

EDRN BIOMARKER DEVELOPMENT

Sam Hanash

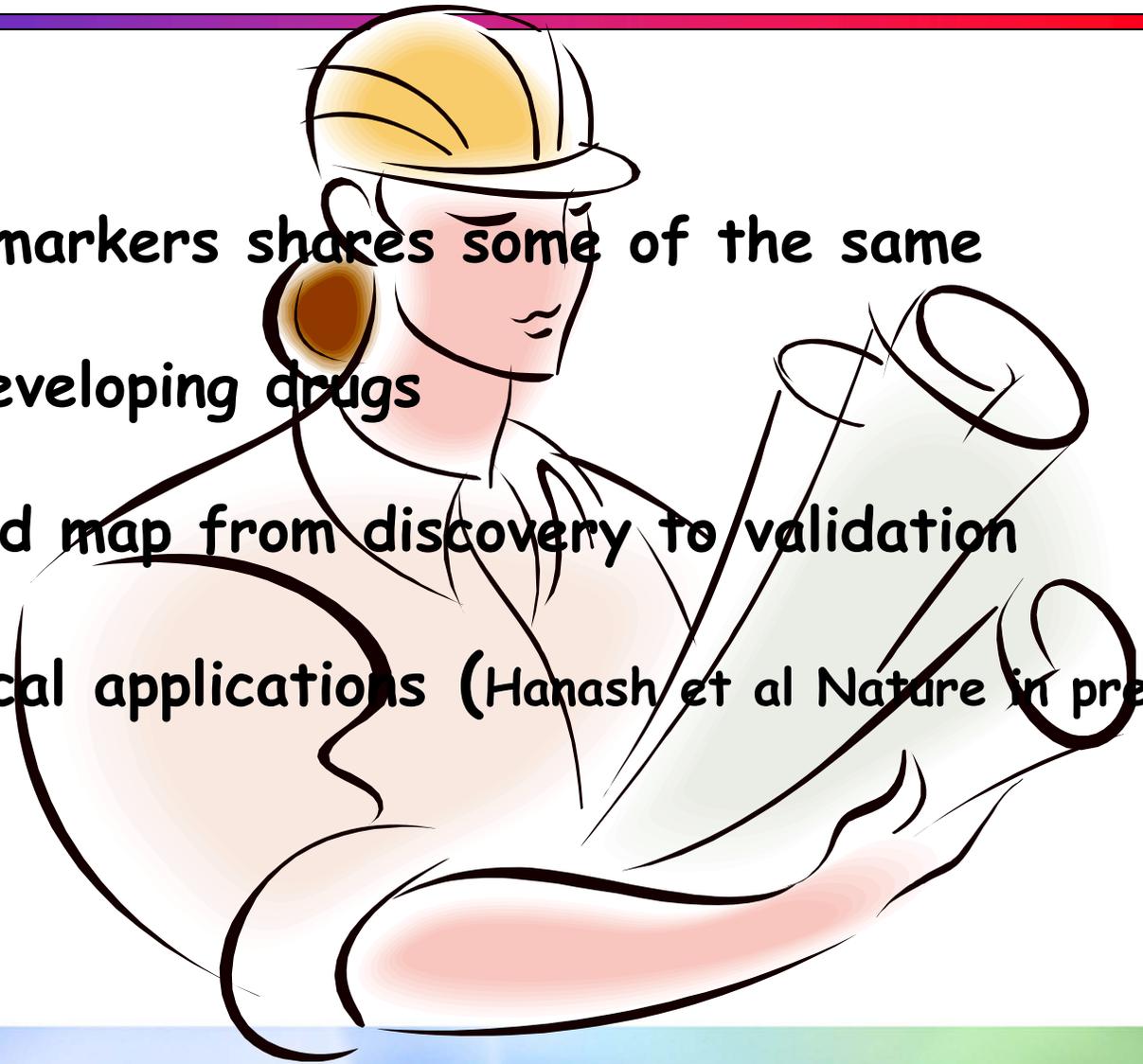
Fred Hutchinson Cancer Research Center

CONTENTS

- 1- Two studies that progressed from discovery to blinded validation in a context of early detection**
- 2- Collaborative work within and outside of EDRN Development laboratories**
- 3- Leveraging resources outside of EDRN**

Why so few biomarkers to date?

