

# Assessing the Quality of Tissue

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I. Standards for assessing preservation of macromolecules (RNA, protein) in tissue

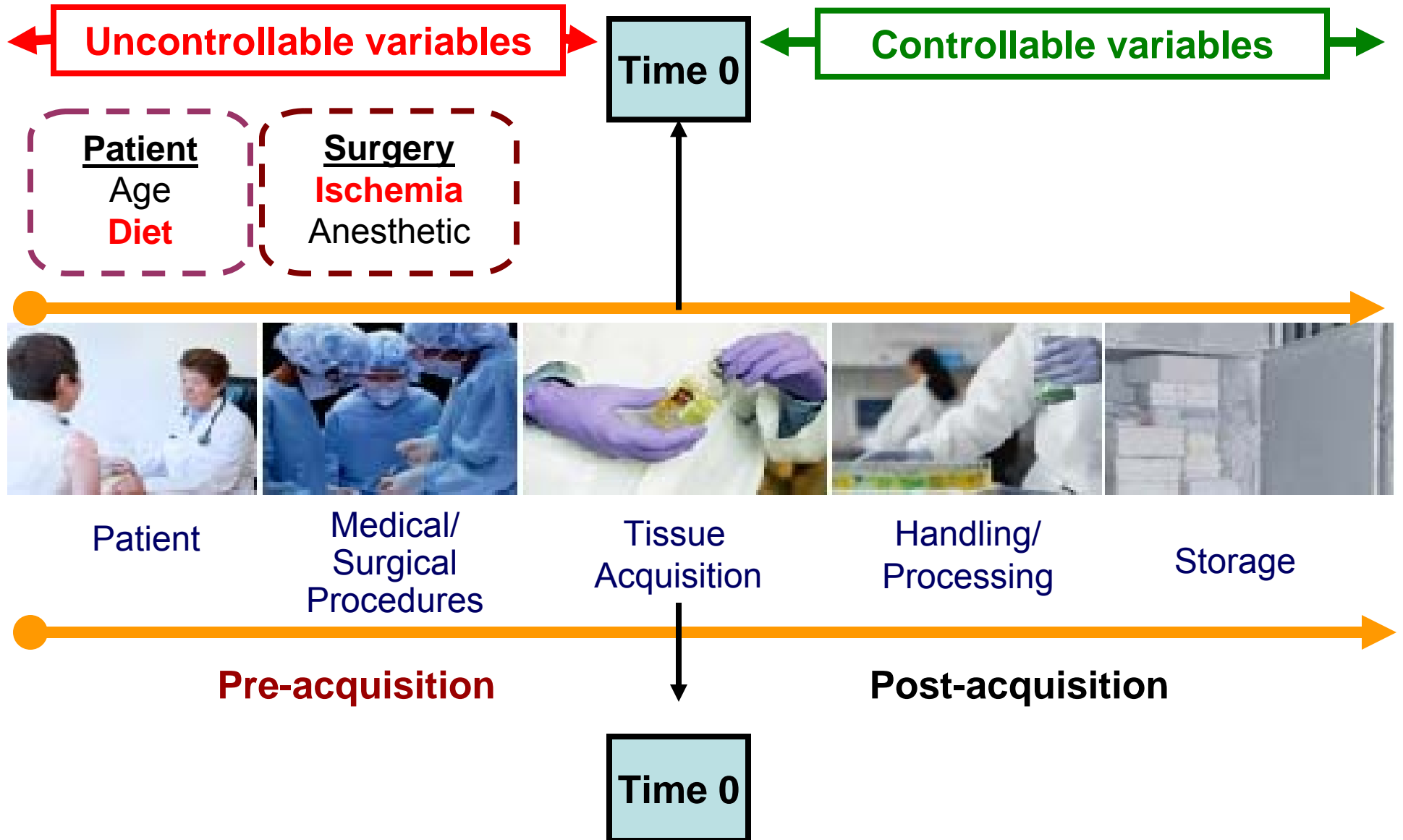
II. A specification for tissue localization experiments (MISFISHIE)

