

New Tools in Biomedical Research: Tissue Arrays

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Getting to the Core of Tissue Microarrays

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Goals

- Define tissue microarrays (TMAs)
- Describe TMA construction methods
- Discuss advantages of and uses for TMAs
- Consider when it is appropriate to use TMAs
- Discuss design considerations & analysis options
- Provide website addresses for TMAs

What is a TMA?

- An array of cores taken from “donor” tissue blocks which are inserted into an “acceptor” block in a defined layout.
- Usually from paraffin embedded tissue.*
- Can hold anywhere from 2 to 2000 cores.
 - Manual or automatic construction.

Manual Construction of TMAs

**Large Corer with Acceptor Mold &
Acceptor Blocks**

**Manual Construction of TMAs:
Small Corer with Acceptor & Donor Blocks**

**Manual Construction of TMAs:
Marked Slides with Donor Blocks
& Sections of TMAs**

Manual Construction of TMAs:

Placing the core in the acceptor block

Other ways to construct TMAs

- **Low Cost, Low Tech** - placement of manually dissected pieces of tissue in a normal paraffin mold.
- **High Cost, High Tech** - commercial automatic and manual arrayers.
<http://www.becherinstruments.com/>
<http://www.chemicon.com/>

Section of a typical TMA

Advantages and Disadvantages of TMAs

- Maximizes tissue use
- Minimizes reagent costs
- Minimizes labor costs
- Maximizes consistency
- Maximizes quality control
- Some damage to donor block
- Specialized equipment
- Requires pathologist's time
- Assumes homogeneity
- Difficult to analyze w/o digital microscopy

Appropriate Use of TMAs

If you can do it with a conventional section, you can do it with a TMA.*

Histology, IHC, FISH, ISH

* THE FINE PRINT - Some Restrictions May Apply

1. Not for heterogeneous tissues
2. Samples may be lost due to core "drop out"
3. Information may be lost due to lack of context.
4. Samples may be difficult to analyze.

Design Considerations

- Adequate representation of tissue vs. Preservation of clinical donor block
 - Small cores vs. Large cores
- Multiple cores from each sample on one array vs. Single cores on duplicate arrays
- Maximize # of sections from each array vs. Preserving optimal antigenicity

Analysis Options

- **Manual, Real-time Analysis** - unwieldy with high density arrays, doable with low density arrays
- **Automated, Digital Analysis** - desirable for high density arrays. Several commercially available platforms (Bacus Labs, Chromavision, Zeiss, Aperio/DAKO, and others)

Informational and Educational Web-links for TMAs

- http://www.the-scientist.com/yr2002/oct/lcprofile1_021028.html
- <http://www.nhgri.nih.gov/DIR/CGB/TMA/>
- http://ccr.cancer.gov/tech_initiatives/tarp/
- <http://pluto3.nci.nih.gov/tissue/default.htm>
- <http://genome-www.stanford.edu/TMA/>
- <http://icg.cpmc.columbia.edu/cattoretto/Protocol/immunohistochemistry/TissueArray.html>

Commercial Web-links for TMAs

- <http://www.molecularhistology.com/microarray07.htm>
- <http://www.ihcworld.com/tissuearray.htm>
- <http://www.chromavision.com/>
- <http://www.baculuslabs.com/>
- <http://www.aperio.com/home.asp>
- <http://www.beecherinstruments.com/>
- <http://www.zymed.com/products/75-xxxx/75-4033.html?emmun010>

