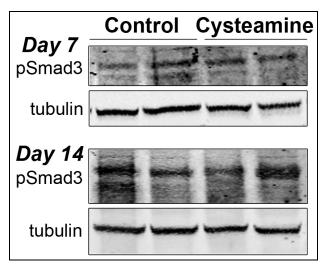
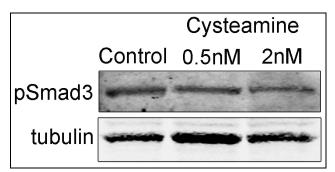


Supplemental Figure A1. Anti-fibrotic dose dependent effects of cysteamine with UUO. A pilot study was performed to determine the efficacy of three different doses of oral cysteamine 200, 400, and 600mg/kg. Mice were placed on cysteamine starting on the day of surgery and sacrificed 14 days after UUO (n=2-4/group).



Supplemental Figure A2. Cysteamine mediated attenuation of fibrosis is independent of TGF- β . Protein levels of phosphorylated Smad3 (pSmad3) measured by Western blotting and normalized to α -tubulin was performed on whole kidney homogenate in control and cysteamine treated mice (n=4/group).



Supplemental Figure A3. TGF- β activity is unaltered by cysteamine with myofibroblast activation. Normal rat kidney fibroblasts were transformed into α SMA+ myofibroblasts by exposure to transforming growth factor beta (TGF- β). Protein levels of phosphorylated Smad3 (pSmad3) measured by Western blotting and normalized to α -tubulin was performed on lysates from cells treated with cysteamine or vehicle alone (n=4/group).

SUPPLEMENTAL TABLES

Gene	Relative expression level (cysteamine:control)
TNF-α	1.31 ± 0.34
TNF-a receptor	0.99 ± 0.22
IL-1β	0.77 ± 0.36
IL-1β receptor	0.91 ± 0.30
Nox4	1.34 ± 0.35
Nox2	0.80 ± 0.21

Supplemental Table A1 No difference in pro-inflammatory and oxidative gene expression profiles after UUO with cysteamine treatment. Semi-quantitative real time RT qPCR was performed on total RNA from whole kidney homogenate from Day 14 UUO kidneys. Genes of interest were normalized to two housekeeping genes, 18S and GAPDH.

Gene	Relative expression level (cysteamine:control)		
	Day 3	Day 7	Day 14
PDGF-Rα	0.97 ± 0.48	1.00 ± 0.27	0.89 ± 0.25
PDGF-Rβ	0.97 ± 0.51	1.47 ± 0.43	0.93 ± 0.31

Supplemental Table A4 No difference in PDGF receptor gene expression profiles after UUO with cysteamine treatment. Semi-quantitative real time RT qPCR was performed on total RNA from whole kidney homogenate from control and cysteamine treated mice UUO kidneys. Genes of interest were normalized to two housekeeping genes, 18S and GAPDH.

Gene	Forward	Reverse
TNF-α	CATCTTCTCAAAATTCGAGTGACAA	TGGGAGTAGACAAGGTACAACCC
TNF-α R	GCTGACCCTCTGCTCTACGAA	GCCATCCACCACAGCATACA
IL-1β	CAACCAACAAGTGATATTCTCCATG	GATCCACACTCTCCAGCTGCA
IL-1β R	CCTCGGGATGGAAATCTGCT	CCAGATTCTGAAGGCTTGCAT
Nox2	GGGAGACTGGACGGAGGGC	ATGCGTGTCCCTGCACAGCC
Nox4	CACCAAACACAGAAGCACAAG	AGAAAGCAAAGCAGGGTATCA
PDGFR-α	ATGGACGCACGCCAGACTGTG	TCTCGACGAAGCCTTTCTCGTGG
PDGFR-β	CACCTTCTCCAGTGTGCTGA	GGAGTCCATAGGGAGGAAGC
Fibronectin	AGACTGCAGTGACCACCATTC	AATGTGTCCTTGAGAGCATAGAC
Procollagen 1	AGAAGTCTCAAGATGGTGGCCG	GGTCACGAACCACGTTAGCATC
Procollagen 3	CAGCTATGGCCCTCCTGATCTT	GTAATGTTCTGGGAGGCCCG
GAPDH	ACTTTGTCAAGCTCATTTCC	TGCAGCGAACTTTATTGATG
18S	GGTGAAATTCTTGGACCGGC	GACTTTGGTTTCCCGGAAGC

Supplemental Table A3: qPCR primer sequences