# Decellularization of rat precision-cut kidney slices — Application of physical and chemical methods

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engineering," Current Opinion in Nephrology and Hypertension, vol. 23, no. 4, 2014.

Animal Tissues," Journal of medical signals and sensors, vol. 11, no. 1, pp. 1–11, 2021.

p. 45005, 2017.





Amount of DNA in native kidney tissue and decellularized PCKS. Chem (n=4), HHP 50 (n=5), HHP 100 (n=4), HHP 200 (n=2). While FTC resulted in a significant reduction in DNA content HHP protocols resulted in relatively similar DNA content and non significant reduction compared to the Chem protocol.  $*P \leq 0.05, **P \leq 0.01, ***P \leq 0.001.$  Data were analyzed with a Mann-Whitney-twosample-test and are given as mean (SD)

> Histology of control and HHP (600 MPa) treated rat kidney tissues (H&E). HHP treated tissue shows huge reduction in interstitial space (red arrows) and a darker color

Number of cell nuclei in kidney tissues after HHP treatment. Control, 10 MPa, 100 MPa, 600 MPa (n=3). The data are normalized to the area of tissue of the control. Data are given as mean (SD). Cell nuclei were counted with QuPath and the tissue area

4 I. Fischer, M. Westphal, B. Rossbach et al., "Comparative characterization of decellularized renal scaffolds for tissue engineering," Biomedical Materials, vol. 12, no. 4,





Histology of decellularized PCKS stained with H&E (20x) shows removal of nuclei in Chem and FTC protocols. Both resulted in an overall preservation of the structures with only minor damage (red arrows)



## Summary

- removal of residual DNA
- In process: further structural analysis with SEM
- epithelial cells (RPTEC/TERT1)



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Amount of GAG in native kidney tissue and decellularized PCKS. Chem (n=4), HHP 50 (n=5), HHP 100 (n=4), HHP 200 (n=3). \*P ≤ 0.05, \*\*P ≤ 0.01. Data were analyzed with a Mann-Whitney-two-sample-test and are given as mean (SD). All Protocols resulted in a significant reduction in GAG amount compared to the native tissue. Chem and FTC protocols showed similar GAG content

representative image of a PCKS decellularized with the Chem protocol and Scanning electron imaged using microscopy (SEM) (2000x). The image shows renal tubular structures. The red arrow points at presumably residual cellular debris and the green arrow at fibrillar ECM proteins (collagen, fibrin etc.)

• FTC resulted in the highest reduction in residual DNA and a better preservation of GAG combined with only minor damage to the ECM • HHP causes compression in kidney tissues leading to ineffective

• In process: Recellularization of PCKS with renal proximal tubular