

Precision Medicine



LUNG CANCER: Precision Medicine Strategy

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Vargas A and Harris CC Nature Review Cancer 16: 525-37, 2016
Robles S and Harris CC Lung Cancer 107:50-58, 2017

Precision medicine

PRECISION MEDICINE



The NEW ENGLAND JOURNAL *of* MEDICINE

A New Initiative on Precision Medicine

Francis S. Collins, M.D., Ph.D., and Harold Varmus, M.D.

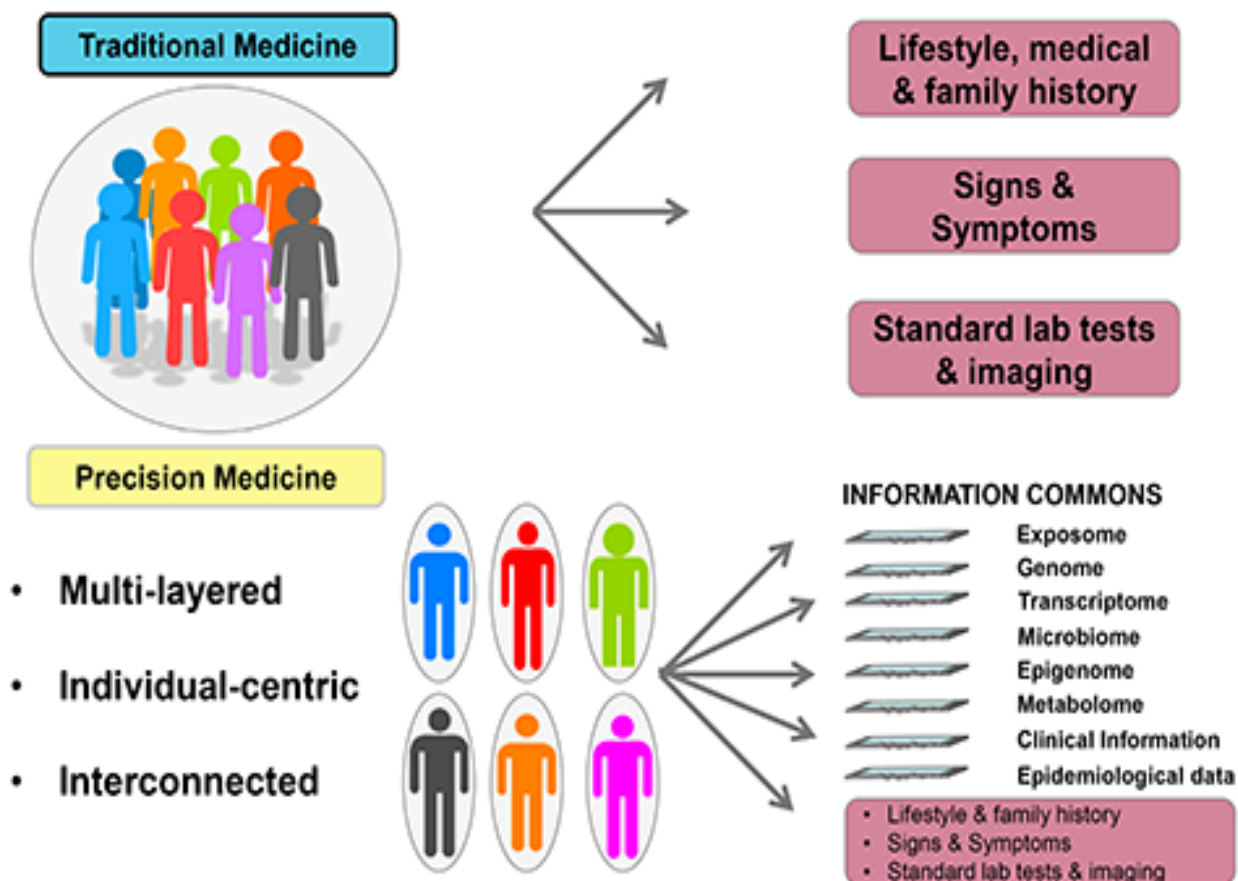
“Tonight, I’m launching a new Precision Medicine Initiative to bring us closer to curing diseases like cancer and diabetes — and to give all of us access to the personalized information we need to keep ourselves and our families healthier.”

— President Barack Obama, State of the Union Address, January 20, 2015

Collins F and Varmus H. NEJM 372: 793-5, 2015

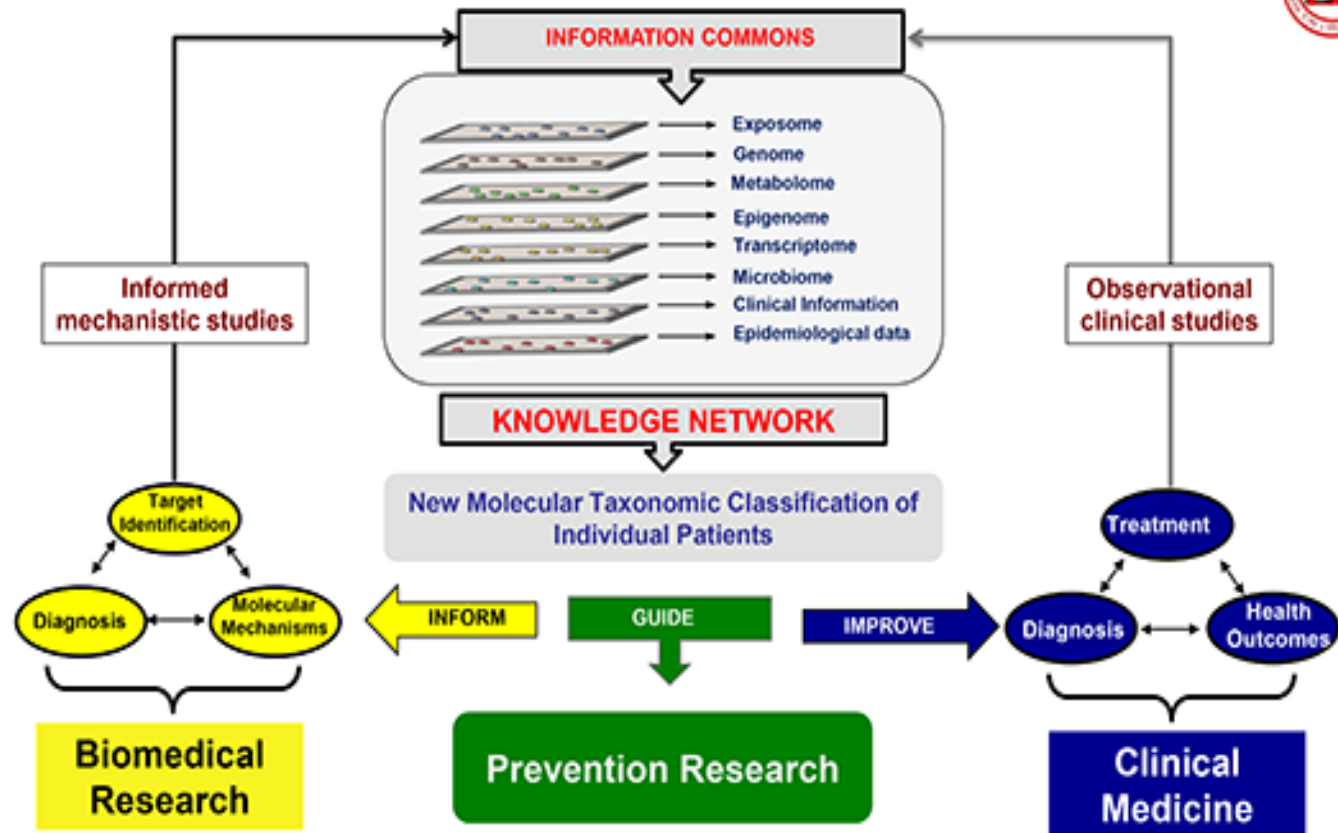
Disease classification

Towards Precision Medicine and a Molecular Taxonomic Classification of Disease



Research strategy

A Precision Medicine Research Strategy



- Modified: "Toward Precision Medicine: Building a Knowledge Network for Biomedical Research and a New Taxonomy of Disease" (National Research Council 2011)
- Vargas, A. and Harris, CC, "Precision Medicine of Cancer", Nature Review Cancer, 18:525-33, 2016

Lung cancer

Precision Medicine: LUNG CANCER



Report of the Surgeon General, 2014

Premature deaths caused by smoking and exposure to secondhand smoke, 1965-2014

Cause of death	Total
Smoking-related cancers	6,587,000
Cardiovascular and metabolic diseases	7,787,000
Pulmonary diseases	3,804,000
Conditions related to pregnancy and birth	108,000
Residential fires	86,000
Lung cancers caused by exposure to secondhand smoke	263,000
Coronary heart disease caused by exposure to secondhand smoke	2,194,000
Total	20,830,000

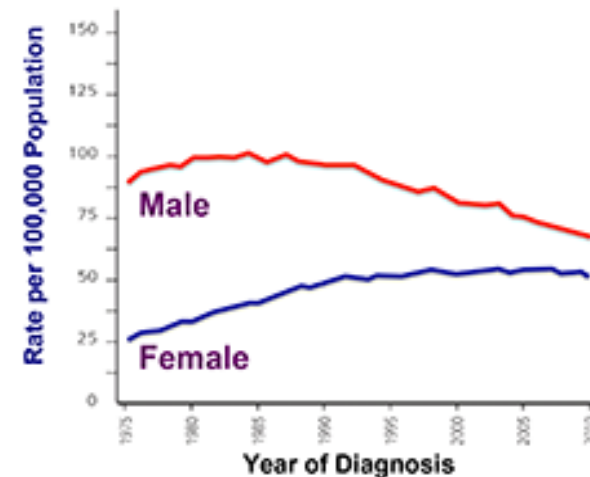
Source: Centers for Disease Control and Prevention, National Center for Chronic Disease Prevention and Health Promotion, Office on Smoking and Health, unpublished data.

Report of Surgeon General, Executive Summary, 2014



Cancer Statistics (ACS), 2014

Lung & Bronchus



Siegel, R., et al., CA Cancer J Clin, 64: 9-29, 2014

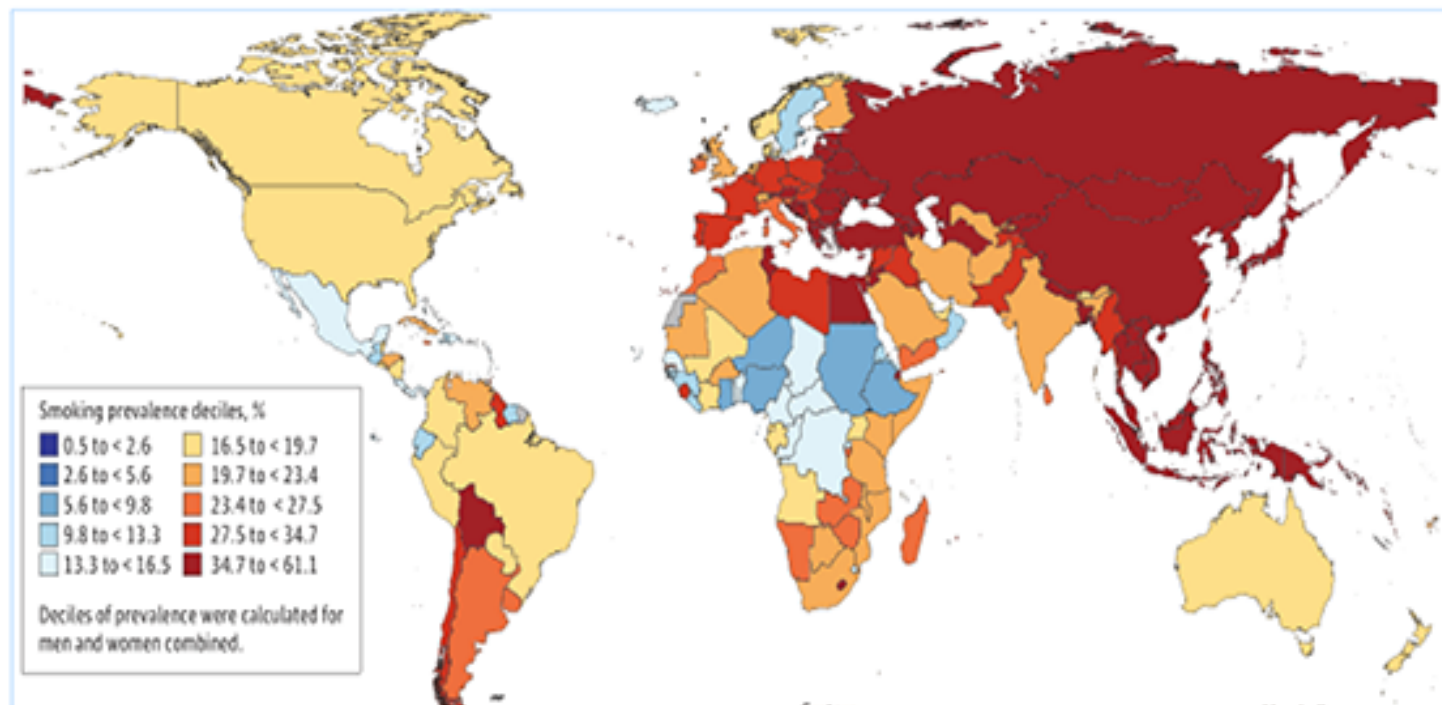
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Geographical map

Smoking Prevalence and Cigarette Consumption in 187 Countries, 1980-2012

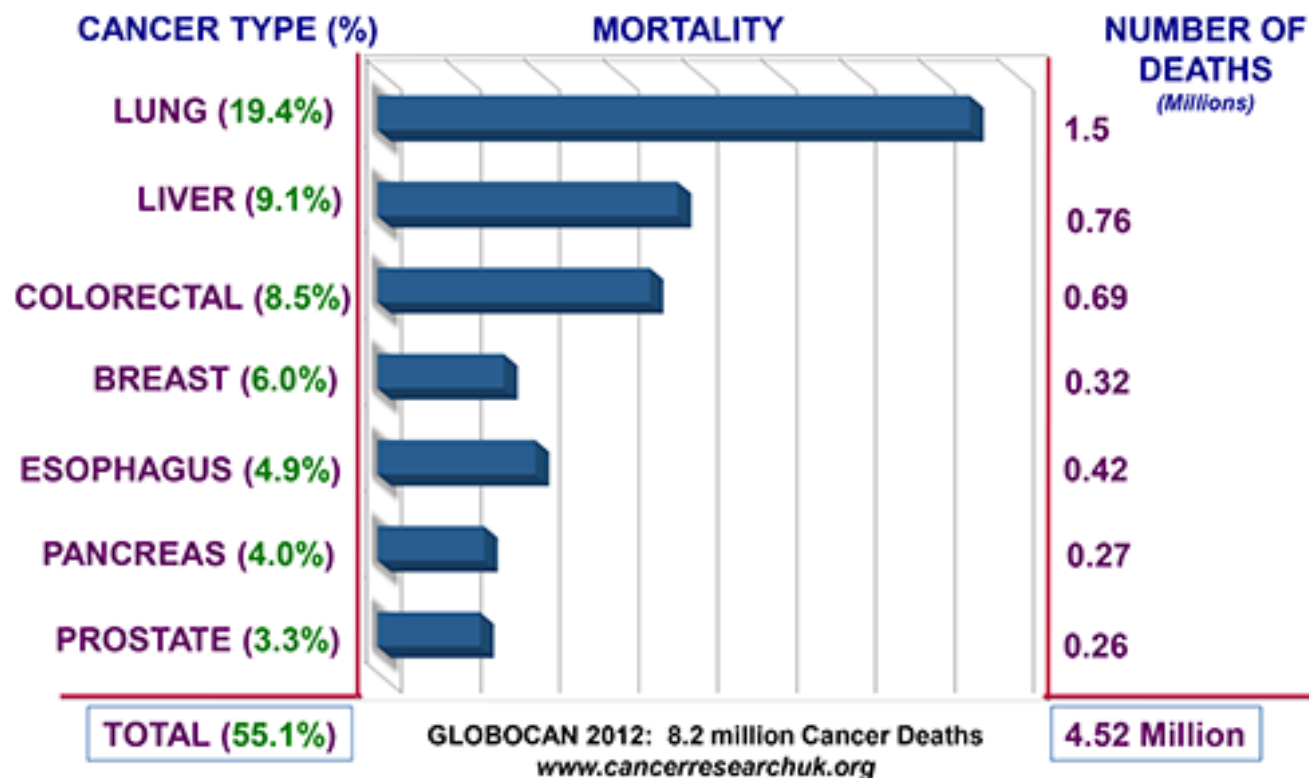


- Estimate Age-Standardized Prevalence of Daily Smoking in 2012 among Men



Worldwide cancers

Major Worldwide Lethal Types of Human Cancer



Smoke exposure

**“EXPOSURE TO SECONDHAND OR ‘ENVIRONMENTAL
TOBACCO SMOKE IS CARCINOGENIC TO HUMANS”**

(Host Factors in Human Carcinogenesis, IARC MONOGRAPH 183)



Some men have constitutions that are like wooded mountains running with springs, others like those with poor soil and little water, still others like land rich in pastures and marshes, and yet others like the bare, dry earth of the plain.

