

Supplemental information

**Renal subcapsular delivery of PGE₂ promotes kidney repair
by activating endogenous Sox9⁺ stem cells**

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SUPPLEMENTAL INFORMATION

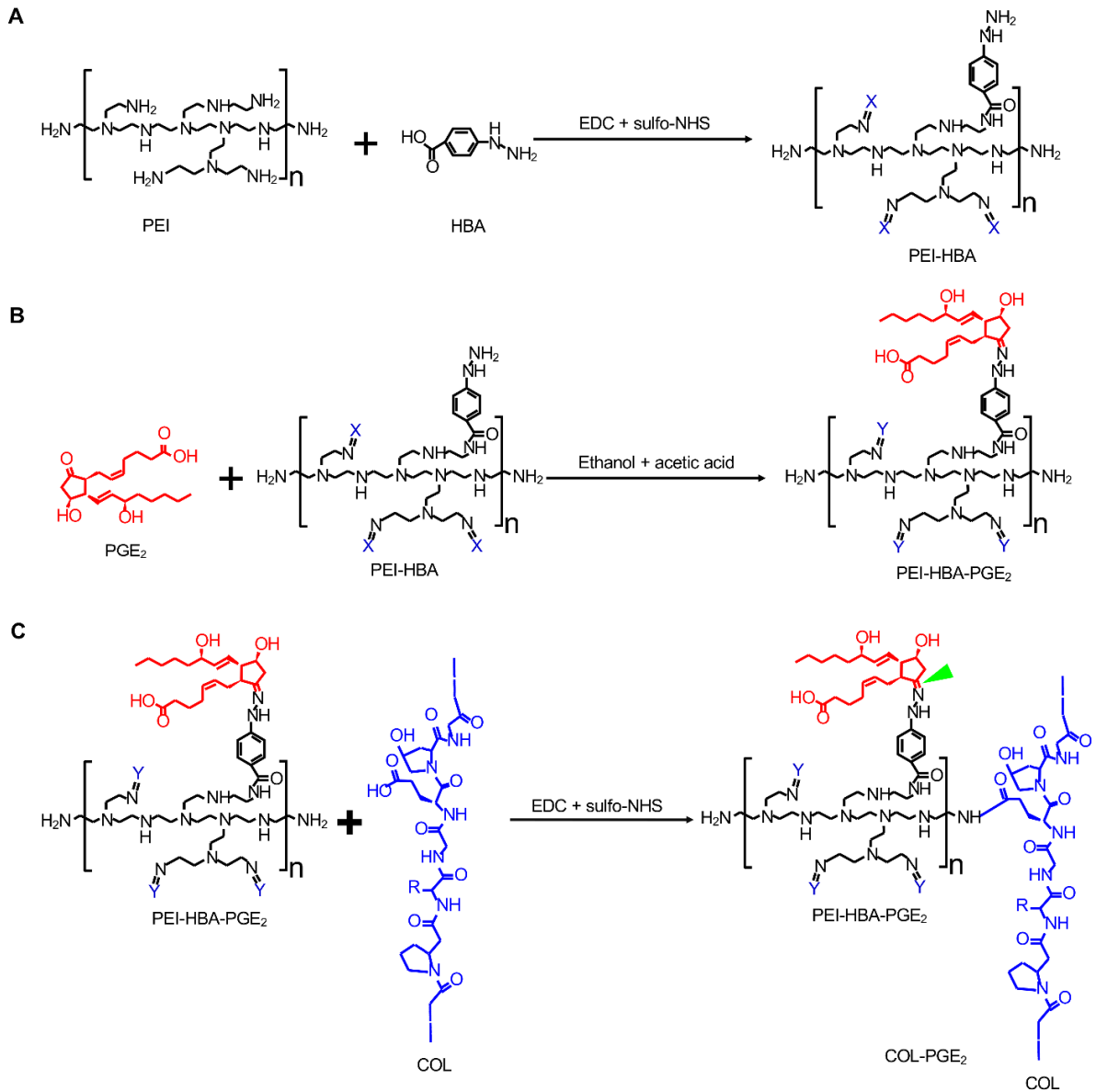


Figure S1. A schematic depicting the method of synthesis of the COL-PGE₂ matrix. (A) The PEI-HBA conjugates were synthesized by the dehydration condensation reaction between the amine group on PEI and the carboxylic group on HBA. **(B)** The ketone group on PGE₂ was reacted with the hydrazine group on PEI-HBA to synthesize the PEI-HBA-PGE₂ conjugates through a dehydration reaction. **(C)** The PEI-HBA-PGE₂ conjugates and collagen were further crosslinked by using dehydration condensation. Mechanism of PGE₂ release: The breakage sites of PGE₂ covalently attached (red) to branched PEI-HBA (black) and collagen (blue) are indicated by green arrows. X= hydrogen or HBA; Y = hydrogen, HBA or HBA-PGE₂. Abbreviations: HBA, 4-hydrazinobenzoic acid; PEI, polyethylenimine.

