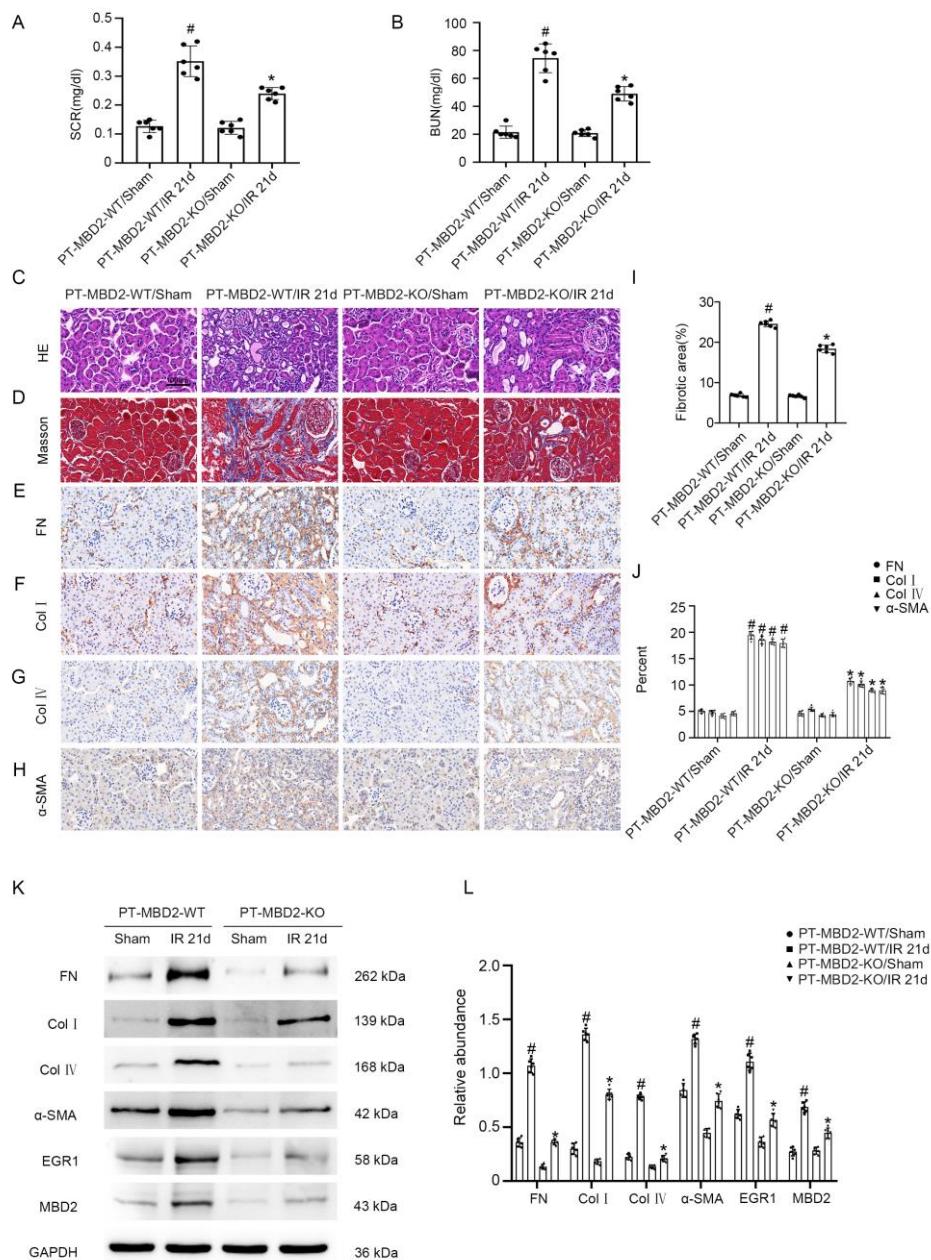


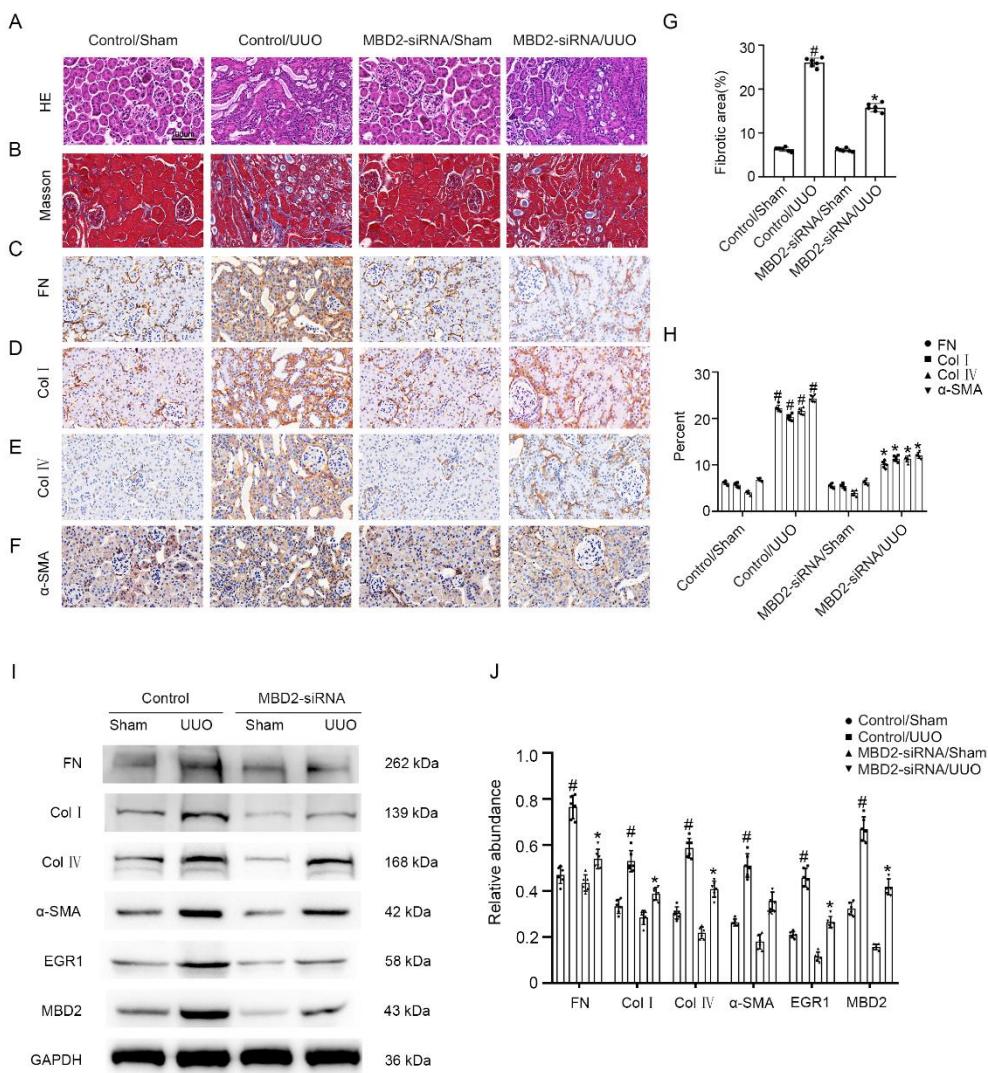
Supplemental information

**Genetic or siRNA inhibition of MBD2
attenuates