

Biospecimen Research Network: Program Update

Helen M. Moore, Ph.D.

2010 BRN Symposium "Advancing Cancer Research Through Biospecimen Science" March 24, 2010



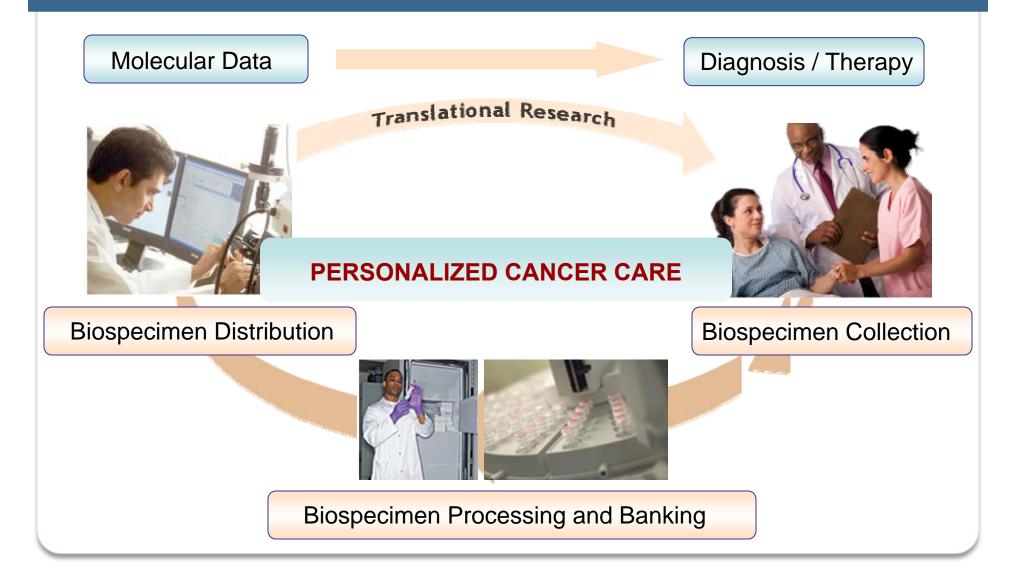


BBR Office of Biorepositories and Biospecimen Research

- To enable better, more reproducible cancer research
- To develop better products from that research
- To do so faster, better, more reliably
- To translate better research to better products for patients

Translational Research Promises to Advance Molecular Medicine for Cancer Patients

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Progress in Cancer Treatment and Research are Dependent on High Quality Human Specimens

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Identification of targets for drug development, treatment and prevention Identify biologic variations that determine drug efficacy and drug toxicity Defining markers for susceptibility, screening and reoccurrence Development of molecular based taxonomy of cancer Elucidation of molecular mechanisms of neoplasia Validation of new therapeutics

> All Depend On High-Quality, Annotated Human Biospecimens

Biospecimen Collection in the U.S.

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•Collection, procession, storage procedures differ

Degree and type of data annotation varies

Scope and type of patient consent differs

Access policies are lacking or unknown to potential users

Materials transfer agreement conditions differ

Supporting IT structures differ in capacity and functionality

 \rightarrow WIDE VARIATION IN QUALITY OF SPECIMENS AND DATA

Biospecimens and Cancer R&D

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- The lack of high-quality biospecimens is the number one obstacle for cancer research
- The accuracy of research data may be compromised when utilizing biospecimens of poor or unknown quality
- This issue must be addressed to move cancer R&D forward
- An issue for us all academics, government, industry

Molecular Research and Analyte Variation

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The lack of standardization of human biospecimens compromises the quality and the utility of molecular research and the advances in clinical medicine dependent on them.



Urgent Needs Being Addressed by OBBR

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BEST PRACTICES: state-of-the-science guidance for biobanking to harmonize procedures for collection, processing, storage and distribution of biospecimens

- The NCI Best Practices for Biospecimen Resources
- RESEARCH to better understand how pre-analytical variables affect the molecular integrity of the biospecimen

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- The NCI Biospecimen Research Network
- and...all of this integrated into a national biobank to facilitate cancer research: caHUB

New Web-based Format of NCI Best Practices ...and Revised Version Coming

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See the poster by Nicole Lockhart for more information

The Cancer Human Biobank (caHUB)

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Envisioned as:

A unique, centralized, non-profit public resource that will ensure the adequate and continuous supply of human biospecimens and associated data of measurable, high quality acquired within an ethical framework.

