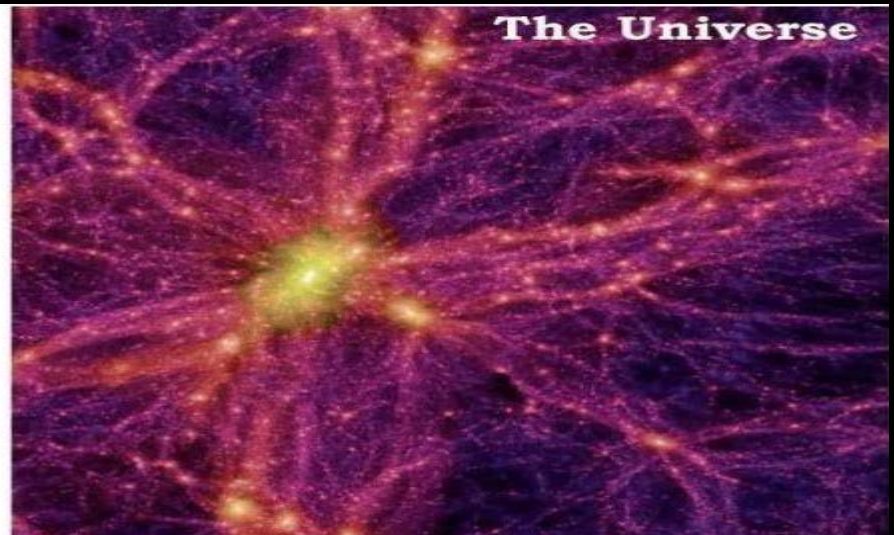
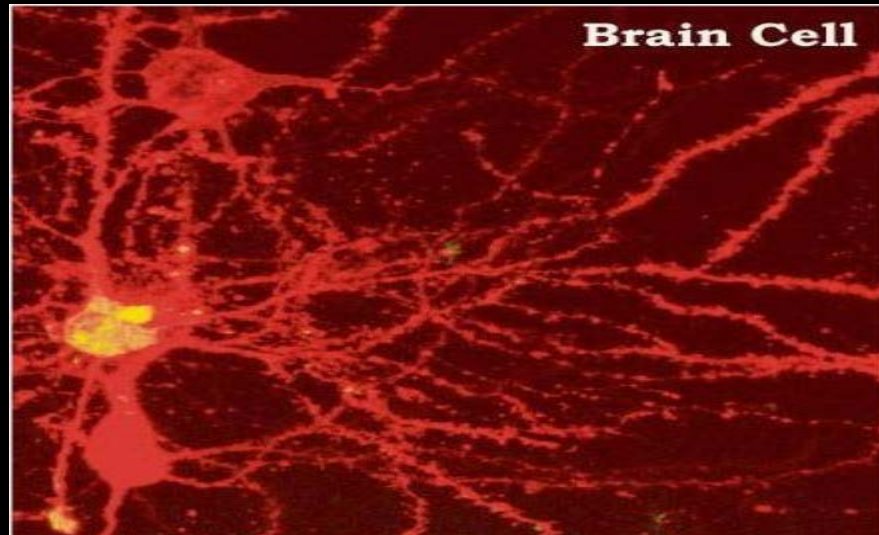


Precision Medicine and the Reclassification of Cancer

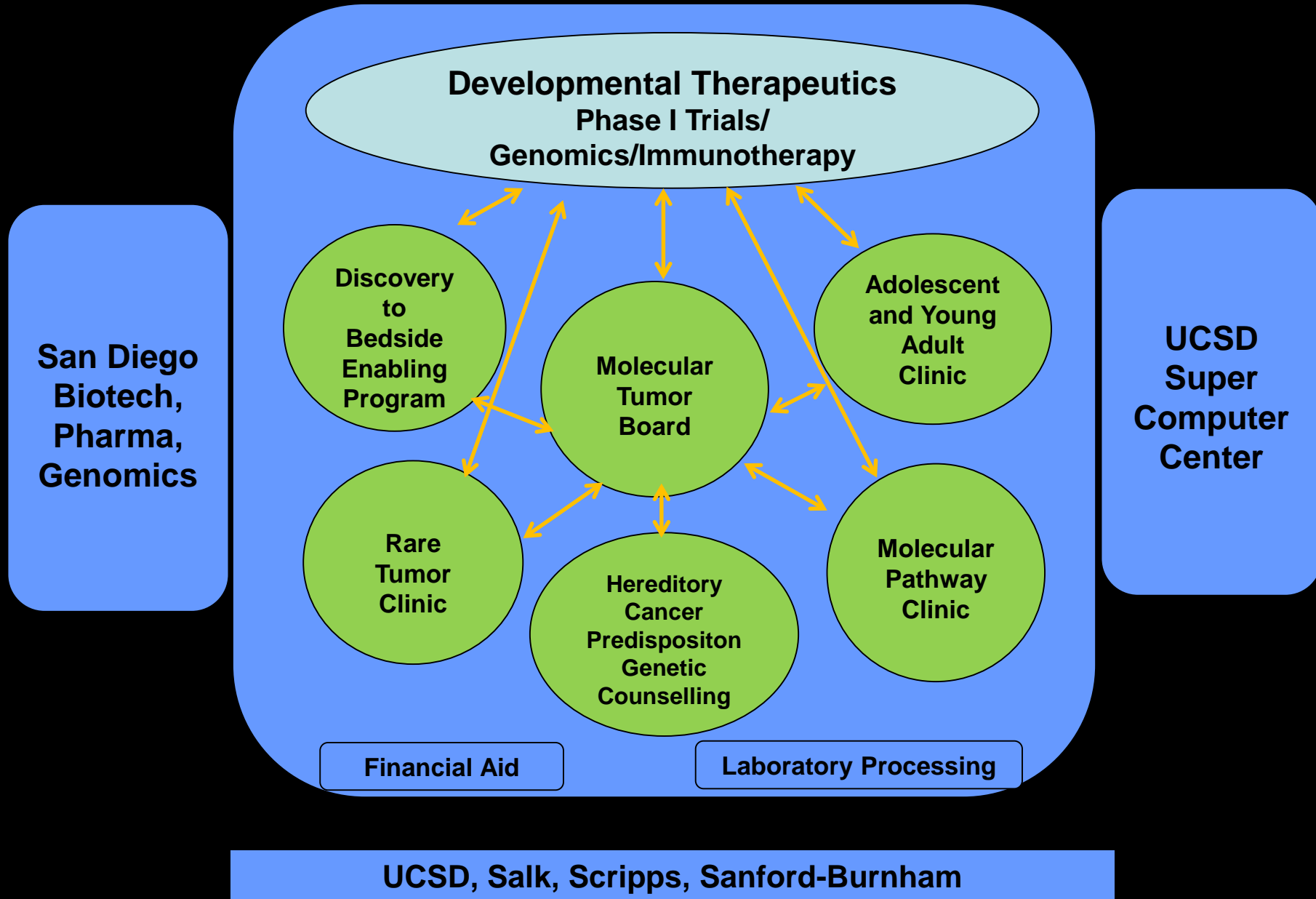
Divide and Conquer

Razelle Kurzrock, MD

Senior Deputy Director , Clinical Science
Director, Center for Personalized Cancer Therapy
Director, Clinical Trials Office
Chief, Division of Hematology/Oncology
UCSD Moores Cancer Center