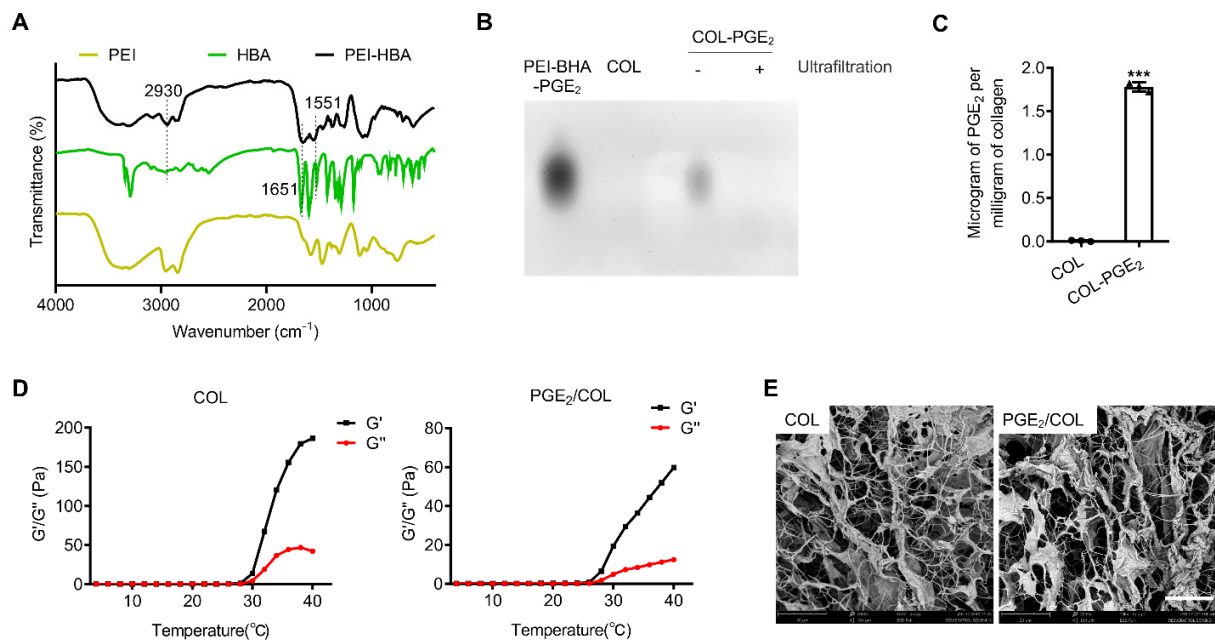


Figure S2. Characterization of collagen and the PGE₂/COL matrix. (A) FT-IR analysis shows the characteristic peaks of PEI, HBA, and PEI-HBA. (B) Tris-borate-EDTA-PAGE was used to detect free PEI-HBA-PGE₂. (C) Quantification of PGE₂ in collagen before and after crosslinking as determined using ELISA. Results are normalized to collagen concentration and expressed as the content of PGE₂ per milligram of collagen. The two-tailed *t* test was used for statistical analysis. Data are expressed as mean ± SD; ****P* < 0.001 versus COL. The experiments were carried out in triplicate. (D) Evaluation of the rheologic profile of collagen (left) and the PGE₂/COL matrix (right) during temperature changes. (E) Scanning electron micrograph image that reveals the morphologic structure of lyophilized collagen (left) and the PGE₂/COL matrix (right). Scale bars, 30 μm.