Please join us for the caHUB session tomorrow afternoon

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The NCI Biospecimen Research Network

 Supporting the development of Best Practices for collection, annotation, processing, and storage, based on <u>scientific evidence</u>

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- What evidence is already available about how different biospecimen collection, processing and storage procedures affect biospecimen molecular integrity?
- What new research is needed?
- *How do we accomplish that research?*

 Evidence-based biospecimen practices will underpin the entire caHUB effort Multiple pre-analytical variables can affect the molecular integrity of the biospecimen

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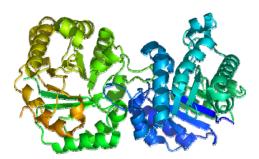
Variables (examples): Variables (examples): Time 0 Time at room temperature **Antibiotics Temperature of room Other drugs** Type of fixative Type of anesthesia Time in fixative **Duration of anesthesia** Rate of freezing Arterial clamp time Size of aliquots Medical/ Restocking Handling/ **Scientific** Surgical Acquisition Storage Distribution Unused Patient Processing Analysis Sample Procedures **Pre-acquisition Post-acquisition**

How Can Changes in Molecular Integrity of Biospecimens Affect Molecular Readout?

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Genomics

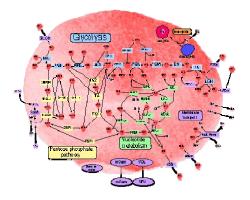




Proteomics

Changes in specific transcript levels based on ischemic time, not disease Lack of reproducibility of protein biomarkers in discovery research
Inconsistent IHC results in Research and Clinical Labs

Metabolomics



Inconsistencies in small molecule readouts, yielding results that point to the wrong pathway

The Biospecimen Research Network: Supporting Collaborative Research

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- Provide a forum for research results on how biospecimen variables affect molecular analysis:
 - The Biospecimen Research Database: Make existing and emerging biospecimen research data more accessible
 - This symposium: "Advancing Cancer Research through Biospecimen Science" <u>http://brnsymposium.com</u>

• Generate new research data:

- Extramural Research Programs
- IMAT Program "Innovative and Applied Emerging Technologies in Biospecimen Science" (RFA)
- Collaborate with other programs to facilitate biospecimen research and evidence-based biospecimen practices