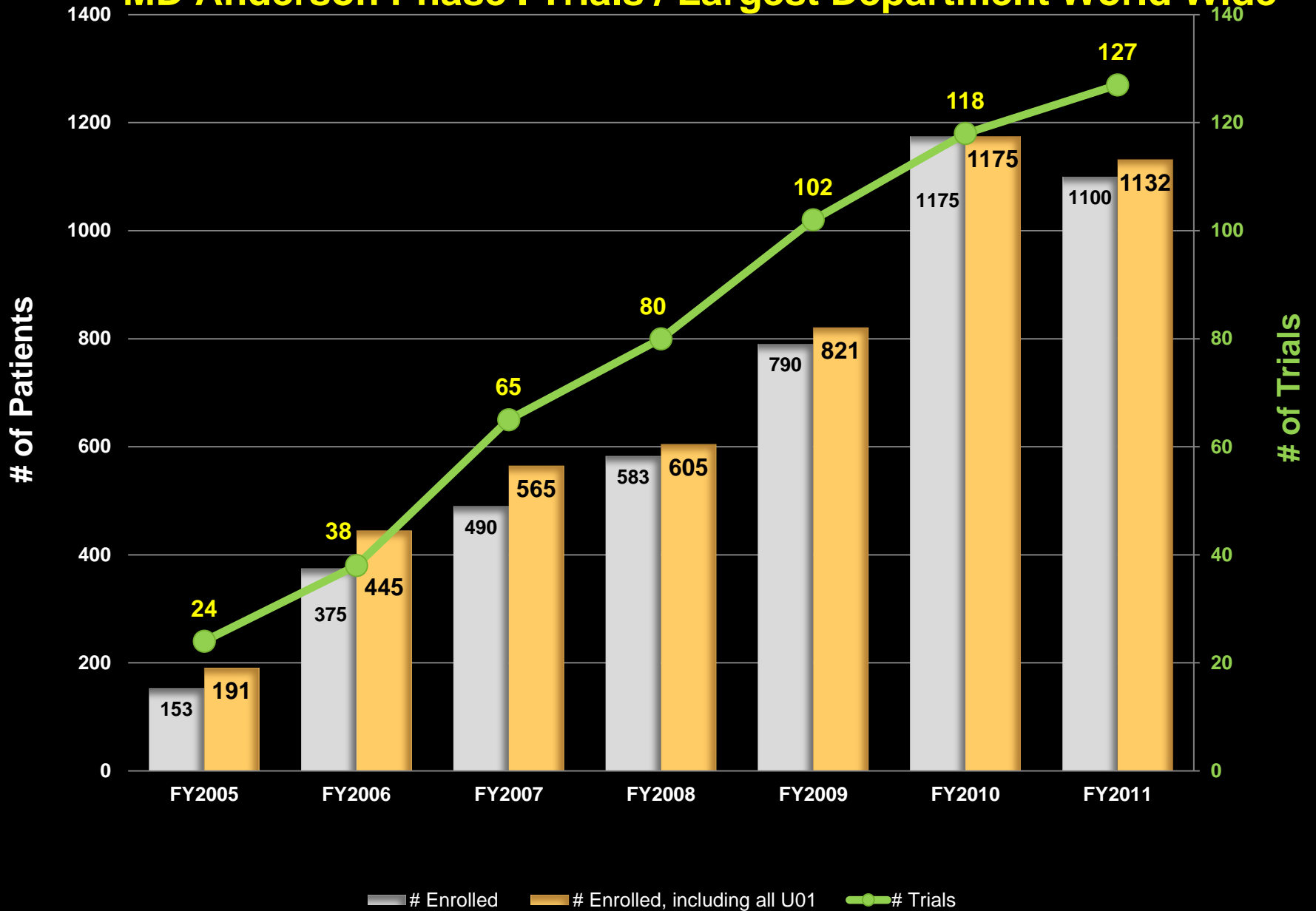


Center for Personalized Cancer Therapy at Moores Cancer Center



Kurzrock Experience

MD Anderson Phase I Trials / Largest Department World Wide



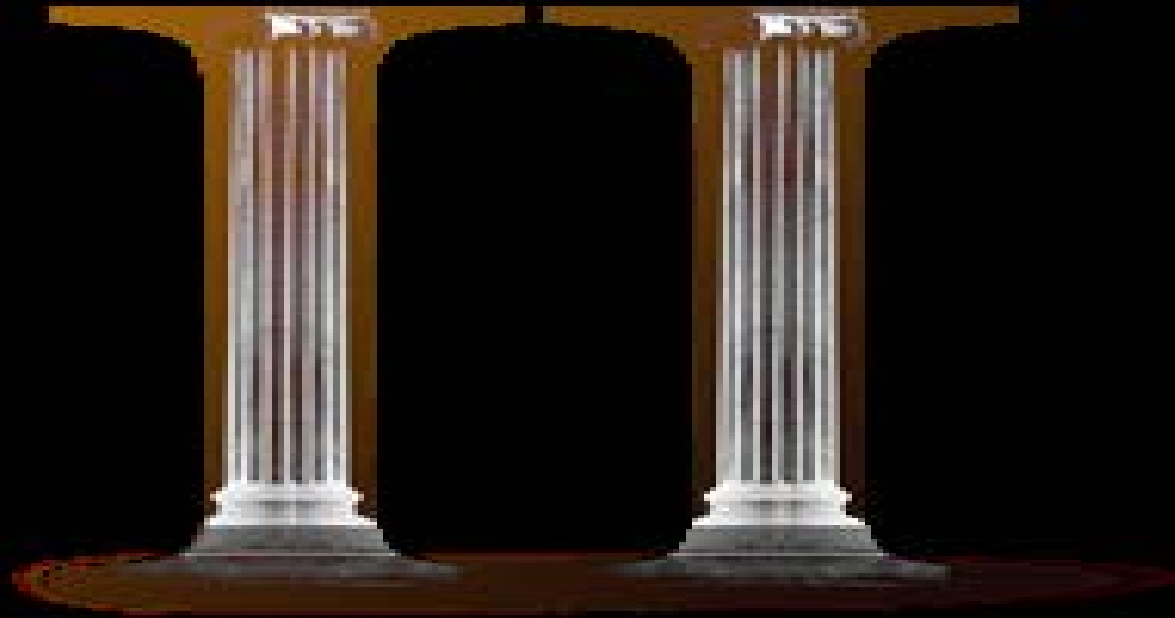
Question: Is it precision medicine or personalized medicine?

Answer: Both

“Precisionalized Medicine”

The Pillars of Precision Cancer Medicine

Genomics Immunotherapy



**The future
is here.**

Precision Medicine

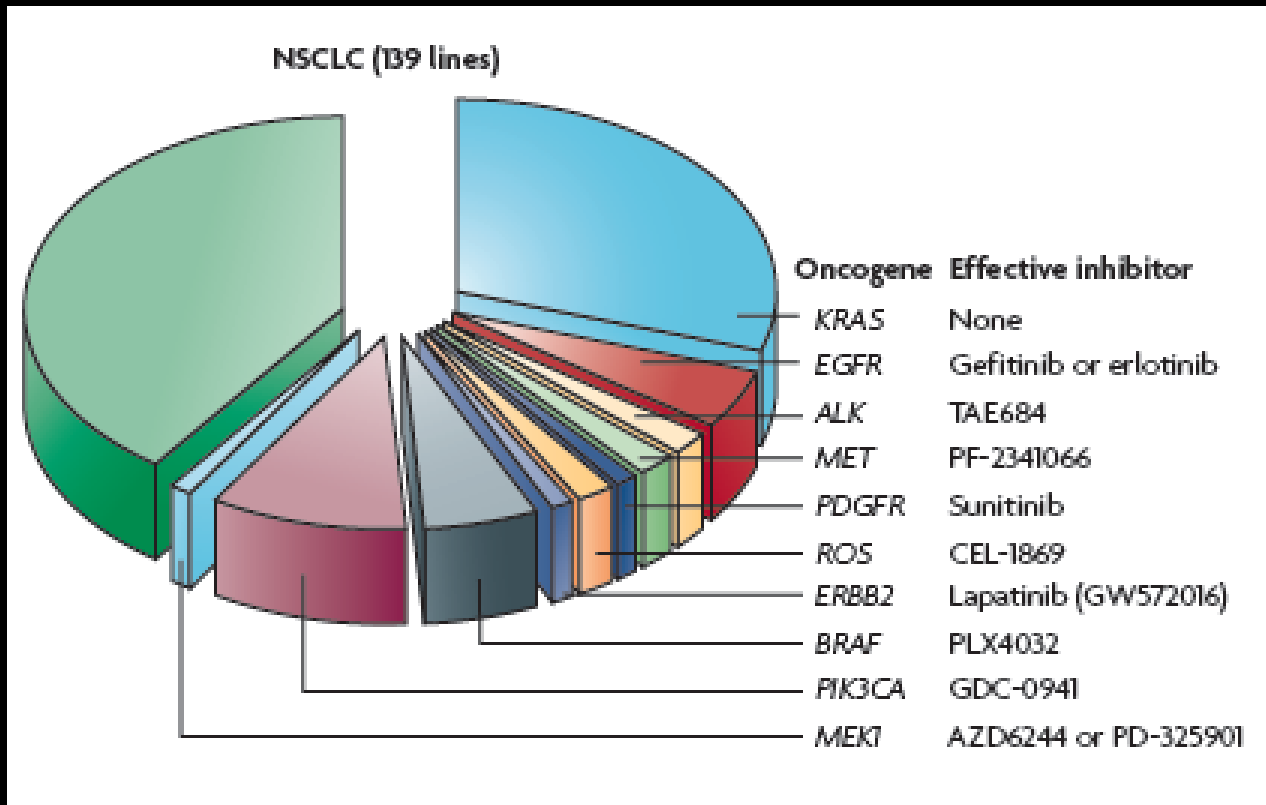
Lessons Learned

- Use combinations of matched drugs for metastatic or complex tumors
- Treat newly-diagnosed patients
- Omics is a disruptive technology; retrofitting the reality unveiled into traditional paradigms is suboptimal
- Harness the immune system
- Transformative changes will require new models for clinical research and practice

Why are cancers difficult to treat?