NCI Biospecimen Best Practices Forums

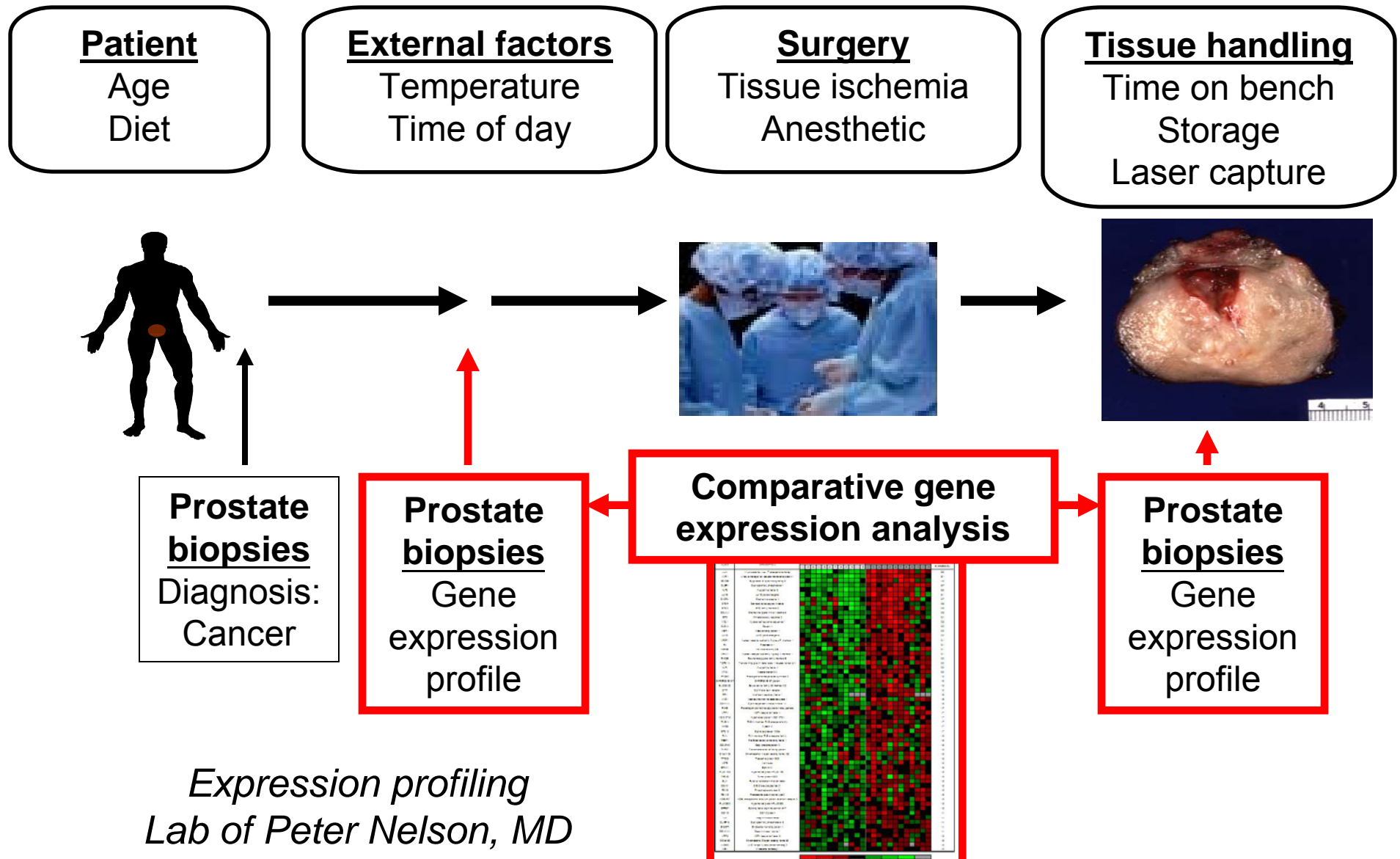
Seattle, WA, January 28, 2008