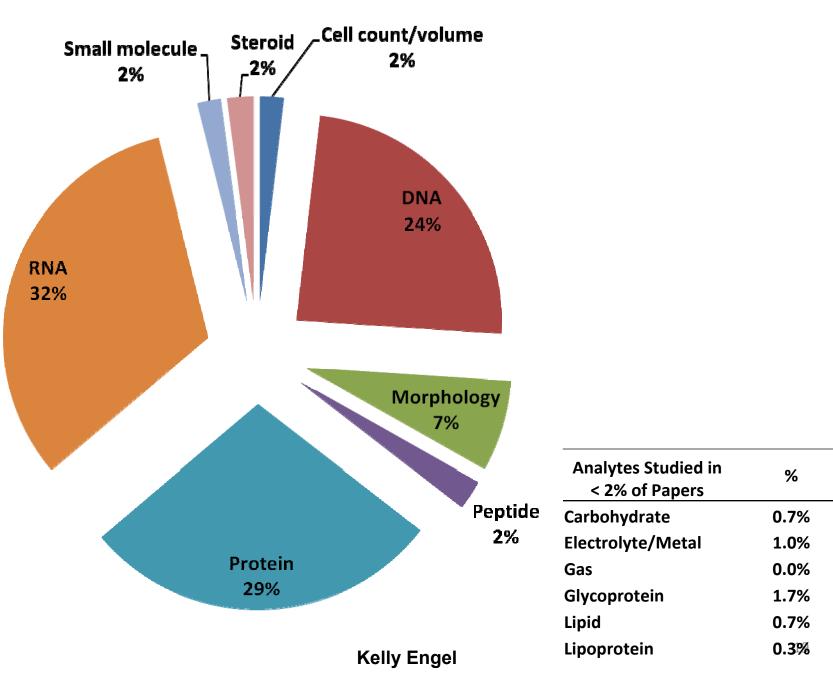


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Learn more	Paper and Study Details					
2009 BIOSPECIMEN RESEARCH NETWORK SYMPOSIUM	PubMed ID: 16288005 Publiced					
March 16-18, 2009	Lips Esther H, Dierrsen Jan WF, van Eijk Ronald, Oosting Jan, Eilers Paul HC, Tollenaar Rob AEM, de Graaf Eelco J, van't Slot Ruben, Wijmenga Cisca, Morreau Hans, van Wenzel Tom					
Home Search			put genotyping and loss-of-heterozygosity detection in formalin- ded (FFPE) tumors using single nucleotide polymorphism arrays.			
<u>Quick Search</u>	Cancer Re	s, 2005, <u>Vol</u>	<u>. 22.</u> Page 10188			
Simple Search	Review Paper?		No			
Advanced Search Experimental Factor Search Curator Login	Purpose of Paper:		To determine if genomic DNA isolated from formalin fixed paraffin embedded (FFPE) tissue is applicable for genotype and loss of heterozygosity (LOH) analysis via single nucleotide polymorphism (SNP) arrays.			
0	Conclusion of Paper:		Results of SNP and LOH analyses were nearly identical among FFPE and frozen specimens. The authors conclude that genome wide genotyping using FFPE tissue is reliable, producing results that are reproducible with frozen specimens.			
	Studies	;				
	Detail	Specimen:	Tissue / Colorectal / Formalin / Normal			
	<u></u>	_	DNA - SNP assay /			
		Findings :	The occurrence of SNPs in FFPE and frozen matched controls were nearly identical (99.9%). Of the 8 LOH regions examined only 2 were not identical among FFPE and frozen specimens (located on chromosomes 3 and 8). Genotyping results were identical among the two SNP arrays (BeadArray and GeneChip).			
	<u>Detail</u>	Specimen:	Tissue / Colorectal / Formalin / Normal			
		Platform:	DNA - DNA Sequencing /			
		Findings :	DNA sequence analysis confirmed SNP results.			
	1	-				

Biospecimen Research Database Update: 300 papers now curated

Journals with 10 or More Papers Identified for BRD Inclusion 0 10 20 30 40 60 70 80 90 100 50 **Clin Chem Clin Chem Lab Med J Clin Pathol Diagn Mol Pathol Am J Clin Pathol Cancer Epidemiol Biomarkers Prev** J Histochem Cytochem Appl Immunohistochem Mol Morphol Mod Pathol **Am J Pathol** J Clin Microbiol J Mol Diagn Proteomics Lab Invest Number of Papers in the BRD Number of Papers Awaiting Curation

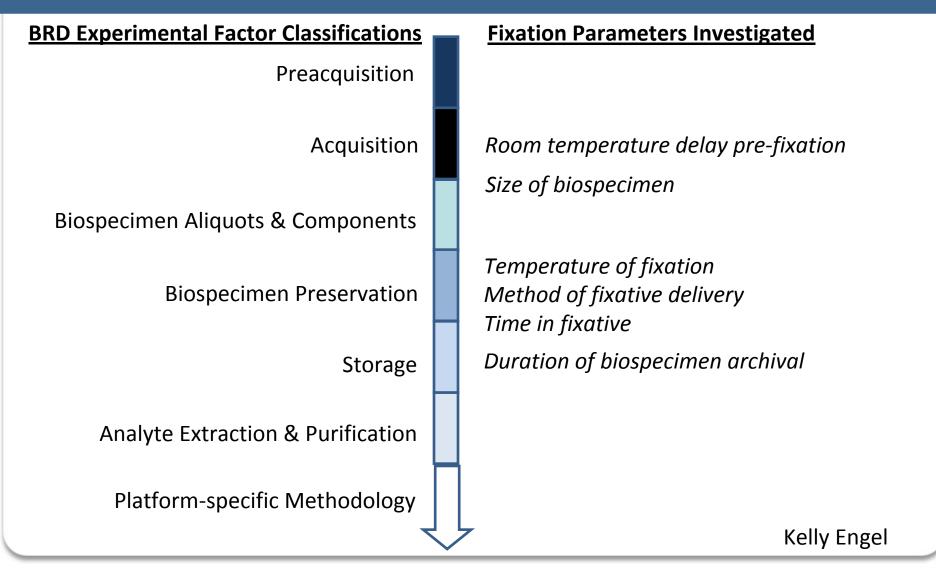
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Analyte Distribution of BRD Paper Entries