- Developing biomarkers shares some of the same challenges as developing drugs
- Requires a road map from discovery to validation for defined clinical applications (Hanash et al Nature in press)



Autoantibodies as biomarkers for early cancer detection

Immune response to tumor antigens

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- Occurs early during tumor development: may allow early cancer detection
 - Is not limited to mutated proteins
 - May involve aberrantly expressed proteins eg oncofetal antigens
 - Epitopes may result from post-translational modifications eg glycosylation

ANTI-ANNEXINS I & II ANTIBODIES IN LUNG CANCER

	# subjects	Annexin I Antibody +	Annexin II Antibody +
<u>Lung Cancer</u>	54	16	18
Adenocarcinoma	30	12	11
Squam cell carcinoma	18	3	4
Small cell carcinoma	4	1	2
Large cell carcinoma	2	0	1
<u>Other cancer types</u>	60	6	0
<u>Other controls</u>	61	0	0
Healthy subjects	51	0	0
Chronic lung disease	10	0	0

CARET Validation strategy

Blinded validation study

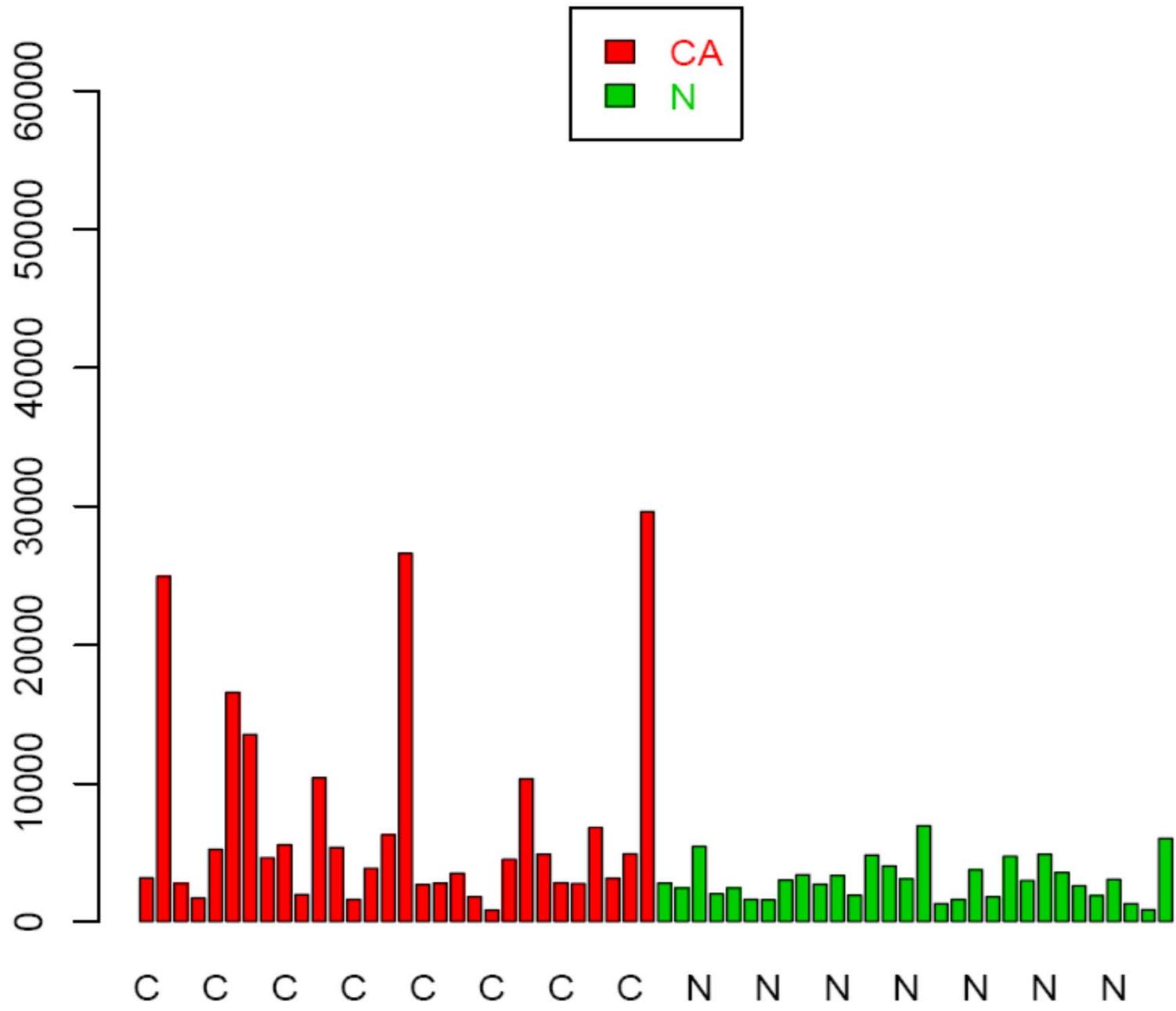
Approach: Protein microarrays

Contents: Natural proteins derived from tumor cell line(s)

Samples: Collected ~1 yr prior to lung cancer dx from
100 cases and 100 matched controls