# Lifecycle of a Biospecimen

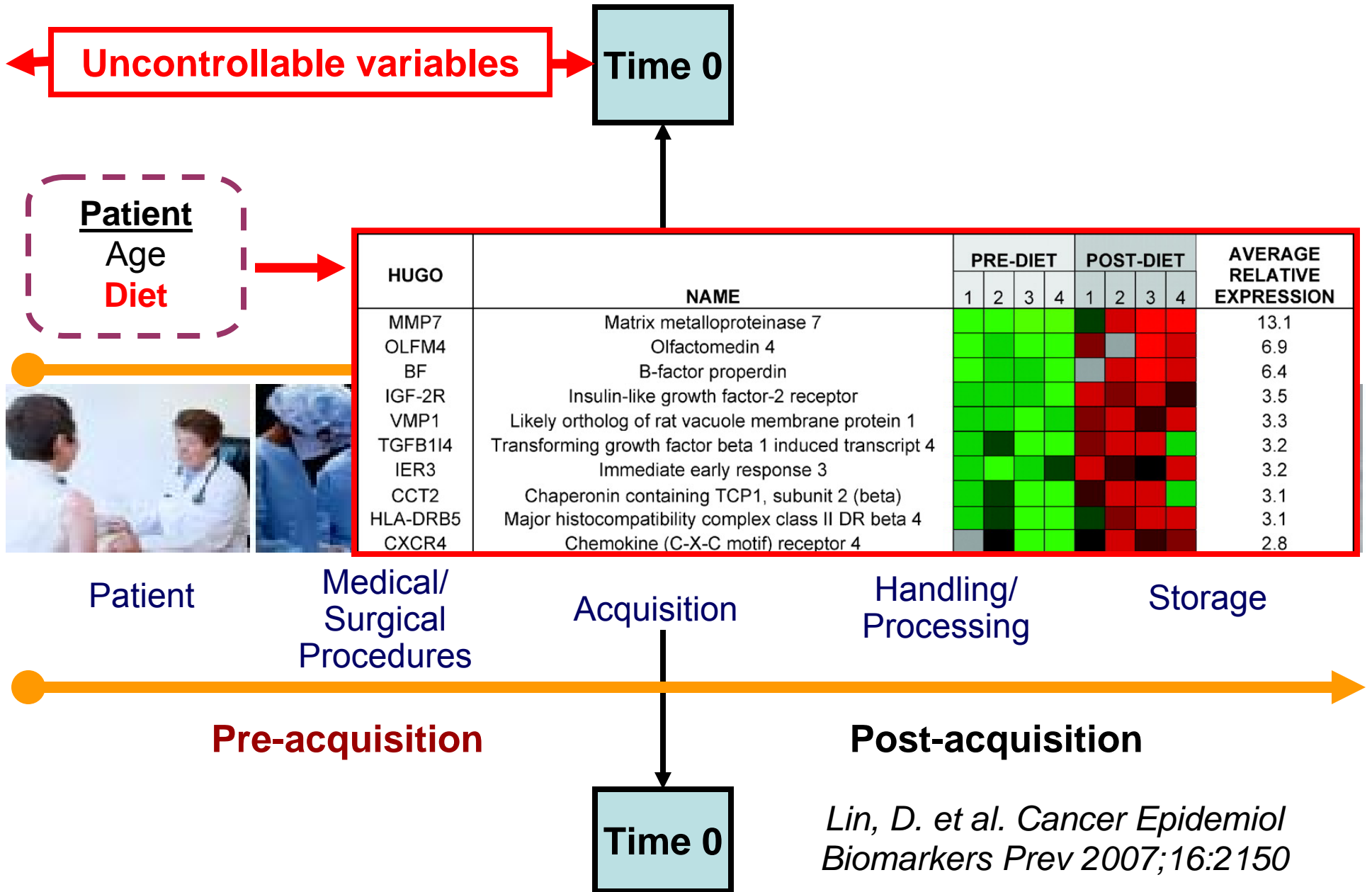
Factors potentially having significant effect on gene expression