Divide and Conquer

Agents work only
in those with
a sensitizing
aberration




Braiteh....Kurzrock, MCT 2007

Munoz J, Swanton C, Kurzrock R, Molecular Profiling and the Reclassification of Cancer;
Am Soc Clin Oncol Educ Book. 2013:

Sharma, Nat Rev Cancer 2010

What can patients expect from traditionally approved drugs ?

Drug	Tumor	Survival Gain	Complete remission
gemcitabine	pancreas	1.5 months	≈ 0%
bevacizumab	colon	2.2 months	≈ 0%
erlotinib	pancreas	11 days	≈ 0%
bevacizumab	NSCLC	2 months	≈ 0%
sorafenib	renal	2 months	≈ 0%
temozolamide	glioblastoma	2.5 months	≈ 0%
docetaxel	prostate	2.4 months	≈ 0%
cetuximab	colon	1.5 months	≈ 1-2 %



Master Protocol

Profile-Related Evidence Determining
Individualized Cancer Therapy



PREDICT

- Histology-Independent targeted approach
- Multiple molecular aberrations assessed
- Patients matched with targeted agents

The Reclassification of Cancer

PIK3CA mutations were found in 10% of 1,000 patients with advanced cancers

- Endometrial cancers (29%)
- Breast cancers (24%)
- Colon cancers (17%)
- Ovarian cancers (14%)
- Lung cancer (13%)
- Head and neck squamous cell cancers (13%)
- Pancreatic cancers (13%)

Molecular aberrations do not segregate well by organ of origin

Matching patients with targeted drugs increases response rates

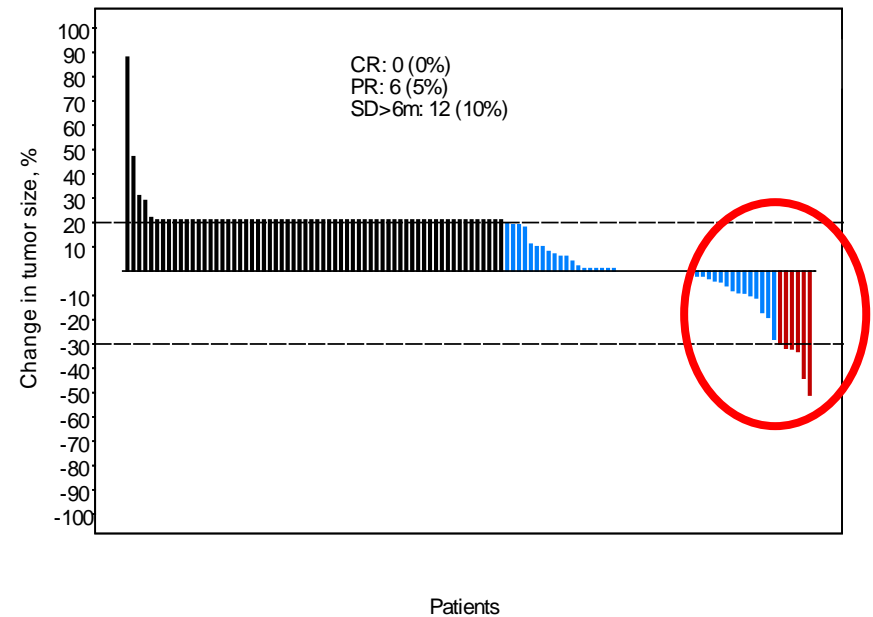
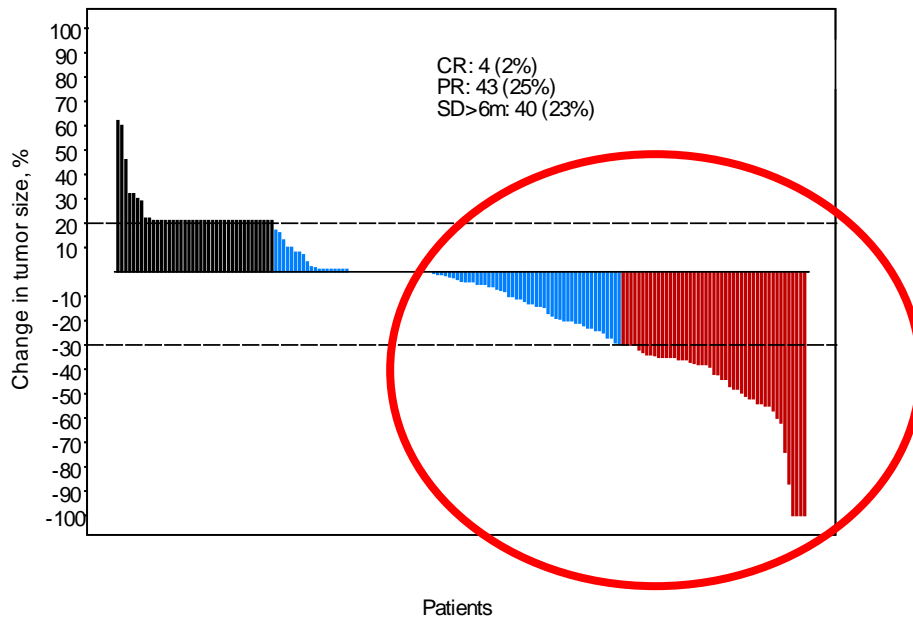
Matched therapy
N=175

Complete/Partial Response = 27%

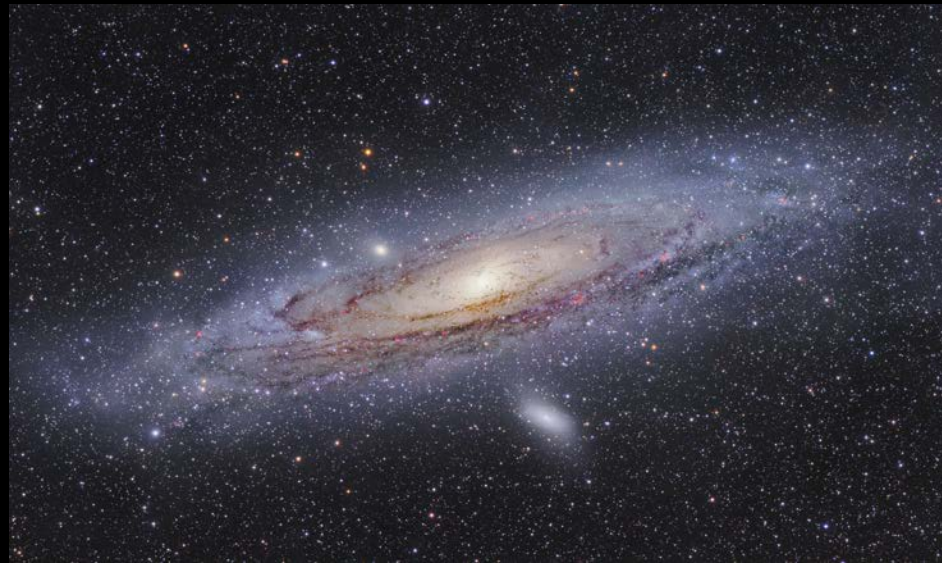
$p < .0001$

Therapy without matching
N=116

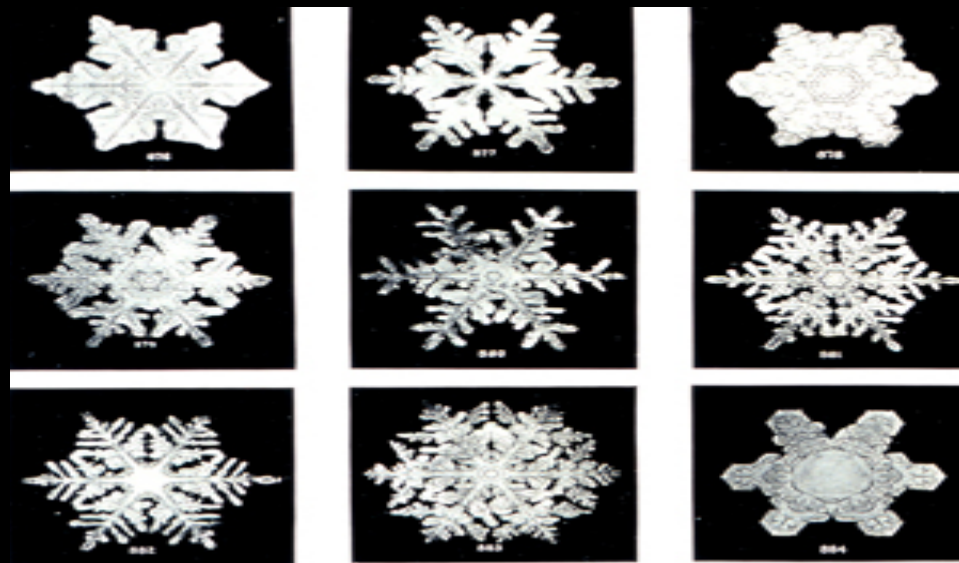
Complete/Partial Response = 5%



Partnering with the UCSD SuperComputer Center



What if every patient with metastatic disease is different?