the UUO- and I/R-induced renal
fibrosis via downregulation of EGR1**

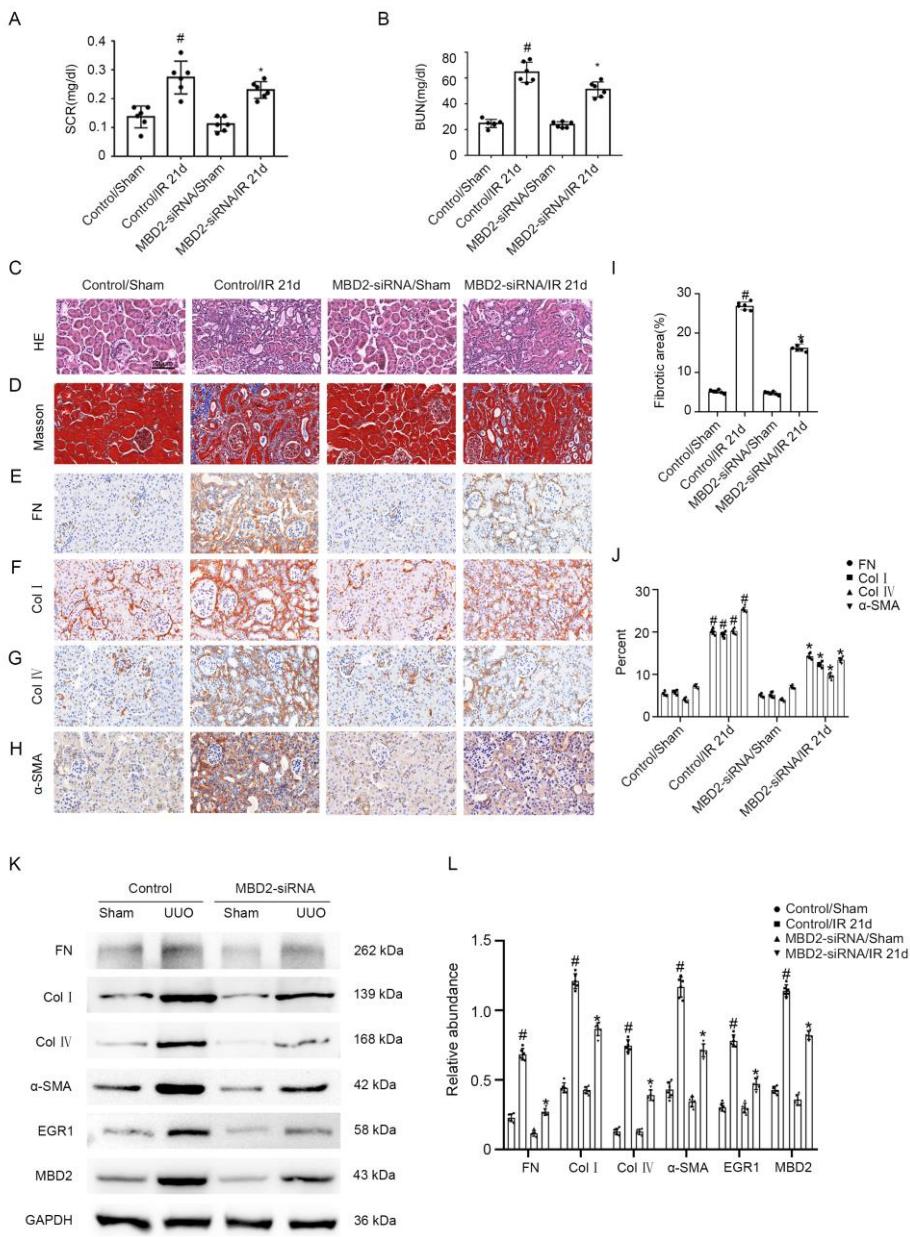
Kai Ai, Xiaozhou Li, Pan Zhang, Jian Pan, Huiling Li, Zhibiao He, Hongliang Zhang, Lei Yi, Ye Kang, Yinhuai Wang, Junxiang Chen, Yijian Li, Xudong Xiang, Xiangping Chai, and Dongshan Zhang



