

# Cancer Drug Development: Where We Came From & Where We're Headed

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YaleNewHaven**Health**  
Smilow Cancer Hospital

Yale **CANCER**  
CENTER  
A Comprehensive Cancer Center Designated  
by the National Cancer Institute

## MAJOR NCI MILESTONES

1937

President Franklin D. Roosevelt signs legislation to establish NCI.

1960

NCI begins funding government-supported cancer centers.

1973

NCI establishes its **Surveillance, Epidemiology, and End Results (SEER)** program to collect and analyze U.S. cancer incidence, survivorship, and mortality data.

1982

PDQ® (Physician Data



1955

The **Clinical Trials Cooperative Group Program** begins testing anticancer agents identified in the NCI drug development program.

1971

President Nixon signs the **National Cancer Act of 1971**, authorizing the NCI director to coordinate the National Cancer Program and establishing important entities of the program.

1976

NCI's **Cancer Information Service (1-800-4-CANCER)** launches to provide the public with the latest and most accurate cancer information.

**December 23, 1971**



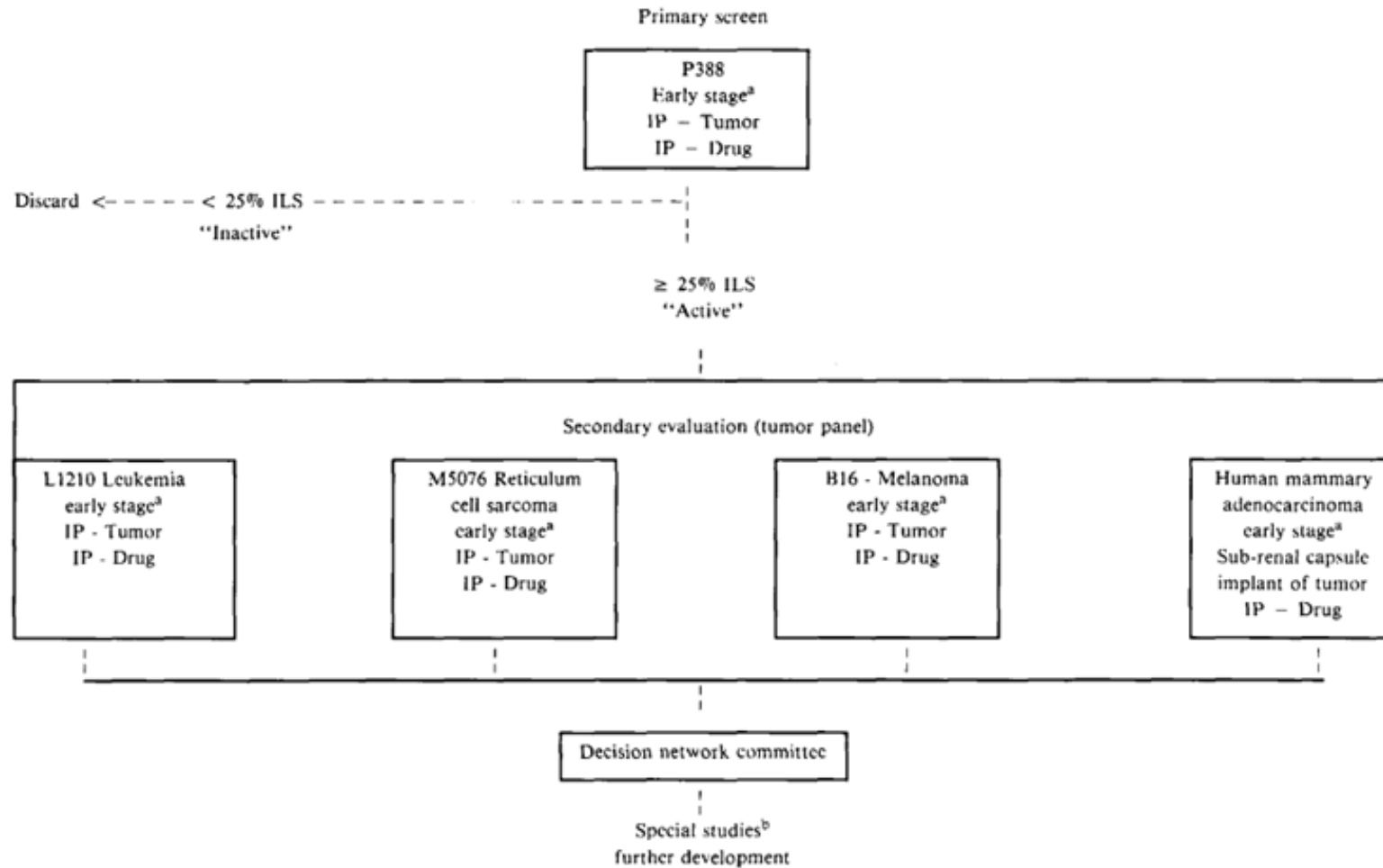
# Conceptual

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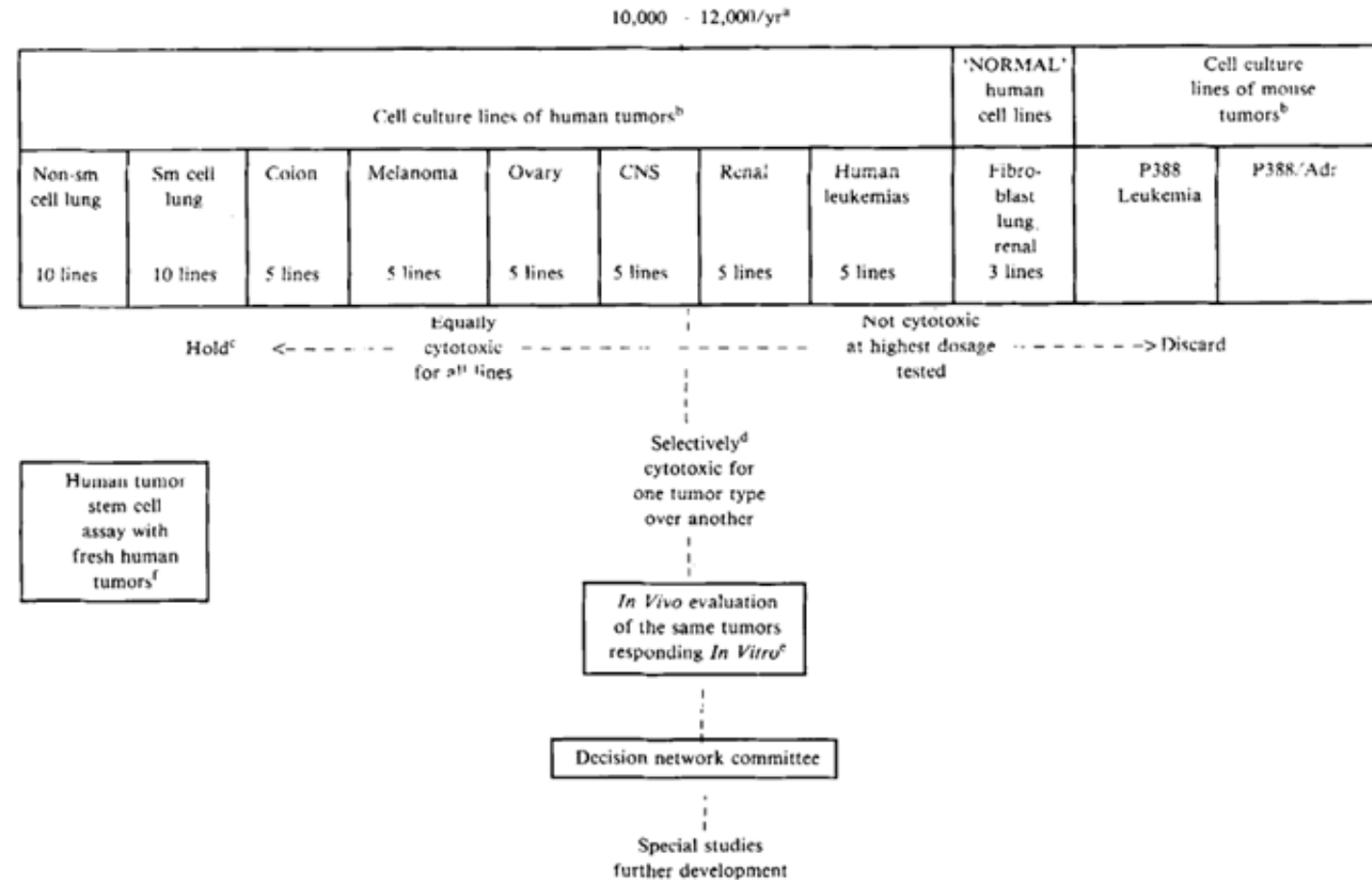