# Design of experiments to evaluate effect of variables on gene expression



# Effect of 6 weeks of high-caloric diet

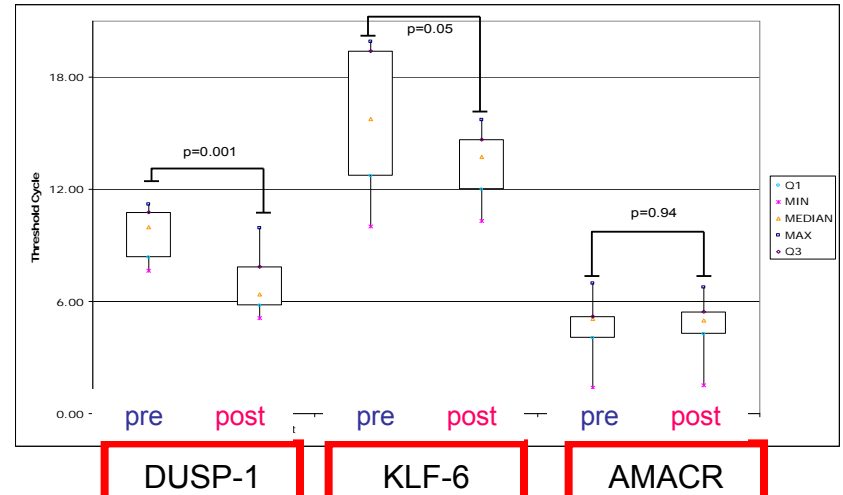
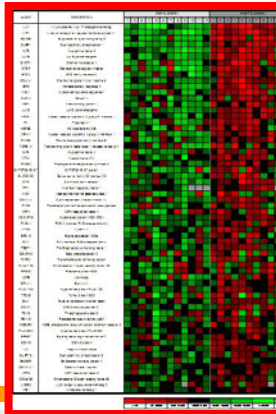