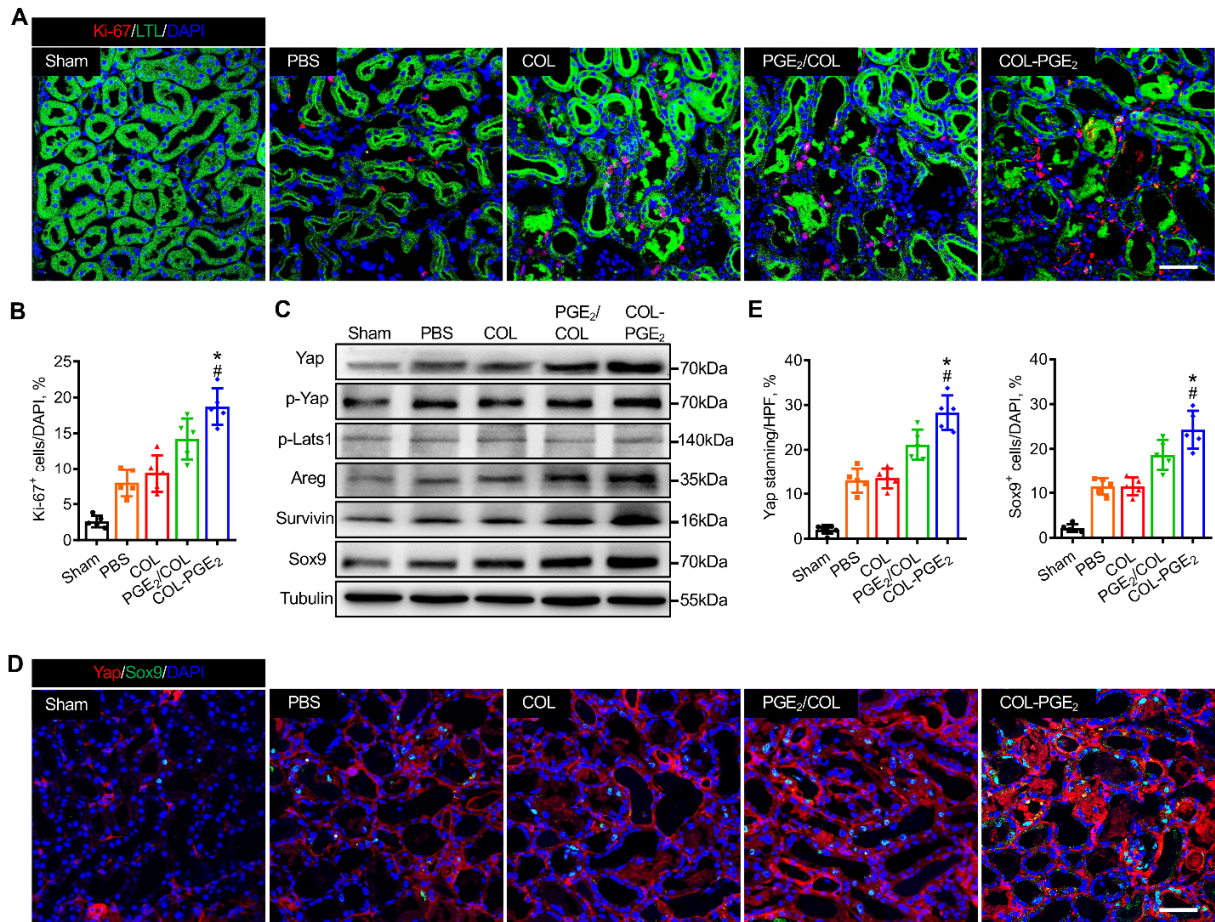


Figure S3. COL-PGE2 promoted the proliferation of proximal tubular epithelial cells and the activation of Sox9 through Yap mediation. (A) Representative images show Ki-67 immunostaining (red) detected in proximal tubular epithelial cells (LTL, green) on day 3 after AKI. (B) Quantification of Ki-67⁺ staining in (A). (C) Immunoblot analysis of the YAP, p-YAP, p-Lats1, Areg, Survivin, and Sox9 protein in the kidney on day 3 of PBS, COL, PGE₂/COL and COL-PGE₂. (D) Immunostaining analysis of YAP (red) and Sox9 (green) on day 3 after AKI. (E) Quantification of YAP⁺ (left) and Sox9⁺ (right) staining. HPF, high-power field. Scale bars, 50 μ m. One-way repeated measures ANOVA with Tukey post hoc tests (B and E) was used for statistical analysis. Data are expressed as mean \pm SD; n=5, * P < 0.05 versus COL; # P < 0.05, versus PGE₂/COL.