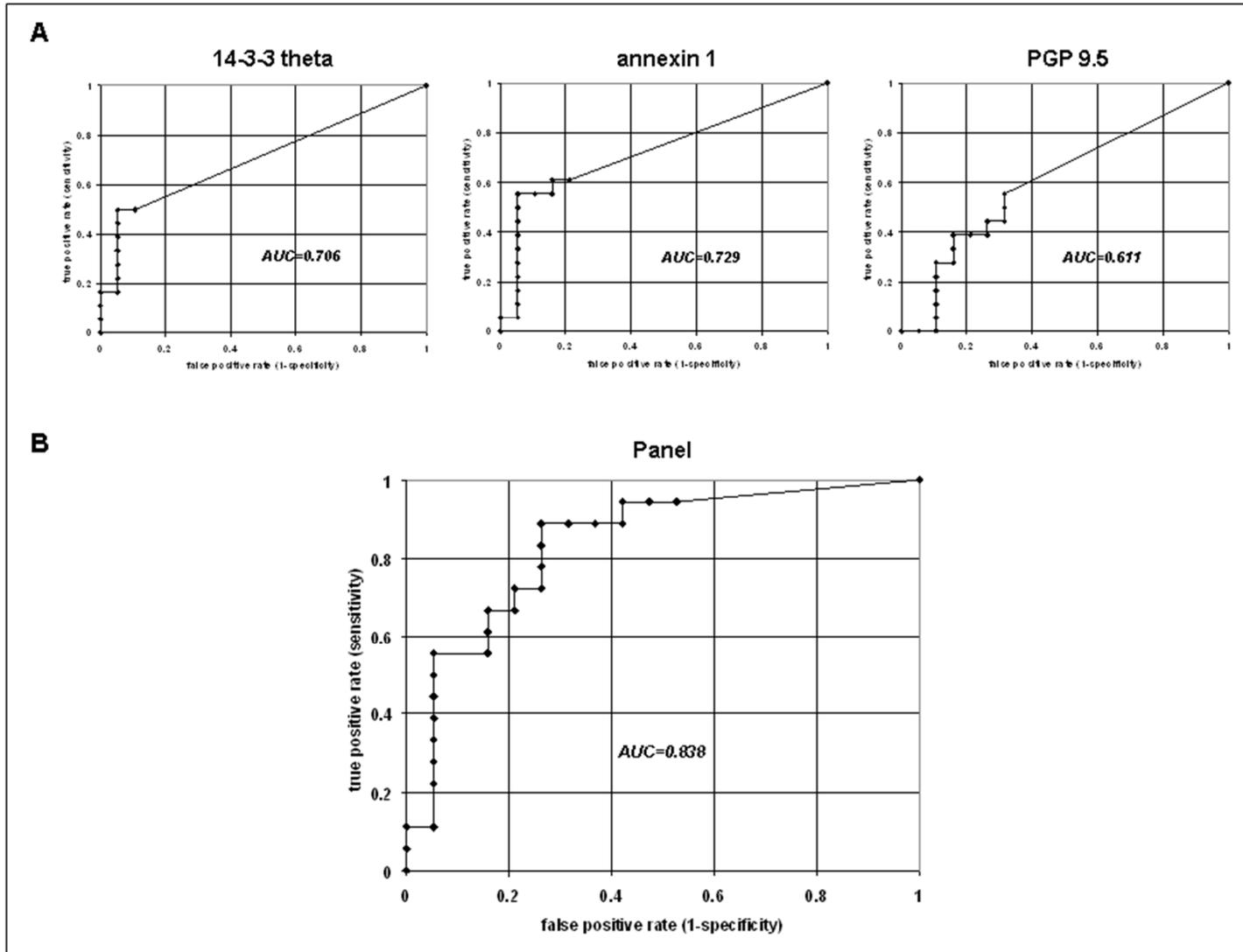
Targets: Annexin, PGP9.5, 14-3-3 theta

Data analysis: NCI EDRN Data Management Center



Identification of 14-3-3 theta as an antigen that induces a humoral response in lung cancer

Sandra R. Pereira-Faca et al (Cancer Research '07)



Validation Phase 2

- **Validate intended clinical application:
Blood test in combination with CT scanning**