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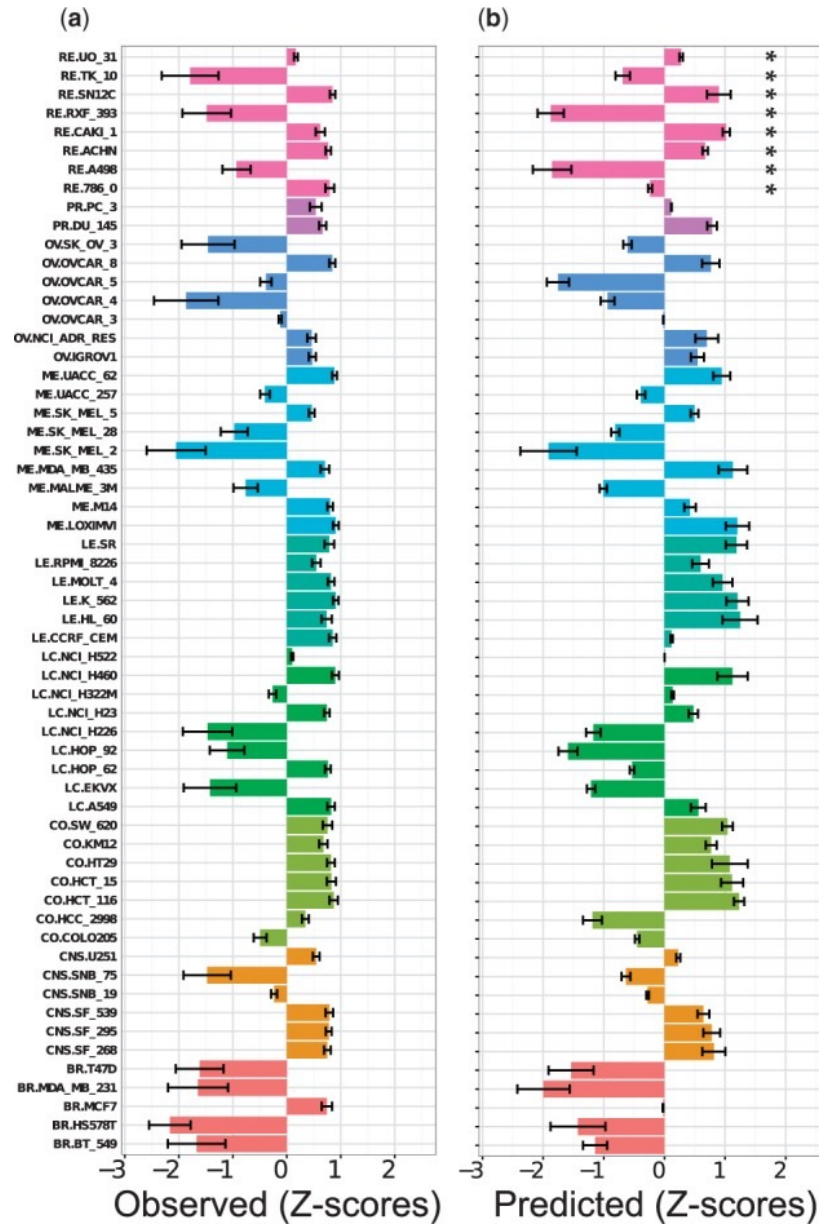
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CENTER  
A Comprehensive Cancer Center Designated  
by the National Cancer Institute

# NCI Drug Screening Schema (1985-1986)



# Proposed NCI "Disease Oriented" Drug Screening Schema (1987)





Cortés-Ciriano I, van Westen GJ, Bouvier G, Nilges M, Overington JP, Bender A, Malliavin TE. Improved large-scale prediction of growth inhibition patterns using the NCI60 cancer cell line panel. *Bioinformatics*. 2016 Jan 1;32(1):85-95. doi: 10.1093/bioinformatics/btv529. Epub 2015 Sep 8. PMID: 26351271; PMCID: PMC4681992.