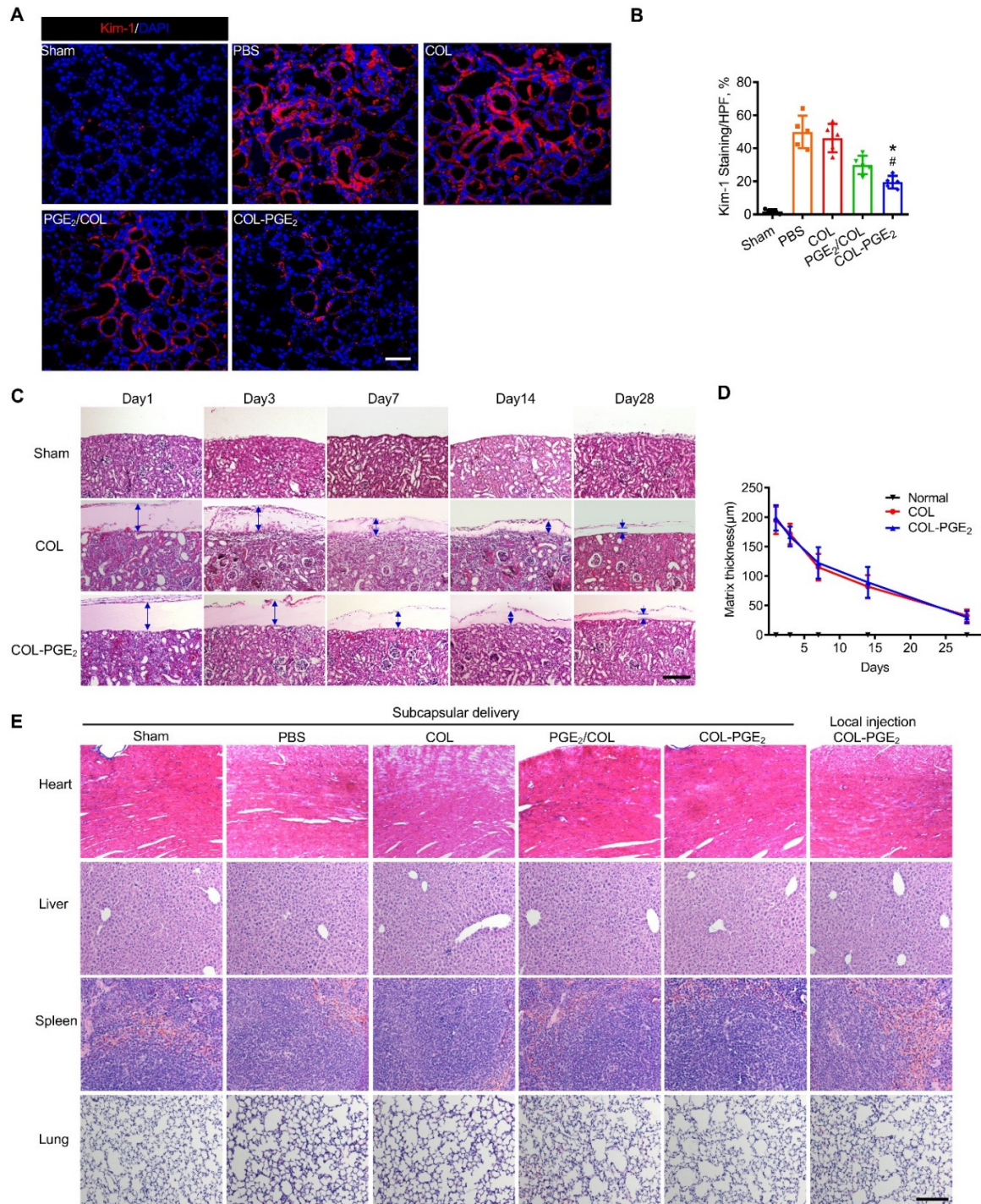


Figure S4. The COL-PGE₂ matrix attenuates kidney injury without organ toxicity. (A) Histological analysis of kidney injury by anti-Kim-1 immunostaining (red) on day 3 post AKI. Scale bars, 50 μ m. (B) Quantification of Kim-1⁺ staining. (C) Analysis of the matrix degradation profile by measuring the thickness of the matrix under the kidney capsules on day 1, 3, 7, 14, 28 post AKI. Scale bars, 200 μ m. (D) Quantification of the thickness of the matrix. (E) H&E staining did not show histological changes in the heart, liver, spleen and lung. One-way repeated measures ANOVA with a Tukey post hoc test (B) and Two-way multivariate analysis of variance (two-way MANOVA) with Tukey post hoc test (D) were used for statistical analysis. Data are expressed as mean \pm SD; n=5, * P < 0.05 versus COL; # P < 0.05, versus PGE₂/COL.

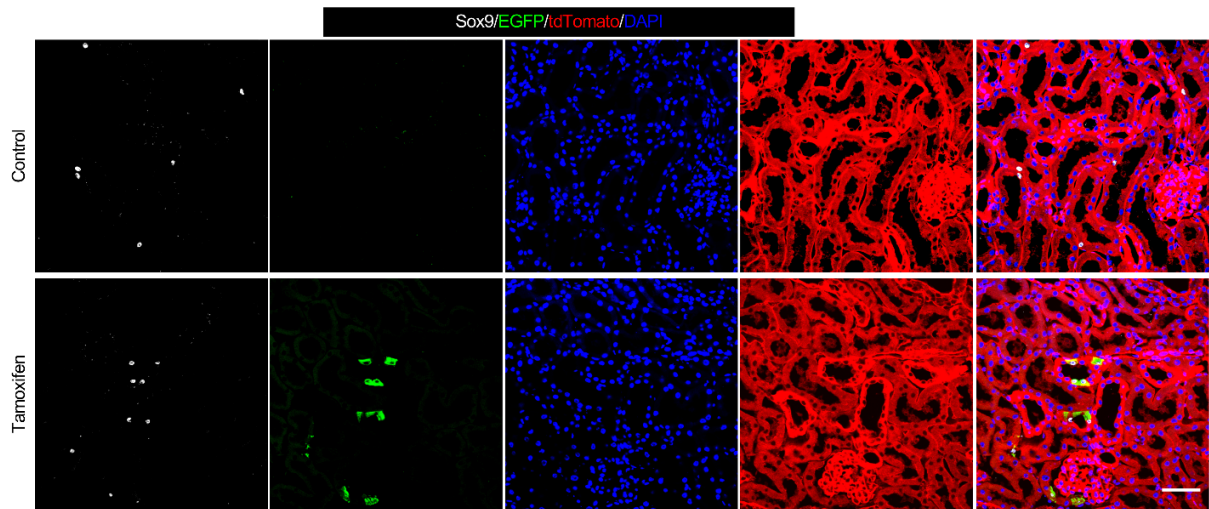


Figure S5. Validation of the Sox9-Cre^{ERT2}; R26^{mTmG} mouse genetic model

To verify the mouse genetic model, tamoxifen (corn oil was used as a control) was injected three times in succession before AKI, and Sox9⁺ cells and EGFP fluorescence colocalization in the kidney were detected by anti-Sox9 immunostaining.