- **Demonstrate increased specificity and sensitivity of CT scans when combined with an autoantibody marker panel for high risk subjects**
- **Retrospective component**
- **Prospective component**

Pancreatic cancer markers from discovery in the mouse to blinded validation study in pre-diagnostic sera

Pancreatic cancer mouse model

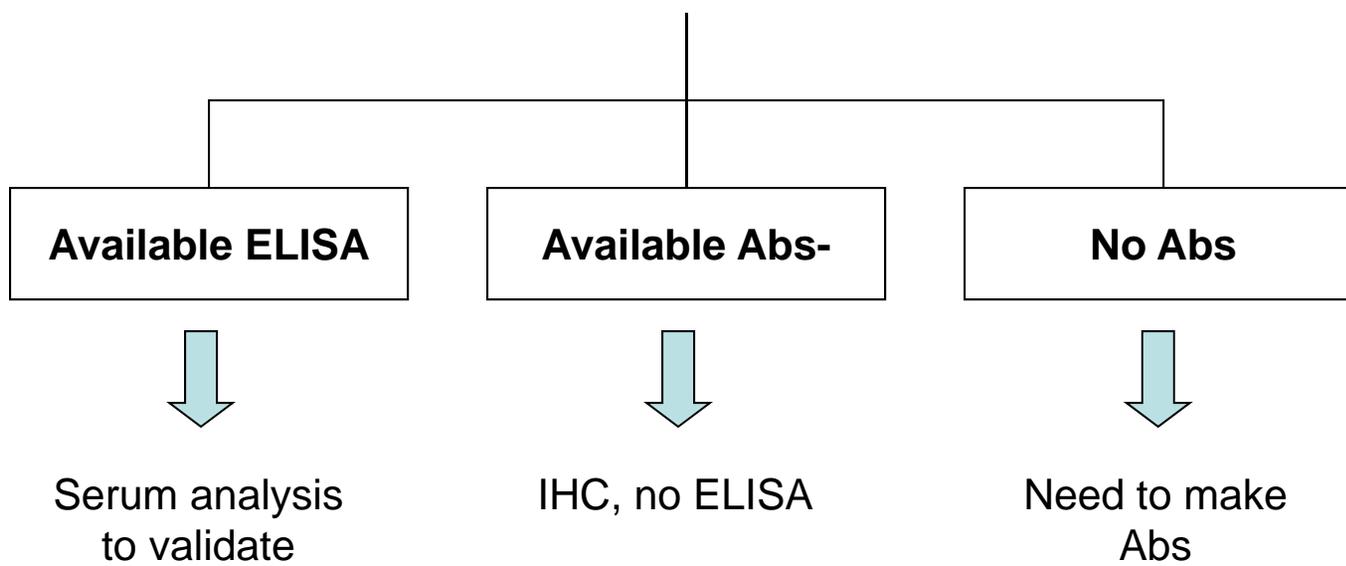
K-ras activation + Ink4a/Arf for pancreatic cancer
R. DePinho and N. Bardeesy