Hippocrates

Airs, Waters, Places

Hirayama, T., Cape Sunion, Greece, 1981

Never smokers

Molecular Epidemiology of Lung Cancer in Never Smokers



Hypothesis:

- Childhood exposure to second-hand smoke and genetic alterations in innate immunity increase lung cancer risk in never smoking adults

Conclusions:

- Parental secondhand smoke exposure during childhood is associated with dose-dependent increased lung cancer risk among never smokers in two cohorts.
 - Especially among those with an *MBL-2* haplotype with a hyperactive innate immune system
 - Early age of onset of lung cancer



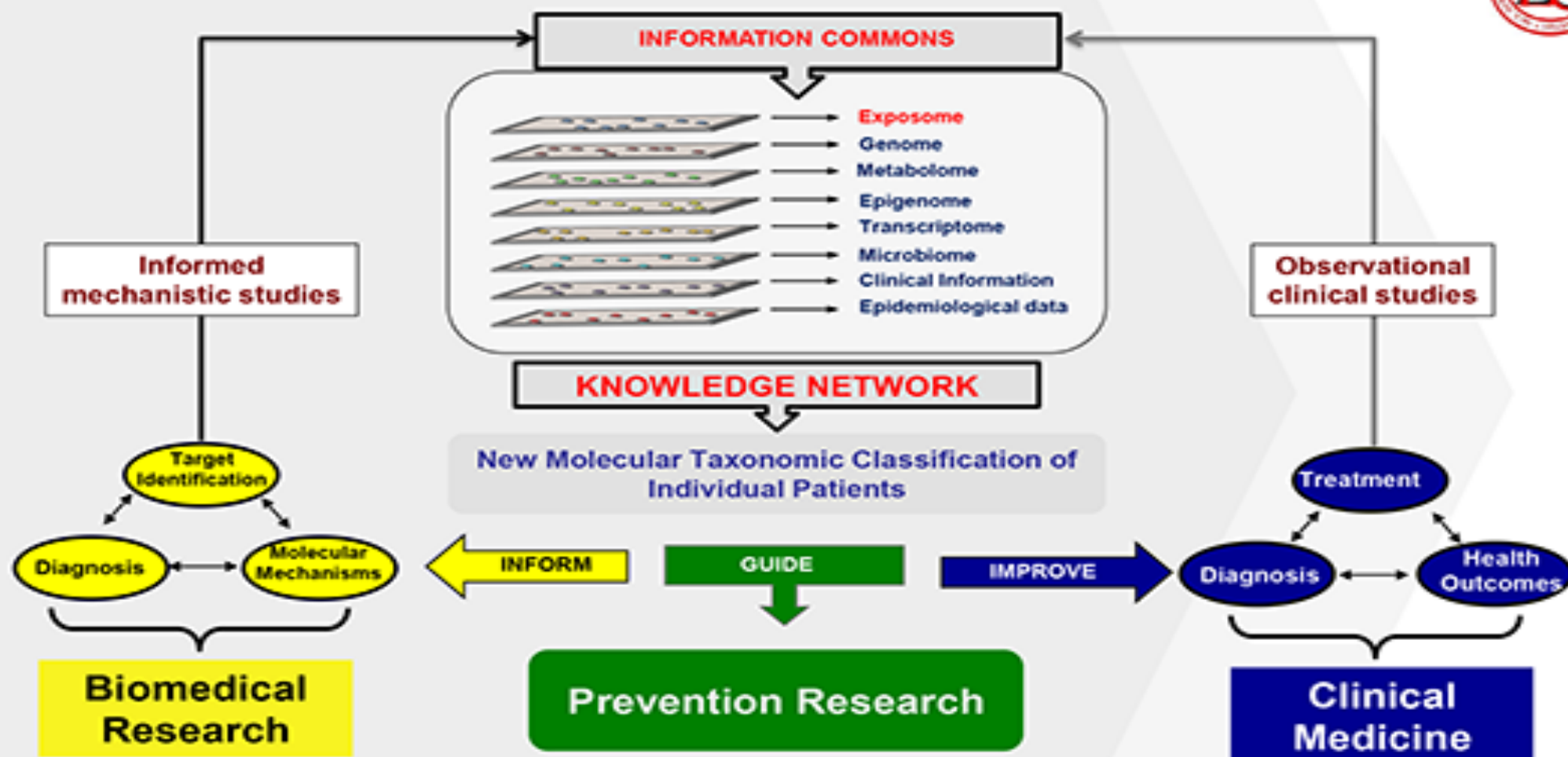
Susan Olivo-Marston

Cancer Epi Bio Prev.,
18: 3375-83, 2009

Collaboration with Jen Jin
and Ping Yang, Mayo Clinic

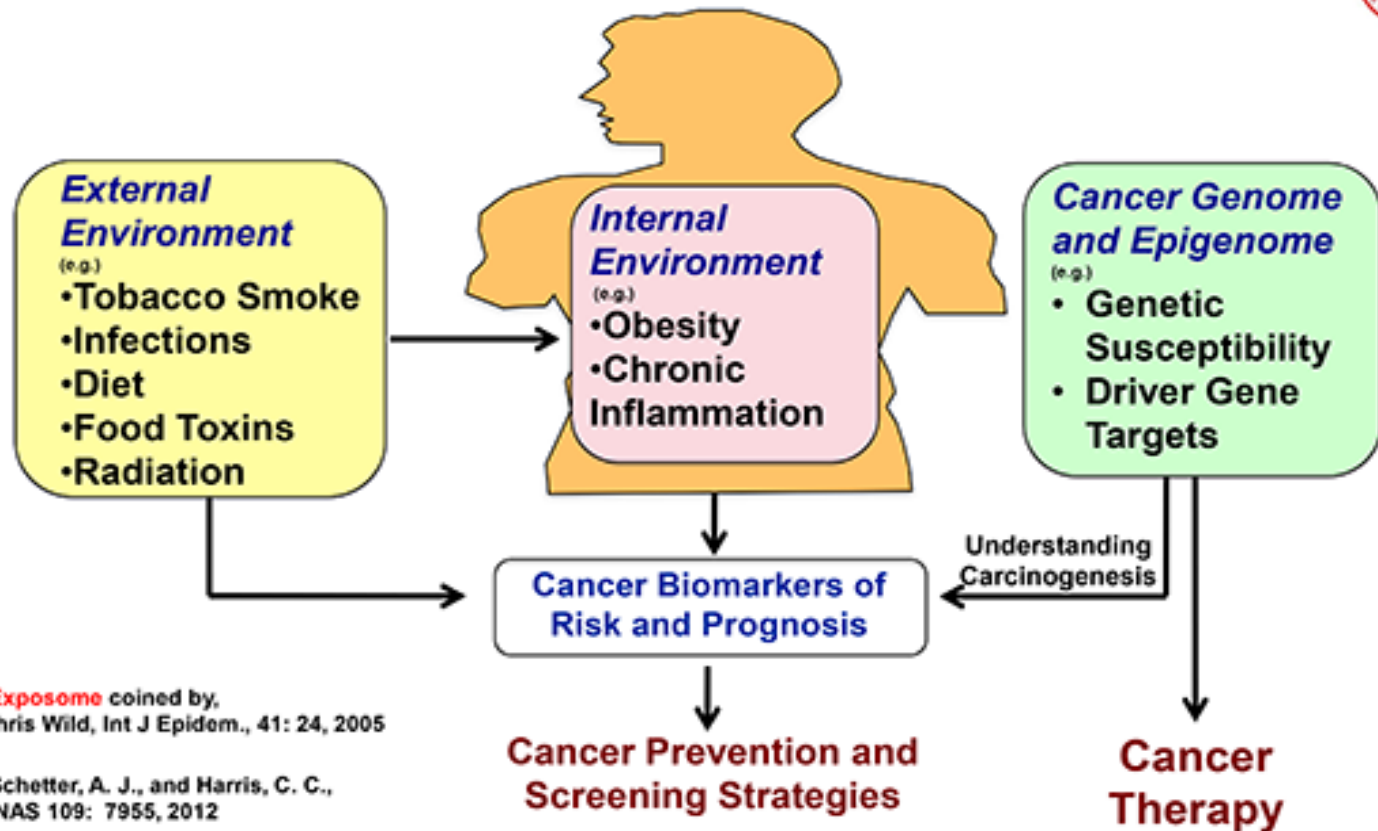
Exposome

A Precision Medicine Research Strategy



Modified: "Toward Precision Medicine: Building a Knowledge Network for Biomedical Research and a New Taxonomy of Disease" (National Research Council 2011)

Exposome¹: p53 Tumor Suppressor is at the Crossroads of the Exposome and Cancer Genome²



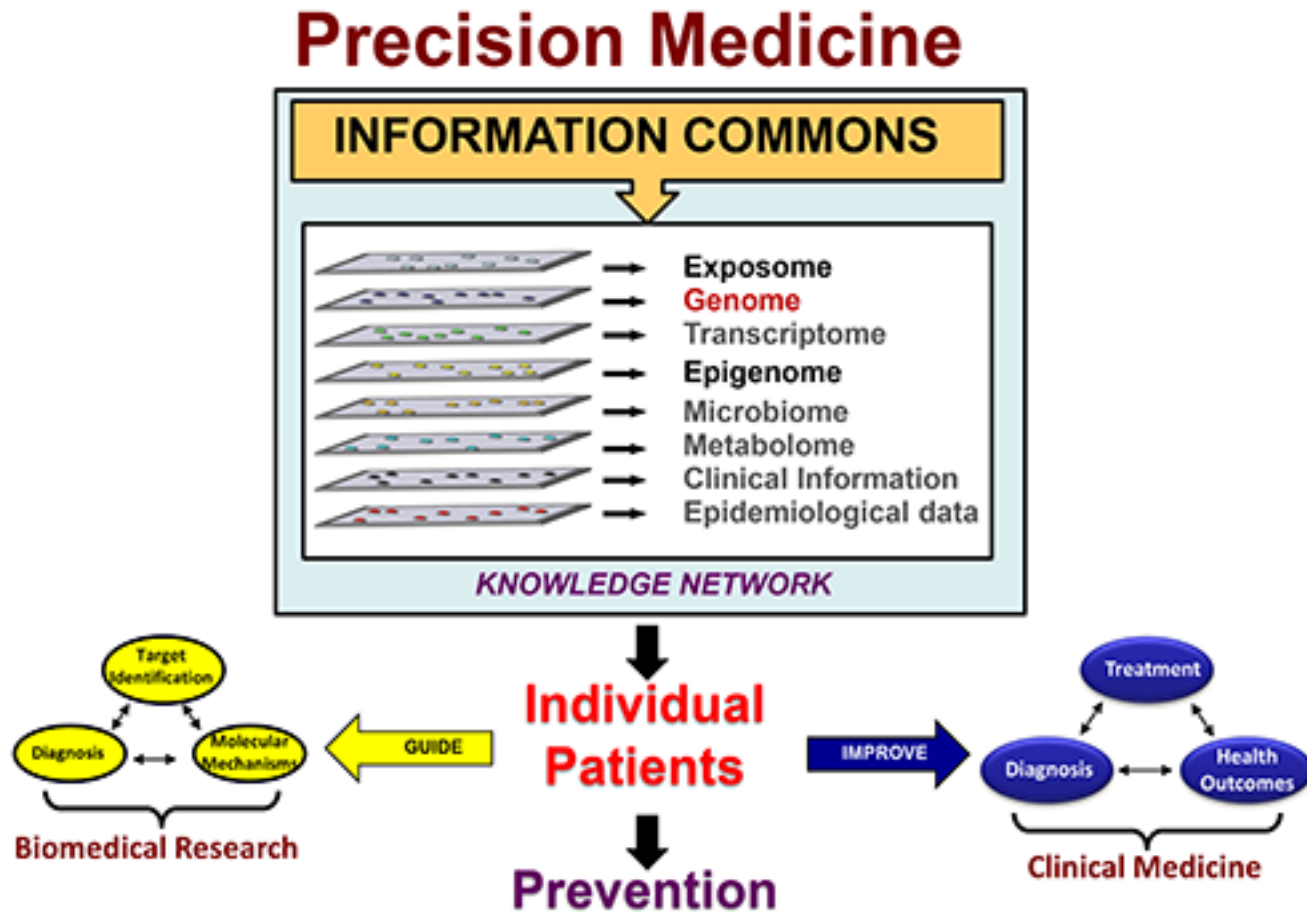
¹Exposome coined by, Chris Wild, Int J Epidemiol., 41: 24, 2005

²Schetter, A. J., and Harris, C. C., PNAS 109: 7955, 2012

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Genome

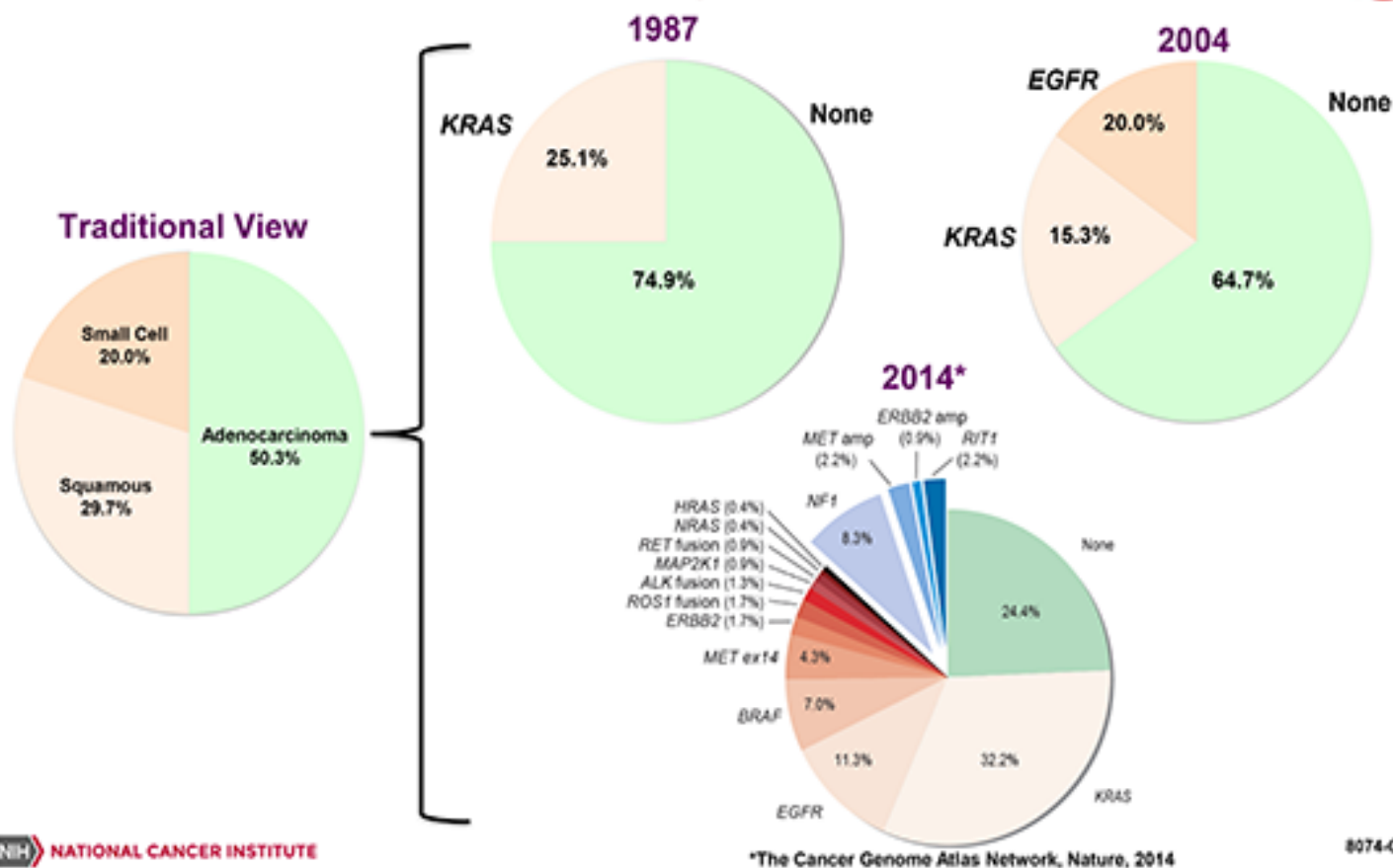


NIH NATIONAL CANCER INSTITUTE Modified: Toward Precision Medicine, National Research Council, National Academy of Science, 2011

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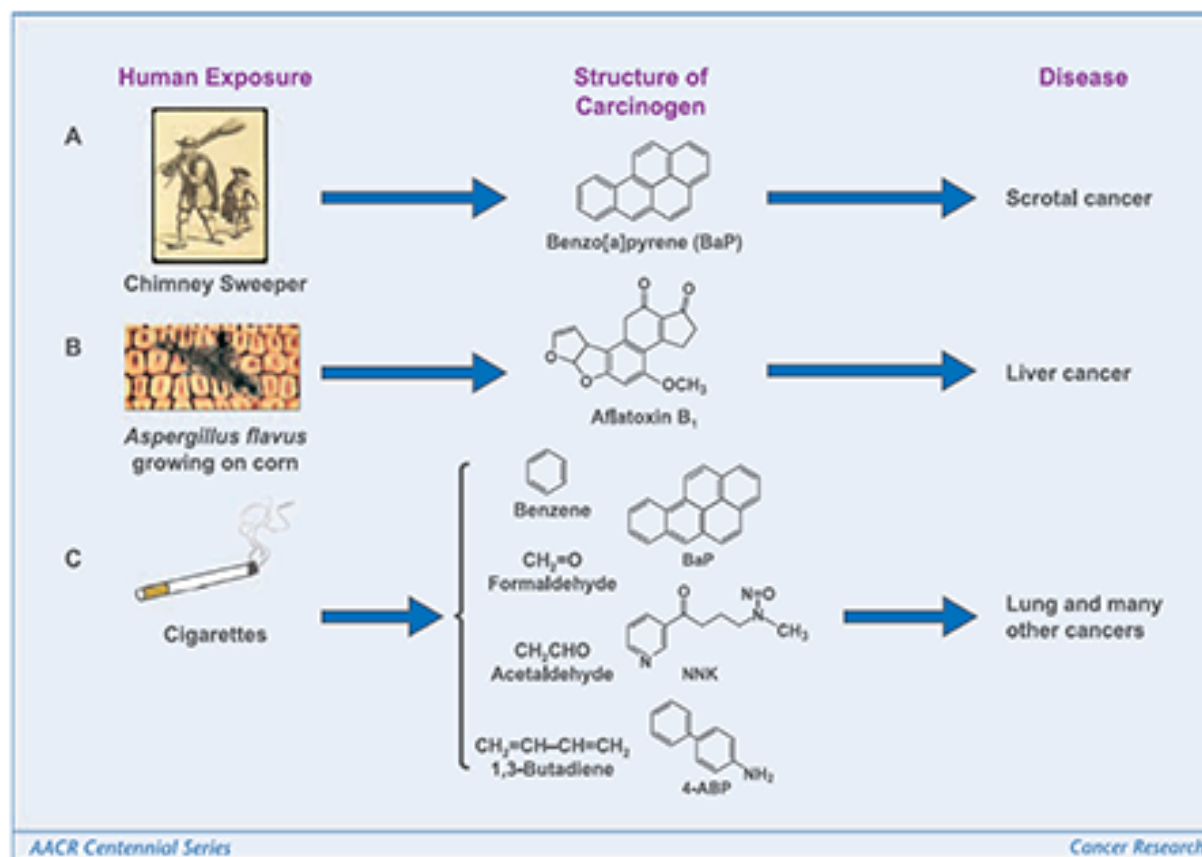
NSCLC evolution

