



OBBR

Office of Biorepositories
and Biospecimen Research

NCI Best Practices for Biospecimen Resources

**Technical and Operational Best Practices
(Real-World Perspective)**

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- **Introduction and Perspective**
- **Examples**
- **Issues Encountered and NCI Best Practices**
- **Conclusion**



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Introduction and Perspective

- **The Cancer Genome Atlas (TCGA) pilot project**
 - 3 year pilot project of the NCI and NHGRI to comprehensively catalog the molecular changes associated with cancer.
 - Three different cancers: brain, ovarian and lung
 - Biospecimens obtained from a network of retrospective collections at multiple academic medical centers.
 - Large scale molecular analysis – 10 platforms, each doing every case in common.
(RNA and micro-RNA profiling, copy number variation, translocation analysis, epigenetics, and sequencing.)
 - Clinical data integrated with molecular data.
 - Integrated data sets made available to the broad research community.



Introduction - Not in scope for this presentation

- **Other important factors that impact biospecimen access:**
 - Human subjects policies, IRB approvals, HIPAA
 - Re-contacting and re-consenting of living patients
 - Material Transfer Agreement, Intellectual Property, Authorship
 - Informatics
 - Extraction and transfer of associated clinical data
 - Process data
 - Standards compliance (caBIG™)
 - Costs



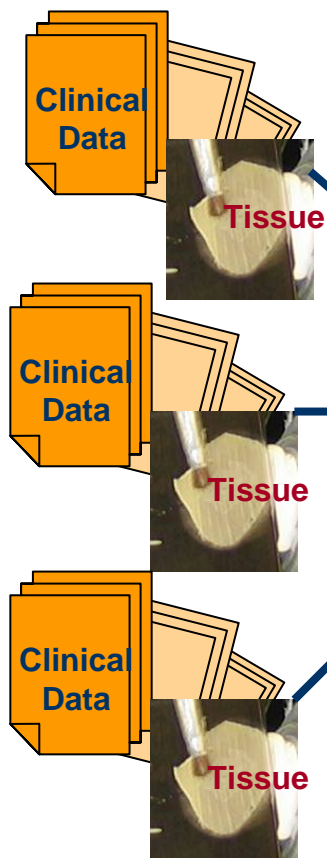
Introduction - TCGA Goals & Biospecimen Quantity and Quality

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- **500 individual cases successfully yielding molecular profiles**
 - Estimated 35% histological + molecular QC failure = 760 cases
 - Statistical power vs. financial constraints
 - Preferably from 2 collections to minimize variability
- **Germline DNA source for every case**
- **Frozen samples with at least 200 mg of tumor tissue per case**
 - DNA + RNA from each sample, enough for all 11 sites
 - No WGA
- **At least 80% of each sample composed of viable tumor cells on histologic assessment**
 - No LCM

Introduction - TCGA Operational Overview

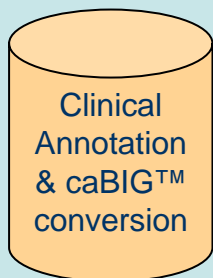
Clinical site biorepositories



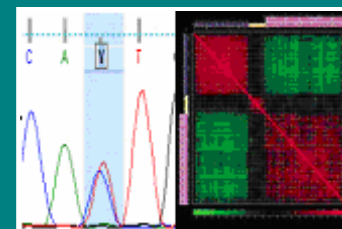
Biospecimen Core Resource

Single central facility

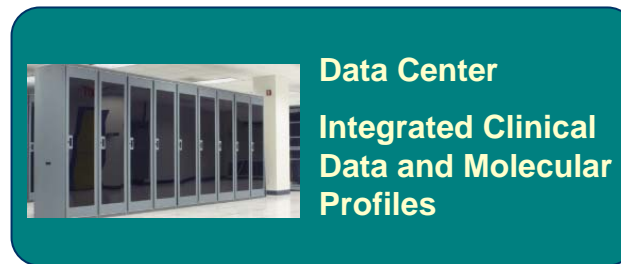
- Biospecimen Inventory
- Pathology Verification
- Molecular Analyte Production & QC



Aliquots

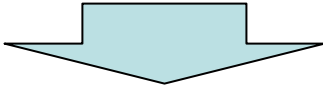


Analytical
Platforms
11 sites





Introduction - Sample Selection Process – on paper

- **Request for Information issued to identify interested biorepository custodians**
 - ~75 responses
 - **Follow-up phone calls to clarify/verify data provided**
 - **Site visits to institutions with estimated sufficient sample sets**
 - Did not include “audit” level review – i.e. going into freezers or databases
 - **Determination of source’s willingness to donate samples and participate in TCGA**
- 
- **Conclusion: TCGA needs could be met by 2 collections per cancer**



Outline

- Introduction and Perspective
- **Examples**
- Issues Encountered and NCI Best Practices
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Examples - Real numbers from beginning ops

- All logged sample number dropped 95 – 99%

	Repository 1	Repository 2
# Frozen samples logged in collection	5000+	1200+
# Samples meeting spec upon detailed (non-physical) review	1392	120
# Samples meeting physical specs	174	18