# Effect of ischemia during surgery on tissue

Uncontrollable variables

Time 0

Surgery  
Ischemia  
Anesthetic



Patient



Medical/  
Surgical  
Procedures



Acquisition



Handling/  
Processing



Storage

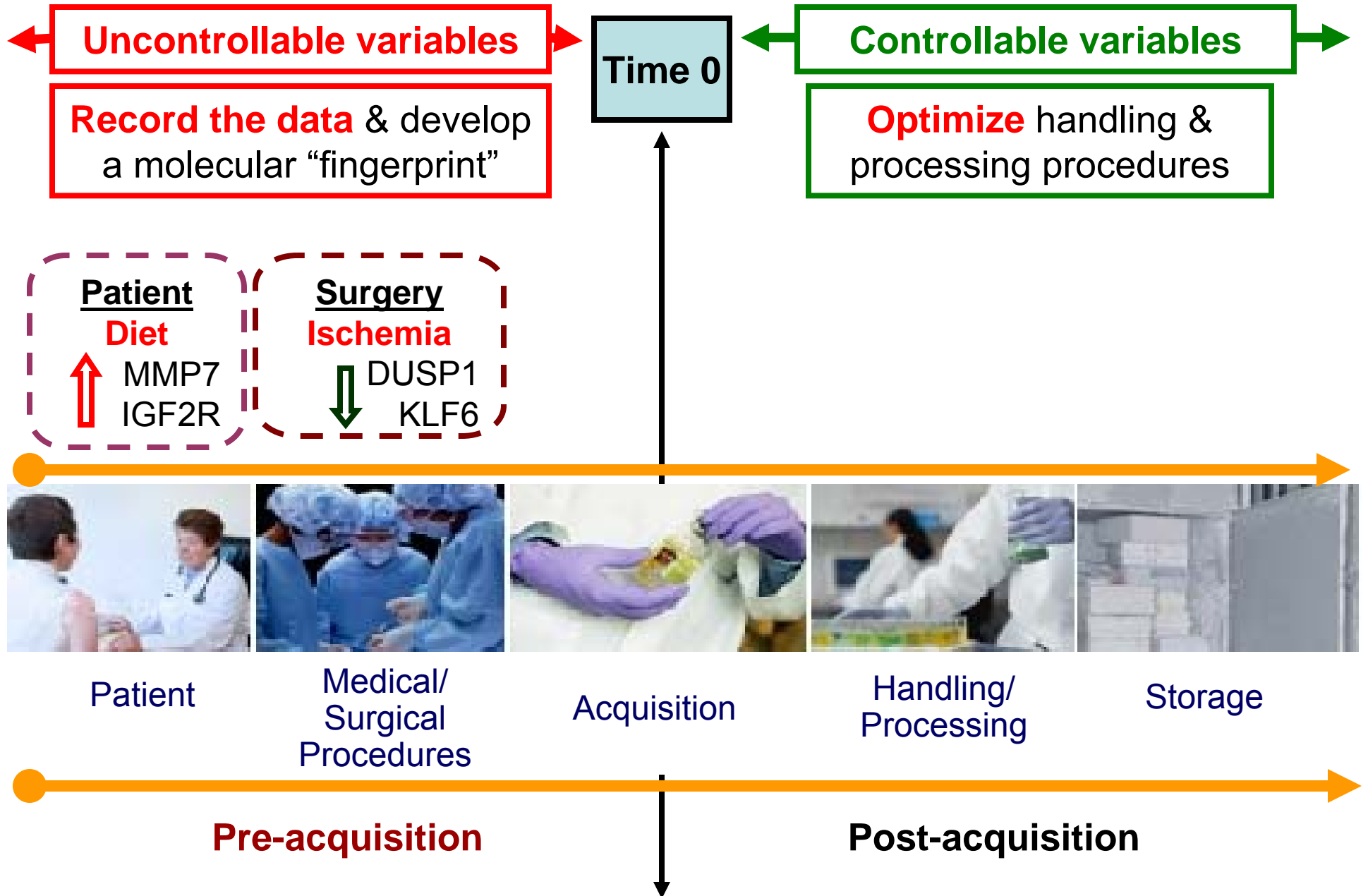
Pre-acquisition

Post-acquisition

Time 0

Lin, Coleman, Hawley..JCO '06;  
10;24(23):3763-70

# Lifecycle of a Biospecimen



# Lifecycle of a Biospecimen

Type of surgery	RRP	Laparoscopy	Cystoprostatectomy	
Serum PSA	6.92	_____	_____	_____
Prior therapy (circle)	None	Hormone?	ChemoRx ?	Other
Nerve sparing (circle if yes)	Right ? no	Left ? no	_____	_____
Times of day tissue	Out of pt	First sample frozen	Last sample frozen	_____
Time	6:40	7:15	7:20	_____
Nodular parenchyma	None	Rare Multiple	Small Medium	Large
Clinical stage?	T1a	T1b	T1c	T2