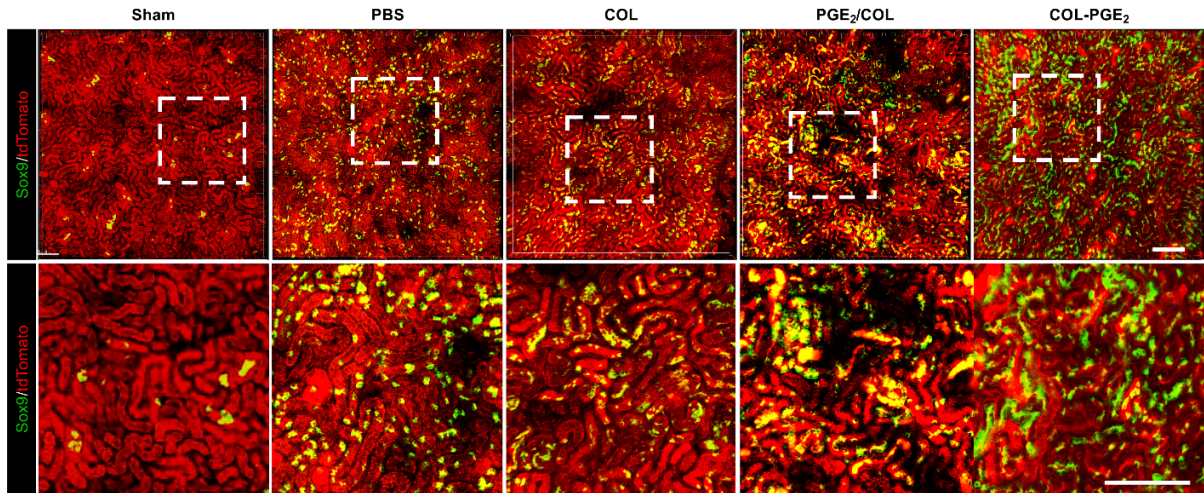


Figure S6. Two-photon living imaging of Sox9⁺ cell expansion following PGE₂ treatment. Representative images of two-photon intravital tracing showed that Sox9⁺ cell-derived cells were abundantly expanded by the administration of COL-PGE₂ matrix on day 7 after AKI. Scale bar, 200 μ m. See also Video S1-4.