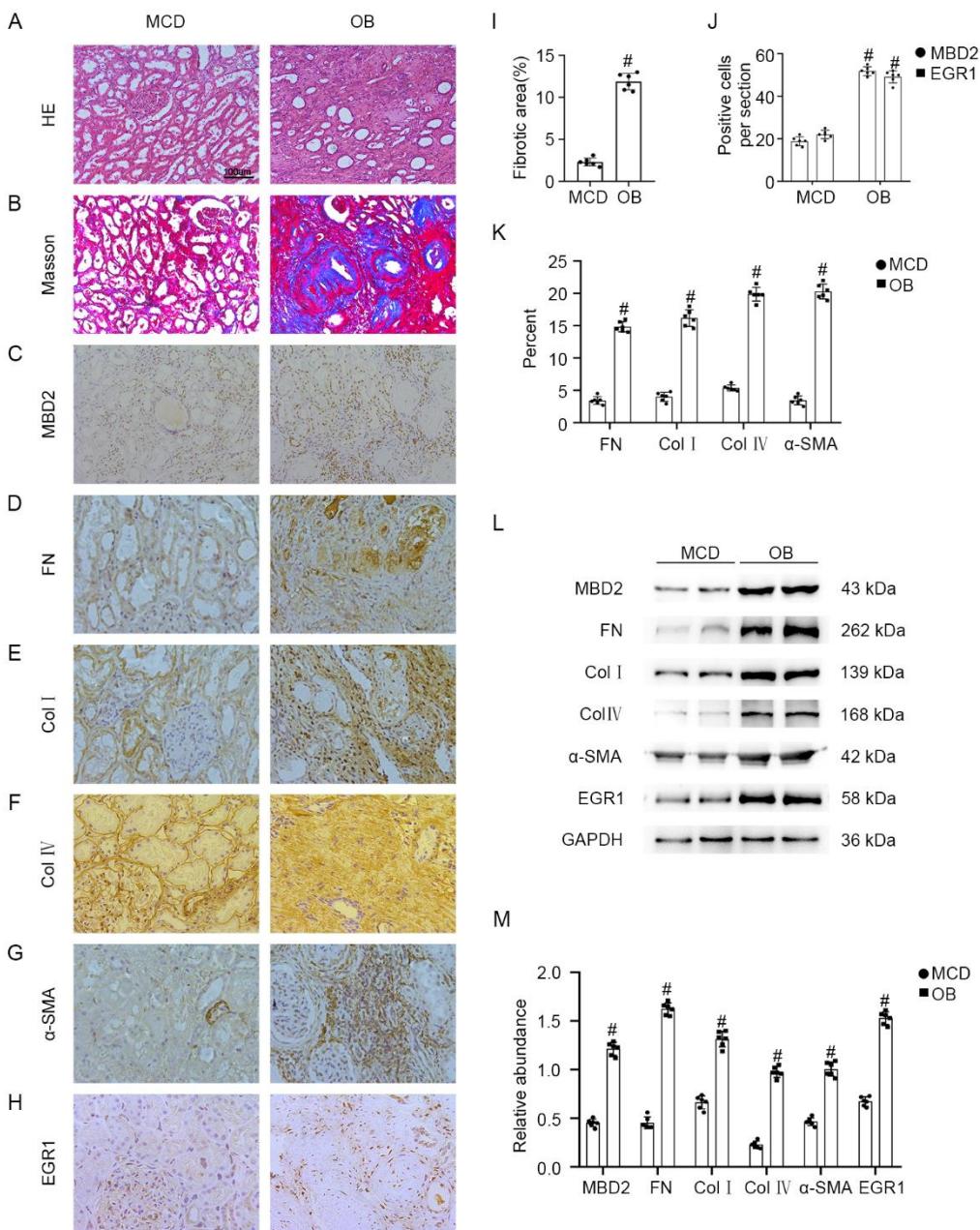
Supplemental Figure 1. PT-MBD2-KO mice attenuated IR-induced renal fibrosis. The littermate mice of the PT-MBD2-WT and PT-MBD2-KO were subjected to 28 minutes of bilateral renal ischemia followed by 21 days of reperfusion. (A) Blood SCR level of mice. (B) Blood BUN level of mice. (C) Representative hematoxylin and eosin staining. (D) Masson trichrome staining shows interstitial collagen deposition (blue). (E-H) Immunohistochemistry of FN, Col I&IV, and α-SMA. (I) Quantify tubulointerstitial fibrosis in the kidney cortex. (J) Quantification of immunohistochemistry staining. (K) Representative immunoblots for the expression of FN, Col I&IV, α-SMA, EGR1 and GAPDH. (L) Densitometry analysis of proteins signals, and normalized to internal control of GAPDH. Data are expressed as means ± sd (n=6). # P<0.05 versus sham group. * P<0.05 versus PT-MBD2-WT IR group. Original magnification, x400. Scale Bar: 100μm.



Supplemental Figure 2. MBD2 siRNA attenuated UUO-induced renal fibrosis. The C57BL/6 mice with UUO operation were injected with MBD2-siRNA or control at 15mg/kg via the tail vein once a week. (A) Representative hematoxylin and eosin staining. (B) Masson trichrome staining shows interstitial collagen deposition (blue). (C-F) Immunohistochemistry staining of FN, Col I&IV, and α-SMA. (G) Quantify tubulointerstitial fibrosis in the kidney cortex. (H) Quantification of immunohistochemistry staining. (I) Representative immunoblots for the expression of FN, Col I&IV, α-SMA, EGR1, MBD2, and GAPDH. (J) Densitometry analysis of proteins signals, and normalized to internal control of GAPDH. Data are expressed as means ± sd (n=6). # $P < 0.05$ versus sham group. * $P < 0.05$ versus Control with UUO group. Original magnification, x400. Scale Bar: 100μm.