Knowledge of Non-Small-Cell Lung Cancer has Evolved Substantially in Recent Decades



Chemical agents

Examples of Chemical Agents Causing Cancer



Three decades

Three Decades(**1970s, 1990s and 2010s**) of Examples of Initial Seminal Advances in Exposure of Environmental Carcinogens Being Molecularly Linked to Mutagenesis



- Development of a Rapid Mutagenicity Testing of Chemical Carcinogens and Metabolic Activation of Chemical Carcinogens
 - McCann J Ames BN; Detection of Carcinogens as Mutagens in the Salmonella/Microsome Test: Assay of 300 Chemicals. P.N.A.S 72:5135-39, **1975**
 - Heidelberger, C. Chemical Carcinogenesis, Annual Rev Biochemistry 44:79-121, **1975**
 - Harris CC et al., Human Bronchus-Mediated Mutagenesis of Mammalian Cells by Carcinogenic Polynuclear Aromatic Hydrocarbons. PNAS 75:2003-7, **1978**.
- **Discovery of TP53 Mutations Linked to Environmental Carcinogen Exposure**
 - Hsu IC Harris CC. p53 Mutational Hotspot in Hepatocellular Carcinoma from Qidong, China. Nature 350: 427-8, **1991**
 - Bressac B Ozturk M. Selective G to T mutations of p53 gene in Hepatocellular Carcinoma from Southern Africa. Nature 350:429-31, **1991**.
 - Holstein M, Sidransky D, Vogelstein B and Harris CC: p53 Mutations in Human Cancer. Science 253: 49-53, **1991**
- Computational Analysis of Genome-Wide DNA Sequencing Data Identifies Exogenous and Endogenous Induced Mutations including those Caused by Chemical and Physical Carcinogens, Inflammation, DNA Repair Defects and Aging
 - Alexandrov LB.....Stratton MR. Signatures of Mutational Processes in Human Cancer. Nature 500:415-421, **2013**

TP53 mutations

Exposome: Examples of Chemical and Physical Agents Causing Cancer and related TP53 Mutations in Human Cancer

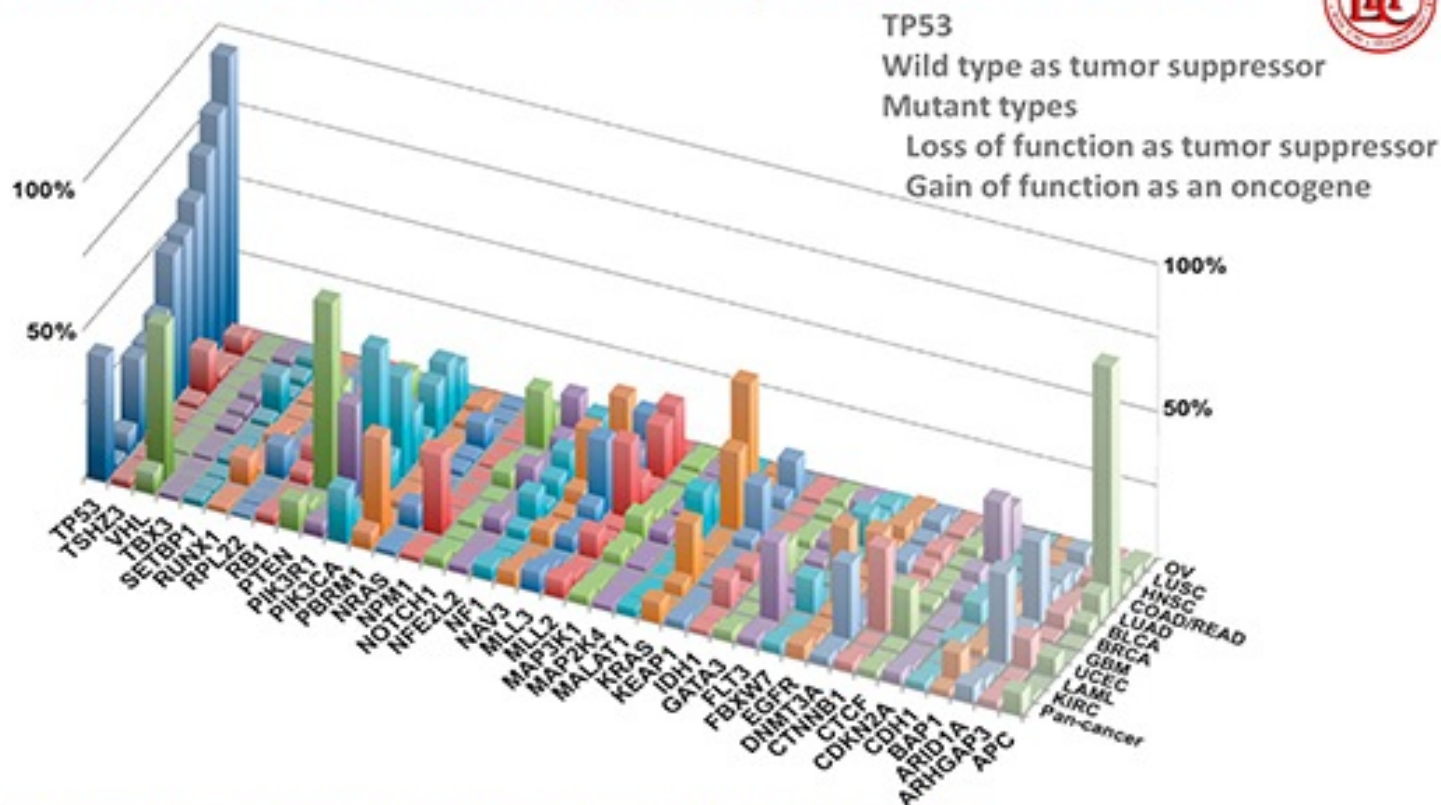


Carcinogen/ (exposure)	Target organ	Base substitution		Distinctive
		IARC database	In Vitro Human Cell Assays	Hotspots (codons)
<i>Aflatoxin B1</i> (dietary contam.)	Liver	G to T	G to T	249 (3 rd base)
<i>PAH (B[a]P)</i> (smoking)	Lung	G to T with Strand bias	G to T with Strand bias	157, 158, 273
<i>UV radiation</i> (sunlight)	Skin	CC to TT	CC to TT	248, 278
<i>Aristolochic Acid</i> (dietary contam.)	Urothelium	A to T with Strand bias	A to T with Strand bias	131, 209, (280)

- Hsu, I.C.....Harris, C.C., Nature, 350: 427-428, 1991
- Bressac, B.....Ozturk, M., Nature, 350: 429-430, 1991
- Hollstein, M., Sidransky, D., Vogelstien, B., and Harris, C.C., Harris, Science, 253: 49-53, 1991
- Greenblatt, M.....Harris, C.C., Cancer Res., 54: 4855-78, 1994

TP53 mutations

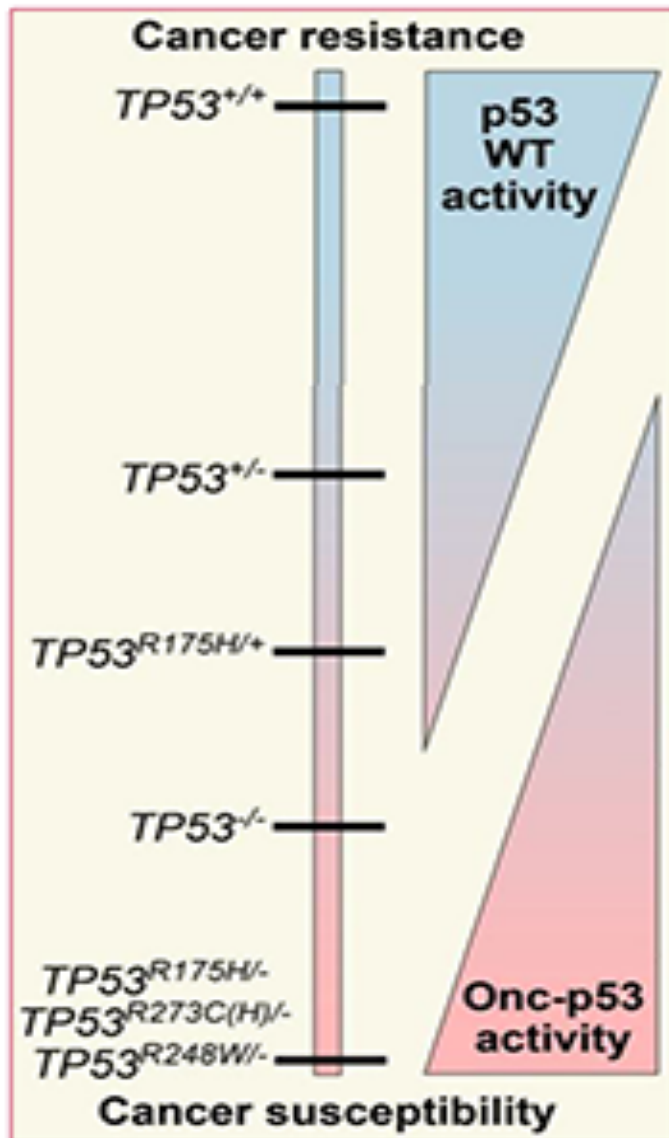
TP53 is the Gene most frequently Mutated in Cancer



Genes most frequently mutated in various types of cancer in the TCGA Pan-Cancer study.
Data were generated by analysis of the mutations released by Kandoth et al. Nature 2013.

Soussi & Wiman, 2015

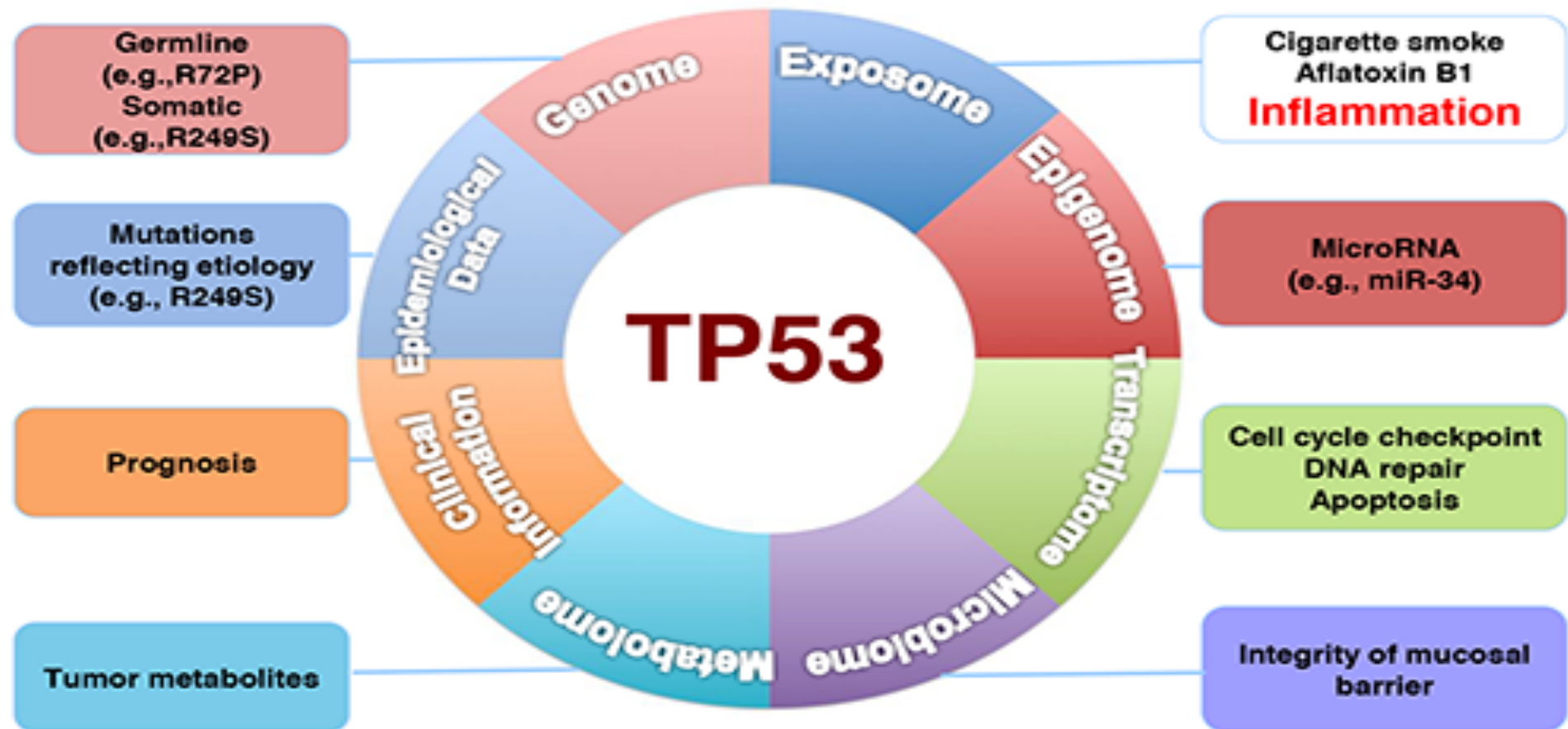
p53



The Spectrum of Protection Against Cancer Provided by WT p53. As Copies of WT p53 ($TP53^{+/+}$) are Lost, Cancer Protection Decreases. When Oncomorphic Mutations are Acquired, Cancer Susceptibility is Increased

TP53 functions

TP53 and its Functions Affect Multiple Layers of “-OMICS” Data in the Precision Medicine Paradigm



Three decades

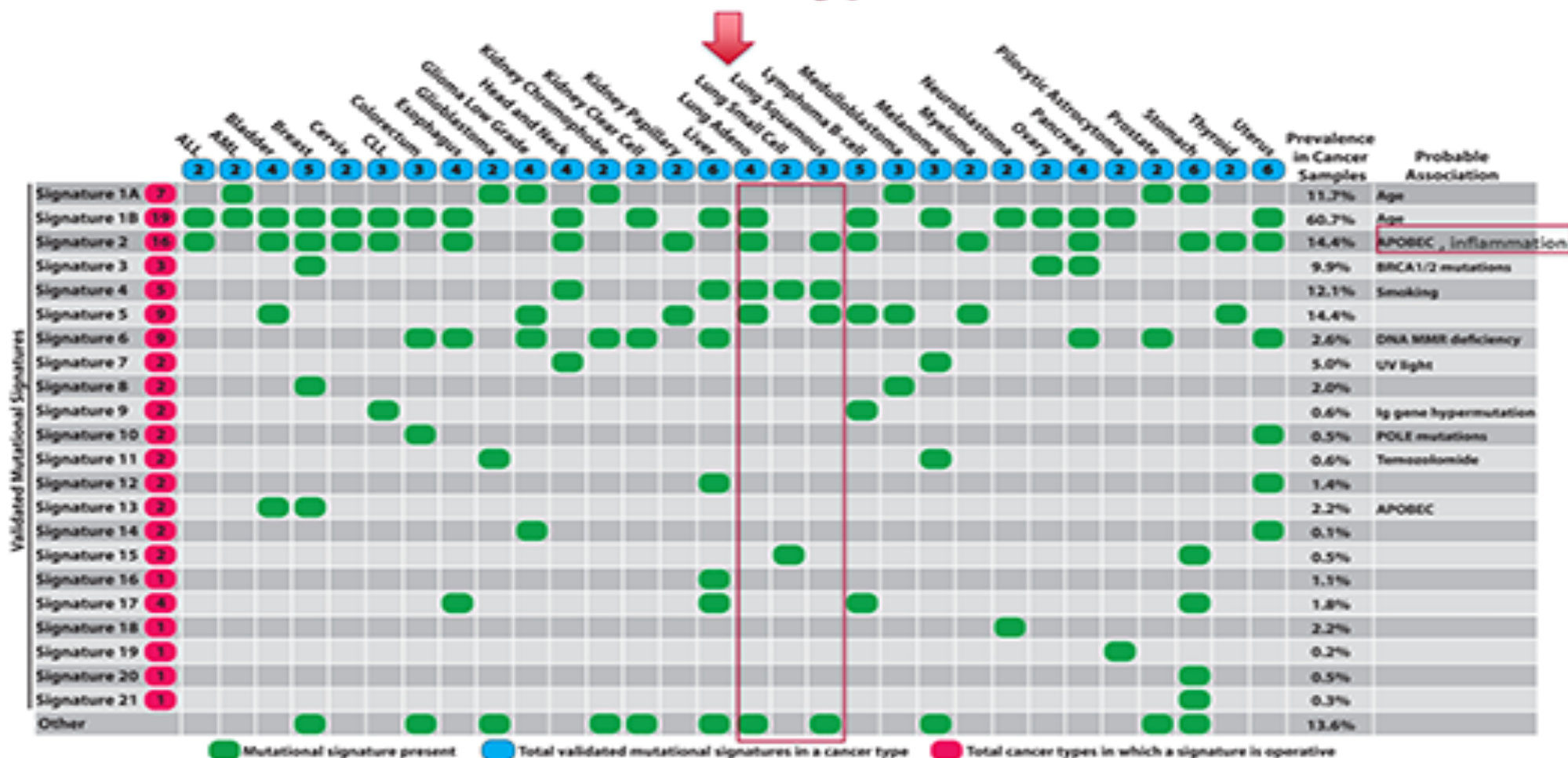
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 - Heidelberger, C. Chemical Carcinogenesis, Annual Rev Biochemistry 44:79-121, **1975**
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- Discovery of TP53 Mutations Linked to Environmental Carcinogen Exposure
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 - Holstein M, Sidransky D, Vogelstein B and Harris CC: p53 Mutations in Human Cancer. Science 253: 49-53, **1991**
- **Computational Analysis of Genome-Wide DNA Sequencing Data Identifies Exogenous and Endogenous Induced Mutations including those Caused by Chemical and Physical Carcinogens, Inflammation, DNA Repair Defects and Aging**
 - Alexandrov LB.....Stratton MR. Signatures of Mutational Processes in Human Cancer. Nature 500:415-421, **2013**