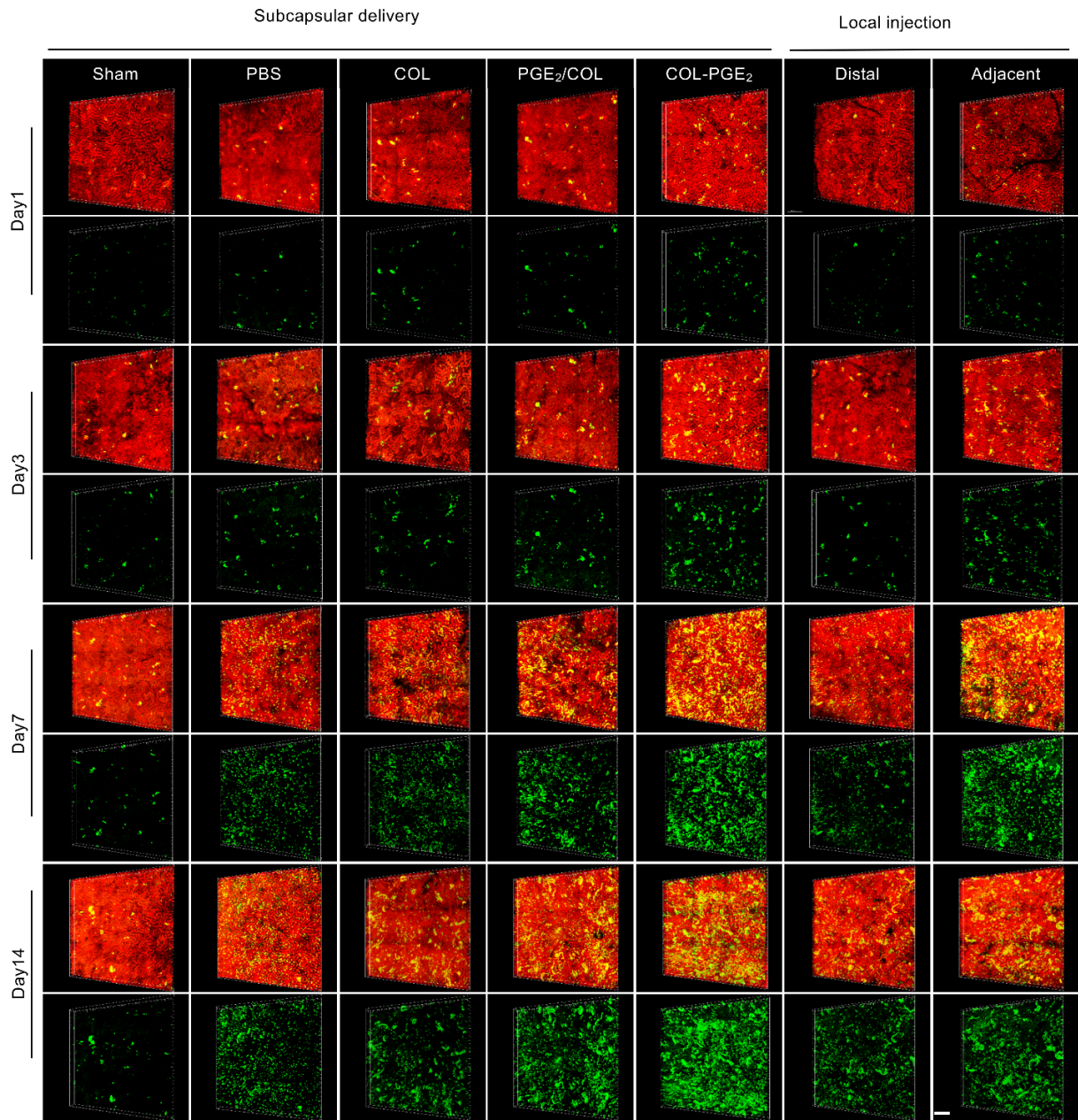


Figure S7. The COL-PGE2 matrix promoted the formation of functional renal tubules in descendants of Sox9⁺ cells. Representative images of two-photon intravital microscopy showed Sox9⁺ derived cells with the administration of PBS, COL, PGE₂/COL and COL-PGE₂ matrix at different time points after AKI. Scale bar, 200 μ m.