Test-Driving the Biospecimen Research Database: FFPE

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Conclusions supported by 3 or more papers populating the BRD

	No. of			
Analyte	BRD Papers	Consensus		
Fixation Parameter: Biospecimen size				
DNA	3	PCR results of small biospecimens (2-10 mm diameter) were		
		favorable to larger biospecimens.		
Fixation parameter: Time in fixative				
DNA 6	6	PCR analysis was optimal in biospecimens fixed for 2-48 h, with		
		adverse effects reported after fixation for ≥72 h.		
RNA 3 3	3	Evidence of RNA degradation was observed in specimens fixed for		
		1-72 h.		
	3	mRNA transcript stability was analyte-specific, with fixation		
		thresholds potentially influenced by platform sensitivity and		
	amplicon length. Quantified analytes included: COX-1, beta-actin,			
		MART, MMP-1, MMP-1, VEGF, p21, EGFR, C-BCR.		
Protein 5	5	Protein immunoreactivity was stable in biospecimens fixed for 6 h-		
	8 d. The antigens investigated included: p27Kip1, ER, PR, AR, c-erb-			
	B2, HER/neu, EGFR, MMR-1, VEGF, p53, PCNA, Ki-67.			
Fixation pa	rameter: Archiv	al of formalin-fixed paraffin tissue blocks		
DNA 4	4	PCR success and efficiency (of 90-435 bp fragments) were not		
	impacted by paraffin block archival at room temperature for 1 wk-			
		8 y.		
RNA 3	3	RNA degradation was more extensive in blocks stored for 3.5-17 \ensuremath{y}		
		compared to those stored for 1 y or less.		
	3	RT-PCR success rate decreased by 0-20% after 1-10 y, 30-50% after		
		10-30 y, and 60% after 40 y of paraffin block archival compared to		
		fresh blocks, although amplicon length also influenced RT-PCR		
3	success. The analytes investigated included hepaptitis C, beta-			
		actin, C-BCR.		
	3	Real-time qRT-PCR analysis of paraffin blocks archived for 1-8 y		
		was successful; while analysis was impaired for blocks stored for		
		11 y or longer. The analytes investigated included LDHA, RPL32,		
		beta-actin, RPL13A, RPLO, CYP1, GUS, TBP, TFRC		
Fixation parameter: Archival of formalin-fixed, paraffin-embedded, slide-mounted sections				
Protein 3	3	Immunostaining was altered in slides stored for 3 mon-3 y at room		
		temperature compared to freshly cut sections. Alterations in		
		immunostaining intensity and duration threshold were antigen-		
		specific. The antigens investigated included ER, PR, HER-2,		
		Chromagranin, CD3, Vimentin, EGFR.		

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Central Themes for BRN Research

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- "Bridging the gap" between existing clinical practice for biospecimens and emerging technologies for personalized diagnostics and therapies
 - Tissue preservation variables and their impact on downstream applications (e.g., HER2)
 - Robotic surgery vs. manual surgery for prostate are tissues harvested from robotic surgery suitable for advanced biomarker detection?
- Defining the most significant variables for prospective collection of tissues, blood, and body fluids
 - Effects of pre-acquisition variables and biomolecule extraction methods on biomolecule analysis results in blood
- Developing evidence-based biospecimen quality indicators for specific analytical platforms
 - How to assess whether a banked specimen is suitable for a specific molecular analysis approach?

