## Supplementary Information

## Four-dimensional Hydrogel Dressing Adaptable to the Urethral Microenvironment for Scarless Urethral Reconstruction

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## **Supplementary Figures**



**Supplementary Fig. 1.** <sup>1</sup>H NMR spectral characterization of gelatin methacryloyl (**GM**), fluorophenylboronic acid (FPBA)-modified gelatin methacryloyl (**GMP**), and *cis*-diol-modified gelatin methacryloyl (**GMD**).



Supplementary Fig. 2. A) Representative strain sweep rheological plots of GMPD-hv hydrogels. B) Viscosity and shear-thinning behavior of GMPD-hv hydrogels. n = 4 independent samples.



Supplementary Fig. 3. A) The C(1s) *XPS* regions of GMPD+hv hydrogel samples. Intensity (cps): count per second. Red line: C(1s) characteristic speak; green line: C=O (288.02 eV); orange line: C-O (286.26 eV); blue line: C-C (284.82 eV); blue arrows represent C(1s) regions. B) The B(1s) *XPS* regions of GMPD+hv hydrogel samples. Red line: B(1s) characteristic speak; blue line: B-O (191.03 eV); green line: B-C (189.46 eV); red arrows represent B(1s) regions.



Supplementary Fig. 4. The statistical data of standard lap shear (A) and incision sealing (B) tests showing the *ex vivo* adhesive performance after 0 and 12 hours. n = 4 independent samples. Data are presented as mean  $\pm$  SD. All error bars represent SD. p values calculated using one-tailed unpaired t-test. ns = no significance. Source data are provided as a Source

## Data file.



**Supplementary Fig. 5.** Representative photographs showing the resistance to water washing at approximately 10 kPa water pressure in the urethral defect model. Red arrows represent the adhesive hydrogels stained by green fluorescence. n = 3 independent samples.



Supplementary Fig. 6. The *in vitro* degradation curves of GMPD-hv and GMPD+hv hydrogels in neutral (pH = 7.4) or acidic solutions (pH = 4.5-6.5). n = 4 independent samples. Data are presented as mean  $\pm$  SD. All error bars represent SD. Source data are provided as a Source Data file.



Supplementary Fig. 7. A, B) Representative photographs (A) and histological examinations of H&E and Masson's trichrome staining (B) of the remaining GMPD hydrogels in the urethral defect model after 3-, 7-, 14-days surgery. Blue circles and red arrows represent the remaining hydrogels. n = 3 biologically independent samples.



**Supplementary Fig. 8.** The *in vitro* swelling ratio of **GMPD**+hv hydrogels in different pH values ranging from 4.5-8.5. n = 4 independent samples. Data are presented as mean  $\pm$  SD. All error bars represent SD. p values calculated using one-tailed unpaired t-test. Source data are provided as a Source Data file.



**Supplementary Fig. 9. A)** Observation of the endothelial cell scratch assay after 0, 6, 12, and 24 hours in the V-GM, TI-PLGA, and control (Ctrl) groups. B) Statistical analyses of the corresponding wound healing rates. n = 3 biologically independent samples. Data are presented as mean  $\pm$  SD. All error bars represent SD. p values calculated using one-tailed unpaired t-test. ns = no significance. Source data are provided as a Source Data file.



Supplementary Fig. 10. Immunofluorescence staining of COL1 (red), COL3 (green), and cell nuclei (DAPI, blue) for fibroblast marker expression after 4- and 7-days culture in the V-GM and TI-PLGA groups. n = 3 biologically independent samples.



**Supplementary Fig. 11. A)** Immunofluorescence staining of CD31 (red), VWF (green), and cell nuclei (DAPI, blue) for endothelial cell marker expression after 4- and 7-days culture in the **V-GM** and **TI-PLGA** groups. **B, C)** Comparative endothelial cell expression levels (*CD31* and *VWF*) after 4- and 7-days culture in the **V-GM**, **TI-PLGA**, and control (**Ctrl**) groups. n = 3 biologically independent samples. Data are presented as mean  $\pm$  SD. All error bars represent SD. p values calculated using one-tailed unpaired t-test. ns = no significance. Source data are provided as a Source Data file.



Supplementary Fig. 12. A, B) The CCK-8 data of 1% w/v extract solution of TOR

microspheres, **GMPD**-*hv*, **GMPD**+*hv*, and **GMPD**+*hv*/**TOR** hydrogels after 24-hours culture for both fibroblasts (**A**) and HUVECs (**B**). n = 4 biologically independent samples. Data are presented as mean  $\pm$  SD. All error bars represent SD. p values calculated using one-tailed unpaired t-test. ns = no significance. Source data are provided as a Source Data file.



Supplementary Fig. 13. A, B) Volcano plot (A) and statistical data (B) of endothelial cell

DEGs analyzed between the E1 and E4 groups. **C**, **D**) The Gene Ontology (GO, **C**) and Kyoto Encyclopedia of Genes and Genomes (KEGG, **D**) enrichment analyses of endothelial cell DEGs after mRNA sequencing in **TOR**-functionalized hydrogels between the E1 and E4 groups, including the top 15 representative upregulated or downregulated signaling pathways.



Supplementary Fig. 14. The magnified H&E staining of the rabbit urethral canal in the GMPD, GMPD-V, GMPD-TI, GMPD-V/TI, and control (Ctrl) groups after 8-weeks surgery. n = 3 biologically independent samples.



Supplementary Fig. 15. The semi-quantitative data of Masson's trichrome staining of the scar thickness in the GMPD, GMPD-V, GMPD-TI, GMPD-V/TI, and control (Ctrl) groups

after 8-weeks surgery. n = 3 biologically independent samples. Data are presented as mean  $\pm$  SD. All error bars represent SD. p values calculated using one-tailed unpaired t-test. ns = no significance. Source data are provided as a Source Data file.



Supplementary Fig. 16. The magnified immunofluorescence staining of the rabbit urethral canal for evaluating epithelialization (AE1/AE3) after different treatments (i.e., GMPD, GMPD-V, GMPD-TI, GMPD-V/TI, and Ctrl groups) for 8 weeks. n = 3 biologically independent samples.



**Supplementary Fig. 17. A, B)** Immunofluorescence staining (A) and quantitative expression level (B) of the rabbit urethral canal for evaluating the M2 macrophages (CD206) after different treatments (i.e., GMPD, GMPD-V, GMPD-TI, GMPD-V/TI, and Ctrl groups) for 8 weeks. n = 3 biologically independent samples. Data are presented as mean  $\pm$  SD. All error bars represent SD. p values calculated using one-tailed unpaired t-test. Source data are provided as a Source Data file.