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Liver Tissue Tissue Microarray (TMA) Embedding and Sectioning -- University of Minnesota TMCs

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Cellular Senescence Net...



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Protocol status: Working

We use this protocol and it's working

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Abstract

Construct TMAs compatible for multiple spatial transcriptomic and proteomic platforms.

Guidelines

Ethics Statement:

This protocol requires prior approval by the users' local Institutional Animal Care and Use Committee (IACUC), Institutional Review Board (IRB) or equivalent ethics committee(s), depending on the source of the tissue.

Materials

FFPE tissue blocks with sufficient tissue (width and depth) for TMA construction

Beecher Manual Tissue Arrayer (MTA-1)

Biopsy punch (2 mm)

Troubleshooting

TMA dimensions are determined based on the area requirements of all planned spatial platforms.

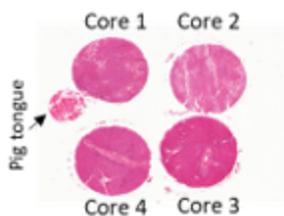
- 1 The smallest capture dimension of all the spatial platforms for the multiplatform experiment was 6.5× 6.5 mm. It was determined that four x 2 mm tissue cores with 2.8 mm spacing could successfully fit within these dimensions.

Selection of tissue core from each FFPE tissue block

- 2 H&E of each tissue block for the TMA was examined by pathologist and 2 mm area determined and annotated on the FFPE block.
- 3 2 mm cores removed from FFPE block using biopsy punch

TMA construction

- 4 The Beecher Manual Tissue Arrayer (MTA-1) was used for TMA construction.
 - 4.1 Four tissue cores were embedded in a paraffin block in a column/ grid format.
 - 4.2 To aid in orientation, a 0.6mm core of pig tongue tissue was included as a marker.
 - 4.3



Example Liver TMA H&E