BRN Research Funding Program: "Biospecimen Research for Molecular Medicine"

Program aims:

- (1) Develop innovative approaches to the control, monitoring and assessment of biospecimen quality
- Investigator-driven Contract Research
- "Biospecimen Molecular Integrity" RFP issued 10/08, four research contracts awarded, research is in progress
- Second issue under ARRA funds, proposals were due 12/09, contracts in negotiation

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- (2) Systematically define the impact of key pre-analytical variables in human biospecimens of specific type on downstream molecular data generated from specific molecular analysis platforms
- Program-directed Research: Contracts in negotiation

Biospecimen Molecular Integrity: Program Approach

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- Encourage the best research from the field
 - RFP modeled on a Broad Agency Announcement
 - The "right" solutions to the many research problems out there are many
 - Different technology platforms may be applied
 - They do not all cost the same

• This can be very challenging research to perform

- Biospecimen collection and molecular analysis procedures must be known and reproducible
- Annotation and Quality Control are paramount

Biospecimen Molecular Integrity: Research Contracts

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- "Investigations into the Effects of Blood Specimen Handling Procedures on Protein Integrity"
 - Chris Becker, PhD, PPD Biomarker Discovery Sciences
- "Credentialing Plasma and Serum Biospecimen Banks for Proteomics Analyses"
 - Katy Williams, PhD, University of California San Francisco
- "Intrinsic and Extrinsic controls for FFPE tissue"
 - David Rimm, MD, PhD, Yale
- "Effects of Biospecimen Integrity, Intratumoral Heterogeneity, and Analytical Variance on Microarray-Based Pharmacogenomics Tests of Breast Cancer"
 - W. Fraser Symmans, MD, MD Anderson (PI), Christos Hatzis, PhD, Nuvera Biosciences (Co-PI)
 - More to be announced soon

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(1) Develop innovative approaches to the control, monitoring and assessment of biospecimen quality.

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(2) Systematically define the impact of key pre-analytical variables in human biospecimens of specific type on downstream molecular data generated from specific molecular analysis platforms.

Tissue Acquisition Variables Project

Tissue Acquisition Variables Project: Directed Research

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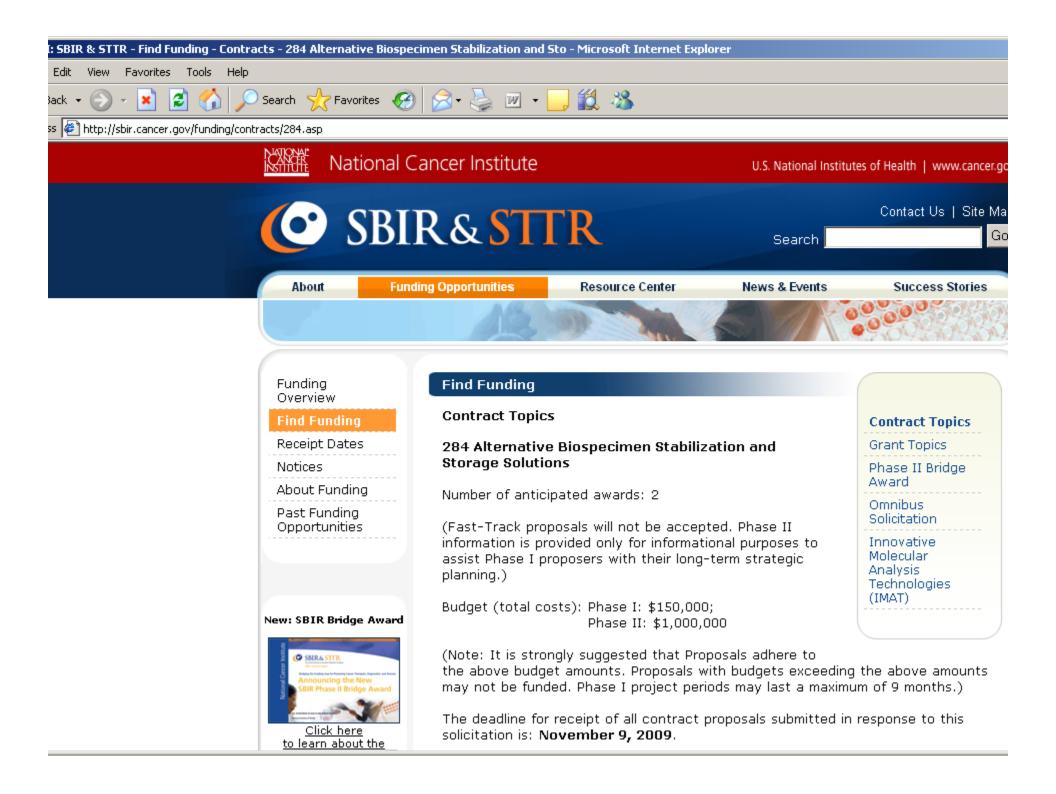
- Aimed at understanding the variability introduced by different tissue fixation and processing procedures
- Multi-site study:
 - Establish the state of practice in cancer tissue collection, processing and storage
 - Study the effect of different such practices on biospecimen molecular integrity
 - Hone in on the procedural variations that cause the most significant variations in molecular integrity
 - Use this data to inform the development of evidencebased Standard Operating Procedures (SOPs)