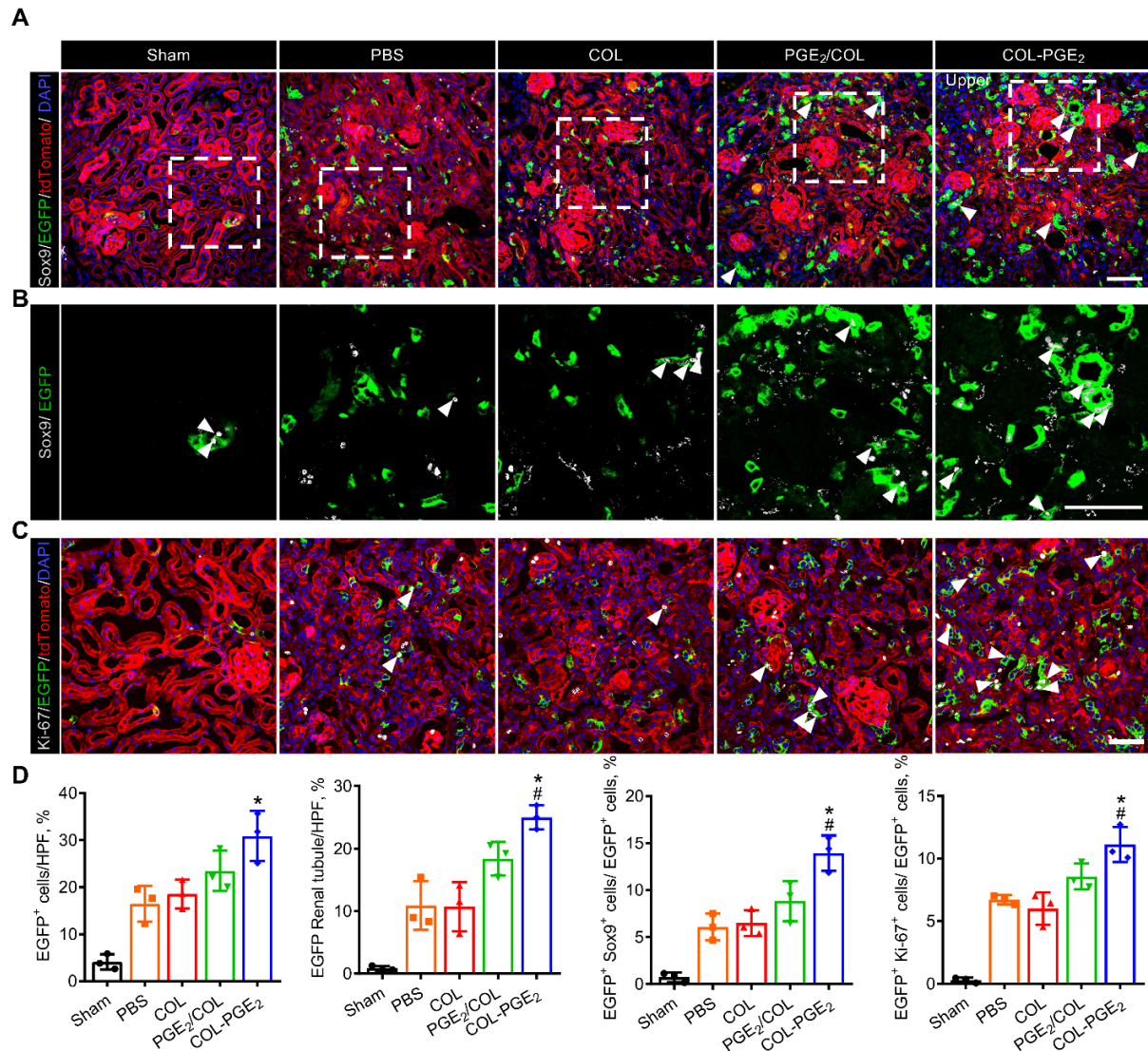


Figure S8. The COL-PGE₂ matrix results in increased activation of Sox9 expression and Sox9⁺ cell proliferation (A) Confocal images and (B) local zoomed-in images for colocalization analysis of anti-Sox9 immunostaining (gray) and Sox9-Cre^{ERT2}-activated EGFP fluorescence in kidneys with the administration of the PBS, COL, PGE₂/COL and COL-PGE₂ matrix on day 14 post AKI. White arrowheads highlight Sox9-EGFP-labeled cells, and white arrowheads highlight Sox9⁺/EGFP-colabeled cells. Scale bar, 50 μ m. (C) Confocal images for the colocalization analysis of anti-Ki-67 immunostaining (gray) and Sox9-Cre^{ERT2}-activated EGFP fluorescence in kidneys with the administration of PBS, COL, PGE₂/COL and COL-PGE₂ matrix on day 14 after injury. White arrowheads highlight Ki-67⁺/EGFP colabeled cells. Scale bar, 50 μ m. (D) Quantification of EGFP⁺ cells, EGFP renal tubules, Sox9⁺/EGFP colabeled cells and Ki-67⁺/EGFP colabeled cells in the kidneys of mice on day 14 after delivery of the COL-PGE₂ matrix post AKI. One-way repeated measures ANOVA with a Tukey post hoc tests (D) were used for statistical analysis. Data are expressed as mean \pm SD; n=3, **P* < 0.05 versus COL; #*P* < 0.05, versus PGE₂/COL.

