCACAGTTTC	

Supplemental Table 2. Antibodies used for experiments

Vendor	Catalog	Application	Working dilution
	number		for WB or (IF/IHC)
Cell Signaling	11816	WB	1:2000
Technology			
Santa Cruz	sc-398169	WB,IHC	1:2000 (1:100)
Biotechnology			
Cell Signaling	4858	WB, IHC,IF	1:2000 (1:200)
Technology			
Santa Cruz	sc-74459	WB	1:2000
Biotechnology			
Santa Cruz	sc-7985-R	WB	1:2000
Biotechnology			
Cell Signaling	2920	WB	1:2000
Technology			
Cell Signaling	4511	WB	1:2000
Technology			
Cell Signaling	8690	WB	1:2000
Technology			
Cell Signaling	9129	IF	1:200
Technology			
Cell Signaling	9532	WB	1:500
Technology			
Sungene	KM9002T	WB	1:5000
C 11 C: 1:	4270	WD	1 2000
	4370	WB	1:2000
	4605	WD	1 2000
	4695	WB	1:2000
1	4660	IVD	1 2000
	4668	WB	1:2000
	0.600	IIID	1.2000
	8690	WB	1:2000
	12107	****	1.2000
Abclonal	A3185	WB	1:2000
Santa Cruz	sc-377009	IF	1:50
Biotechnology			
	Cell Signaling Technology Santa Cruz Biotechnology Cell Signaling Technology Santa Cruz Biotechnology Santa Cruz Biotechnology Cell Signaling Technology Cell Signaling	Cell Signaling Technology Santa Cruz Biotechnology Cell Signaling Technology Santa Cruz Biotechnology Santa Cruz Biotechnology Santa Cruz Biotechnology Cell Signaling Technology Abclonal A3185	Cell Signaling Technology11816WBSanta Cruz Biotechnologysc-398169WB,IHCCell Signaling Technology4858WB, IHC,IFTechnologyWBSanta Cruz Biotechnologysc-74459WBSanta Cruz Biotechnologysc-7985-RWBCell Signaling Technology2920WBCell Signaling Technology4511WBCell Signaling Technology9129IFCell Signaling Technology9532WBCell Signaling Technology4370WBCell Signaling Technology4695WBCell Signaling Technology4668WBCell Signaling Technology4668WBCell Signaling Technology4668WBTechnologyCell Signaling Technology4668WBTechnologyCell Signaling Technology4668WBTechnologyCell Signaling Technology4668WBSanta Cruzsc-377009IF

Supplemental Table 3. List of 46 genes with significant alteration in the mRNA level in the renal cortex of the *DEPTOR*-cKO mice *vs* that of the control mice under physiological condition by transcriptomic sequencing and screening assay.

Gene	log ₂ (Fold change)	Description
Cyp2b10	-4.01	cytochrome P450, family 2, subfamily b,
Сур2010	-4.01	polypeptide 10
Ccl28	-3.68	chemokine (C-C motif) ligand 28
Rnf180	-3.41	ring finger protein 180
Rn7sk	-3.25	RNA, 7SK, nuclear
Rmrp	-3.02	RNA component of mitochondrial RNAase P
Hsd11b1	-2.81	hydroxysteroid 11-beta dehydrogenase 1
Ctxn3	-2.80	cortexin 3
Slco1a1	-2.73	solute carrier organic anion transporter family, member 1a1
Srd5a2	-2.61	steroid 5 alpha-reductase 2
Acsm3	-2.59	acyl-CoA synthetase medium-chain family member 3
Cyp4a12a	-2.56	cytochrome P450, family 4, subfamily a, polypeptide 12a
Rpph1	-2.48	ribonuclease P RNA component H1
Hapln1	-2.36	hyaluronan and proteoglycan link protein 1
Kif20b	-2.34	kinesin family member 20B
Eri2	-2.32	exoribonuclease 2
Pik3c2g	-2.29	phosphatidylinositol 3-kinase, C2 domain
F1K3C2g		containing, gamma polypeptide
Mir6236	-2.25	microRNA 6236
Hprt	-2.17	hypoxanthine guanine phosphoribosyl transferase
Zpld1	-2.16	zona pellucida like domain containing 1
Mfsd2a	-1.97	major facilitator superfamily domain containing 2A
Odc1	-1.96	ornithine decarboxylase, structural 1
Frzb	-1.93	frizzled-related protein
Ces2b	-1.87	carboxyesterase 2B
Slc22a7	-1.80	solute carrier family 22 (organic anion transporter), member 7
Lipo1	-1.72	lipase, member O1
Nr4a1	-1.70	nuclear receptor subfamily 4, group A, member 1
Azgp1	-1.66	alpha-2-glycoprotein 1, zinc
I134	-1.66	interleukin 34
Egr1	-1.63	early growth response 1
Cxcl12	-1.57	chemokine (C-X-C motif) ligand 12
Ttbk1	-1.55	tau tubulin kinase 1
Cd36	-1.54	CD36 antigen
Ces2c	-1.35	carboxylesterase 2C
Tnfsf10	-1.26	tumor necrosis factor (ligand) superfamily, member 10

B4galt5	-1.26	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 5
C1qtnf3	-1.20	C1q and tumor necrosis factor related protein 3
Scd1	1.35	stearoyl-Coenzyme A desaturase 1
Apoe	1.39	apolipoprotein E
Pdk4	1.49	pyruvate dehydrogenase kinase, isoenzyme 4
Cish	1.49	cytokine inducible SH2-containing protein
Tma7	1.88	translational machinery associated 7
B4galnt1	1.94	beta-1,4-N-acetyl-galactosaminyl transferase 1
Kynu	2.30	kynureninase (L-kynurenine hydrolase)
Cyp4a14	2.76	cytochrome P450, family 4, subfamily a, polypeptide 14
Prlr	3.14	prolactin receptor
Gpr112	3.70	adhesion G protein-coupled receptor G4

Supplemental Figure 4. Network for protein interactions among the 46 genes.

