

SUPPLEMENTAL MATERIAL

TEAD1 Prevents Necroptosis And Inflammation In Cisplatin-Induced Acute Kidney Injury Through Maintaining Mitochondrial Function

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Supplementary Figure 1. Generation of proximal-tubule specific TEAD1 knockout mice (TEAD1^{PKO}). (a) Schematic representation of genetic crosses to conditionally delete TEAD1 in proximal tubules of kidneys (b) Representative photomicrographs of kidney sections immunostained for TEAD1 (brown) and counterstained with hematoxylin (blue) in TEAD1^{CON} and TEAD1^{PKO} mice. Proximal tubules are indicated by black arrows. Scale bar: 50 μ M.

a

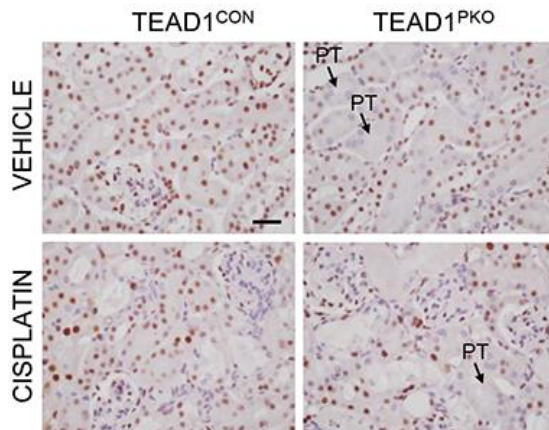
♂ PEPCK-Cre^{-/-}TEAD1^{flox/flox} X ♀ PEPCK-Cre^{+/-}TEAD1^{flox/flox}



PEPCK-Cre^{-/-}TEAD1^{flox/flox} (proximal tubule specific KO; TEAD1^{PKO}) (~ 50 % offspring)

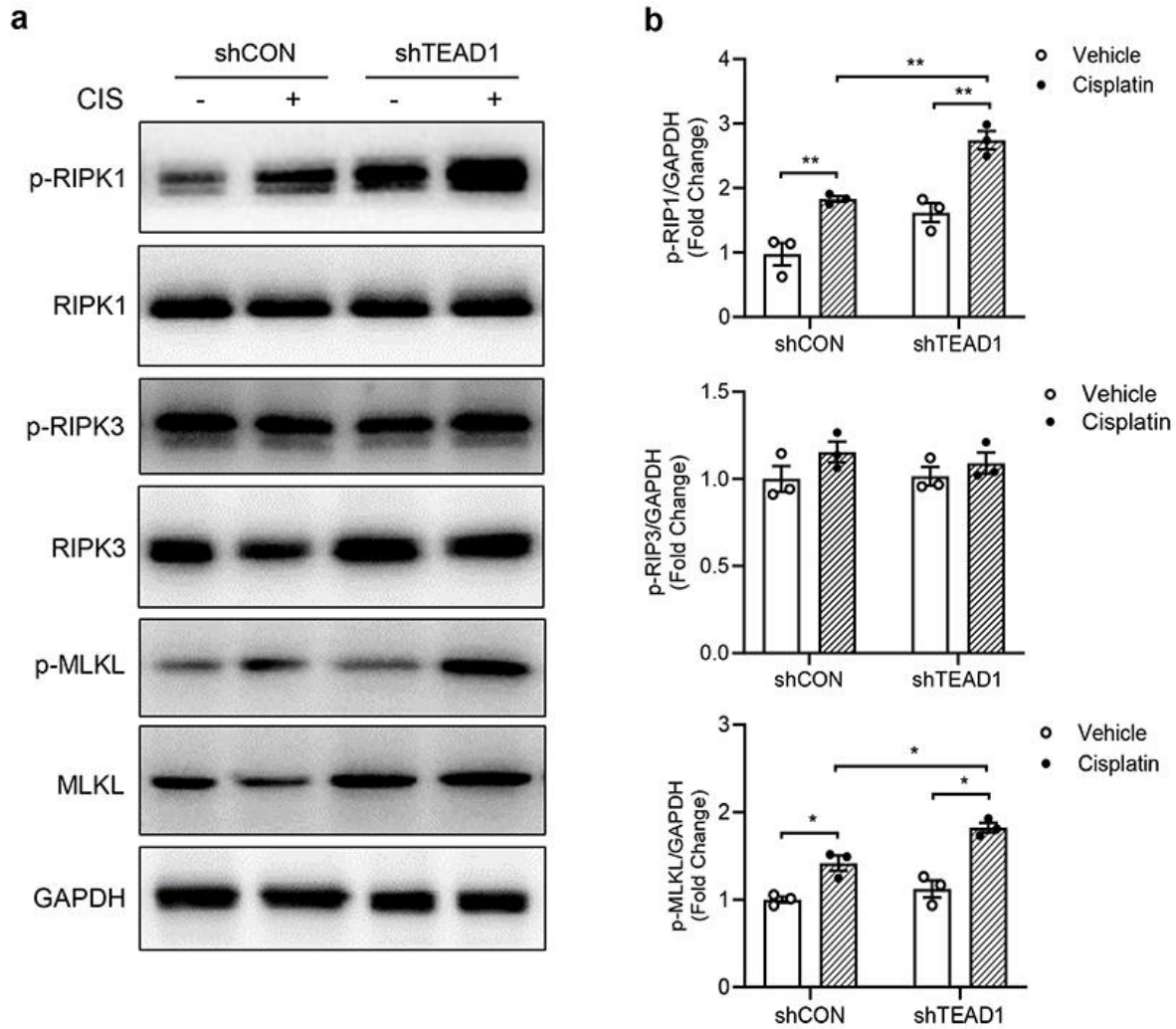
PEPCK-Cre^{+/-}TEAD1^{flox/flox} (WT control; TEAD1^{CON}) (~ 50 % offspring)