Somatic mutations

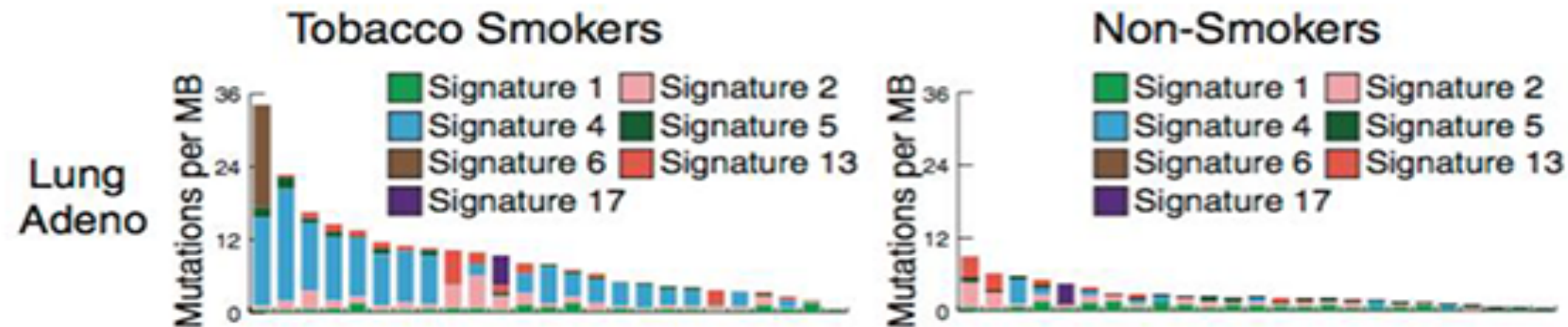
The Prevalence of Somatic Mutations Across Human Cancer Types



Mutations and tobacco smoking



Mutational Signatures Associated with Tobacco Smoking



Alexandrov, L.B., et al., Science, 354(6312): 618-772, 2016

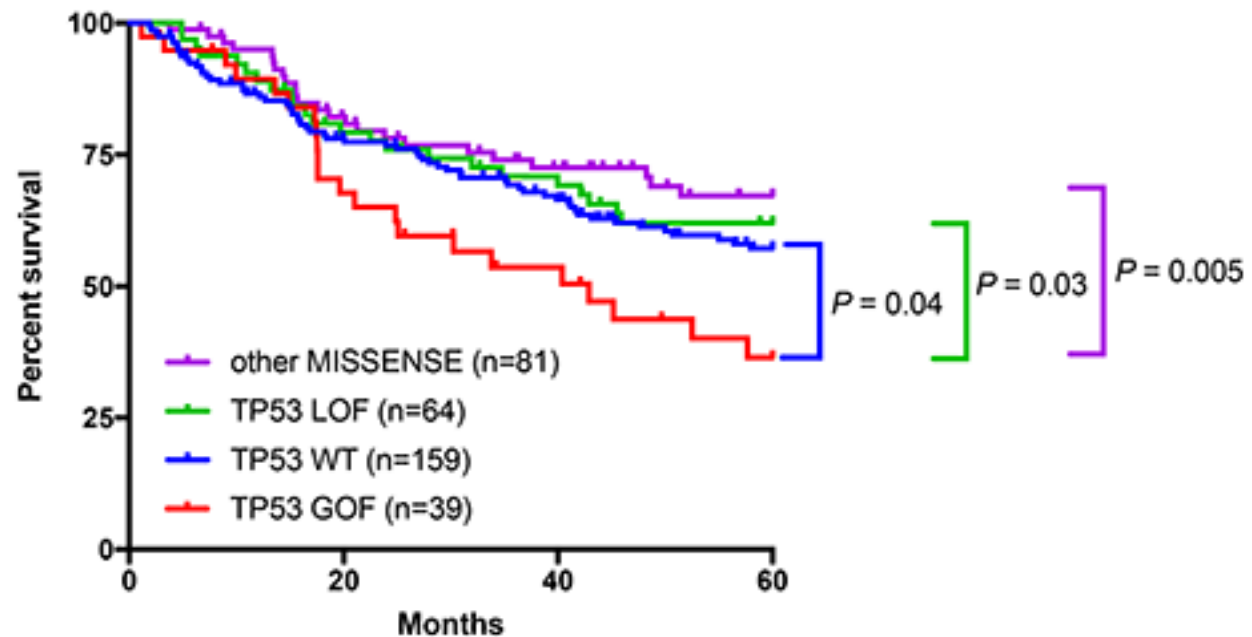
Carcinogens fingerprint

A Carcinogen's Fingerprint in Human Lung Cancer DNA Can be Reproduced in Experimental Systems: Benzo(a)pyrene



Patient prognosis

Gain of Function(GOF) TP53 Mutations are Associated with Poor Prognosis of Lung Cancer Patients



$HR_{GOFvsallothers}: 1.8; CI, 1.0-3.0; P= 0.034$

Adjusted for patient age, race, sex, and smoking history, and tumor histology, and stage.

Dataset of 352 surgical cases of NSCLC analyzed by TP53 capture sequencing

Chronic inflammation and infection



Chronic Inflammation and Infection Can Increase Cancer Risk

Inherited > Acquired

Disease	Tumor Site
Hemochromatosis	Liver
Crohn's Disease	Colon
Ulcerative Colitis	Colon
Familial Pancreatitis	Pancreas

Global Impact

- 2 million human cancers per year are related to infection
- Other causes of inflammation are associated with many more cancers per year (e.g. smoking 6 million cancers/year)

Acquired > Inherited

Disease Agent	Tumor Site
---------------	------------

Viral

Hepatitis B	Liver
Hepatitis C	Liver

Bacterial

Helicobacter Pylori	Gastric
PID	Ovary

Parasitic

S. hematobium	Urinary Bladder
japonicum	S. Colon
Liver Fluke	Liver

Chemical, Physical, and Metabolic Examples

Acid reflux	Esophagus
Obesity	Multiple sites & Liver
Smoking	Multiple sites

Serum cytokines

Increased Expression of Serum Pro-Inflammatory Cytokines is Associated with Lung Cancer Risk, Diagnosis and Survival



Increased Risk

Diagnosis

Poor Survival



↑ **IL-8**, ↑ **CRP** (EA)

JNCI 103: 1112, 2011

↑ **IL-8**, ↑ **IL-6** (AA, EA)
↑ **IL-1B & IL-10** (AA)

JNCI 103: 1112, 2011
J. Thoracic Oncol, 1494, 2014

↑ **IL-6 & -8** (AA, EA), ↑ **MBL-2** (EA)
↑ **TNF α** (EA), ↑ **IL-10 & IL-12** (AA)

CEBP: 215, 2010
JNCI 99: 1401, 2007
J. Thoracic Oncol, 1494, 2014
CEBP: 488, 2016

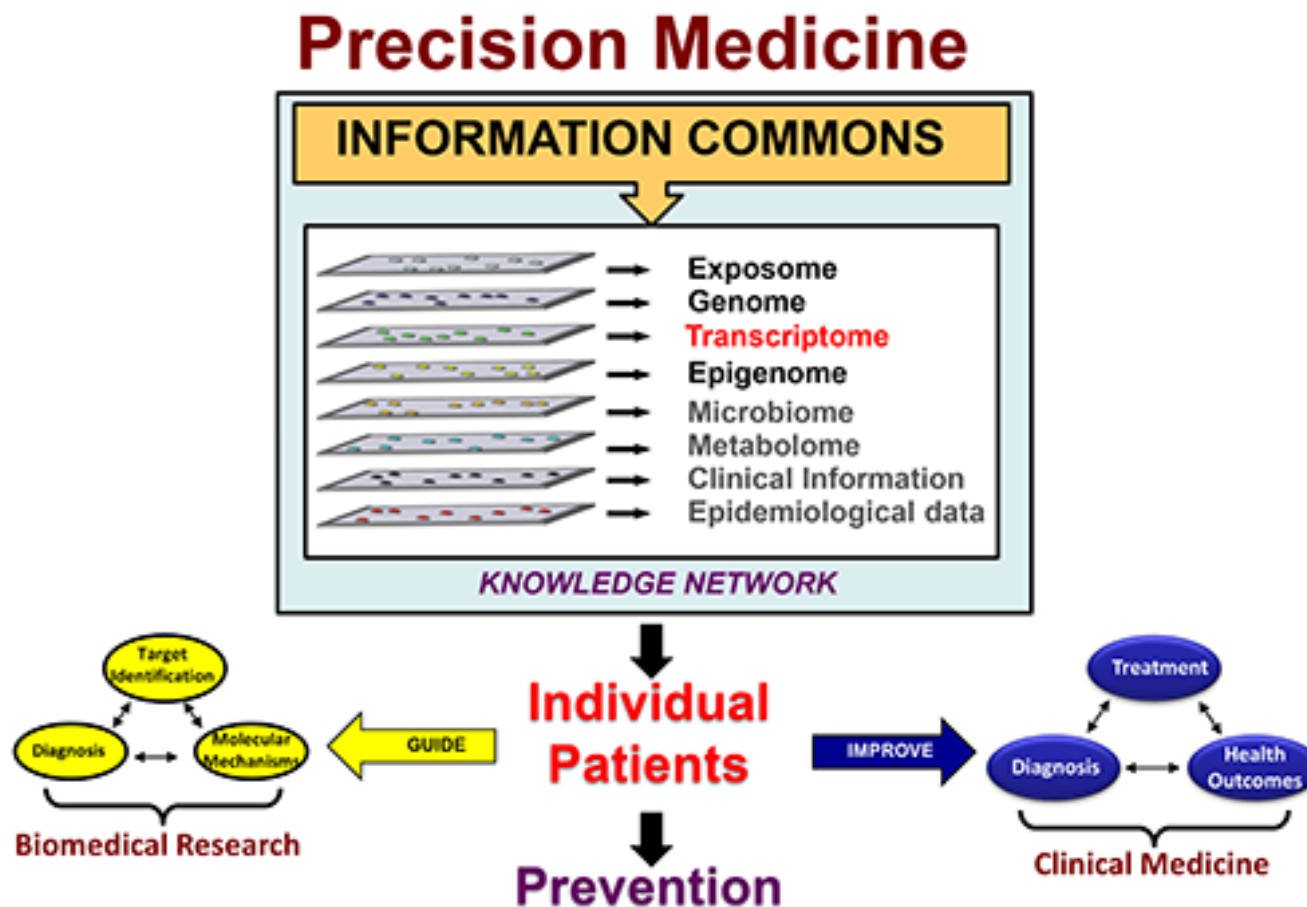
Health Disparity:

EA, European-American

AA, African-American

Collaboration: Ann Schwartz

Transcriptome



NIH NATIONAL CANCER INSTITUTE Modified: Toward Precision Medicine, National Research Council, National Academy of Science, 2011

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MicroRNA

MicroRNA and Cancer



- **MicroRNA**

- Small non-coding RNAs that are evolutionarily conserved and regulate gene expression.
- Protein output of hundreds of genes are repressed by each microRNA destabilizing mRNA and to a lesser extent inhibiting translation of mRNA.



Victor Ambros



Gary Ruvkun

- **Human Cancer**

- MicroRNAs are differentially expressed in human cancers.
- MicroRNAs can predict risk, diagnosis, prognosis and therapeutic outcome.

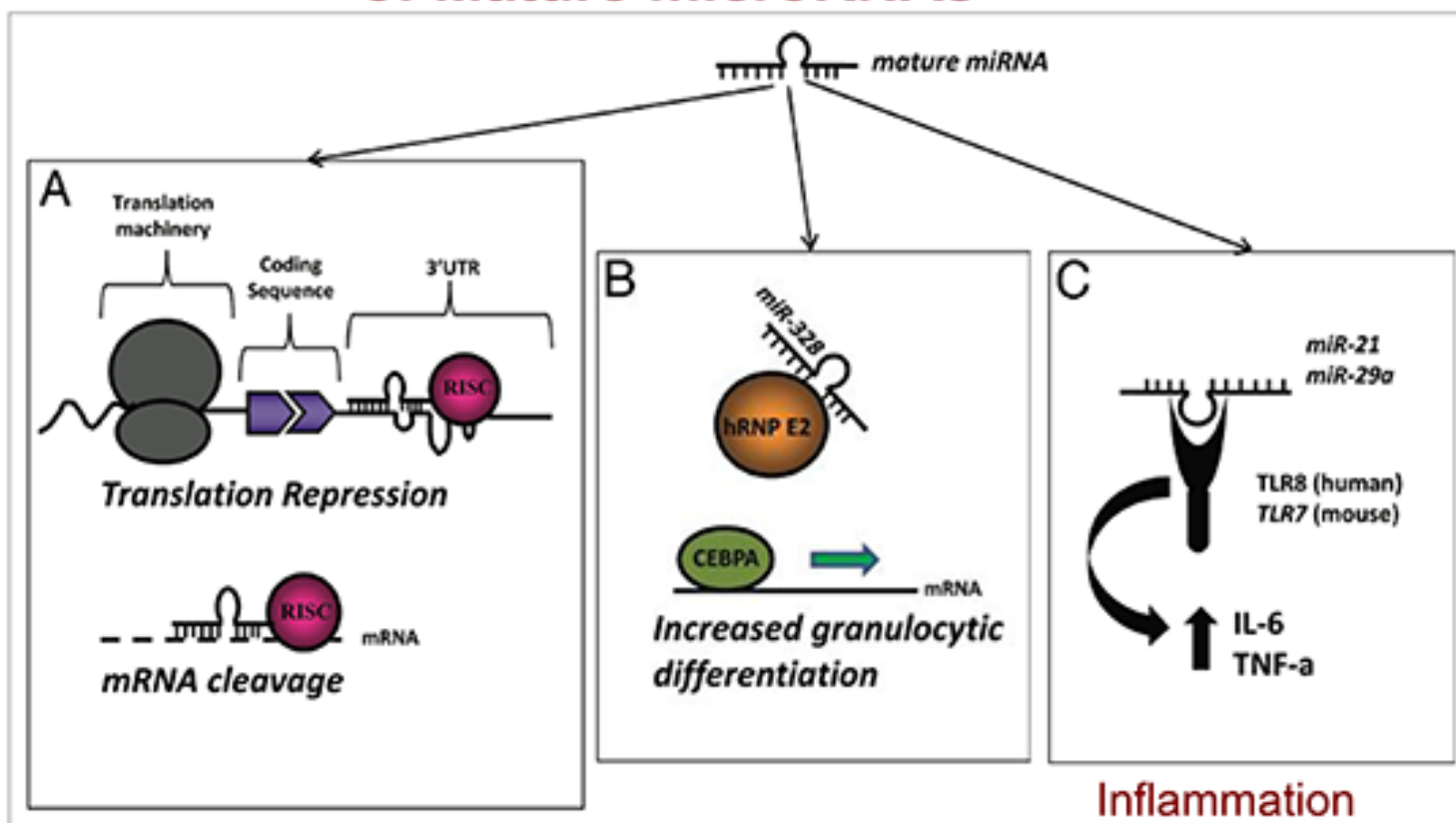


Carlo Croce

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microRNA mechanisms

Different Mechanisms of Action of Mature microRNAs

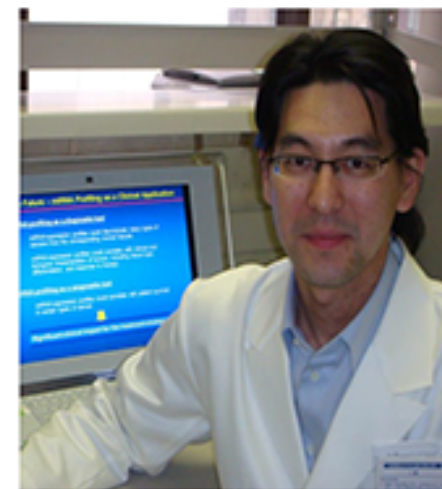


Hypothesis

Hypothesis: MicroRNAs are Associated with Lung Cancer Diagnosis and Prognosis



- miRNAs profiles were significantly different both between primary lung cancers and corresponding non-cancerous lung tissues and among different histological types of lung cancer.
- **Increased mir-21, mir-155 and mir-106b, and decreased let-7** were each associated with diagnosis and prognosis including **stage 1 lung cancer.**