← Before full pathology review



Examples - Top 5 Sources of GBM Failure

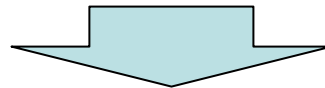
- **Matched normal germline DNA controls (blood or other) lacking**
- **Insufficient tumor cellularity in samples**
 - Tumor cellular composition too low
 - % necrosis too high
- **Specimen size too small**
 - Insufficient tissue to generate minimum required amount of DNA/RNA for all analyses
- **Molecular quality insufficient**
 - QC failure of DNA or RNA
 - Insufficient amount
- **Tumor not primary disease**
 - Samples derived from recurrent, i.e. previously treated GBMs (confounding issue: Rx-related effects)



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Best Practices - Lessons Learned

- **Quality of existing sample sets are typically overestimated by biobanks**
- **Collection of control samples is not routine in existing protocols**
- **Anatomic site-matched normal controls may be impossible to acquire**
- **Histologic quality does not guarantee molecular quality**
- **Data are lacking to define quality parameter cut-points accurately**
 - How does cellular composition affect genomics profiling?
 - Necrosis?
 - Yields of DNA, RNA per weight by tissue?



- **Biospecimen research is needed to understand effects of tissue variables on analysis data from different platforms**



Best Practices - Specimen Collection and Processing

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- ❑ **Tissue collection protocols need to start at the beginning**
 - ❑ Surgical /OR staff, biopsies, pre-op, consent
- **Handling appropriate for specimen type and study design**
- ✓ **Minimize collection and processing time**
- **Standard Operating Procedures**
 - Quality management system
 - Document all protocols
 - Training programs
- **Tag all specimens with human + machine readable labels**
 - Alphanumeric code
 - Barcode / RFID



Best Practices - Collecting and Managing Clinical Data

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- **Relevant clinical data**
 - Longitudinal data, clinical follow-up, outcomes
 - CTMS: patient tracking, study calendars, electronic data capture
- **Relevant epidemiologic data**
- **Informatics system for tracking all aspects of collection, processing and distribution**
 - caBIG™
- ✓ **Comply with privacy rules and human subjects regulations**



Best Practices - Monitoring and Storage

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- **Inventory tracking system**
 - Check in / check out, log all handlings
- **Store specimens in a “stabilized state”**
 - Appropriate temperature
 - Aliquots
 - Minimize thawing and refreezing
- **Disposal according to SOPs**
- Monitor and document storage equipment**
 - Temperature tracking critical



Best Practices - Record Keeping

- ✓ **IRB protocol governing collection**
 - ✓ Informed consent
 - ✓ Version year, tiered, permitted uses, re-contact OK
 - Exemptions
- ☐ **Subject vital status**
 - **Material Transfer parameters**
 - **Date of Collection**
 - Archived specimens prior to the era of molecular medicine
 - Prior to HIPAA Privacy Rule (April 14, 2003)



Best Practices - Packaging and Shipping

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- **Packaging procedures**
- **Records of specimen distribution**
- **Shipping procedures**
 - Appropriate temperature
 - Length of time
- ✓ **Shipping container electronic tagging**
 - ✓ Temperature, orientation
- **Train personnel**



Best Practices - Biosafety Procedures

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- **Identify risks and hazards**
 - Infectious, Radiation, Chemical
- **Record exposure incidents**
- **Provide treatment**
- **Indemnification agreements**



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Conclusions

- **Garbage In -> Garbage Out**
- **Make the up-front investment in samples – it will be worth it**
 - Tissue Banking is the protocol.
 - Too often it is considered a sideline
 - Treat your donors like clinical trial participants.
 - Track participants over time to get data: clinical follow-up / outcomes
- Do histopathology review (and molecular QC) prior to deposition
 - Categorize your samples
 - Discard what should be discarded

Print Close

Ardais RNA Pathology Verification Report			
Case ID	C10000013500	Prep Number	1
Case Information			
Primary Diagnosis from Donor Institution Pathology Report	Adenocarcinoma of breast, ductal		
Sample Information			
Diagnosis and Tissue			
Sample Pathology from Ardais Pathology Verification	Adenocarcinoma of breast, ductal		
Tissue of Origin of Diagnosis	Breast		
Site of Finding	Breast		
Sample Composition: Microscopic cell distribution by %			
Normal Cells (NRM):	0%		
Non-Neoplastic Lesional Cells (LSN):	0%		
Tumor Cells (TMR):	85%		
Tumor Cellular Stroma (TCS):	15%		
Tumor Hypo-/Acellular Stroma (TAS):	0%		
Necrosis (NEC):	0%		
Extended Composition/Comments	Tumor Stroma (Cellular): Desmoplastic reaction, Inflammatory cells		
Microscopic Appearance	Tumor		

Images

Pathology Image (4x)	Pathology Image (20x)	Gel Electrophoresis	RT-PCR

Bioanalyzer Ratio (28S/18S): 1.38

Fluorescence

Time (seconds)

Print Close

Inbox - Microsoft Outlook Ardais BIGR(R) - Microsof... RNA Pathology Verific...



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