Malignant Snowflakes

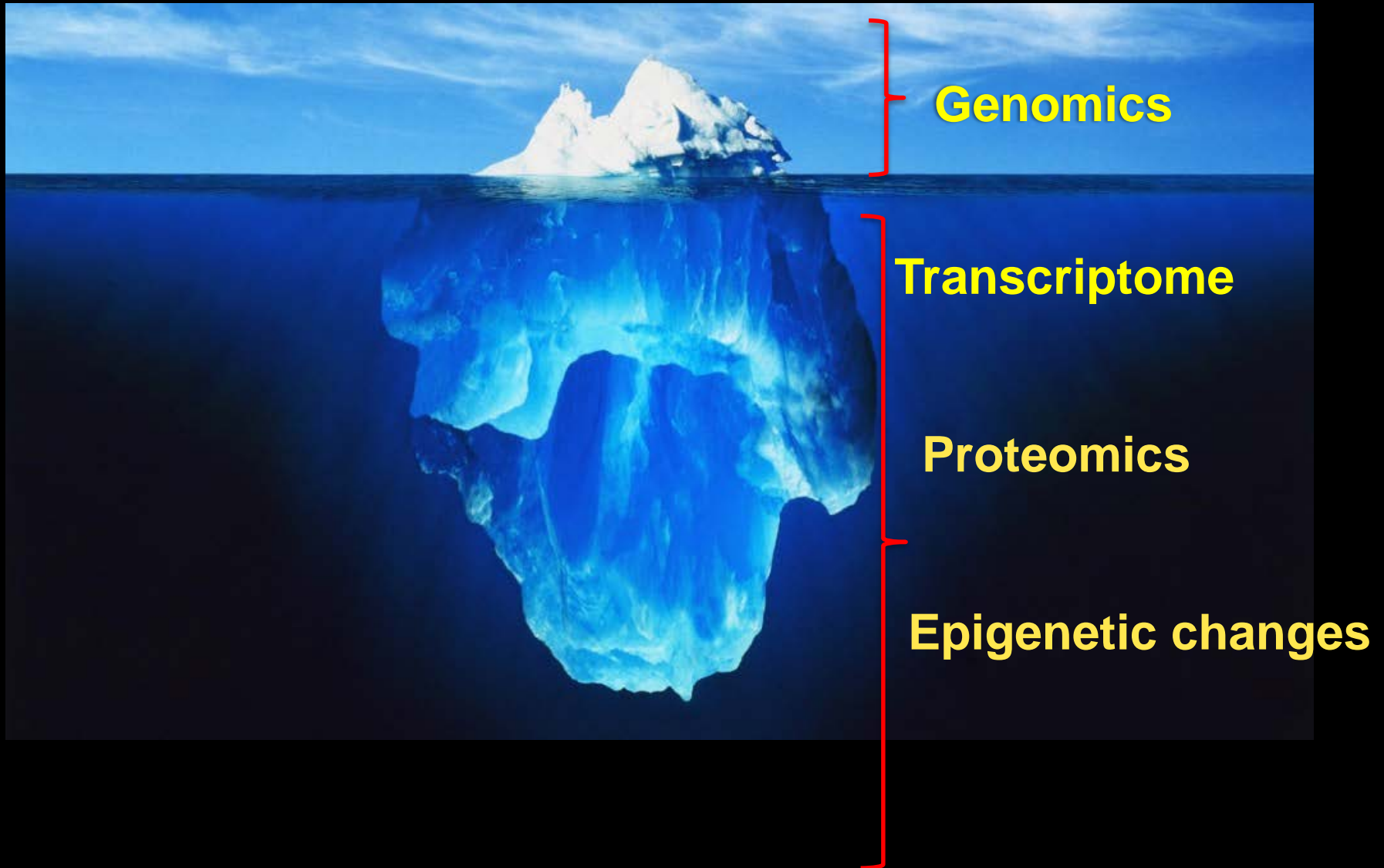


Pt number

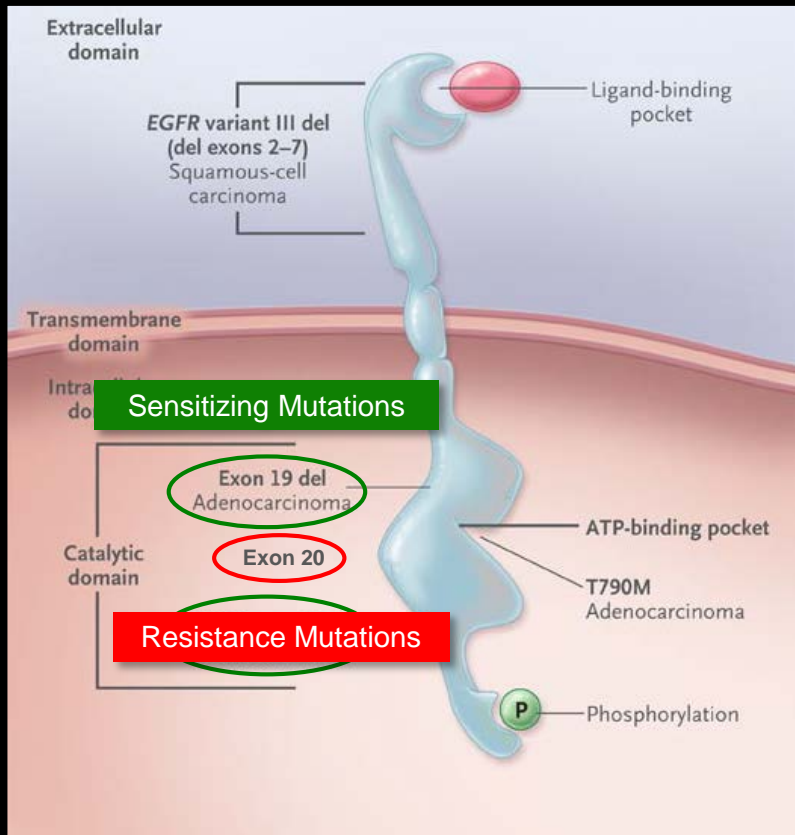
Molecular Results (Foundation Medicine)

1	PIK3CA amplification, SOX2 amplification, TP53 G302fs*42, FLT3 L260*
2	AKT1 (E17K)
4	EGFR amplification, CCND1 amplification, CDKN2A/B loss, FGFR1 amplification, MYC amplification, TP53 P151A
42	ERBB2 amplification, PIK3CA H1047L, AURKA amplification, TP53 R342P, CREBBP P858S, ZNF217 amplification
25	ERBB2 amplification, MYC amplification, CDK6 amplification, TP53 R213*
7	ESR1 Y537S
13	GATA3 *445fs*2+
16	RET C634R, GATA3 P436fs*11+
18	AKT3 amplification, MYC amplification, MYCL1 amplification, TP53 R248Q
54	NF1 R1276Q

Tip of the Iceberg



Epidermal Growth Factor Receptor (EGFR) In Silico Modelling in Lung Cancer



3D In Silico Modeling (SDSC SAN DIEGO SUPERCOMPUTER CENTER)

The 3D model shows the EGFR protein structure with a large 'X' over the extracellular domain, indicating that the binding site is blocked or altered. An arrow labeled **Cetuximab** points to the extracellular domain. Mutations are highlighted in boxes:

- Exon 20:** D770_P772del_insKG, D770>GY
- Exon 19:** ΔE764-A750

Below the model are two axial CT scans of a lung. The left scan shows a tumor (red outline) with a black arrow pointing to it. The right scan shows the same tumor after treatment, with a red outline and a black arrow pointing to it, indicating a reduction in size.