-Plasma from mice with early stage tumor and matched controls

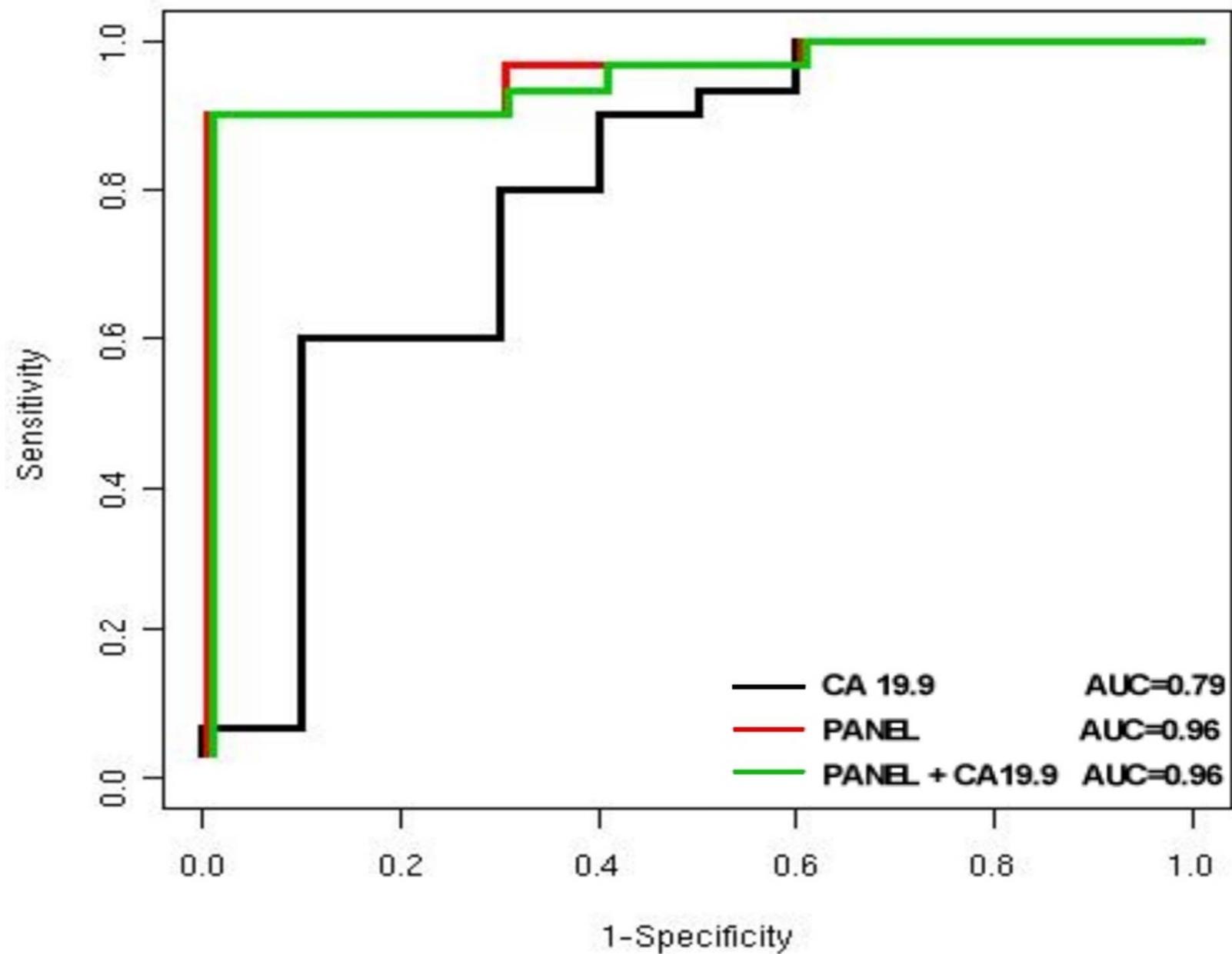
-Plasma from mice with advanced stage tumor and matched controls