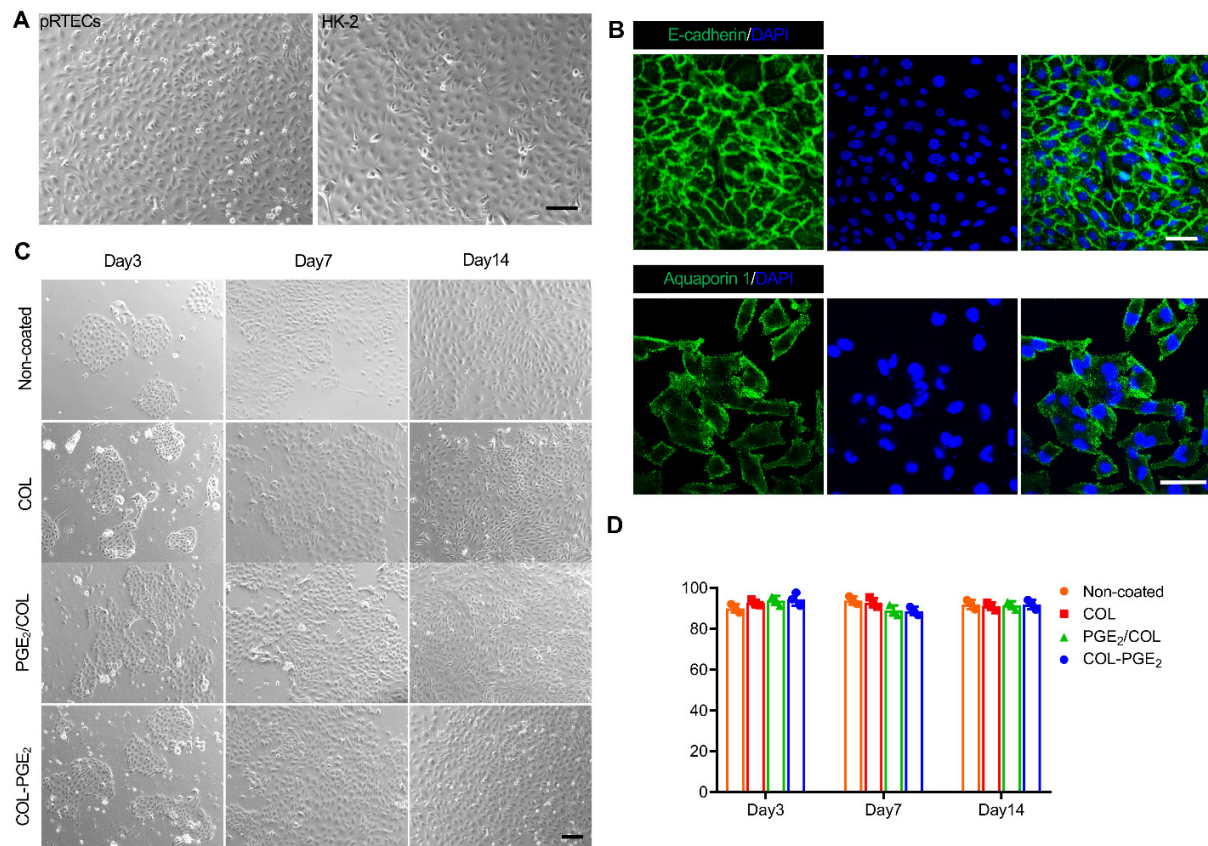


Figure S9. Renal tubular epithelial cell morphology and matrix biocompatibility. (A) Comparison of the morphology between isolated primary renal tubular epithelial cells (pRTEC) and HK-2 human renal proximal tubule cells. Scale bar, 100 μ m. (B) Immunofluorescence staining with antibodies against E-cadherin (top) and Aquaporin-1 (bottom) shows the isolated pRTECs. Scale bar, 50 μ m. (C) The morphology and colony formation capacity of pRTEC cultured on noncoated or COL, PGE₂/COL or COL-PGE₂ matrix for 3, 7, and 14 days. Scale bar, 100 μ m. (D) Quantification of cell viability by trypan blue staining and calculation of the percentage of live cells cultured on noncoated or COL, PGE₂/COL or COL-PGE₂ matrix-coated plates for 3, 7, and 14 days. Two-way MANOVA with Tukey post hoc test was used for statistical analysis. Data are expressed as mean \pm SD; All data are representative of three independent experiments and are shown as mean \pm SD.

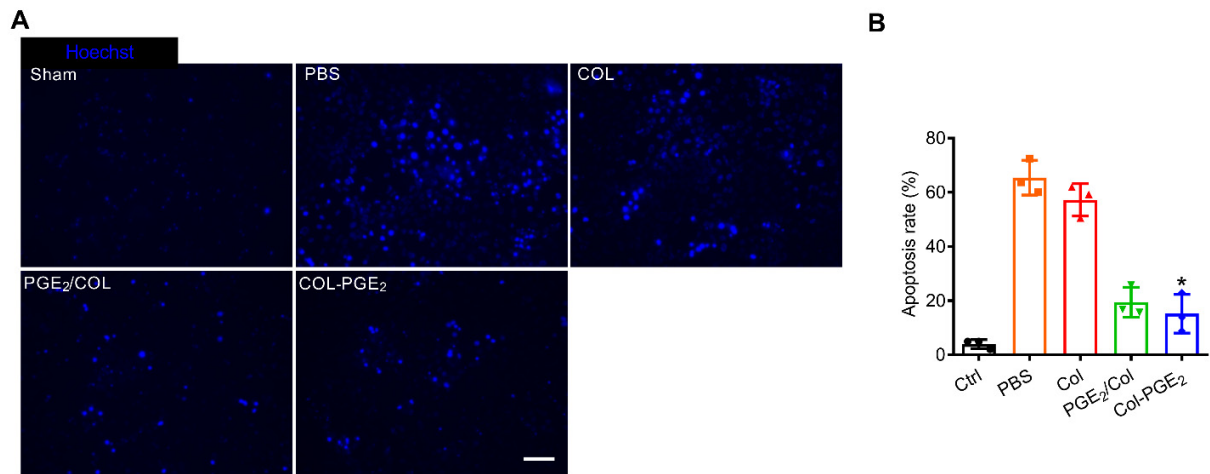


Figure S10. COL-PGE₂ matrix exerts an anti-apoptosis effect *in vitro*. (A) Hoechst 33342 staining showed the survival of pRTECs cultured on noncoated, PBS, COL, PGE₂/COL or COL-PGE₂-coated plates upon exposure to 150 μ M H₂O₂ for 6 h. Scale bar, 100 μ m. (B) Quantification of the apoptosis index of pRTECs under H₂O₂-induced stress. One-way repeated measures ANOVA with a Tukey post hoc test (B) was used for statistical analysis. Data are expressed as mean \pm SD, n=3, **P* < 0.05 versus COL.