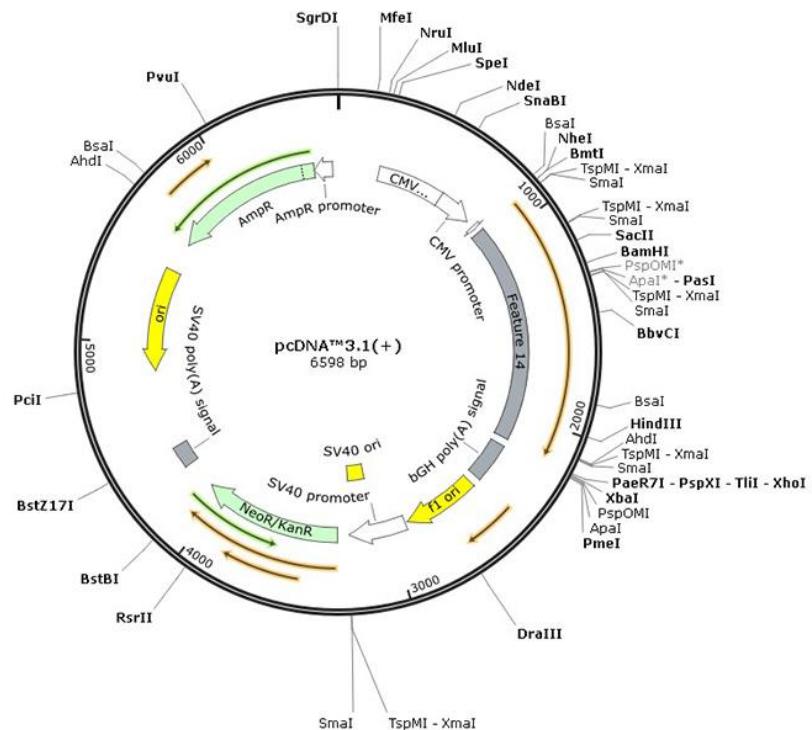
Supplemental Figure 3. MBD2 siRNA attenuated IR-induced renal fibrosis. The C57BL/6 mice with IR operation were injected with MBD2-siRNA or control at 15mg/kg via the tail vein once a week. (A) Blood SCR level of mice. (B) Blood BUN level of mice. (C) Representative hematoxylin and eosin staining. (D) Masson trichrome staining shows interstitial collagen deposition (blue). (E-H) Immunohistochemistry of FN, Col I&IV, and α-SMA. (I) Quantify tubulointerstitial fibrosis in the kidney cortex. (J) Quantification of immunohistochemistry staining. (K) Representative immunoblots for the expression of FN, Col I&IV, α-SMA, EGR1 and GAPDH. (L) Densitometry analysis of proteins signals, and normalized to internal control of GAPDH. Data are expressed as means ± sd (n=6). # P < 0.05 versus sham group. *P < 0.05 versus Control with IR group. Original magnification, x400. Scale Bar: 100μm.



Supplemental Figure 4. MBD2-induced downstream genes expression in patients with OB.

The samples of kidneys from OB patients were gathered. (A) Representative hematoxylin and eosin staining. (B) Masson trichrome staining shows interstitial collagen deposition (blue). (C-H) Representative immunohistochemistry staining of MBD2, ECM related genes, and EGR1. (I) Quantification of the tubulointerstitial fibrosis. (J&K) Quantification of immunohistochemistry staining (L) Representative immunoblot of MBD2, ECM related genes and EGR1. (M) Densitometry analysis of proteins signals, and normalized to internal control of GAPDH. Data are expressed as means \pm sd ($n=6$). # $P<0.05$ versus MCD group. Original magnification, $\times 400$. Scale Bar: 100 μ m.

MBD2 plasmid map



Supplemental Figure 5. The Map of MBD2 plasmid. The sequences listed as below:

GACGGATCGGGAGATCTCCGATCCCCTATGGTGCAGTCTCAGTACAATC
TGCTCTGATGCCGCATAGTTAAGCCAGTATCTGCTCCCTGCTTGTGTTG
GAGTCGCTGAGTAGTGCAGCAGAAATTAAAGCTACAACAAAGGCAAG
GCTTGACCGACAATTGCATGAAGAATCTGCTTAGGGTTAGGCCTTGC
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