Nozumu Yanaihara

let-7: confirmed studies by Takahashi and Slack

Yanaihara et al., Cancer Cell,
9:189, 2006

Upregulated microRNA



Commonly Up-Regulated microRNAs in Carcinomas

miR	N	Tumor Type
miR-21	6	Breast Colon Lung Pancreas Prostate Stomach
miR-17-5p	5	Breast Colon Lung Pancreas Prostate
miR-191	5	Colon Lung Pancreas Prostate Stomach
miR-29b-2	4	Breast Colon Pancreas Prostate
miR-223	4	Colon Pancreas Prostate Stomach
miR-128b	3	Colon Lung Pancreas
miR-199a-1	3	Lung Pancreas Prostate
miR-24-1	3	Colon Pancreas Stomach
miR-24-2	3	Colon Pancreas Stomach
miR-146	3	Breast Pancreas Prostate
miR-155	3	Breast Colon Lung

- microRNAs shared by the signatures of six solid cancers*

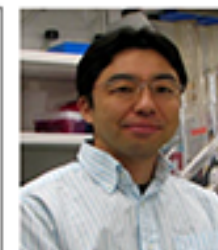
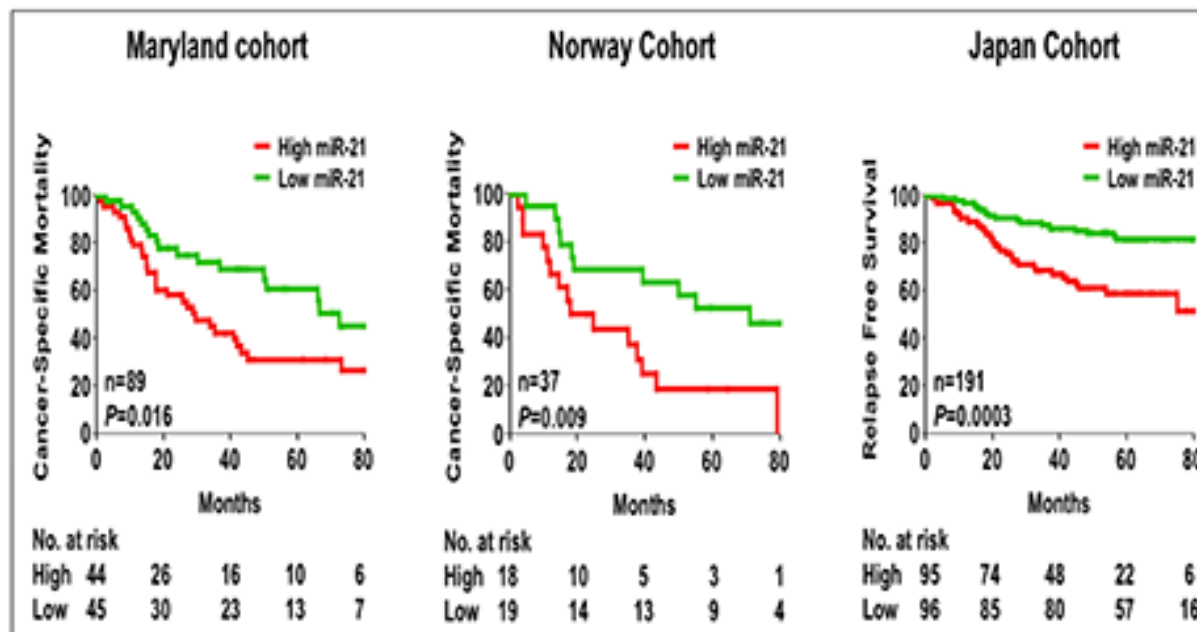
- **miR-21** is both up-regulated in 18 major cancers and a biomarker of poor survival in 14

*The list includes 21 commonly up-regulated microRNAs in three or more (N) types of solid cancers ($p\text{-value}=2.5 \times 10^{-3}$).

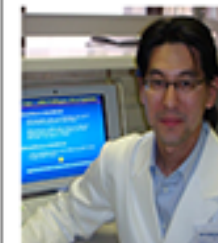
Volinia et al., PNAS 103: 1-5, 2006

Increased miR-21 expression

Increased *miR-21* Expression in Human Lung Cancer is Associated with Poor Prognosis in Three Cohorts



Motonobu Saito



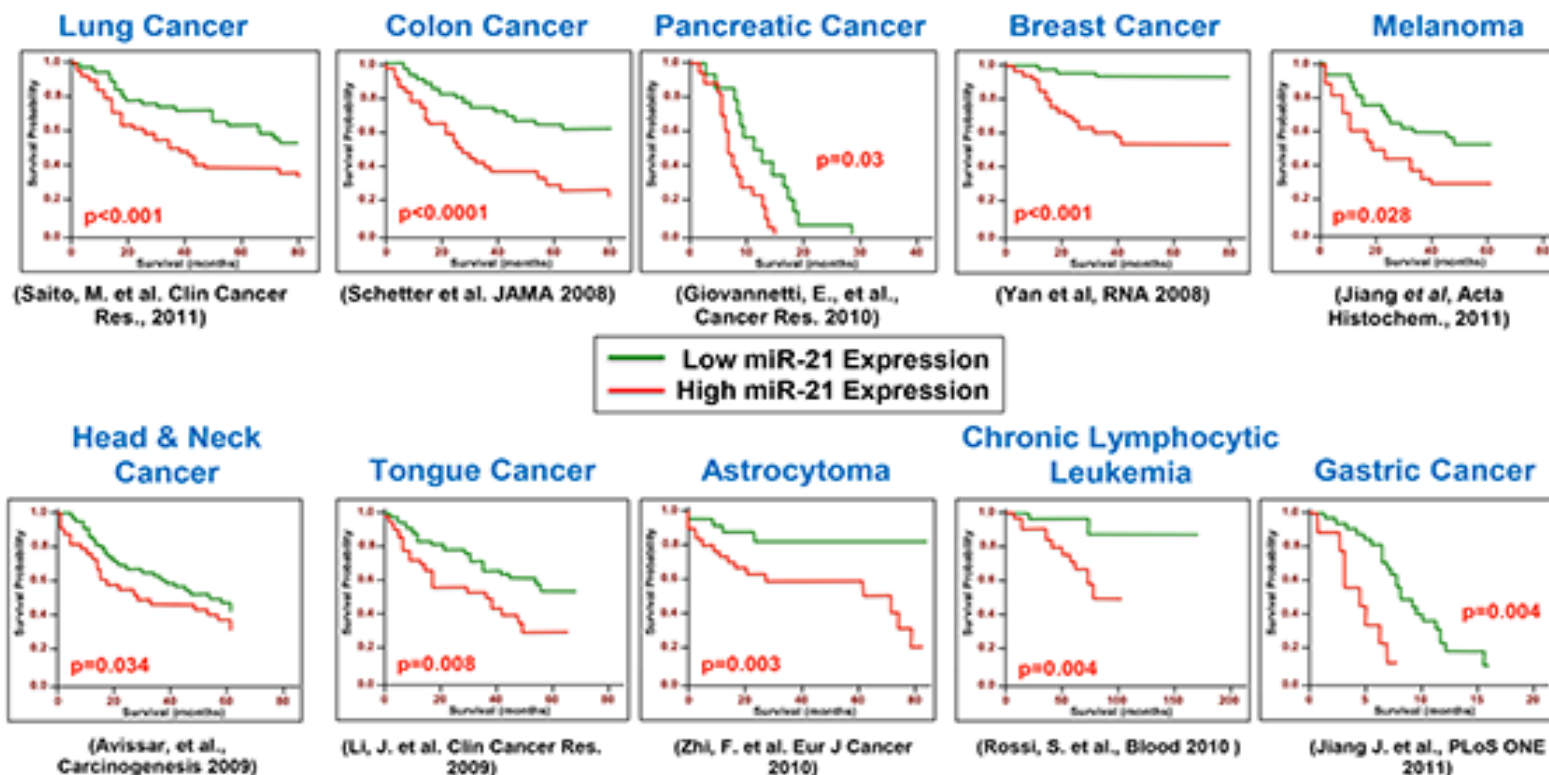
Nozumu Yanaihara

Yanaihara *et al.* Cancer Cell 9:189, 2006

Saito *et al.* Clinical Cancer Res 17:1875, 2011

miR-21 expression

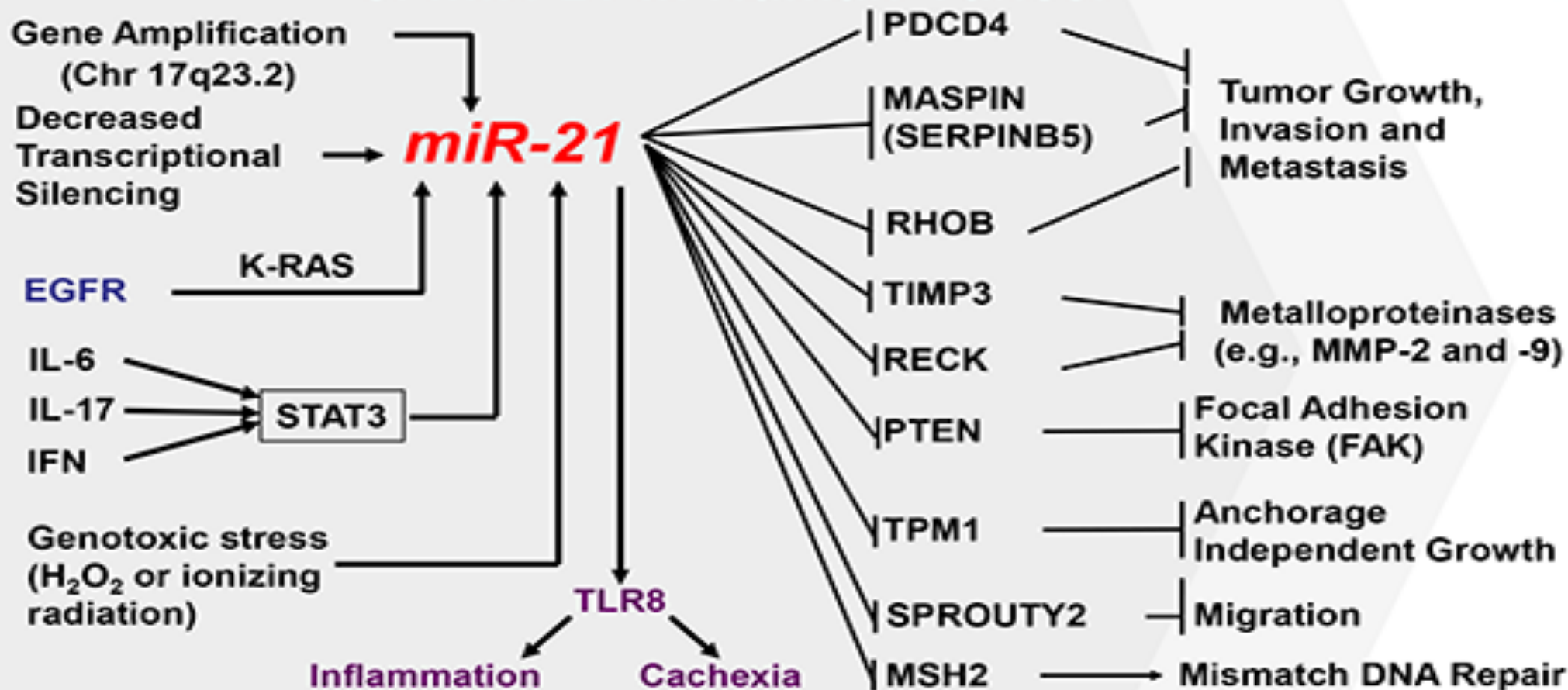
Increased *miR-21* Expression is Associated with Poor Prognosis in Multiple Types of Cancers



miR-21



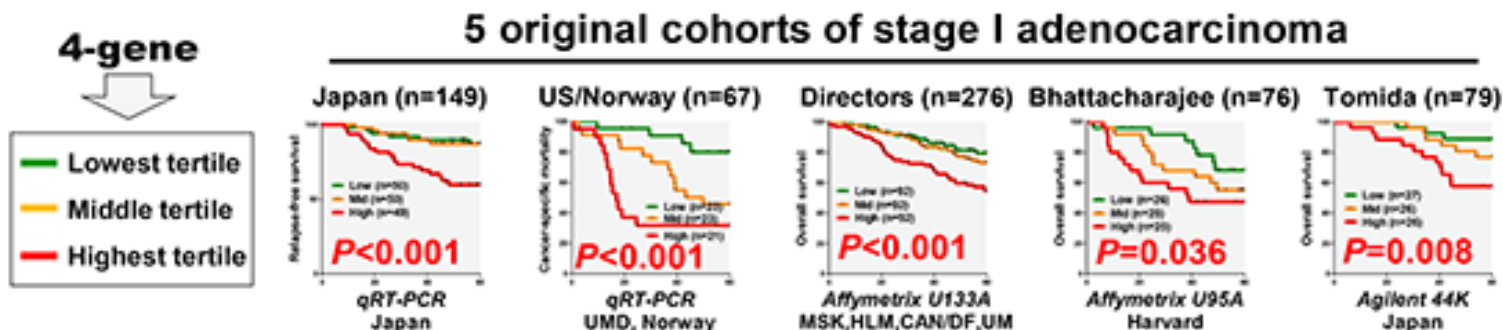
Examples of the Mechanistic Underpinning of *miR-21* in Human Cancer



SeikeHarris, PNAS 106: 12085-90, 2009
Schetter and Harris, Carcinogenesis 31: 37-49, 2010
Fabbri and Croce, RNA Biology 10: 169-174, 2013
He and Croce, PNAS 111: 4525-29, 2014

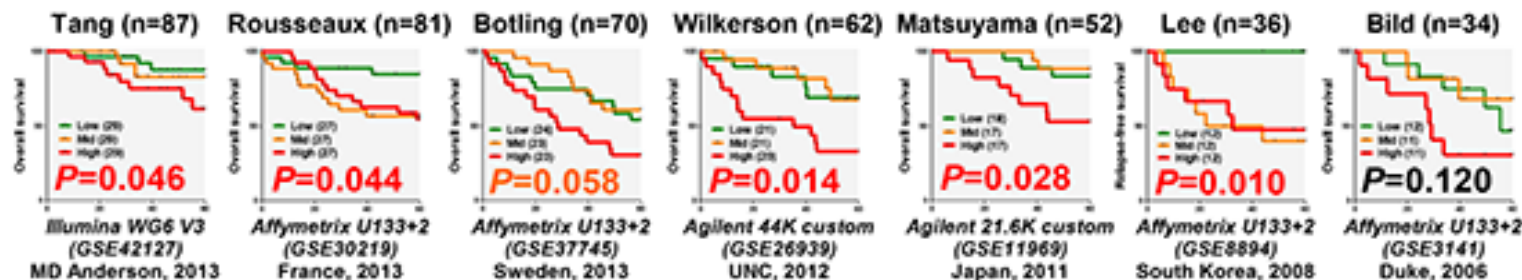
4-gene classifier

The Performance of the 4-gene Classifier (Increased XPO1, BRCA1 and HIF1a and Decreased DLC1) in 12 Independent Cohorts of Stage I ADC



• Akagi, I, Okayama, H et al, Cancer Res 73:3821-32, 2013

7 new cohorts of stage I adenocarcinoma



NIH NATIONAL CANCER INSTITUTE • Okayama, H, Schetter, A ...Harris CC Cancer Epi Prev Biomarker 23:2884-94, 2014.