Strategies

```
graph TD; A[Strategies] --> B[Customized Combinations and Immunotherapy for Advanced Disease]; A --> C[Treat Newly-Diagnosed Disease];
```

Customized
Combinations and
Immunotherapy
for Advanced
Disease

**Treat
Newly-Diagnosed
Disease**

Transforming Outcomes

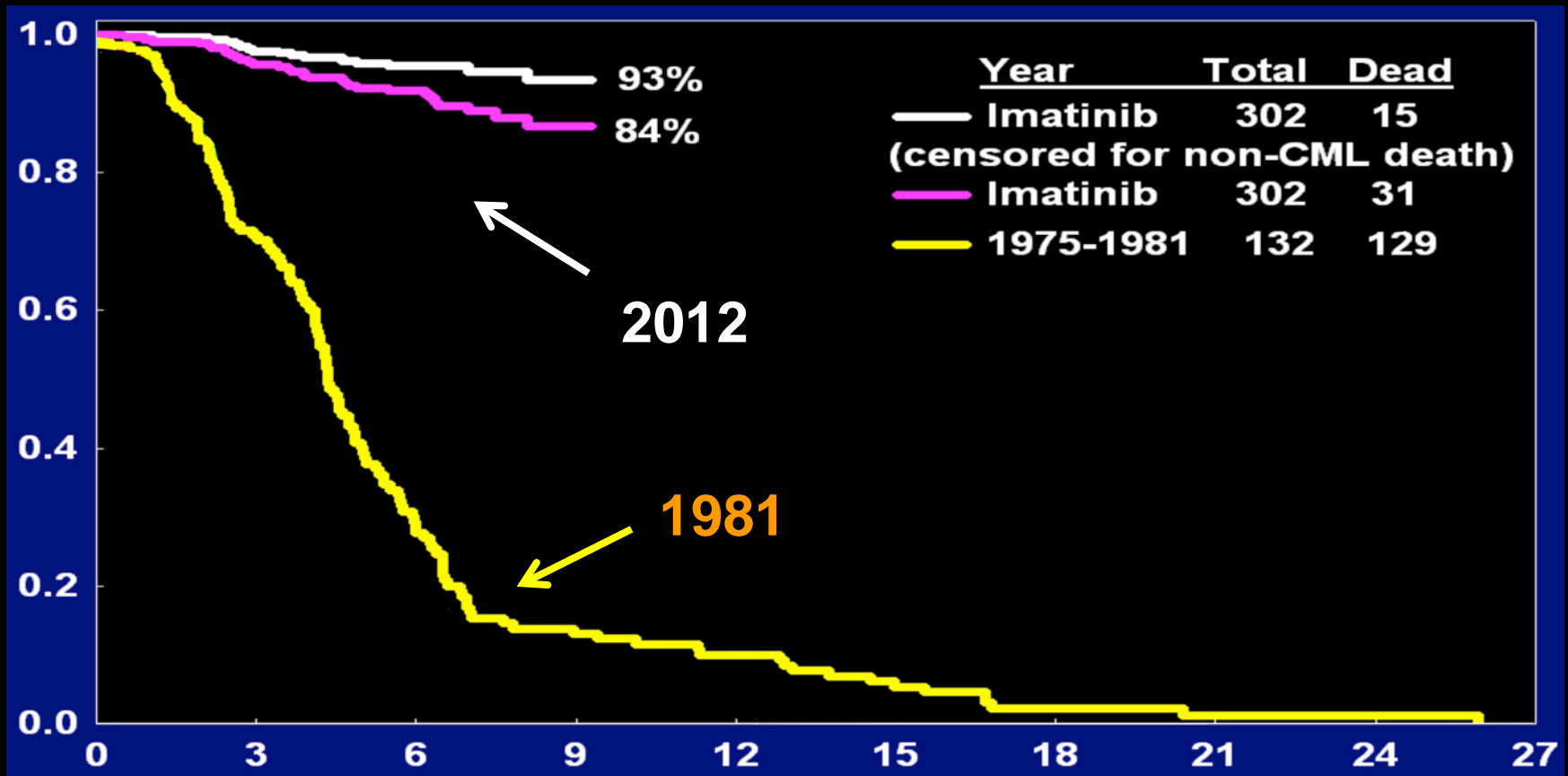
in Solid Tumors?

Is It About Time?

Lessons from the Chronic Myelogenous Leukemia (CML) Story

A Fatal Disease Transformed

- Median survival in 1980s was about 4 years
- Median survival in 2012 is 20+ years



Treatment of Medulloblastoma with Hedgehog Pathway Inhibitor GDC-0449

N ENGL J MED 361:12 NEJM.ORG SEPTEMBER 17, 2009

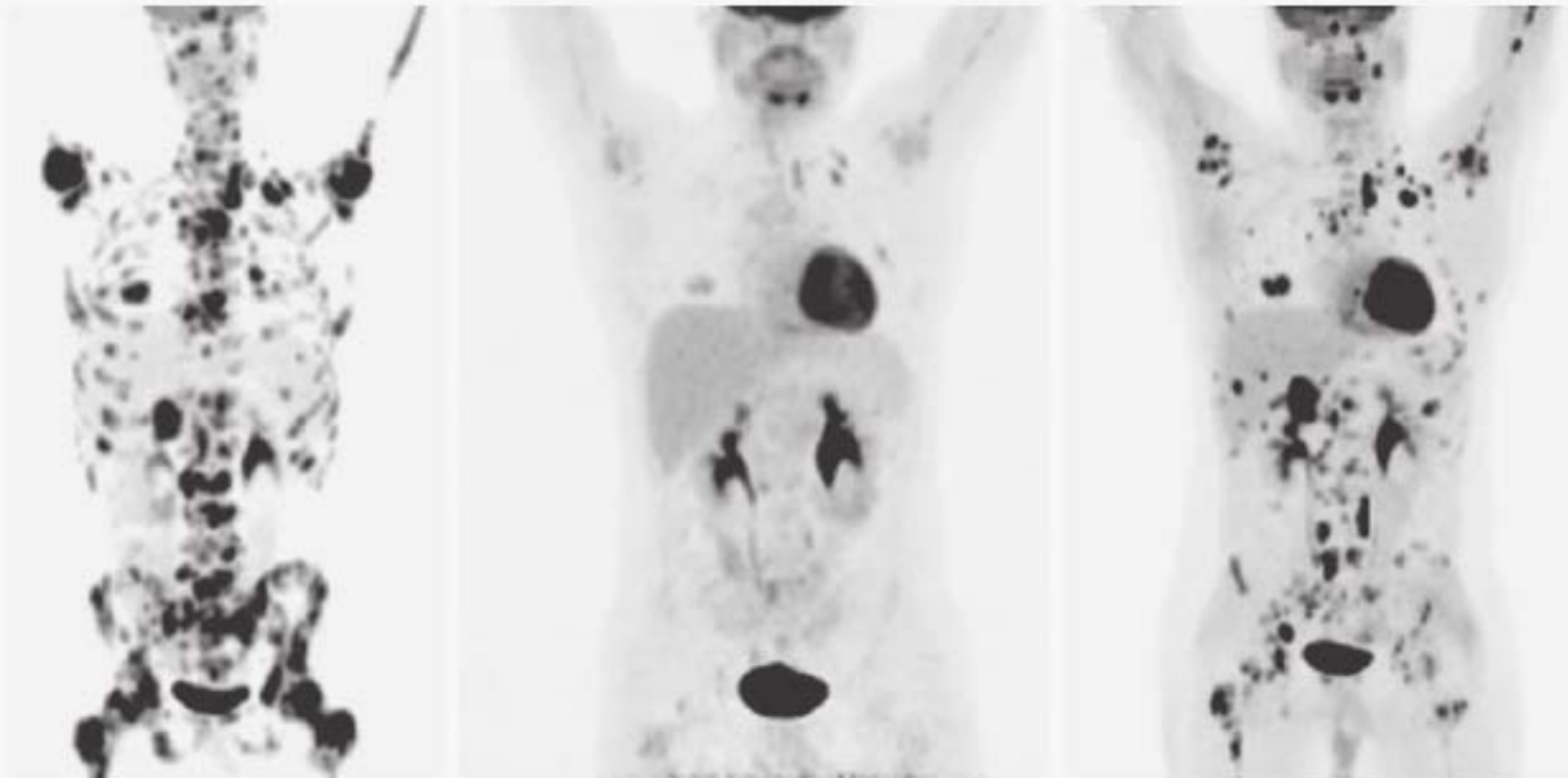
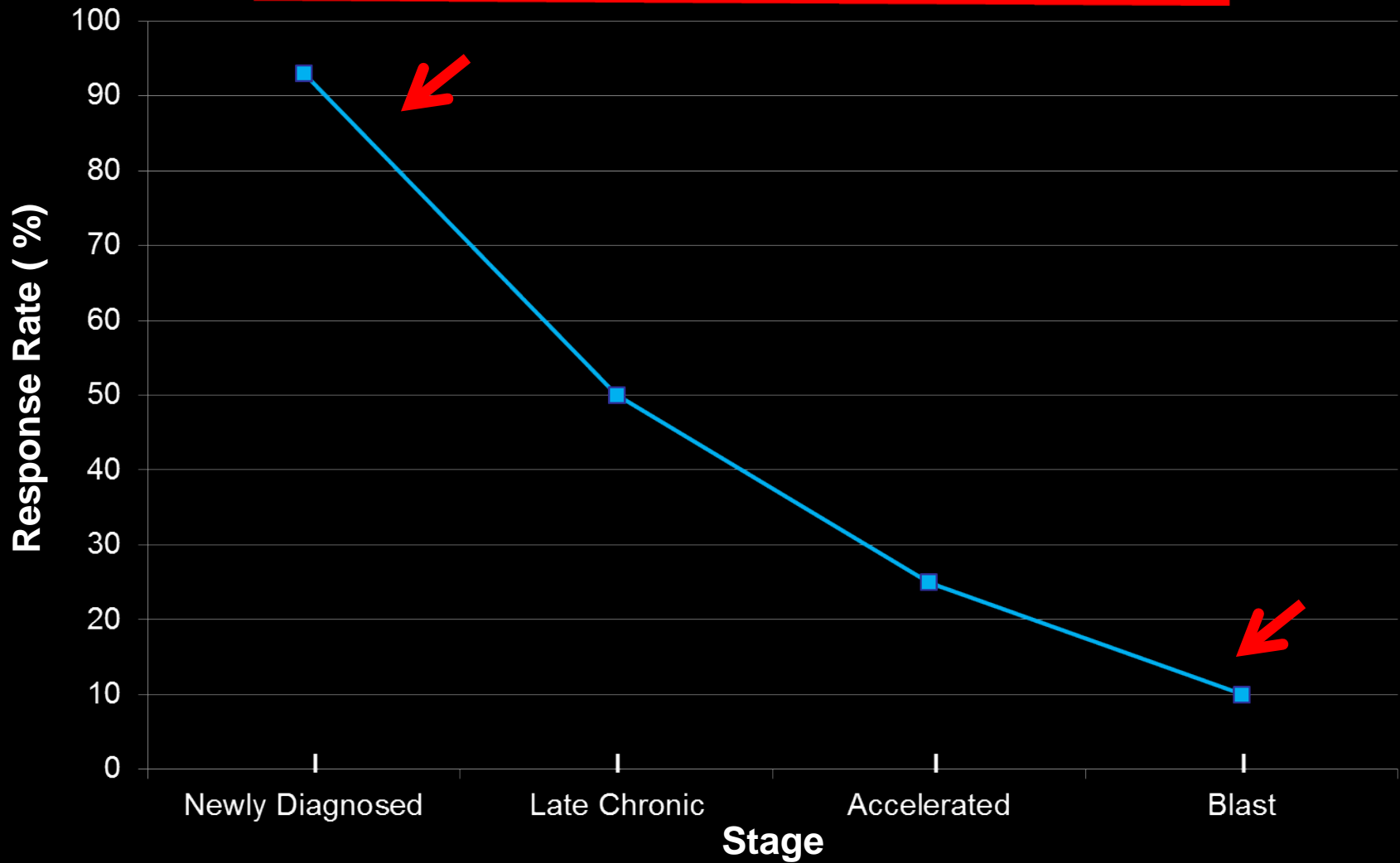


