USP25 Regulates KEAP1-NRF2 Anti-Oxidation Axis and Its Inactivation

Protects Acetaminophen-Induced Liver Injury in Male Mice

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Su	pplen	ientarv	Table	1. Anti	body	info	mation.

Antibody	Cat No.	Supplier	Purpose
USP25	A7975	ABclonal	1:1000 for WB; 2 µg for IP
KEAP1	10503-2-AP	Proteintech	1:2000 for WB; 2 µg for IP
GAPDH	60004-1-Ig	Proteintech	1:5000 for WB
β-Actin	66009-1-Ig	Proteintech	1:5000 for WB
Lamin B1	12987-1-AP	Proteintech	1:1000 for WB
CYP2E1	19937-1-AP	Proteintech	1:3000 for WB
USP28	17707-1-AP	Proteintech	1:1000 for WB
NRF2	Sc-722	Santa Cruz Biotechnology	1:1000 for WB
UB	Sc-8017	Santa Cruz Biotechnology	1:200 for WB
NRF2	ab62352	Abcam	1:1000 for WB
MKK4	9152S	CST	1:1000 for WB
P-MKK4 (Ser257)	4514P	CST	1:1000 for WB
JNK	ab208035	Abcam	1:2000 for WB
P-JNK (Thr183/Tyr185)	4668P	CST	1:1000 for WB
Rabbit IgG	3900S	CST	2 µg for IP
НА	H6908	Sigma-Aldrich	1:2000 for WB
FLAG	F1804	Sigma-Aldrich	1:2000 for WB
Goat anti mouse (H+L)	115-035-003	Jackson ImmunoResearch Laboratories	1:5000 for WB
Goat anti rabbit (H+L)	111-035-003	Jackson ImmunoResearch Laboratories	1:5000 for WB

Gene	Sequence
β-Actin-F	5-GGCTGTATTCCCCTCCATCG-3
β-Actin-R	5-CCAGTTGGTAACAATGCCATGT-3
Keap1-F	5-TGCCCCTGTGGTCAAAGTG-3
Keap1-R	5-GGTTCGGTTACCGTCCTGC-3
<i>Nrf2</i> -F	5-TCTTGGAGTAAGTCGAGAAGTGT-3
<i>Nrf2</i> -R	5-GTTGAAACTGAGCGAAAAAGGC-3
Nqo1-F	5-AGGATGGGAGGTACTCGAATC-3
Nqo1-R	5-AGGCGTCCTTCCTTATATGCTA-3
Gclc-F	5-GGGGTGACGAGGTGGAGTA-3
Gclc-R	5-GTTGGGGTTTGTCCTCTCCC-3
Gclm-F	5-AGGAGCTTCGGGACTGTATCC-3
Gclm-R	5-GGGACATGGTGCATTCCAAAA-3
Usp25-F	5-CAGAAGCACCAGCAGACATTT-3
Usp25-R	5-TGGCATTCTTTGCAGTGAGGA-3

Supplementary Table 2. Primer sequences used for RT-PCR.

Supplementary Table 3. The sequences used for shRNA.

Gene	Sequence
negative control	5-TTCTCCGAACGTGTCACGT-3
shUSP25-1 (human)	5-TCGATGGTGTTCCTACCT-3
shUSP25-2 (human)	5-GGGAGTACTTGAAGGTAAA-3
shUsp25-1 (mouse)	5-GCACAGAAATAGAGAAATA-3
shUsp25-2 (mouse)	5-GAAGAAACGCTCCGAGTGA-3

Supplementary Table 4. Cloning primers used in the present study.