Validation

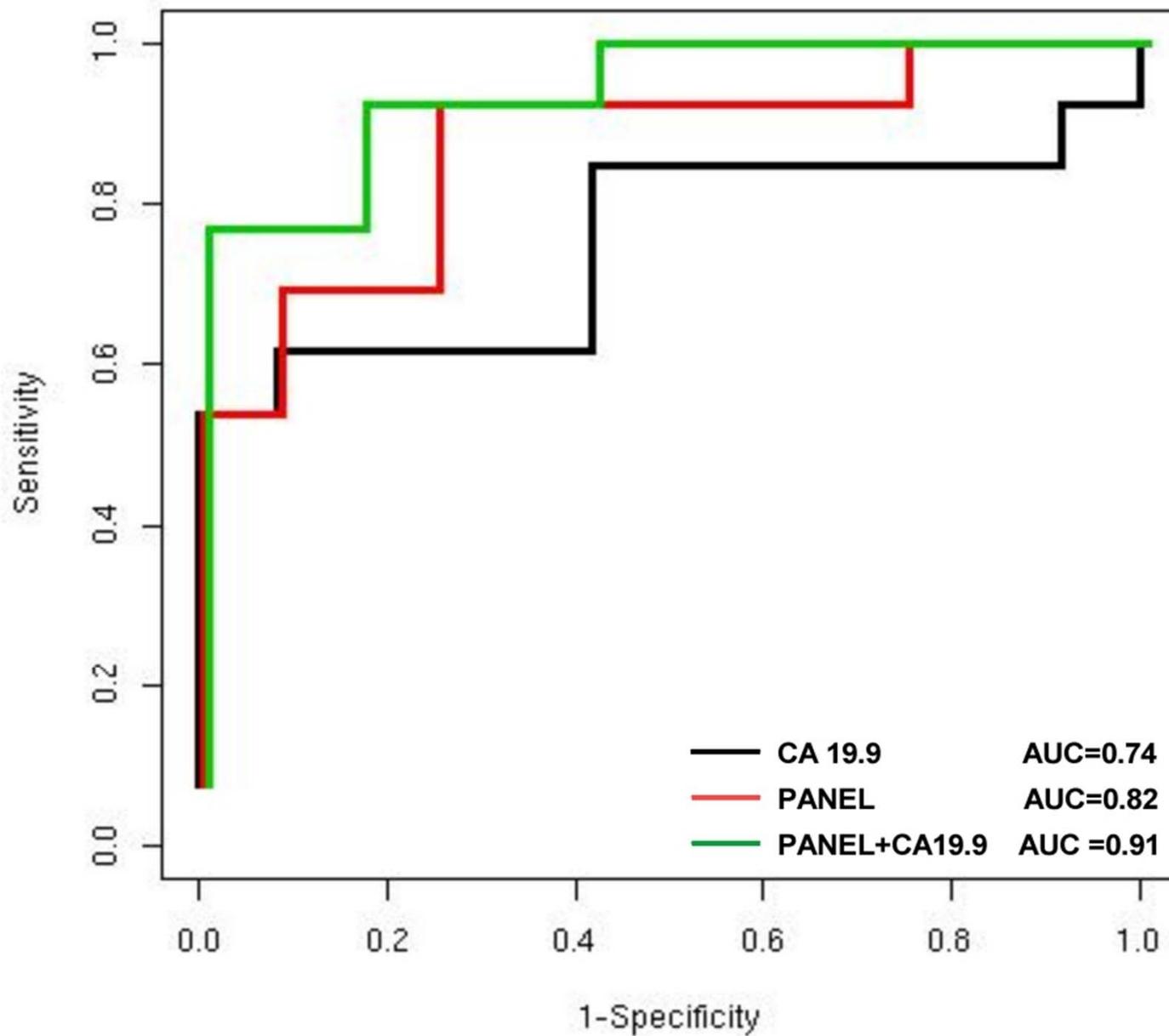
54 potential biomarkers



CANCER VS. PANCREATITIS



PRE-DIAGNOSIS PANCREATIC CANCER
VS.
MATCHED CONTROLS



NHLBI-WHI

**Markers
for early
detection of
colon Cancer**

**10 Academic
Institutions**

NCI-EDRN

NHLBI-NCI-WHI-EDRN Collaborative study

S. Hanash & R. Prentice
Co-PIs

Early
Detection
Research
Network 

100 colon cancer cases that occurred 6 – 18 m following yr 3 blood draw + 100 matched controls

10 teams applied a variety of proteomics approaches to aliquots from the same blood draws