Figure 1. Tumor Response on Positron-Emission Tomographic (PET) Scanning.

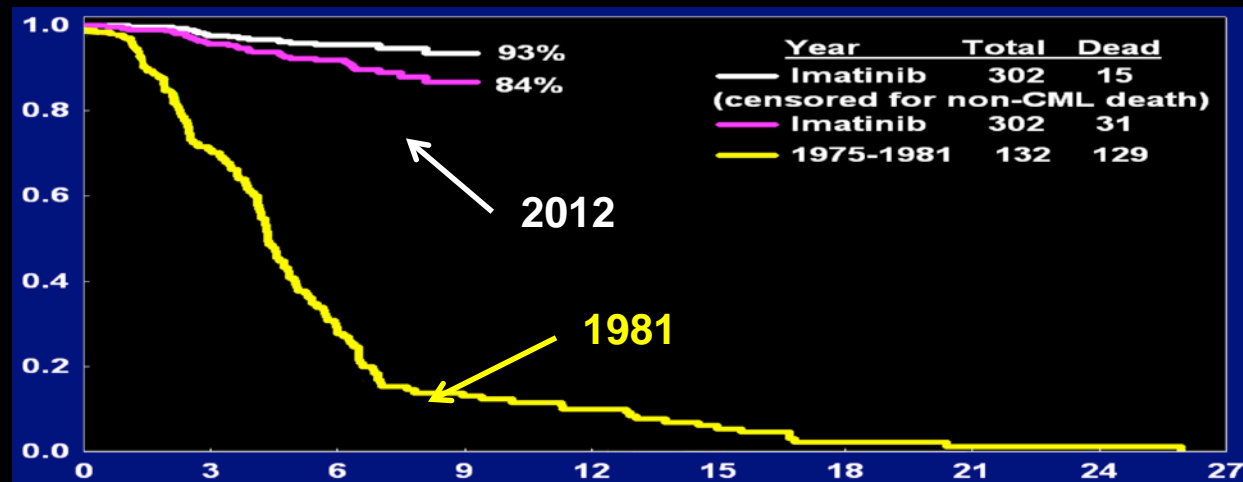
Whole-body projections from ^{18}F -fluorodeoxyglucose (FDG)-PET scans are shown. Panel A shows the pretreatment scan; Panel B, the repeat scan after 2 months of therapy with the hedgehog pathway inhibitor GDC-0449; and Panel C, the repeat scan after 3 months of therapy.

Response Rate of Chronic Myelogenous Leukemia Rises Rapidly in Newly Diagnosed Disease



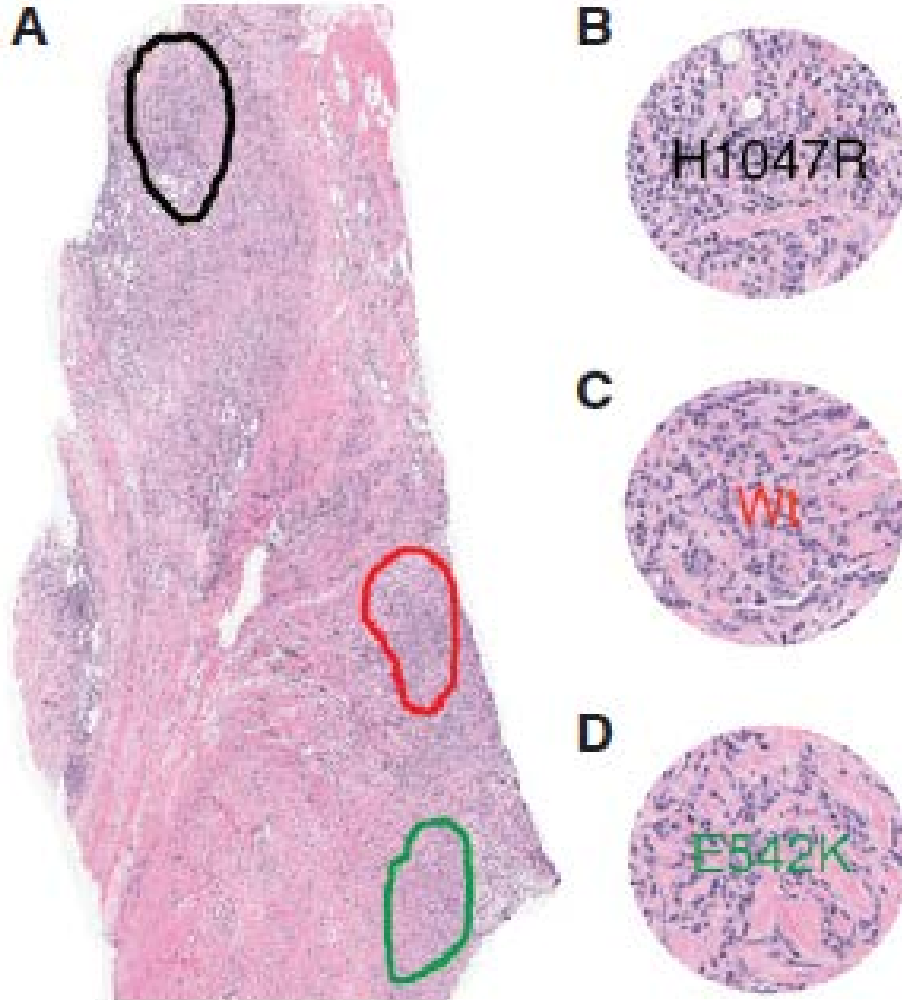
Key factors leading to the revolution in outcome of chronic myelogenous disease