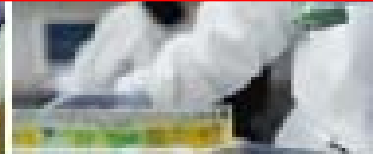
Patient



Medical/  
Surgical  
Procedures



Acquisition



Handling/  
Processing



Storage

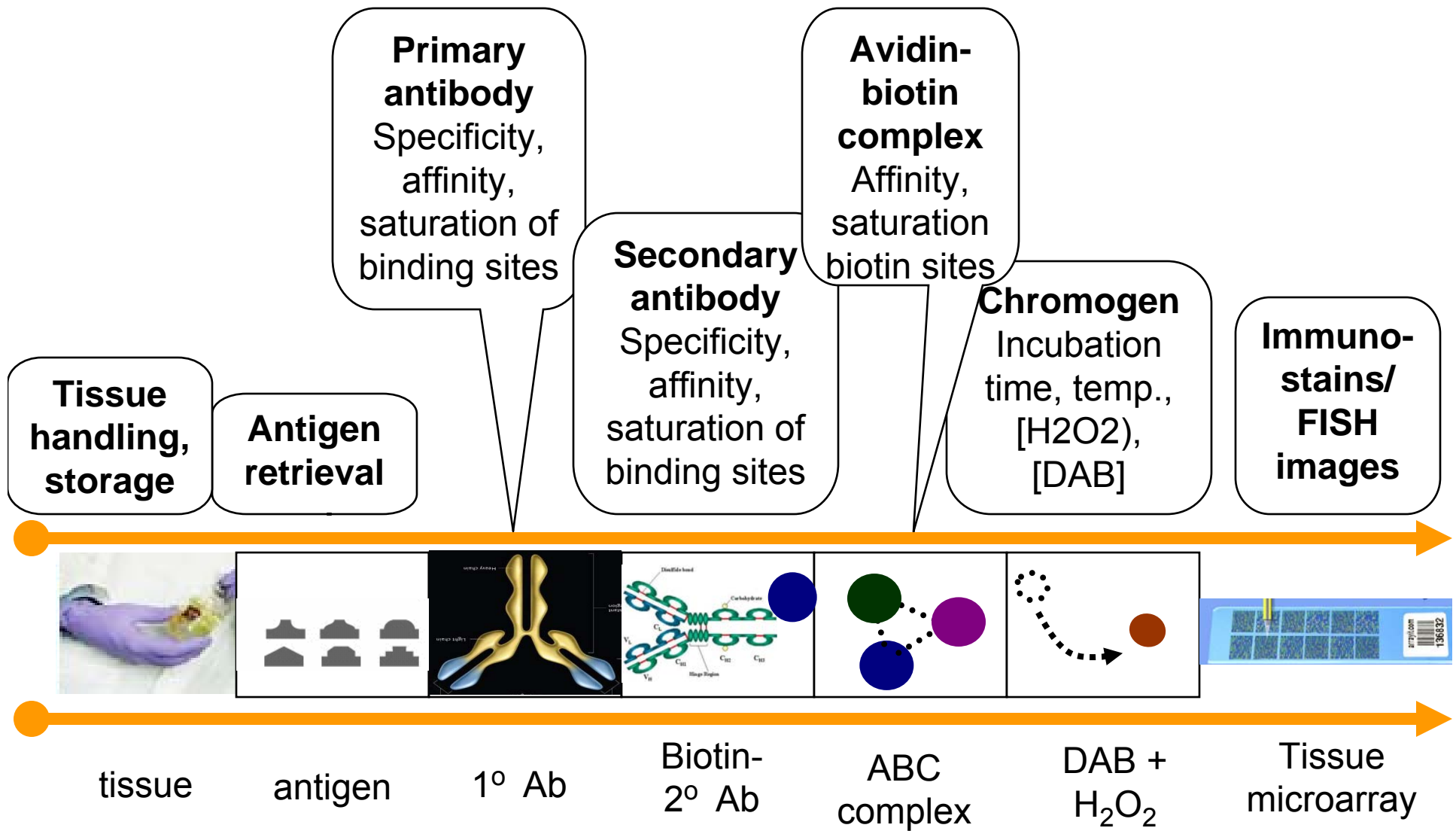
Pre-acquisition

Post-acquisition

Time 0

# Factors affecting immunohistochemical stains

## Controllable variables





## **MISFISHIE** – *report all data in your experiments*

- **M**inimum **I**nformation **S**pecification **F**or *In Situ* **H**ybridization and **I**mmunohistochemistry **E**xperiments (MISFISHIE)
- The purpose is to "**Insure that the minimum information that a researcher at a different lab needs to reproduce or evaluate the experiment is provided.**"
- MISFISHIE does **not** specify the data format, merely **what** information must be communicated
- Based on principles of **MIAME** (specification for reporting expression microarray data)

*Deutsch E, et. al. OMICS. 2006;10(2):205-8 (PMID: 16901227 ); Nature Biotechnology (in press)*

Project Information

Resources

Software Links

Minimum Information Specificat  
Immunohistochemistry Experim

## Major Sections:

1. Experiment Design
2. Specimens, Treatments
3. Probes or Antibodies
4. Staining  
Protocols/Parameters
5. Imaging Data and  
Parameters
6. Image Characterizations

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Web [Images](#) [Video](#) <sup>New!</sup> [News](#) [Maps](#) [more »](#)

MISFISHIE

Google Search

I'm Feeling Lucky

[Advanced Search](#)  
[Preferences](#)  
[Language Tools](#)

**32 articles from:**

Am J Clin Path  
Am J Surg Path  
Cancer Res  
J Clin Path  
Lab Invest  
Modern Path  
Molecules & Cells  
Nature  
Science



## Section 3. Reporters (Antibodies, Probes)

Unambiguous reporter identification, ideally genomic

Full sequence or clone id of the reporters

Protocol for obtaining exact reporter (purchase from..., create, etc.)

