

# MALDI Mass Spectrometry Imaging of Human Penile Tissue Scaffolds following Organ Decellularization to Evaluate Extracellular Matrix Preservation Caitlin M. Tressler<sup>1</sup>, Allister Suarez<sup>2,3</sup>, Yu Tan<sup>2,3</sup>, Wilmina N. Landford<sup>2,3</sup>, Devin Coon<sup>2,3</sup>, and Kristine Glunde<sup>1,4</sup>

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mode (m/z 500 to 3,200) with 100 micron pixel size (200 laser shots) on a Bruker RapifleX MALDI TOF/TOF instrument.









- Methods were successfully developed to image tryptic peptides and glycans from both fresh-frozen and decellularized human penile tissue.
- Segmentation analysis results demonstrate that some ion suppression may be present in the fresh-frozen specimen due to cellular components not observed in the decellularized specimen.
- Unsupervised analysis, including segmentation and component analysis, demonstrated that the ECM of the human penis is unique to each structure within the penis (i.e., cavernosa and tunica). This indicates that individualized reseeding techniques may be necessary for tissue engineering, which are specific to the ECM of each specialized penile structure. \* We are currently working on establishing a pipeline for peak identification using MS/MS and LC-MS/MS to determine the
- identities of proteins and glycans in this tissue. ↔ We intend to employ hematoxylin-eosin (H&E) and immunohistochemistry (IHC) stains to confirm our MALDI MSI findings.