- Key factors:
 - Known driver target (Bcr-Abl)
 - Targeted agent (imatinib)
 - Treat newly-diagnosed patients



Metastases = Blast Crisis in Leukemia

Tumor Microheterogeneity

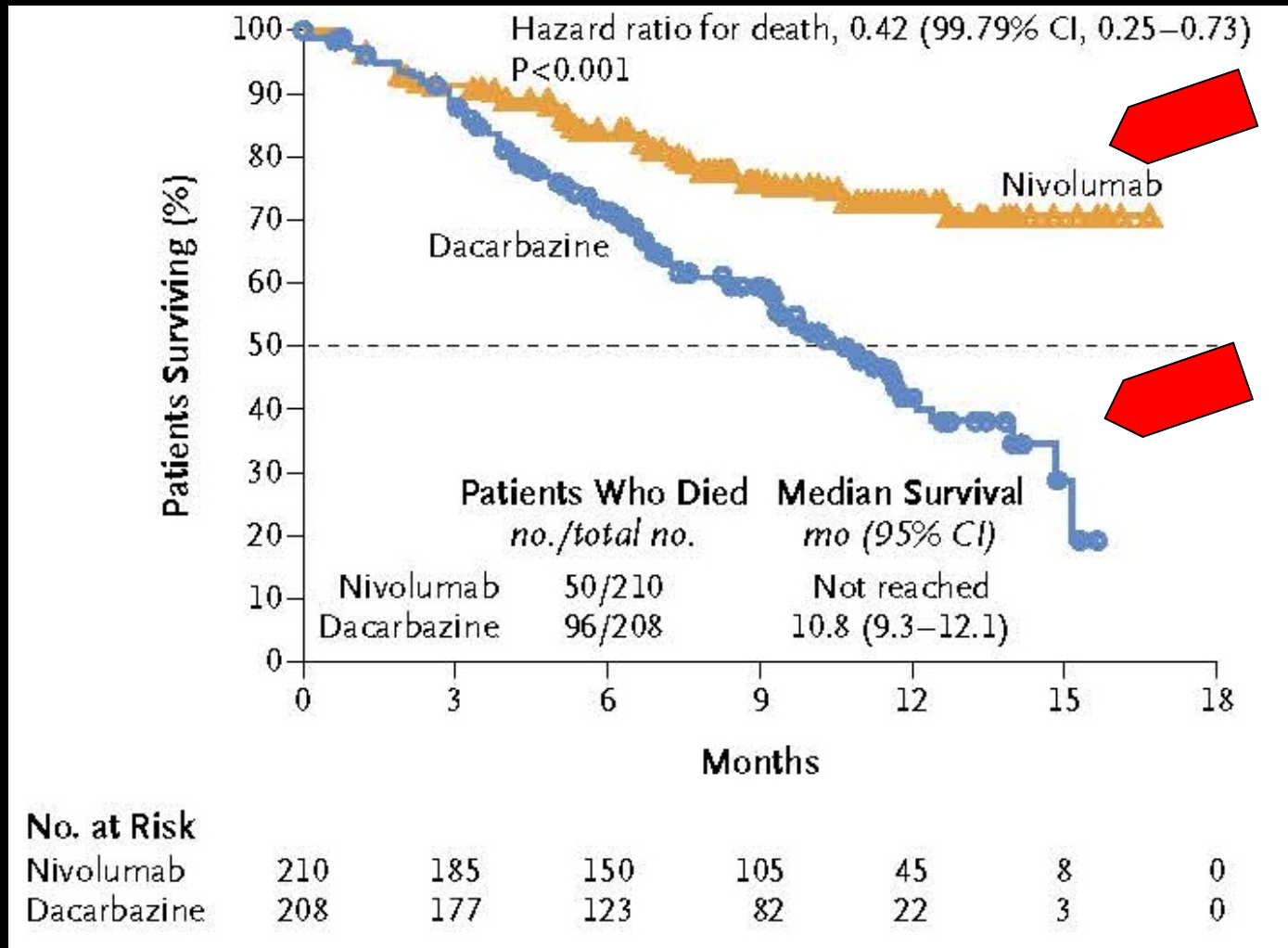


- Molecular profile can differ even within the single lesion
- Discrepancy between molecular profile of primary and metastatic lesion (~20%).

Harnessing the immune system

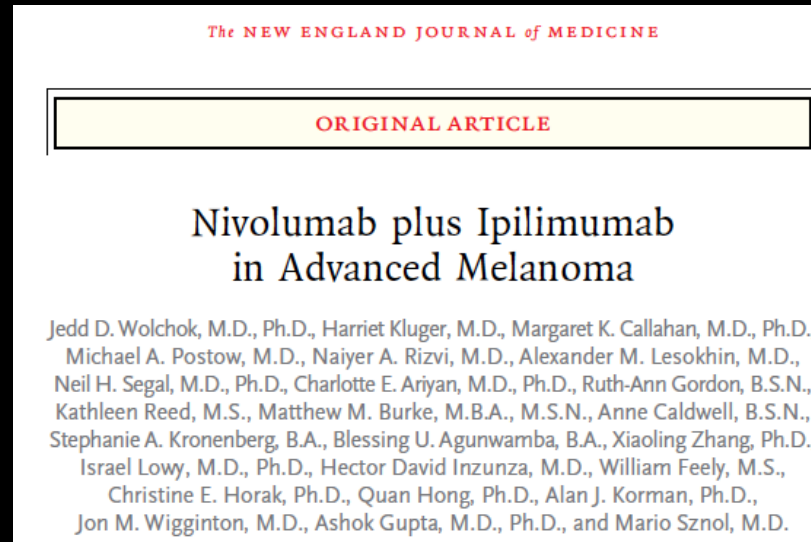
Immunotherapy is revolutionizing cancer care

Metastatic Melanoma: Long-term remissions



Robert et al. Melanoma.....N Engl J Med 2015; 372:320-330

Combinatorial immune blockade is likely the rule, not the exception



- ASCO 2014 update
 - 2 year survival rate- 79%
 - Comparison: dacarbazine monotherapy 2 year survival rate- 18%
 - Prior therapies (1-3+) in 38%

Predicting super-responders to immunotherapy

Biomarker

- PDL-1 negative: 0-17%
- PDL-1 positive: 36-100%

Patel and Kurzrock, MCT 2015

Unique characteristics

- Delayed responses with initial progression
- Subset of patients with advanced disease that have long-term complete remission (?cure)