All data compared and integrated

Promising biomarkers to be validated in a second phase

Participating Institutions

Fred Hutchinson Cancer Research Center

Harvard Medical School

R. Kucherlapati

PNNL

R. Smith

Johns Hopkins University

D, Chan

Northeastern University

W. Hancock

University of Michigan

A. Chinnaiyan

Wayne State University

M. Tainsky

Eastern Virginia University

J. Semmes

Wistar Institute

D. Speicher

University of Pittsburgh

W. Bigbee

Innovative nature of the study

Discovery studies at the pre-clinical stage

Reduced bias due to multi-institutional sample collection

Reduced bias due to asymptomatic status of subjects at the time of blood draw

Analysis of aliquots of the same samples by multiple investigators/platforms

Sample blinding at the time of data collection

Centralized integrated analysis of all data collected

Data Analysis

Data processing: CPAS (M. McIntosh)

Data management: EDRN DMCC (Z. Feng)

Statistical Analysis: WHI (R. Prentice)

Findings

A total of 2,343 high confidence protein groups were identified which corresponds to up to 2,876 distinct gene symbols (compared to HUPO PPP of 889 proteins identified).

A total of 1,846 of these proteins were identified in at least two separate laboratories.

Summary of significance

65 proteins identified with quantitative values and $P < .05$ in one or more labs (41 up, 24 down in pre-diagnostic specimen relative to matches controls).

11/65 proteins showed significance in more than one lab.

Ingenuity Pathways Analysis

File Edit Window Help

Dr. Zhang CLOSE SESSION

NEW Search for Genes or Chemicals Enter gene names/symbols/IDs or chemical/drug names here SEARCH ADVANCED

4IPA_WHI_Colon_PeiJuneLis

Summary Networks Functions Canonical Pathways Lists Pathways Molecules Network Explorer Overlapping Networks

Top Networks

ID	Associated Network Functions	Score
1	View Cell-To-Cell Signaling and Interaction, Hematological System Development and Function, Organismal Functions	51
2	View Cardiovascular Disease, Cancer, Lipid Metabolism	48
3	View Hematological Disease, Cardiac Hemorrhaging, Hematological System Development and Function	46
4	View Cell Signaling, Cell-To-Cell Signaling and Interaction, Connective Tissue Development and Function	41
5	View Immune Response, Organismal Injury and Abnormalities, Infectious Disease	39

Top Bio Functions

Diseases and Disorders

Name	p-value	# Molecules
Cancer	1.60E-22 - 6.3...	223
Hematological Disease	3.33E-20 - 6.3...	101
Inflammatory Disease	4.70E-14 - 6.4...	93
Cardiovascular Disease	6.10E-13 - 7.1...	85
Organismal Injury and Abnormalities	6.25E-11 - 6.3...	86

Molecular and Cellular Functions

Name	p-value	# Molecules
Cellular Movement	6.97E-29 - 6.4...	158
Cell-To-Cell Signaling and Interaction	1.83E-21 - 7.4...	185
Cell Signaling	7.99E-21 - 6.2...	253
Cellular Growth and Proliferation	2.56E-12 - 7.0...	203
Cellular Function and Maintenance	1.67E-11 - 6.9...	59