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Meta-Analysis

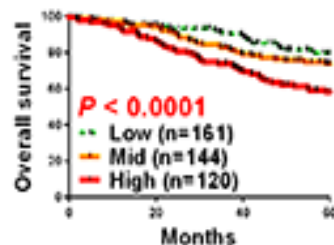
Meta-Analysis of the 4-gene Classifier (↑ XPO1, BRCA1, HIF1a, ↓ DLC1) in 12 Independent Cohorts of Stage I ADC



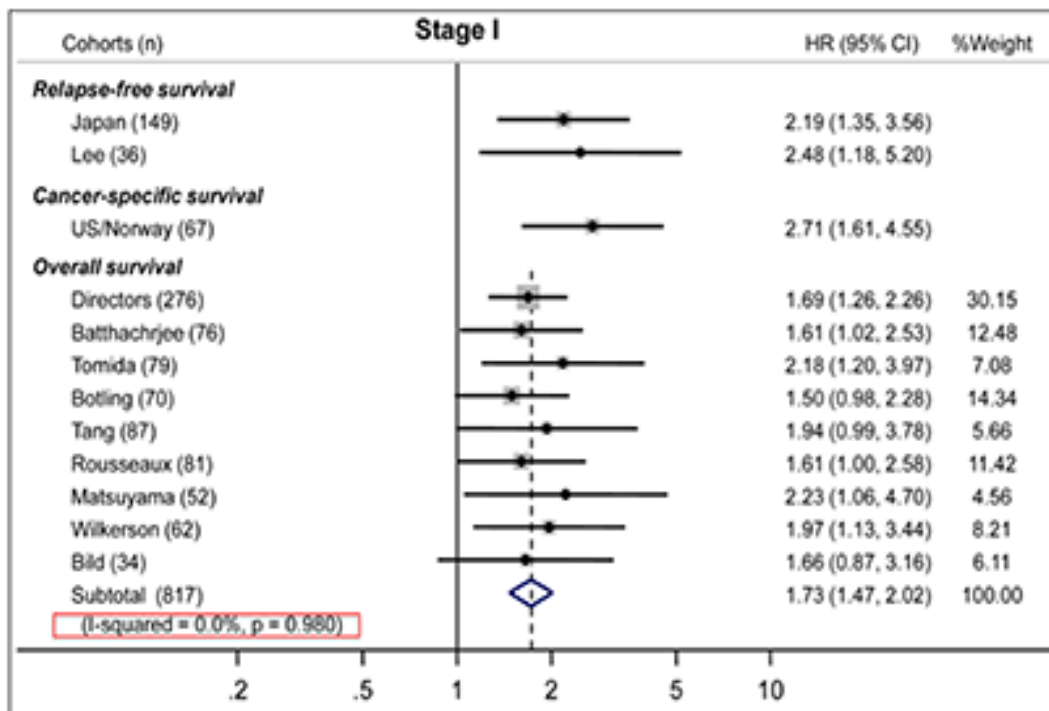
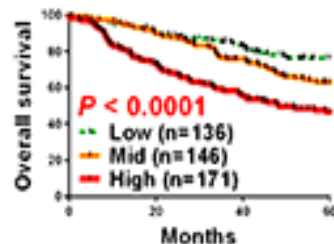
- 883 Stage I Patients

- Combined models included 9 datasets (n=817) with overall survival information

4-gene in Stage IA



4-gene in Stage IB



Okayama H, Schetter A ...Harris CC Cancer Epidemiol Biomarkers Prev 23:2884-94, 2014

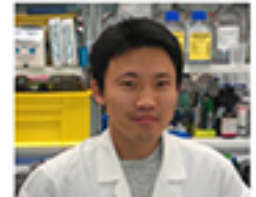
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Mechanistic cancer biomarkers

Mechanistic Cancer Biomarkers: Proof of Principle in Lung Adenocarcinoma



- Hypothesis: The combination of protein-coding genes that are mechanistically related to lung Adenocarcinoma and the non-coding mir-21 is a better prognostic classifier than either alone



Hiro Okayama

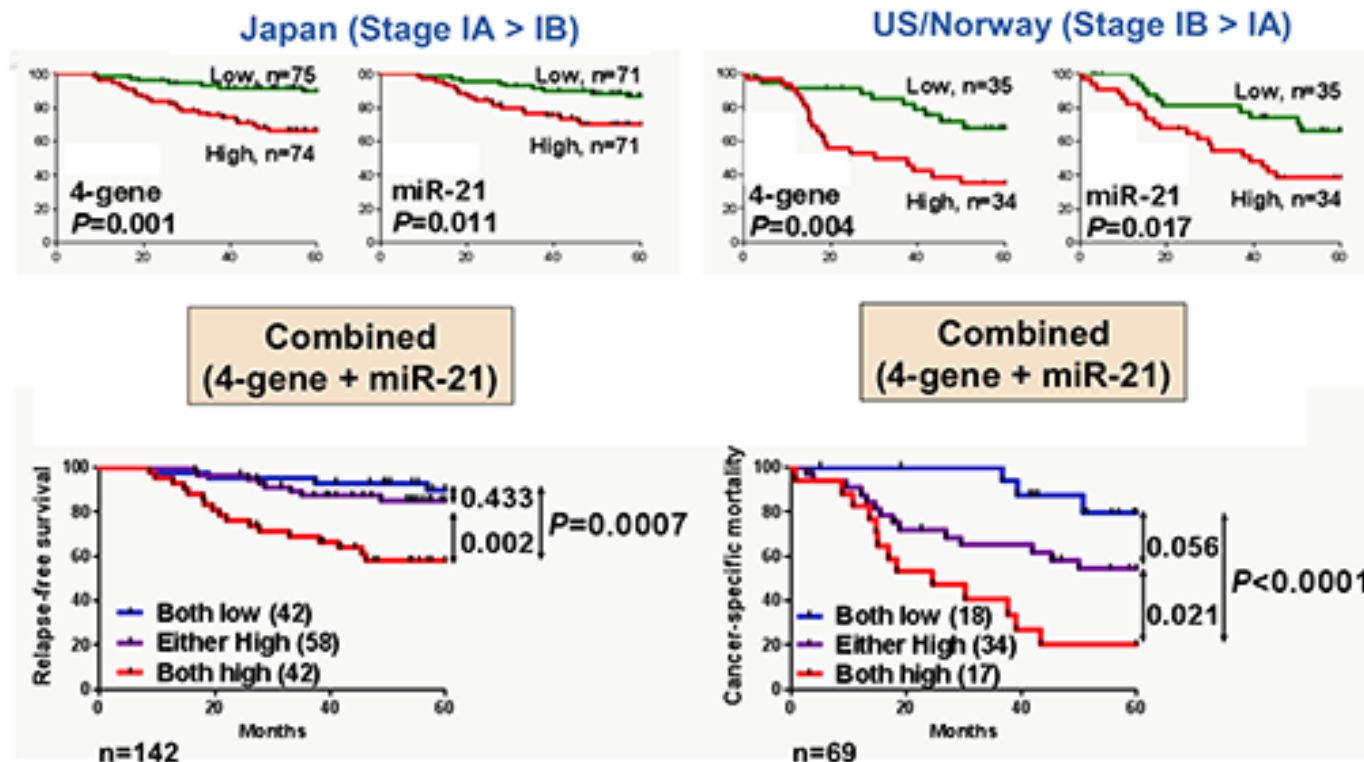


Ichiro Akagi

Akagi et al, Cancer Res. 73: 3821-32, 2013

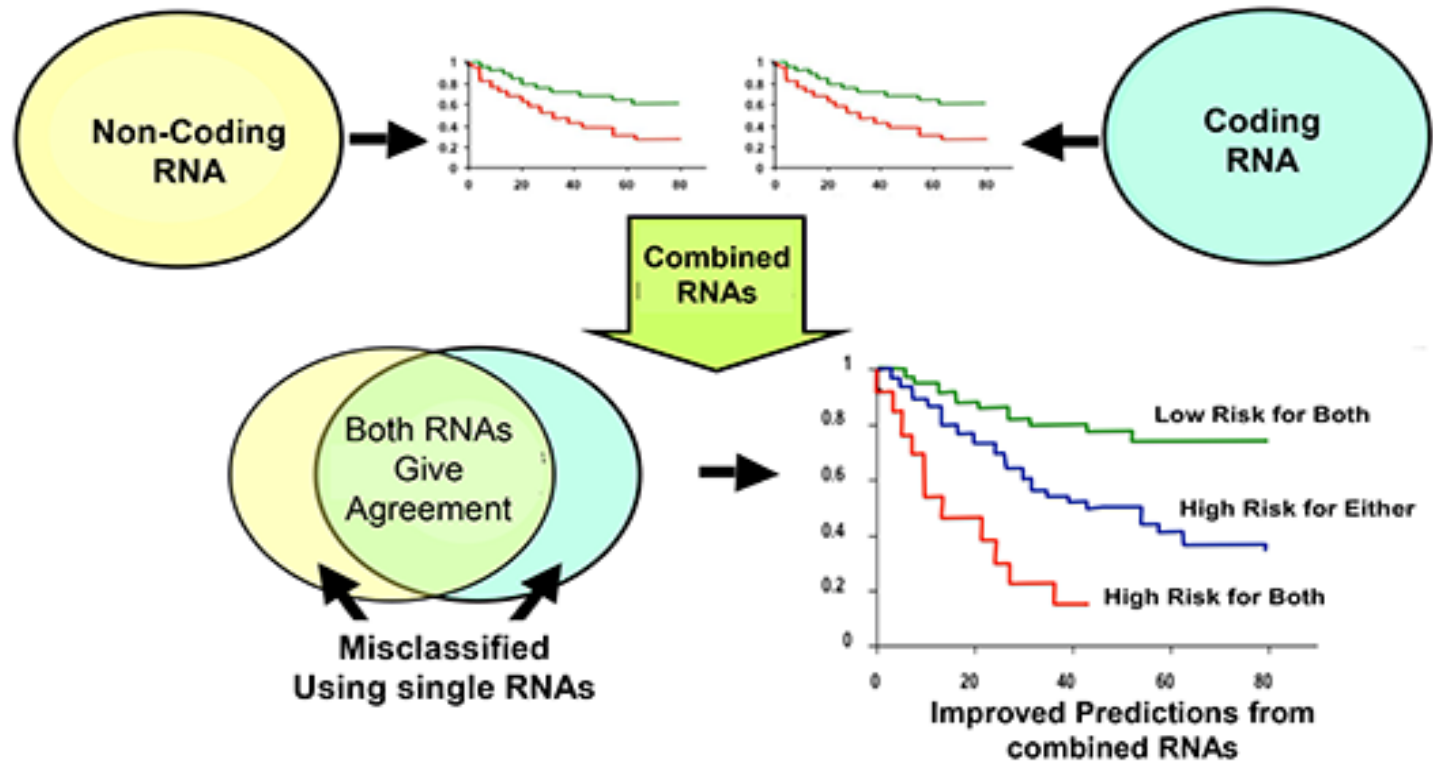
miR-21 and 4 coding genes

Hypothesis: The Combination of miR-21 and Four Coding Genes (Increased XPO1, BRCA1 and HIF1a and Decreased DLC1) Predicts Prognosis of Stage I Lung Adenocarcinoma



Non-coding RNA

Principle: The Combination of Protein Coding and Non-Coding Gene Expression is a more Robust Prognostic Classifier of Early Stage Cancer



Proof of Principle

Further Proof of Principle

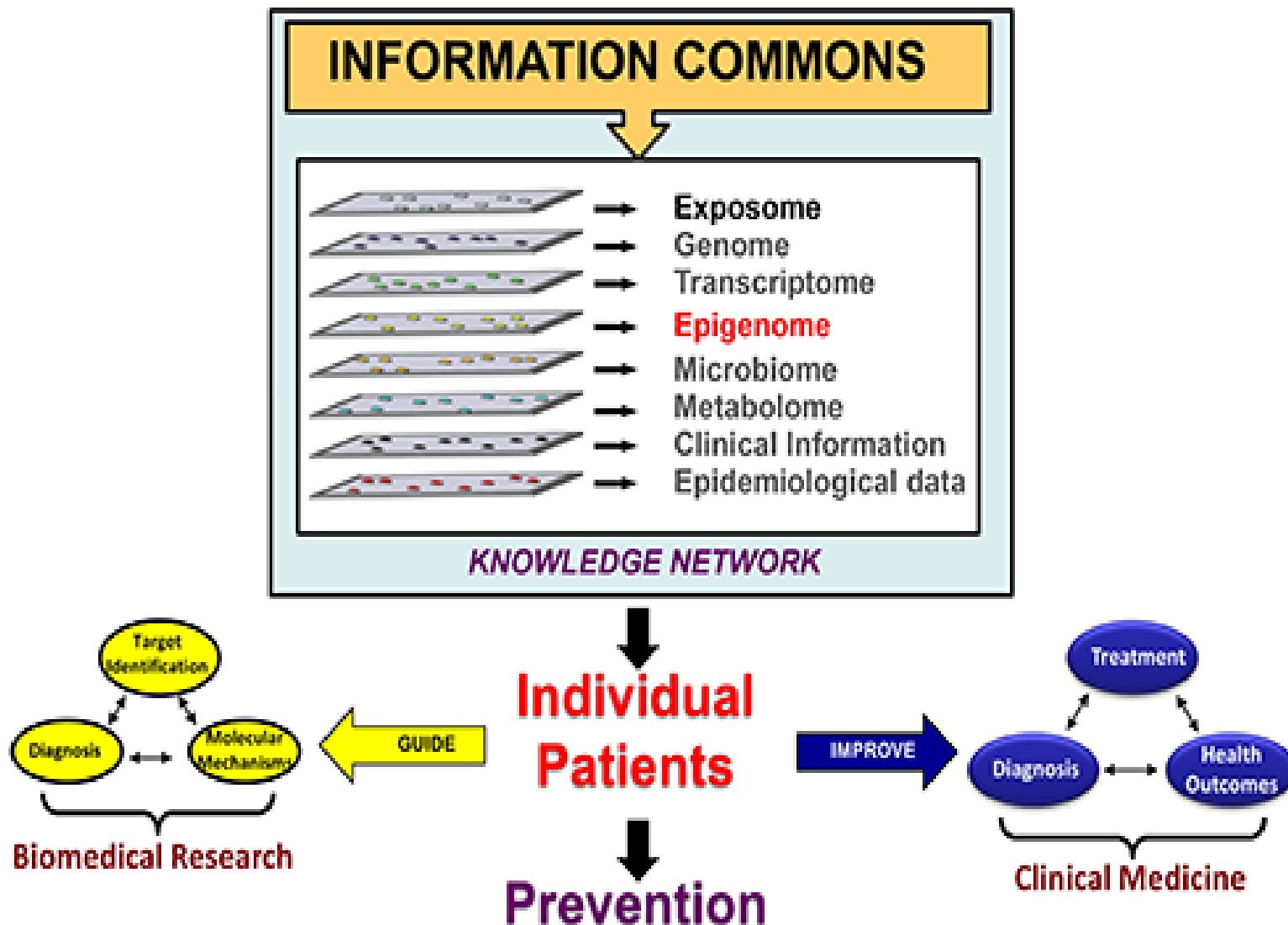


- **Colon Carcinoma**
Schetter et al Clin. Cancer Res 15: 5878, 2009.
- **Esophageal Adenocarcinoma**
Nguyen et al Clin. Cancer Res 16: 5824, 2013
- **Esophageal Squamous Cell Carcinoma**
Zhao et al Int J Cancer 132: 2901, 2013
- **Lung Adenocarcinoma**
Akagi et al Cancer Res 73: 3821, 2013
- **Breast Adenocarcinoma**
Volinia et al P.N.A.S. 110:7413, 2013

Epigenome



Precision Medicine



DNA methylation

EPIGENOME: DNA METHYLATION



- **HYPOTHESIS:** An integrated biomarker classifier of stage I lung adenocarcinoma based on independent mRNA, microRNA and DNA methylation biomarkers, will further improve the prognostic classification