Gene	Sequence $(5' \rightarrow 3')$
USP25-SIM/UIM-F	5-GCTCTAGAATGGACTACAAAGACGATGACGACAAGACCGTGGAGCAGAACGTGCT-3
USP25-SIM/UIM-R	5-ATAAGAATGCGGCCGCTAAACTAT TTAATCCCTCCAAACTTCTGT-3
USP25-USP-F	5-GCTCTAGAATGGACTACAAAGACGATGACGACAAGTCTCGAAACCCTTATGATAG-3
USP25-USP-R	5-ATAAGAATGCGGCCGCTAAACTATTTACTGAGACGCTAAAAGCTT-3
USP25-CTD-F	5-CGGAATTCATGGACTACAAAGACGATGACGACAAGAAATTGAGAGAGA
USP25-CTD-R	5-ATAAGAATGCGGCCGCTAAACTATTTATCTTCCATCAGCAGGAGT-3
KEAP1-BTB-F	5-GCTCTAGAATGGACTACAAAGACGATGACGACAAGCAGCCAGATCCCAGGCCTAG-3
KEPA1-BTB-R	5-ATAAGAATGCGGCCGCTAAACTATTTAGACAGCACCGTTCATGAC-3
KEAP1-IVR-F	5-CCGGAATTCCGGATGGACTACAAAGACGATGACGACAAGGGTGCTGTCATGTACCAGAT-3
KEAP1-IVR-R	5-ATAAGAATGCGGCCGCTAAACTATTTAGTGCAGGGTGAGCTCCTC-3
KEAP1-DGR-F	5-CCGGAATTCCGGATGGACTACAAAGACGATGACGACAAGAAGCCCACGCAGGTGATGCC-3
KEAP1-DGR-R	5-ATAAGAATGCGGCCGCTAAACTATTCAACAGGTACAGTTCTGCTG-3



Supplementary Fig. 1. Induction of liver injury by APAP administration in mice.

(a) Serum ALT and AST levels in control and APAP-treated mice. Male C57BL/6 mice were fasted for 14 h and then treated with APAP (300 mg/kg, i.p.) for 0, 6, and 24 h. At each time point, 6-7 animals were sacrificed, and the blood and liver samples were collected for analysis. 0 h: n = 6 mice, 6 h: n = 7 mice, 24 h: n = 6 mice.

(b) Hematoxylin and eosin staining of the liver sections of the mice in (a). Necrotic areas were encircled, and the percent of necrotic areas per view field was calculated and plotted. 0 h: n = 6 mice, 6 h: n = 7 mice, 24 h: n = 6 mice. Scale bar, 200 µm.

Error bars denote SEM. Two-tailed student's t tests analysis (a and b). Source data are provided as a Source Data file.



Supplementary Fig. 2. The status of *Usp25* does not affect other aspects of APAP-induced responses in the liver.

(a) Western blotting analysis of the indicated proteins in the liver of $Usp25^{+/+}$ or $Usp25^{-/-}$ mice treated with APAP (300mg/kg, i.p., n = 3-6 per group) for 0, 6 or 24 h.

(b) Western blotting analysis of the protein levels of CYP2E1 in the liver samples from (a). (c) HPLC coupled mass spectrometry analysis of APAP-CYS concentration in the liver samples from (a). n = 5 mice per group. Error bars denote SEM. Statistical significance was analyzed with two-tailed student's t tests analysis. Source data are provided as a Source Data file.



HepG2

Supplementary Figure 3

Supplementary Fig. 3. The loss of *Usp28* does not provide protection against APAP-induced liver injury.

(a) $Usp28^{+/+}$ and $Usp28^{-/-}$ mice were treated with APAP (300mg/kg, i.p., n = 6 per group) after 14 h fasting. 24 h after APAP administration, the mice were sacrificed and blood and liver samples were collected for analysis.

(b) Serum ALT and AST levels. n = 6 mice per group.

(c) Hematoxylin and eosin staining of the liver sections of the mice in (a). Necrotic areas were encircled, and the percent of necrotic areas per view field was calculated and plotted. n = 6 mice per group. Scale bar, 200 µm.

(d) Quantification of *Keap1*, *Nrf2* and NRF2 target gene expression via qPCR in the liver from the mice in (a). n = 6 mice per group.

(e) Western blotting analysis of liver proteins from the mice sacrificed at 24 h in (a). The bands with a similar molecular weight to USP28 in the $Usp28^{-/-}$ samples are non-specific.

