

Tissue Micro arrays (TMA)

What is it?

- The use of Multi-tissue blocks was first introduced by H. Battifora in 1986. In 1998, J. Kononen and collaborators developed the current technique, which uses a novel sampling approach to produce tissues drill cores of regular size and shape that can be arranged more densely in an array format.

What can it be used for?

- Tissue microarrays (TMA) can be used for simultaneous analysis of the expression of proteins and mRNAs as well as for ligand binding studies in tissue biopsies/autopsies that are arranged in a high-density format.
- One of the main advantages with the TMA technique is the possibility to use relatively slow methodological approaches such as in situ hybridization and immunohistochemistry in a high throughput format. These qualities make TMAs an attractive screening tool for expression analysis both for the target identification/validation process as well as for toxicological evaluations and autoradiography studies.

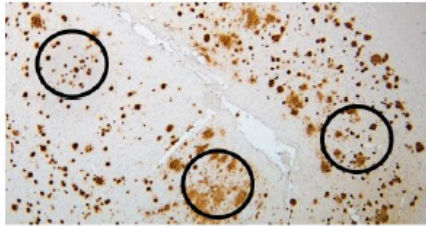
Our services

- The Offspring team has extensive experience constructing TMAs for generation of strong data in tissues from both preclinical animal models of disease and the corresponding human disease state.
- The TMA can be generated from fixed and paraffin embedded human and animal tissues.
- We also use TMA-formatted tissues for tissue cross reactivity (TCR) studies before clinical trial (biologics and in some cases SMDs)
- A typical build of a TMA includes:
 - Pre-evaluation of tissue quality
 - Pathological characterization
 - Defining sites for sampling of drill cores
 - Assembly of tissue cores (diameter = 0,6 - 2mm) from selected tissue areas into a recipient TMA block.
 - Reconfirmation informativity and functionality of the TMA

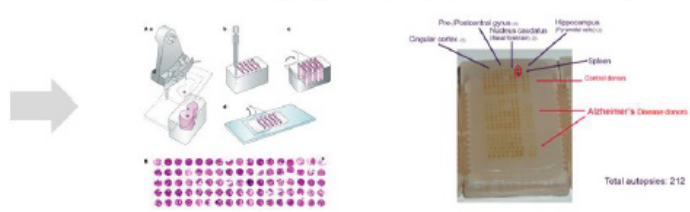




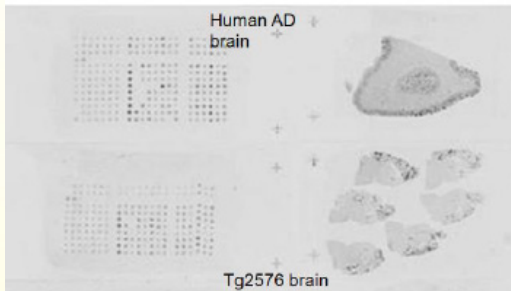
Defining pathology



Constructing high density tissue microarrays



Comparing drug target expression in animal models and human tissues



Characterizing compound binding properties on large sets of tissues