Ana Robles

Robles a et al., J. Thoracic Oncology 10: 1037-48, 2015

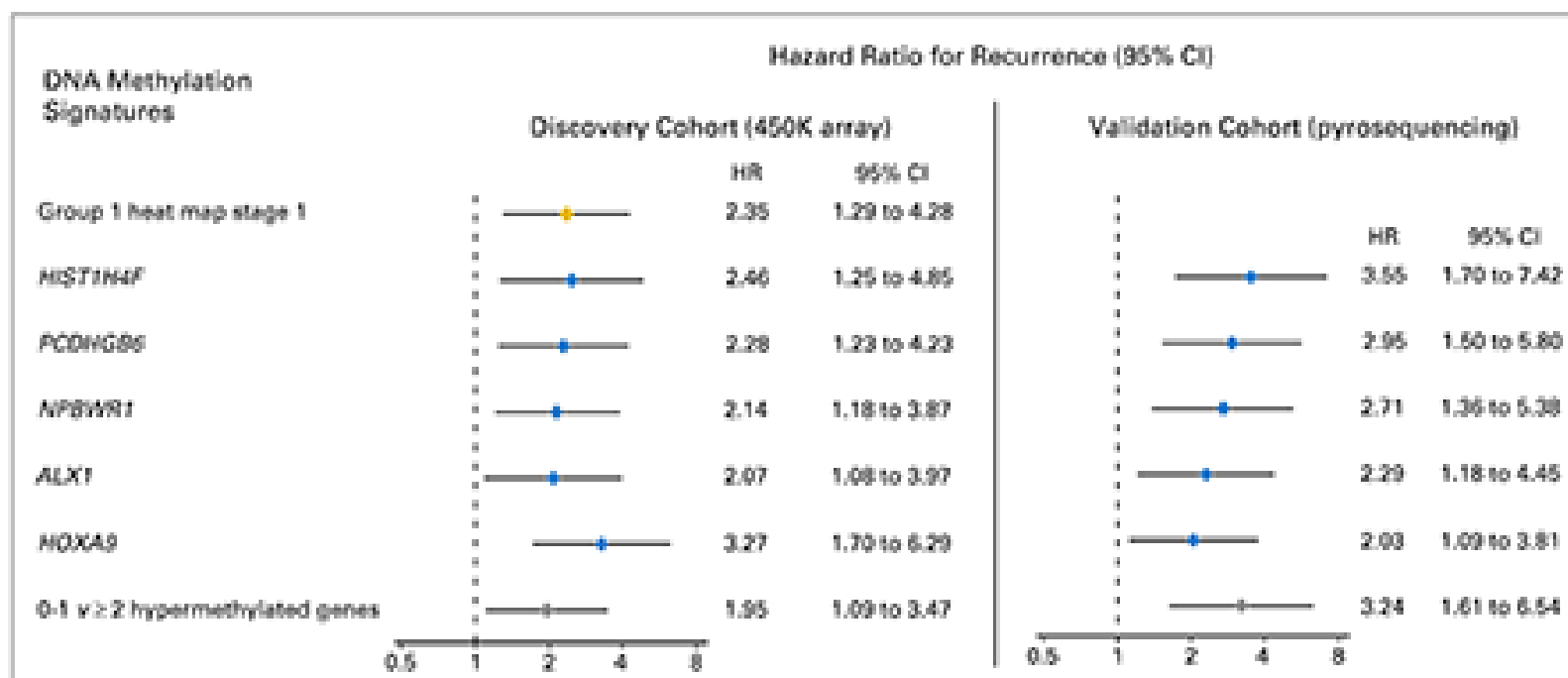
Prognostic epigenetic signature

DNA Methylation in NSCLC is a Prognostic Epigenetic Signature

A Prognostic DNA Methylation Signature for Stage I Non-Small-Cell Lung Cancer

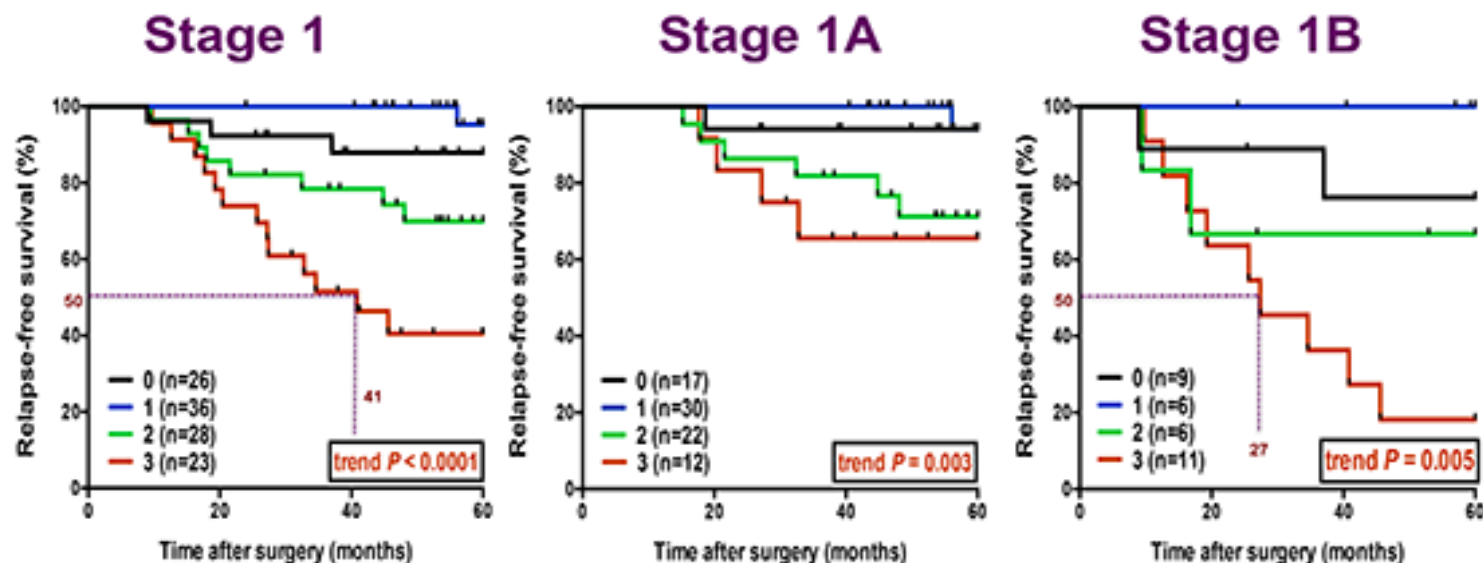
Juan Sandoval, Jesus Mendez-Gonzalez, Ernest Nadal, Guoan Chen, F. Javier Carmona, Sergi Sayols, Sebastian Moran, Holger Heyn, Miguel Vizoso, Antonio Gomez, Montse Sanchez-Carpes, Yassier Avcinor, Fabian Müller, Christoph Bock, Miquel Taron, Josefina Mora, Lucia A. Muscarella, Triantafyllos Liloglou, Michael Davies, Marina Pollan, Maria J. Pajares, Wenceslao Torre, Luis M. Montuenga, Elisabeth Brambilla, John K. Field, Luca Roz, Marco Lo Jacomo, Giorgio V. Scagliotti, Rafael Rosell, David G. Beer, and Manel Esteller

(J Clin Oncol 2013;31(32):4140-7).



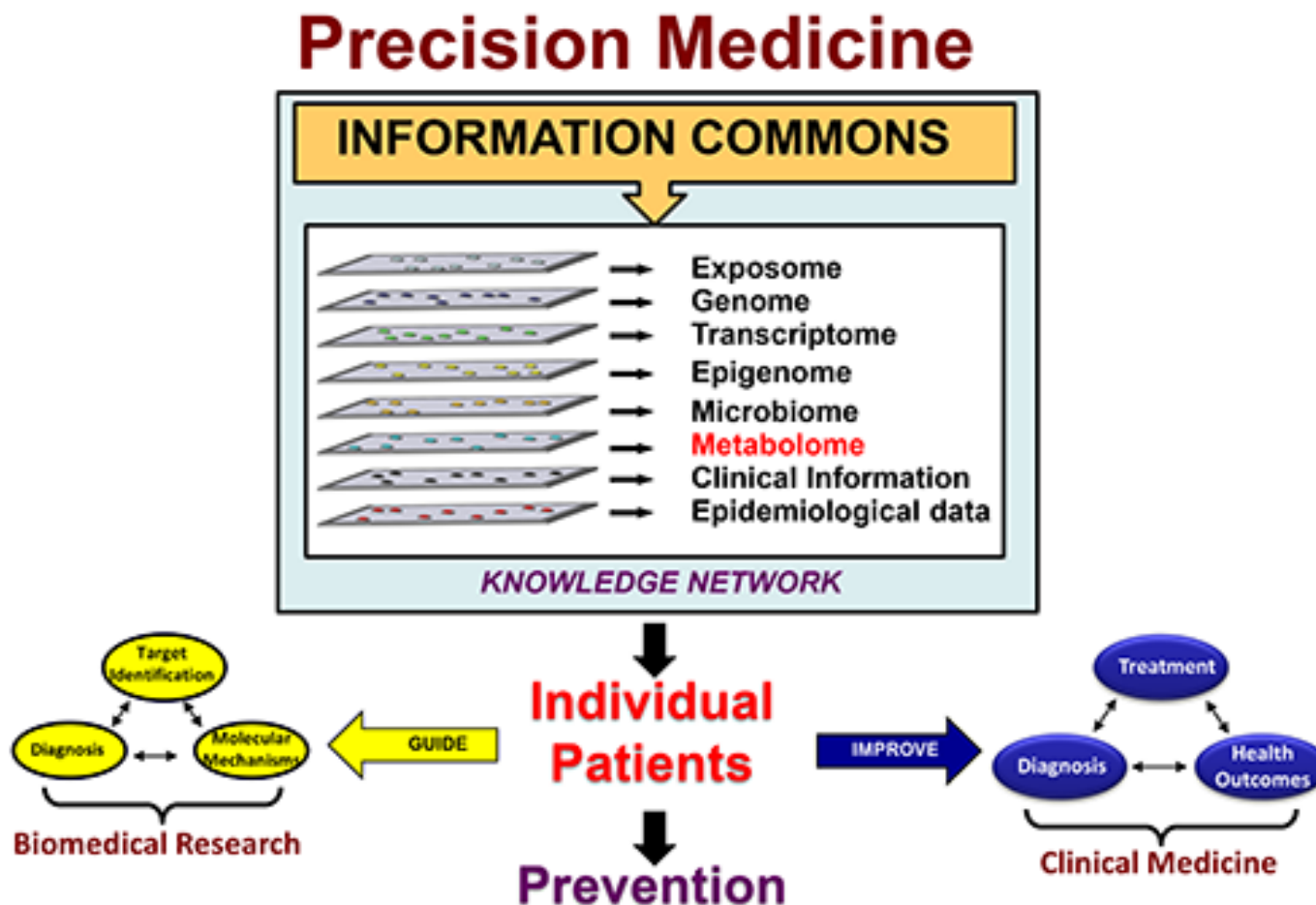
Survival curve

An Increasing Combined Score of Mechanistic Biomarkers Conferred Greater Risk for Poor Outcome in Stage 1 Lung Adenocarcinoma, and Within Stage 1A/1B Subgroup Analysis



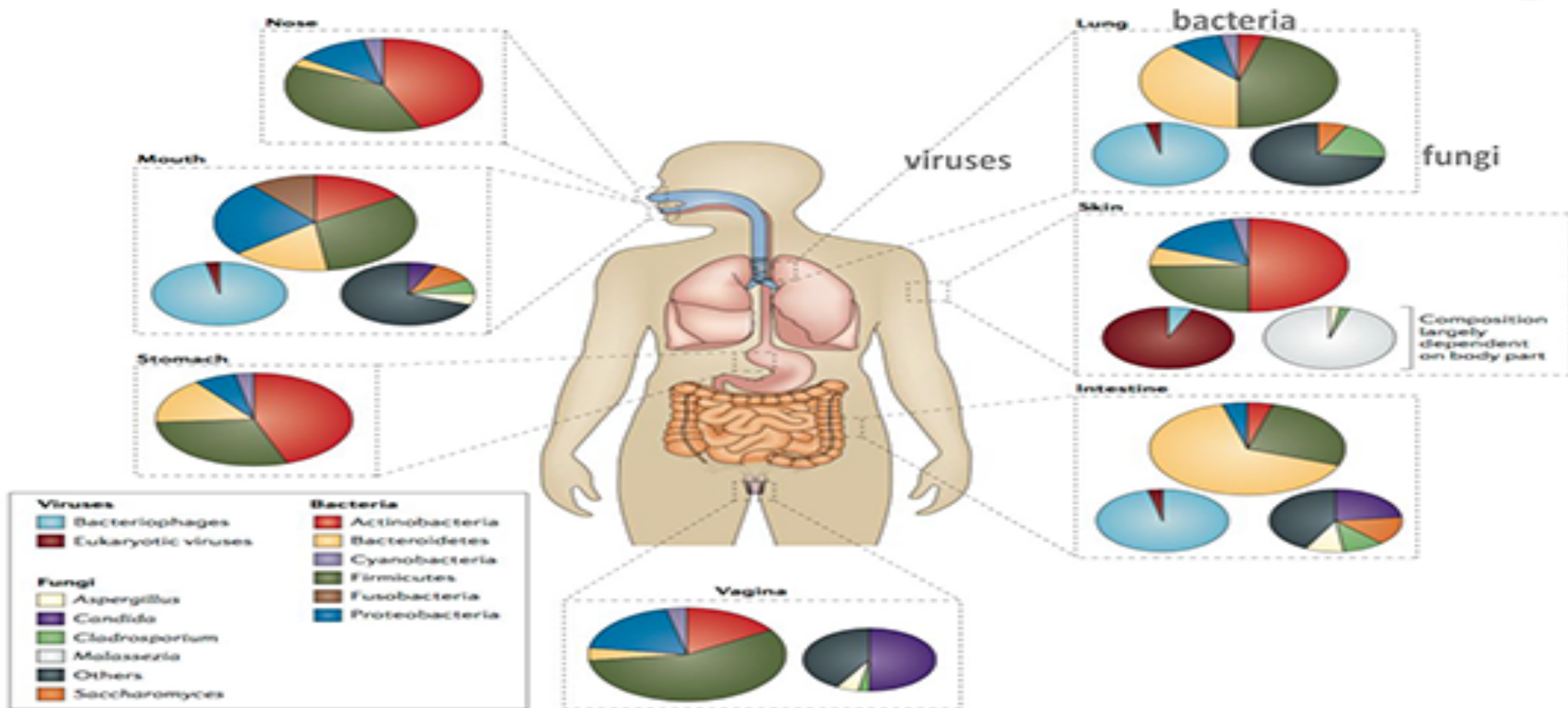
- Cases were categorized according to the combined number of high values for HOXA9 methylation, miR-21 (Clin. Cancer Res. 2011;17:1875-82) and 4-gene signature (Cancer Res. 2013;73:3821-32) and HOXA9 promoter methylation (J. Thoracic Oncology 10: 1037-48, 2015).
- P values calculated by log-rank test for trend.

Microbiome



Heterogeneous microbial habitats

Examples of Heterogeneous Microbial Habitats in Humans



Microbiome and carcinogenesis

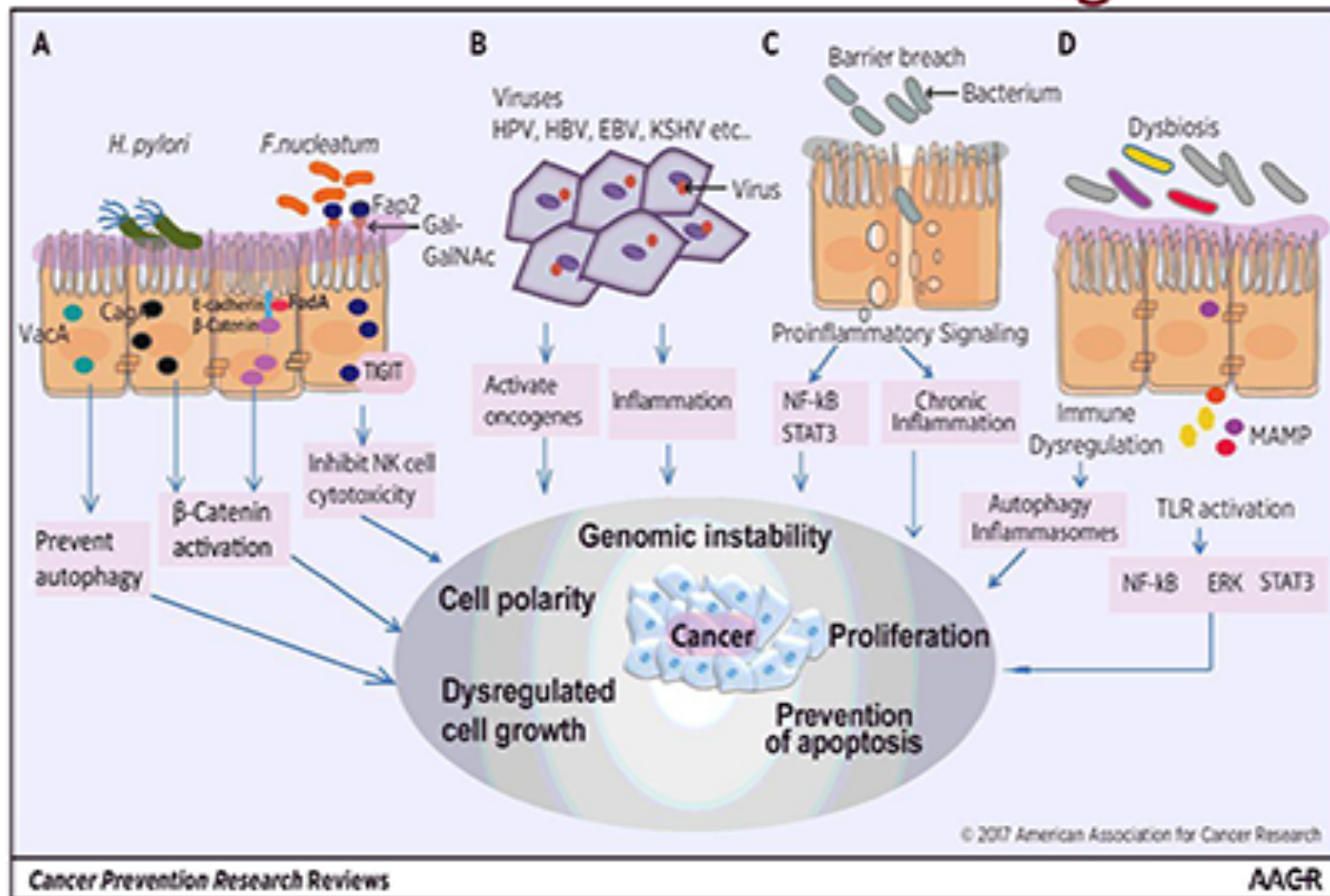
Bacterial and Viral Microbiome Promotes Carcinogenesis

H. pylori
Gastric cancer

F. nucleatum
Colon Cancer

Dysbiosis
Multiple Cancers
Cancer Therapy

HPV
Cervical and Head and Neck Cancer



Microbiome, TP53 and lung cancer

Interaction between the Microbiome, Smoking and TP53 in Human Lung Cancer



- Certain taxa including *Acidovorax* were higher in both adenocarcinoma and squamous carcinoma (SCC) in smokers and former smokers than in non-smokers.
- A group of taxa were significantly associated with SCC, of which *Acidovorax* was enriched in smokers
- SCC cases with *TP53* mutations had a higher abundance of the SCC-associated taxa, including *Acidovorax*

SCC-associated taxa are enriched in tumors with *TP53* mutations, which establishes a microbiome-gene interaction in lung cancer tissue

Metabolome



Precision Medicine

INFORMATION COMMONS



KNOWLEDGE NETWORK

Individual Patients

Prevention



Biomedical Research

GUIDE

IMPROVE



Clinical Medicine

Tumor metabolism

Tumor Metabolism and Metabolome



- **Hypothesis:** Unbiased metabolomics will discover biomarkers associated with the risk, diagnosis, prognosis and therapeutic outcome of lung cancer.



Majda Haznadar



Ewy Mathe

Urinalysis

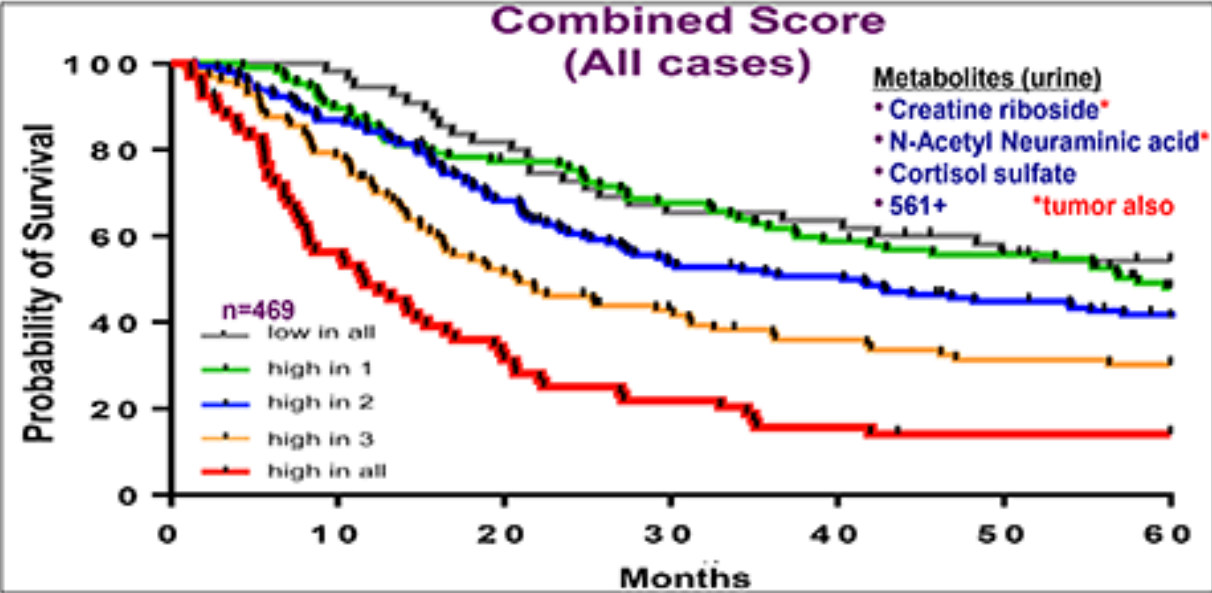


- Hippocrates tasted urine in the diagnosis of disease in his patients. (460-370, BC)
- Physiologist J. A. Armstrong writes, “From a liquid window through which physicians felt they could view the body’s inner workings, urine led to the beginnings of laboratory medicine...”