(f) Western blotting analysis of the indicated proteins in HepG2 cells depleted of USP28 (n = 3 biologically independent experiments).

(g) Western blotting analysis of the ubiquitinated species of endogenous KEAP1 in HepG2 cells depleted of *USP28*. The cells were treated with MG132 (20 μ M) for 6 h before harvesting for analysis (n = 2 biologically independent experiments). The numbers are normalized relative levels of KEAP1 ubiquitination.

Error bars denote SEM. Statistical significance was analyzed with two-tailed student's t tests analysis. Source data are provided as a Source Data file.



Supplementary Fig. 4. USP25 is a deubiquitinase for KEAP1.

(a, b) Western blotting analysis of Keap1 in HepG2 (a) and AML12 (b) cells depleted of *USP25* expression with shRNAs (n = 3 biologically independent experiments).

(c) Determination of the ubiquitination levels of KEAP1 in HEK293T cells depleted of *USP25* expression. FLAG- KEAP1 and siRNAs were co-transfected for 48 h. The cells were treated with MG132 (20 μ M) for 6 h before harvesting for analysis (n = 2 biologically independent experiments).

(d) Determination of the ubiquitination levels of KEAP1 in HEK293T cells overexpressing *USP25* or *USP25*^{C178S}. The plasmids for the expression of FLAG- KEAP1, WT USP25 or USP25-C178S were co-transfected for 48 h. The cells were treated with MG132 (20 μ M) for 6 h before harvesting for analysis (*n* = 2 biologically independent experiments). Source data are provided as a Source Data file.



Supplementary Figure 5

Supplementary Fig. 5. CT1113 treatment could not bring additional protection in *Usp25*-deficient mice.

(a, b) Western blotting analysis of USP25 and KEAP1 in HepG2 (a) or AML12 (b) cells treated with CT1113 for 48 h (n = 3 biologically independent experiments). The relative protein band intensities were quantified and plotted.

(c, d) $Usp25^{+/+}$ and $Usp25^{-/-}$ mice (c) or $Usp28^{+/+}$ and $Usp28^{-/-}$ mice were given vehicle or CT1113 (20 mg/kg, i.g., bid) for 48 h (and continued for the rest of the time until sacrificing), fasted for 14 h, and then given APAP injection (300 mg/kg, i.p.). The animals were sacrificed at 24 h after APAP administration. The blood and liver samples were collected for analyses of serum ALT and AST levels. Usp25 WT with vehicle: n = 5 mice, Usp25 WT with CT1113: n = 5 mice, Usp25 KO with vehicle: n = 4 mice, Usp25 KO with cT1113: n = 4 mice, Usp28 WT with vehicle: n = 4 mice, Usp28 KO with vehicle: n = 4 mice, Usp28 KO with vehicle: n = 4 mice, Usp28 KO with cT1113: n = 4 mice, Usp28 KO with cT1113: n = 5 mice.

(e) Western blotting analysis of the indicated proteins in the liver of the vehicle or CT1113 treated C57BL/6 mice injected with APAP (300 mg/kg, i.p., n = 3-6 per group) for 0, 6 or 24 h. Error bars denote SEM. Two-tailed student's t tests analysis (c and d), non-parametric tests with blue colored (c). Source data are provided as a Source Data file.



Supplementary Fig. 6. Efficacy comparison between CT1113 and RTA402.

(a) Schematic of the experiment. Male C57BL/6 mice were fasted for 14 h, given APAP (300 mg/kg body weight), and CT1113 or RTA402 1 h later for once. Both CT1113 and RTA402 were given through i.p. The animals were sacrificed at 6 h after APAP administration, 6 animals per group.

(b) Serum ALT and AST levels. n = 6 mice per group.

(c) Hematoxylin and eosin staining of the liver sections from mice in (a). Necrotic areas were encircled and quantified. n = 6 mice per group. Scale bar, 200 µm.

Error bars denote SEM. Two-tailed student's t tests analysis (b and c), non-parametric tests with blue colored (b and c). Source data are provided as a Source Data file.