b



Supplementary Figure 2. Knockdown of TEAD1 in TCMK-1 cells promote necroptosis. (a)

Representative Western blots show phosphorylated RIP1, RIP1, phosphorylated RIP3, RIP3, phosphorylated MLKL and MLKL protein expression in TCMK-1 cells transduced with shCON or shTEAD1 and treated with 10 μ M cisplatin or vehicle for 24 h (b) Quantitative analysis of phosphorylated RIP1, phosphorylated RIP3, and phosphorylated MLKL protein expression in TCMK-1 cells treated with 10 μ M cisplatin or vehicle for 24 h. * $P < 0.05$ and ** $P < 0.01$, $n = 3$ per group.



Supplementary Table 1. Sequence of primers used for genotyping and real time RT-PCR.

Primer	Forward primer	Reverse primer
TEAD1 genotyping	GCCTTCTGAGTGCTAGCATTAAAGG	AAGGCAGACTCCTTCATTGGATGG
PEPCK ^{Cre} genotyping	ACCTGAAGATGTTTCGCGATTATCT	ACCGTCAGTACGTGAGATATCTT
IL-1 β	CTTCAGGCAGGCAGTATCACTCAT	TCTAATGGGAACGTCACACACCAG
IL-6	AGGATAACCACTCCCAACAGACCTG	CTGCAAGTGCATCATCGTTGTTCA
MCP-1	TCACCTGCTGCTACTCATTACCA	TACAGCTTCTTTGGGACACCTGCT
TNF α	CATGAGCACAGAAAGCATGATCCG	AAGCAGGAATGAGAAGAGGCTGAG
iNOS	GGCAGCCTGTGAGACCTTTG	GCATTGGAAGTGAAGCGTTTC
MLKL	ACCCTTCAGAGGCACAACAC	TGTCATTGGATTCCGGTGGGG
RIP3	GAAGACACGGCACTCCTTGTA	CTTGAGGCAGTAGTTCTTGGTGG
PGC1 α	AACTTGCTAGCGGTCCTCA	TGGCTGGTGCCAGTAAGAG
GAPDH	CCAATGTGTCCGTCGCGTGGATCT	GTTGAAGTCGCAGGAGACAACC