Survival curve

Lung Cancer Metabolome and Prognosis



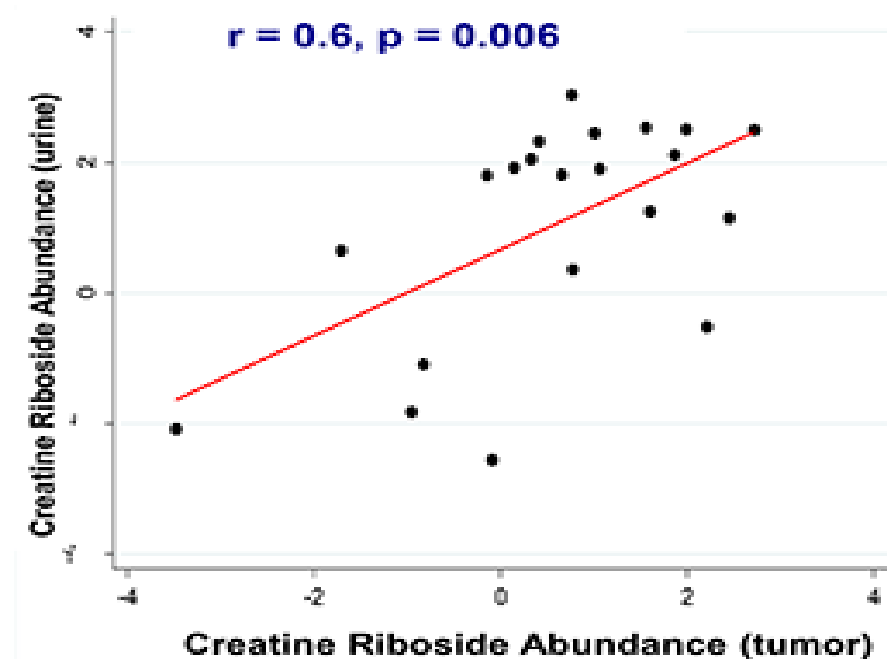
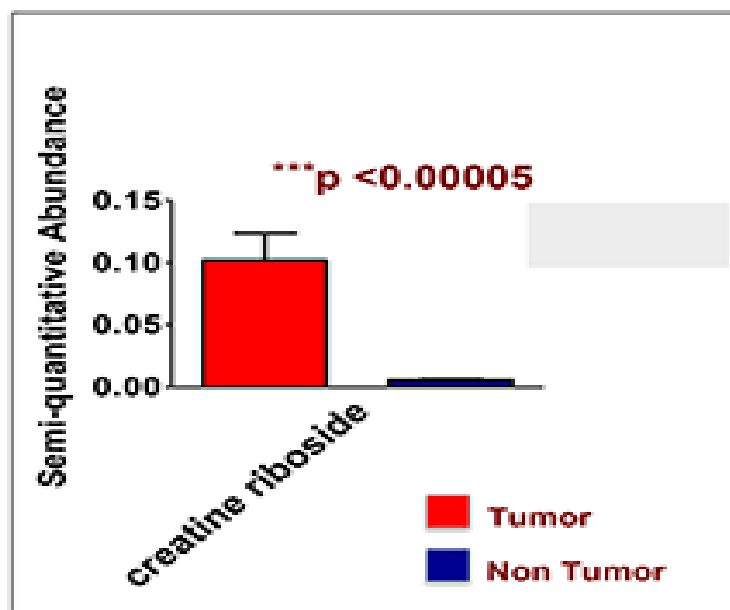
	HR (95% CI)	P
Reference	1	1.00
High in 1	1.10 (0.67-1.75)	0.75
High in 2	1.36 (0.86-2.14)	0.19
High in 3	1.84 (1.15-2.94)	0.01
High in all	3.65 (2.34-6.00)	<0.00001

Creatine ribose

Creatine Ribose is Positively Correlated in Cancer and Urine



Lung Adenocarcinoma
(targeted UPLC-MS/MS)
N = 48 (Tumor), 48 (Non Tumor)



CT screening

Low-Dose CT Screening for Lung Cancer Detects Early Stage and Reduces Mortality



The **NEW ENGLAND**
JOURNAL of MEDICINE

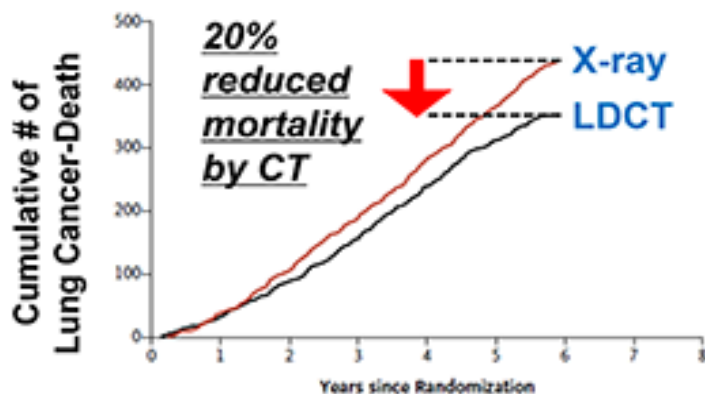
ESTABLISHED IN 1812

AUGUST 4, 2011

VOL. 365 NO. 5

Reduced Lung-Cancer Mortality with Low-Dose Computed Tomographic Screening

The National Lung Screening Trial Research Team*



NIH NATIONAL CANCER INSTITUTE

n=53,454 participants
Current/former heavy smokers, 55-77 y.o.

X-ray

n=26732

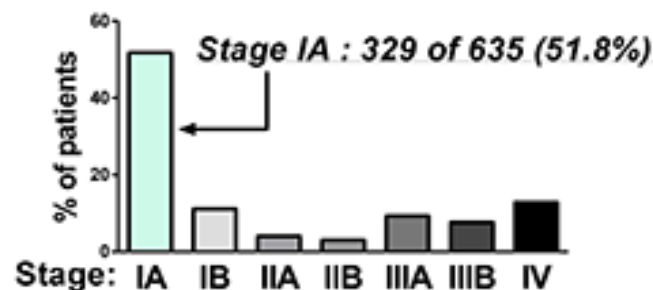
LDCT

n=26722

Low-Dose CT

• False Positive Rate = 96.4%

Stage of Lung Cancers Diagnosed After Positive CT Screening



8167-CH

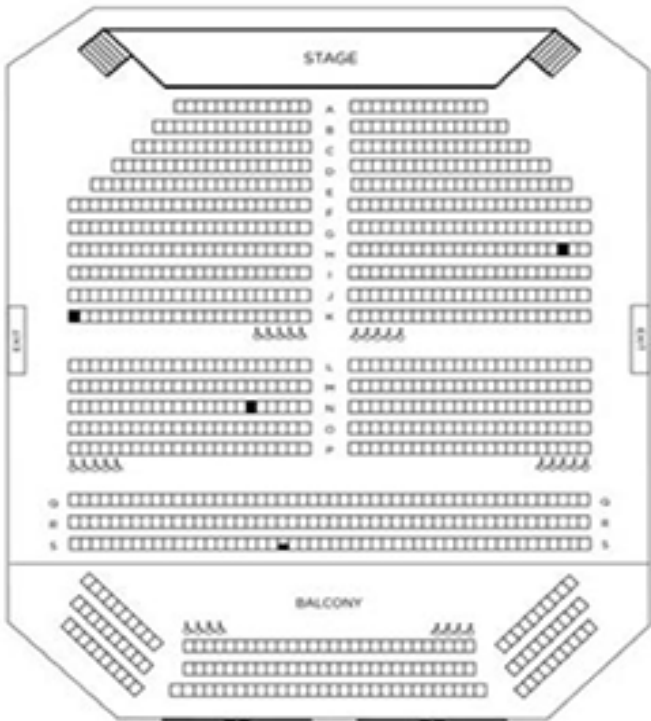
Spiral CT scans

Analogy of the Number of Lung Cancer Deaths Averted with Annual Spiral CT Scans, Compared to the Total Number of Scans Done

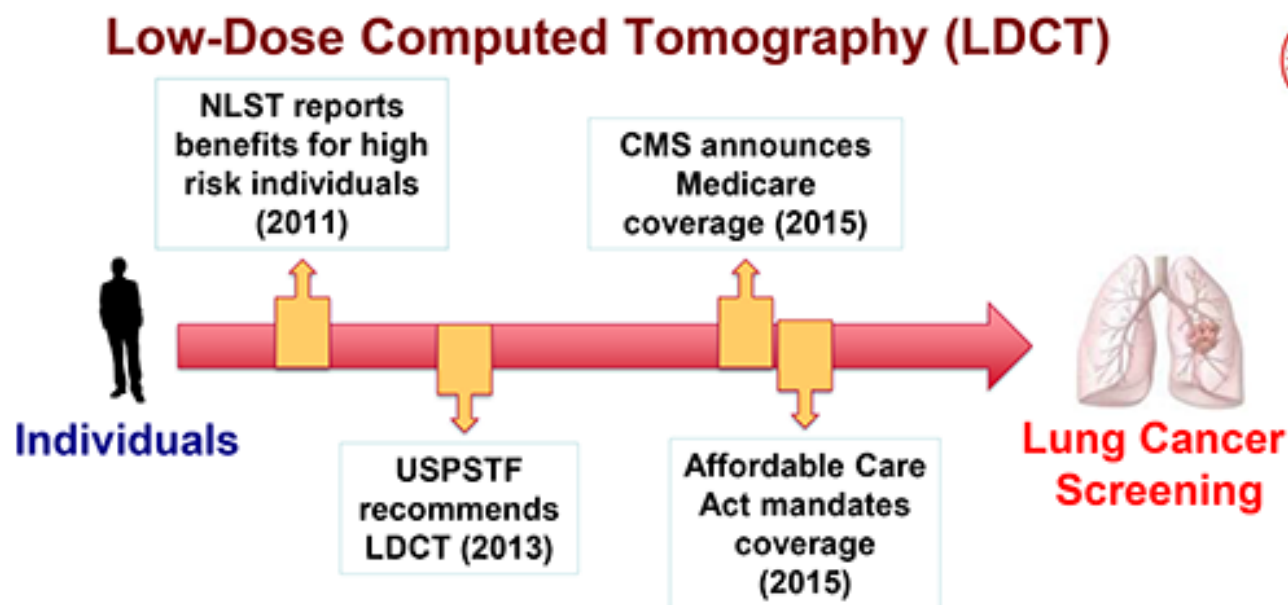


Lung Cancer

False Positive



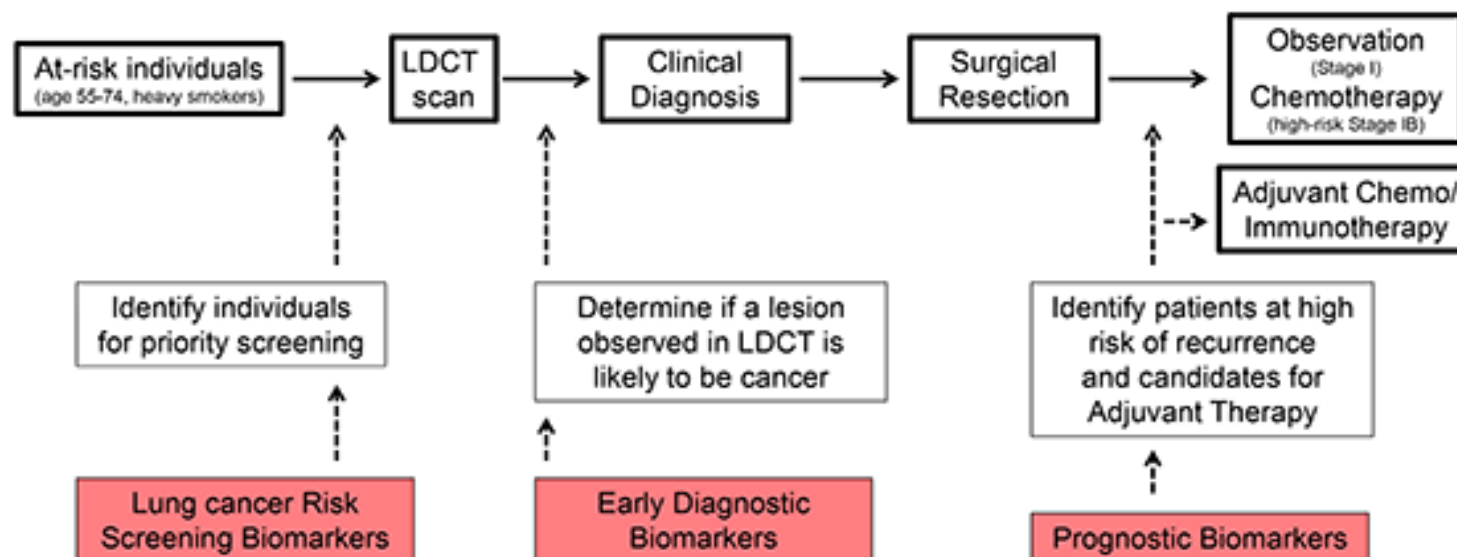
Low dose CT



- High risk: age 55-74, 30 pack-year history, quit within 15 years or current smoker
 - Need to identify high risk individuals outside criteria?
 - Need to identify individuals within the criteria and prioritize those with the greatest risk?
- 8.8 million Americans were LDCT-eligible (2010)

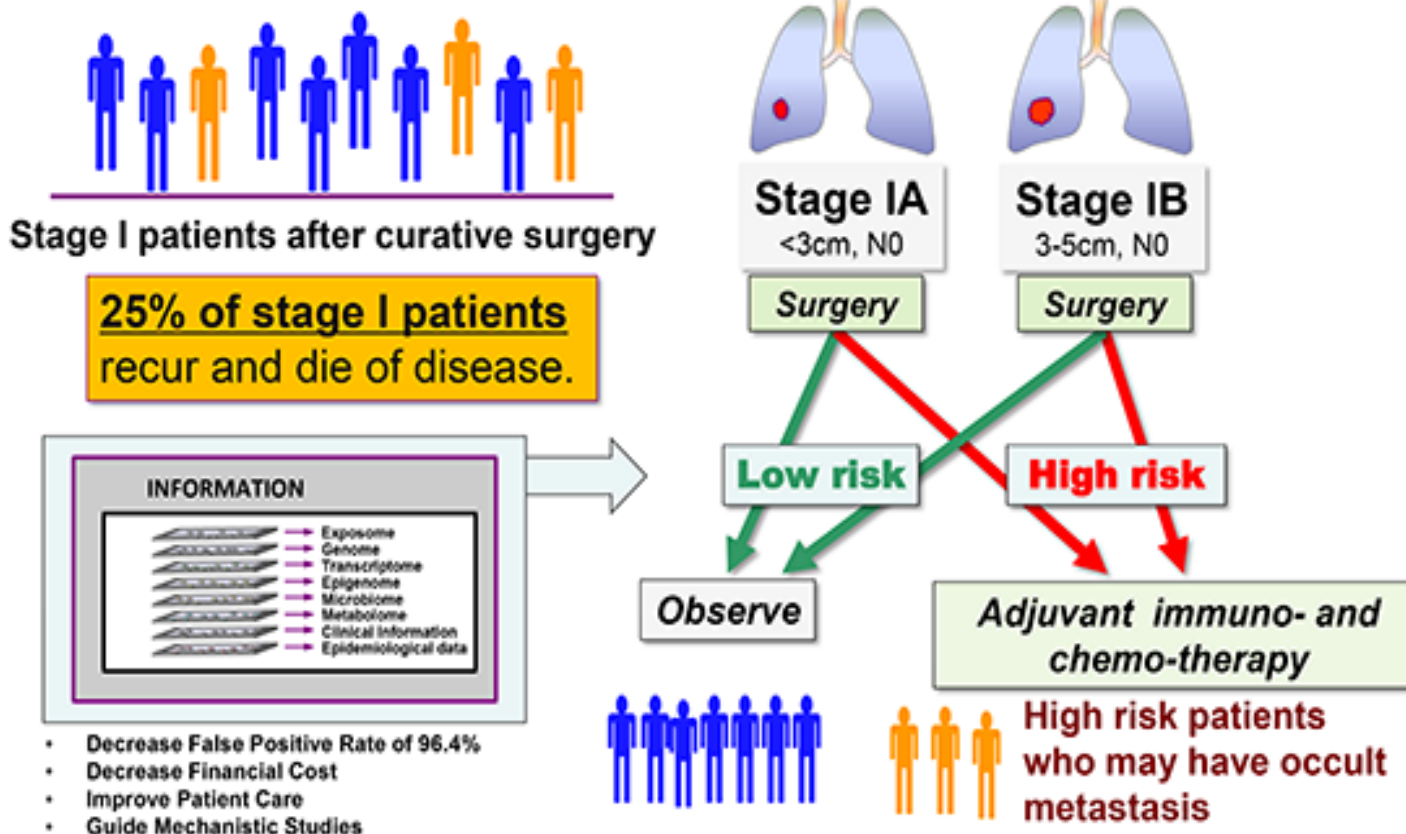
Unmet needs

Early Stage Lung Cancer: Unmet needs



Precision medicine goals

A Goal of Precision Medicine is to Identify 25% of Stage I Lung Cancer Patients whose Cancer will recur



Collaborators

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