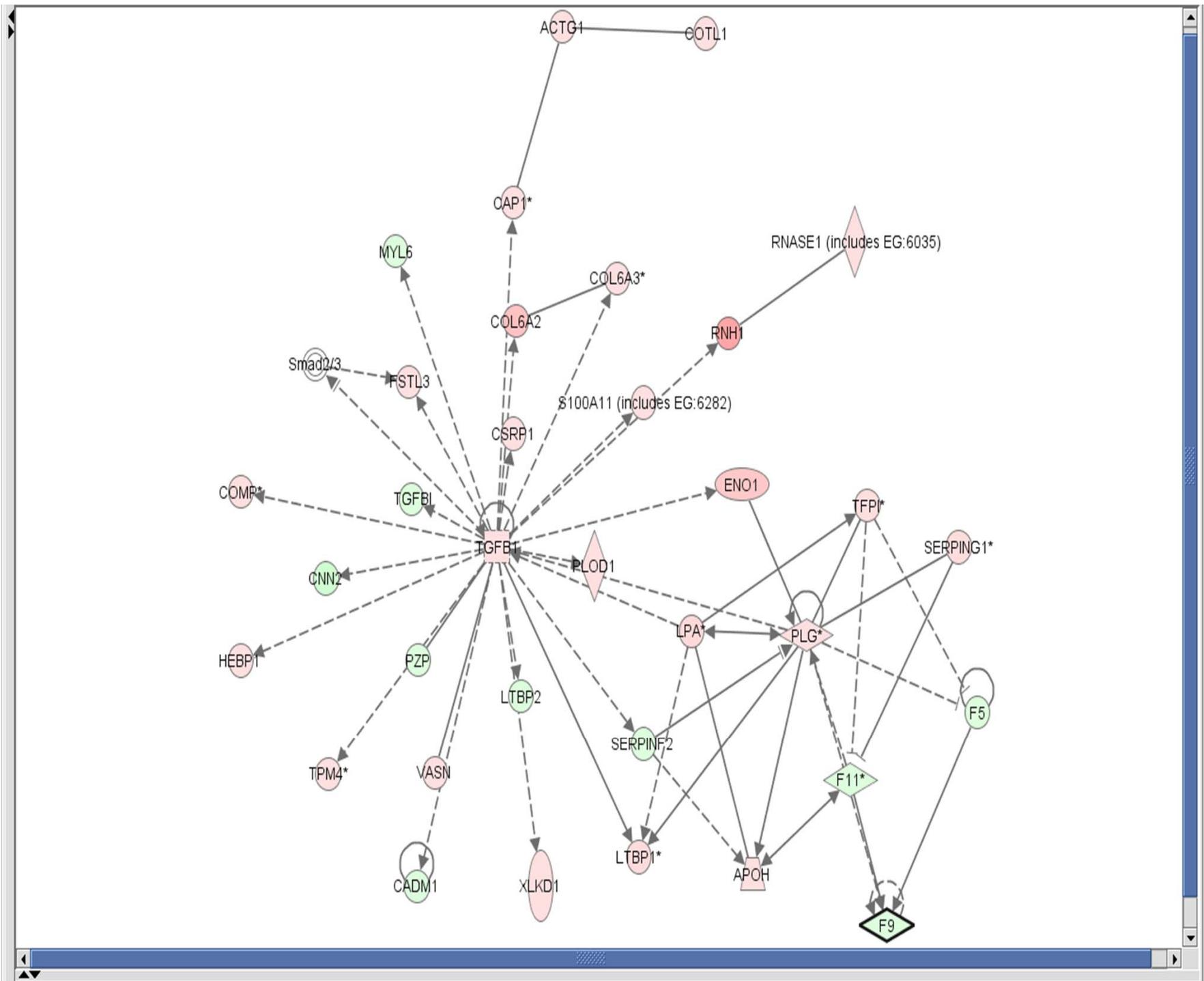
Physiological System Development and Function

Name	p-value	# Molecules
Immune Response	9.86E-23 - 6.9...	151
Tissue Development	1.83E-21 - 4.7...	151
Hematological System Development and Function	1.60E-19 - 6.4...	161
Organismal Functions	6.66E-17 - 1.6...	37
Cardiovascular System Development and Function	2.58E-13 - 6.8...	84

Top Canonical Pathways

Name	p-value	Ratio
Complement and Coagulation Cascades	2.35E-58	51/70 (0.729)
Actin Cytoskeleton Signaling	1.51E-05	24/254 (0.094)
IGF-1 Signaling	2.39E-05	13/90 (0.144)
NRF2-mediated Oxidative Stress Response	3.99E-05	16/142 (0.113)
Leukocyte Extravasation Signaling	6.31E-04	17/186 (0.091)

Top Genes



Phase 2

Validation of candidate markers using a second set of WHI subjects

Further mining of the data for PTMs, Glycan modifications..

Leveraging non-EDRN resources

**NCI: Mouse Models, Glycomics Alliance,
Nanotechnology**

Other NIH Institutes: HUPO PPP

Cohorts: CARET, WHI, PLCO

Foundations

- Lustgarten: Pancreas**
- Labrecque: Lung**
- Avon: Breast**
- Canary: Lung, pancreas, ovary, prostate**

CANARY FOUNDATION

Stopping cancer early...the best possible investment



Canary Lung Project

MISSION: Early detection of lung cancer through a combination of imaging, sputum and/or blood based testing applicable to lung cancer among smokers as well as never smokers.