Tissue Acquisition Variables Project: Directed Research

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- Contracts with Tissue Source Sites are in negotiation
- Additional funding opportunities to be announced
 - To propose robust ways of analyzing the tissues to reveal the molecular readout induced by preanalytical variables such as time to fixation, fixation time, and different processing parameters





How do we change the landscape of research through Biospecimen Science?

Through building a new knowledge base in biospecimen science

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• New research

- Communication of research through publications, meetings, and the Biospecimen Research Database
- Through changing how research using human biospecimens is published
 - Hear Scott Jewell's talk later today about "Biospecimen Reporting for Improved Study Quality (BRISQ),"
 - Effort grew out of a workshop held at this symposium last year
 - The basic idea is that researchers need to say more in their publications about where the biospecimens forming the bass of their study came from and how they were treated
 - Potentially huge impact on making research results more reproducible
- Through this meeting and all of you who are passionate about this area

A Few Program Notes



- Thank you for joining us here today!
- Please sign up for "Meet the Experts" lunch tables
 - See sign up sheets outside and flyers in your program book
 - Thank you to all the "Experts" leading these tables!
- Please join us tonight for a poster session and reception starting at 5:15
- Have a great meeting!

Thanks to...



- The 2010 BRN Symposium planning team:
 - Sherilyn Sawyer, Jim Vaught, Andrea Kelly, Renate Myles, Frank Bajowski, Jennifer Kostiuk
- The two fellows who helped to get this symposium off the ground 2 years ago:
 - Asha Collins and Mark Lim
- The Biospecimen Research Database core team:
 - Kelly Engel, Andrew Breycheck, Ian Fore
- All of our incredible OBBR team, past and present
- Our extended team at SAIC-Frederick who work with us to develop and manage research contracts
- Carolyn Compton our fearless leader!

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3RD ANNUAL BIOSPECIMEN RESEARCH NETWORK SYMPOSIUM

ADVANCING CANCER RESEARCH THROUGH BIOSPECIMEN SCIENCE

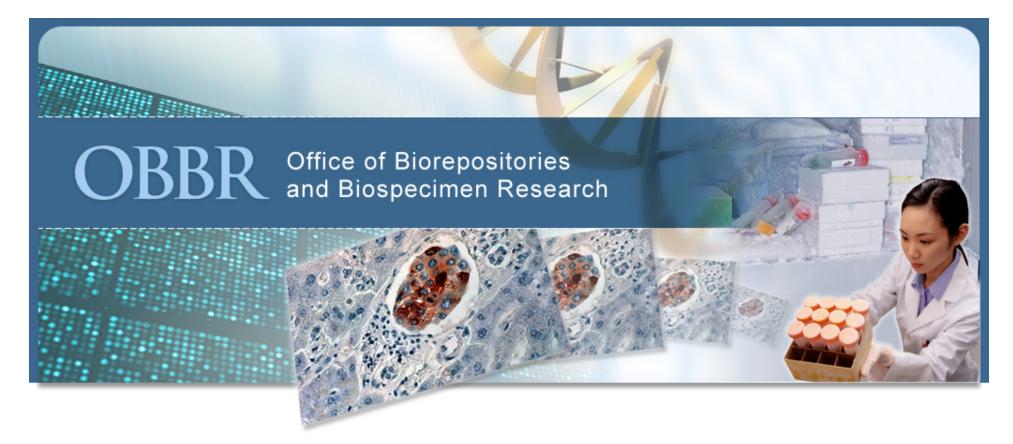
March 24-25, 2010

Bethesda North Marriott Hotel & Conference Center Bethesda, Maryland http://www.brnsymposium.com

For symposium updates, please send your contact information to biospecimens@mail.nih.gov

U.S. DEPARTMENT OF HEALTH AND HUMAN SERVICES

National Institutes of Health



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