Liquid Biopsy Program

Doing genomics on DNA from a small tube of blood sample

No tissue biopsy

~700 patients



Liquid Biopsy Program

→ Blood

→ Urine

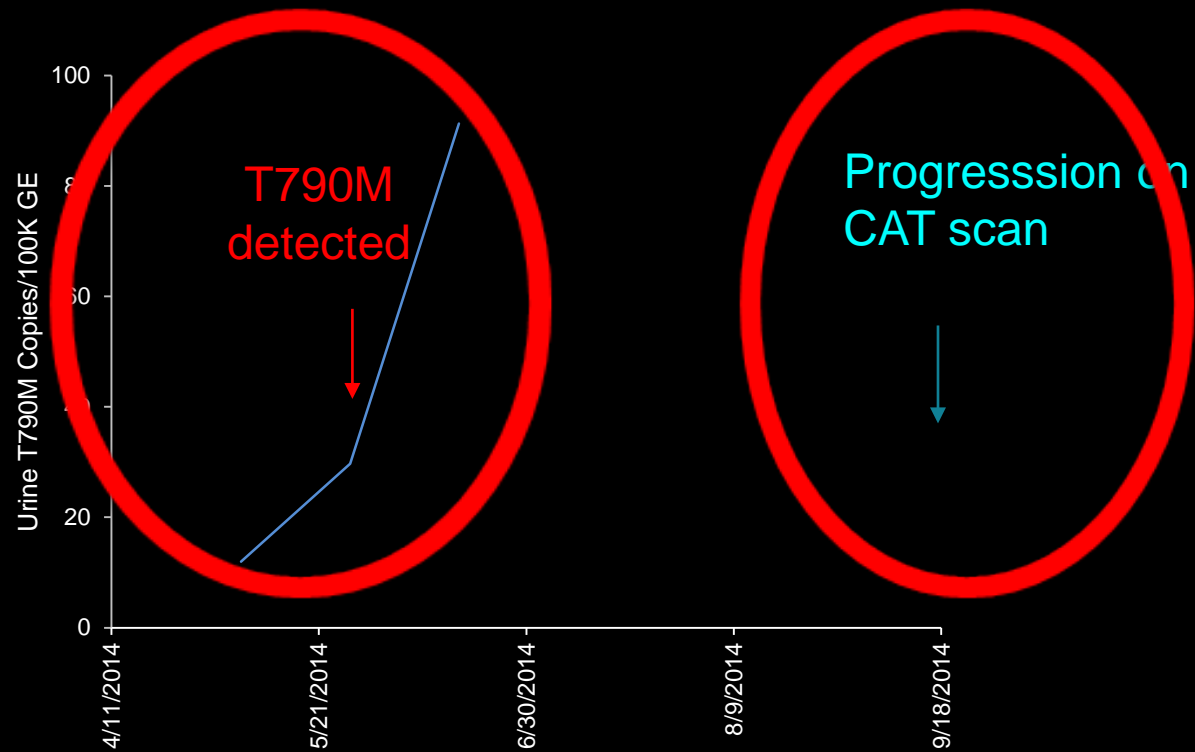
→ Ascites

Theoretically samples shed
DNA from multiple metastatic sites.

Lung Cancer

Early Detection of Progression

Urine



Liquid Biopsy (N = 171 Patients)

Circulatory Tumor DNA
N = 54 genes

	No. of Patients/Total (%)
All	99/171 (58%)
Glioblastoma	9/33 (37%)
Actionable	67/171 (39%)
Healthy Volunteer	1/222 (0.45%) [p53]

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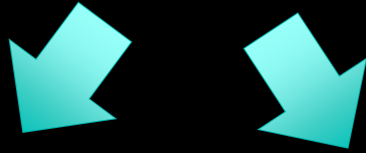
Circulating Tumor DNA
N = 54 genes

	No. of Patients/Total (%)
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Healthy Volunteer	1/222 (0.45%) [p53]

Detecting EGFR Amplifications in Ascites in Lung Cancer

Case No.	Age/Sex	Diagnosis	Tissue NGS ^a	Ascites ctDNA ^b
1	64/man	Adenocarcinoma of the lung, metastasis to pleura and peritoneum	<p>CDK4 amplification^c MDM2 amplification</p> <p>[Also, PCR-based assay (Response Genetics) of primary lung tumor in December 2010 was negative for EGFR, ROS1 and ALK aberrations]</p> <p>(March 2013, Pleural mass)</p>	<p><u>Somatic Mutations:</u> EGFR amplification</p> <p><u>Total CNVs detected:</u> 23</p> <p>(May 2014)</p>

Liquid biopsy applications



Customized combinations for advanced disease.

Need to know all genomic aberrations from multiple metastases

Follow newly-diagnosed disease

Monitor resistance

Precision Medicine
Lessons from meta-analyses
of 70,253 patients

Meta-Analyses Conducted

- 1) Trials leading to FDA approval from trastuzumab (1998) until June 2013
→ 38,104 patients; 112 trials
- 2) Phase II studies published between
→ 32,149 patients; 570 trials

Summary of results

Multivariable analysis: N = 38,104

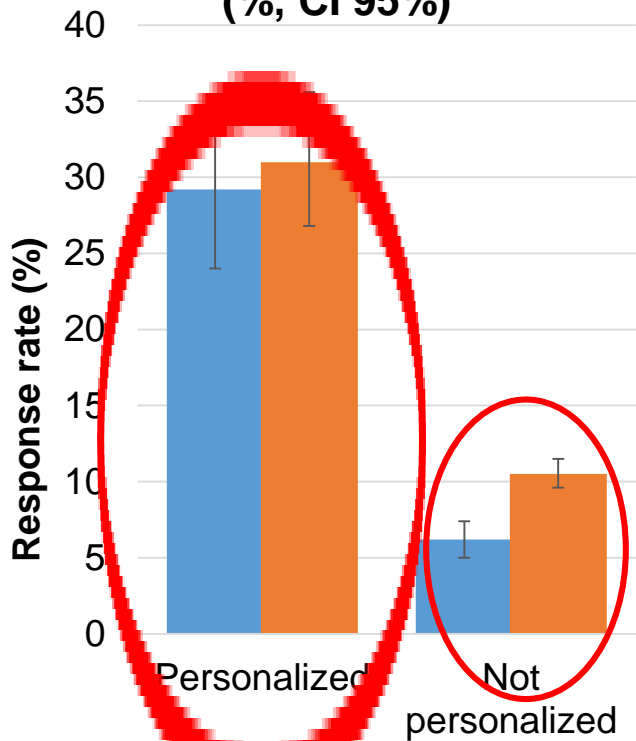
Randomized trials meta regression

Characteristic	P-value
RRR meta-regression	
<i>Personalized therapy strategy</i>	0.03
Progression Free Survival	
<i>Personalized therapy strategy</i>	<0.001
<i>Ctrl arm placebo vs. active drug</i>	<0.001
<i>Hematologic tumor vs solid</i>	0.004
<i>Cross-over allowed</i>	<0.001
Overall Survival	
<i>Personalized therapy strategy</i>	0.07
<i>Hematologic tumor vs solid</i>	0.006

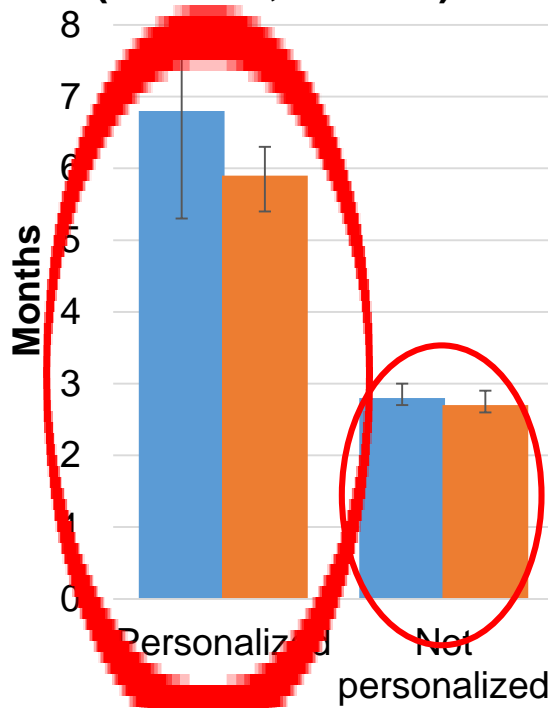
All trials meta-regression (RR) and weighted pooled multilinear regression (PFS/OS)

Characteristic	P-value
Response rate	
<i>Personalized therapy strategy</i>	<0.001
<i>Hematologic tumor vs solid</i>	<0.001
Progression Free Survival	
<i>Personalized therapy strategy</i>	0.002
Overall Survival	
<i>Personalized therapy strategy</i>	0.041

**Response Rate
(%, CI 95%)**



**Median PFS
(Months, CI 95%)**



■ Pooled analysis

■ Meta-analysis

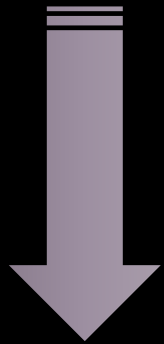
CONCLUSIONS

- **Non-personalized targeted arms led to poorer outcomes than cytotoxics arms**

(All $P < 0.0001$, except $P = 0.048$ for OS meta-analysis).

ARMS type	POOLED Analysis			Meta-analysis		
	RR (%)	PFS (Mos)	OS (Mos)	RR (%)	PFS (Mos)	OS (Mos)
Non-personalized targeted	4	2.6	8.7	7.5	2.5	8.3
Cytotoxic	12	3.3	9.4	16.1	3.3	9.3
Personalized targeted	30	6.9	15.9	31.3	6.1	13.7

Worst outcome



Best outcome

THANK YOU
for your time and interest

Questions??

rkurzrock@ucsd.edu
teoam2011@gmail.com