Other important attributes (e.g., mono- or polyclonal, organism in which antibody generated)

**Information not supplied (25% of articles):**

Antibody clone numbers and/or catalogue numbers

Conditions of antigen retrieval (buffer, heat duration)

Antibody Name	Locus Link ID	Origin	Vendor Details
CD49a	3672	BD PharMingen	<a href="http://www.bdbiosciences.com/external_files/pm/doc/tds/ihc/live/web_enabled/75311E_550568.pdf">http://www.bdbiosciences.com/external_files/pm/doc/tds/ihc/live/web_enabled/75311E_550568.pdf</a>
CD90	7070	BD PharMingen	<a href="http://www.bdbiosciences.com/external_files/pm/doc/tds/ihc/live/web_enabled/74851E_550402.pdf">http://www.bdbiosciences.com/external_files/pm/doc/tds/ihc/live/web_enabled/74851E_550402.pdf</a>

## Section 4. Staining

Detection Method (number of reporters, detection reagent & systems)

Staining protocol (enough to reproduce?)

Details about positive and negative controls

**Information not supplied (25% of articles):**

Was nonspecific protein binding blocked?

Was endogenous peroxidase inhibited?

**Negative controls?**



The screenshot shows the header of a webpage. On the left is the logo for the Institute for Systems Biology, which includes the text 'INSTITUTE FOR Systems Biology' and a circular icon with a grid of dots. To the right of the logo is the text 'SCGAP UESC Protocols'. Below the logo and title is a blue horizontal bar with the text 'SCGAP Consortium Portal' and 'NIH/NIIDDK Main'. To the right of the bar, there is a paragraph of text: 'Below is a listing of the various protocols the UESC group uses to generate the Immunohistochemistry and Microarray data presented here. You can see details of a given protocol by following the link from the name.'

<http://scgap.systemsbiology.net/resources/protocols.php>

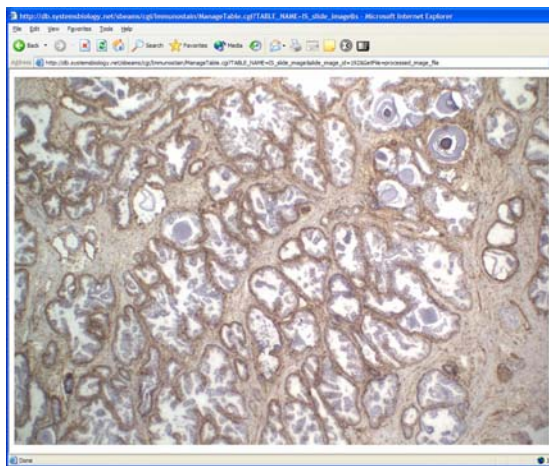
## Section 5. Imaging Data

The digital images for each assay (can the images be downloaded to your computer and evaluated ?)

**Information not supplied (85+ % of articles):**

Images of all specimens (for reader to evaluate for her/himself)

(for example) Was “nonspecific staining” expression of an antigen by cells not expected to stain?



# Specimens # Stains # Images

**Summary by Organism:**

Human	135	687	2222	<a href="#">[Immunostain Summary]</a>
Mouse	11	72	163	<a href="#">[Immunostain Summary]</a>

**Summary by Tissue Type:**

Human Bladder	28	315	1025	<a href="#">[Immunostain Summary]</a>
Mouse Bladder	3	24	75	<a href="#">[Immunostain Summary]</a>
Human Prostate	107	372	1197	<a href="#">[Immunostain Summary]</a>
Mouse Prostate	8	48	88	<a href="#">[Immunostain Summary]</a>

## Conclusions

- To determine if a molecular profile of experimental tissue is abnormal:
  - For **uncontrollable variables** (diet, anesthesia, patient age, duration of surgery, tissue ischemia) we need to **develop a molecular fingerprint** and subtract that fingerprint profile from the experimental gene expression profile
  - For **controllable variables** (rapidity of processing, freezing, storage), we need to **optimize handling** of specimens to minimize the effect of steps that most affect gene expression profiles
- To achieve these goals, we need to
  - Obtain all relevant information from all experimental data sets (MISFISHIE specification)

# Urologic Epithelial Stem Cell Group

# Questions ??

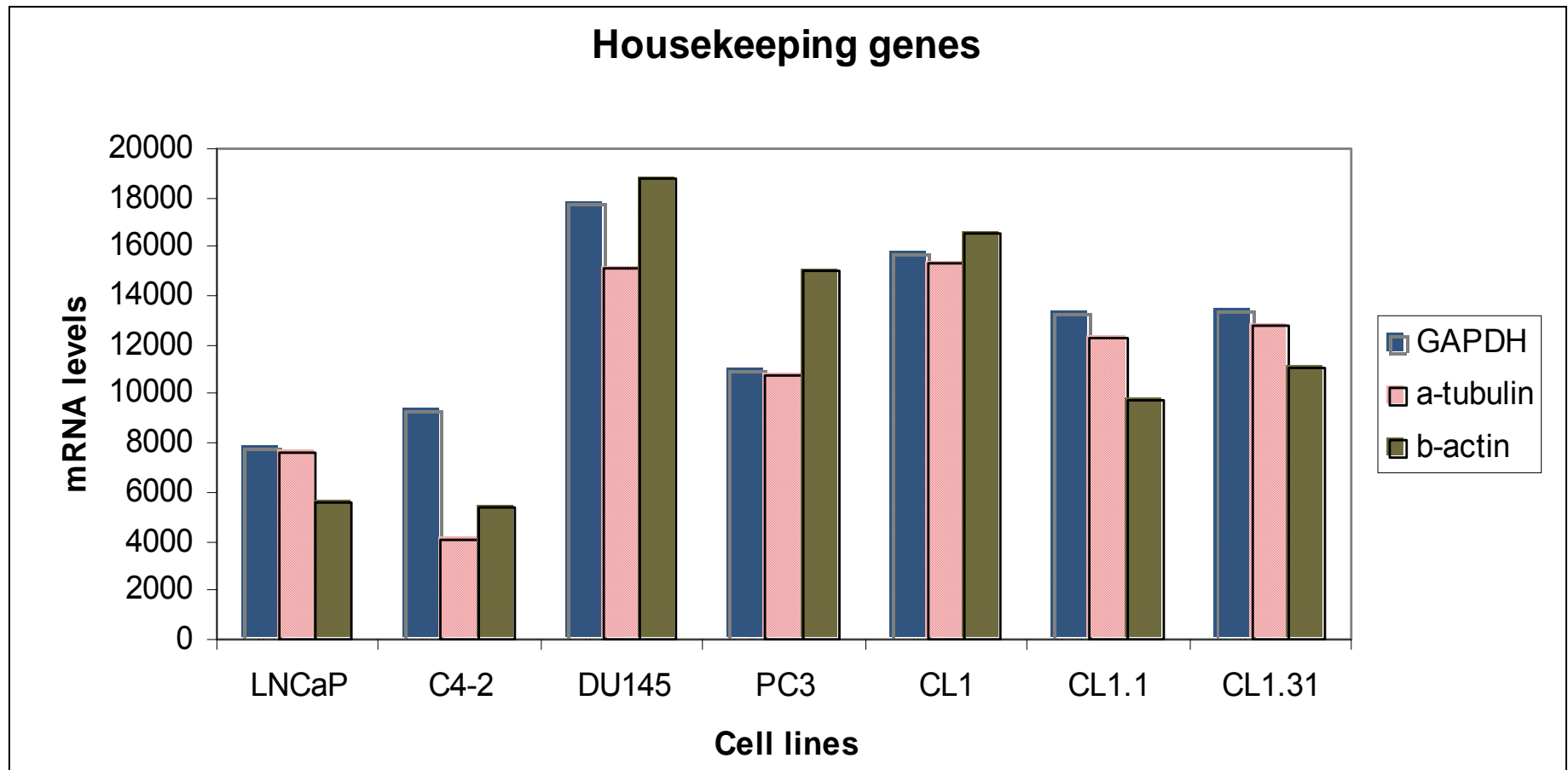
Alvin Liu, PhD: PI,  
Molecular Biologist

Eric Deutsch, PhD:  
Bioinformatics

Laura Pascal, PhD:  
Post-doc, Stem cell biology

NIDDK U01 Stem Cell Consortium (Prostate & Bladder)

Is there a set of “housekeeping” genes/proteins that can be used as a metric for preservation?



**Range of mRNA:**

$\beta$ -actin: 5,500 – 19,000

$\alpha$  tubulin: 4,000 – 15,500

GAPDH: 8,000 – 18,000

There is no single “housekeeping gene” and probably not a generic set of genes that can be used as a general metric for RNA preservation.