# Interleukin-2 (IL-2)

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- Discovered in the supernatant of activated T cells in 1976
- Primary role: induce immune responses by stimulating proliferation & differentiation of effector & memory T cells and NK cells
- High-dose IL-2 found to expand **cytotoxic** lymphocytes
- Tumors do not express IL-2 receptors - any antitumor activity result of IL-2 stimulation of immune cells





# IL-2 First in Human Study

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- Twenty patients were reported in 1985 (23 treated) who received a wide variety of different regimens and doses of rIL-2
- Half-life was ~7 min with a later delayed clearance consistent with a two-compartment model as IL-2 was released from extravascular space into the plasma compartment
- Marked depletion of all lymphoid cells was seen almost immediately after IL-2 administration, which rebounded after IL-2 was discontinued
- Significant toxicities became apparent in these early studies, including fever, chills, malaise, arthralgias, and unexpected capillary leak, which led to weight gain from marked fluid retention

**No tumor regression observed**

# IL-2 at higher dose

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- 25 patients with metastatic cancer treated with increasing doses of IL-2 until toxicity precluded further dose escalation
- Early phase of study – pts received 60,000 IU/kg every 8 h
- Subsequent doses: 180,000 or 600,000 IU/kg
- 4/7 patients with mMelanoma & 3/3 patients with mRCC exhibited regression
- 1st demonstration of IL-2 mediated tumor regression in humans
- Subsequent explosion of HD IL-2 trials against various metastatic cancers
- Eventual FDA approval: mRCC and melanoma in 1992 and 1998

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# Results of Treatment of 255 Patients With Metastatic Renal Cell Carcinoma Who Received High-Dose Recombinant Interleukin-2 Therapy

By Gwendolyn Fyfe, Richard I. Fisher, Steven A. Rosenberg, Mario Sznol, David R. Parkinson, and Arthur C. Louie

***Purpose:*** To determine the efficacy and toxicity of a high-dose interleukin-2 (IL-2) regimen in patients with metastatic renal cell carcinoma.

***Patients and Methods:*** Two hundred fifty-five assessable patients were entered onto seven phase II clinical trials. Proleukin (aldesleukin; Chiron Corp, Emeryville, CA) 600,000 or 720,000 IU/kg was administered by 15-minute intravenous (IV) infusion every 8 hours for up to 14 consecutive doses over 5 days as clinically tolerated with maximum support, including pressors. A second identical cycle of treatment was scheduled following 5 to 9 days of rest, and courses could be repeated every 6 to 12 weeks in stable or responding patients.

***Results:*** The overall objective response rate was 14% (90% confidence interval [CI], 10% to 19%), with 12 (5%) complete responses (CRs) and 24 (9%) partial responses (PRs). Responses occurred in all sites of disease, including bone, intact primary tumors, and visceral metastases, and in patients with large tumor burdens or bulky indi-

vidual lesions. The median response duration for patients who achieved a CR has not been reached, but was 19.0 months for those who achieved a PR. Baseline Eastern Cooperative Oncology Group (ECOG) performance status (PS) was the only predictive prognostic factor for response to IL-2. While treatment was associated with severe acute toxicities, these generally reversed rapidly after therapy was completed. However, 4% of patients died of adverse events judged to be possibly or probably treatment-related.

***Conclusion:*** High-dose IL-2 appears to benefit some patients with metastatic renal cell carcinoma by producing durable CRs or PRs. Despite severe acute treatment-associated toxicities, IL-2 should be considered for initial therapy of patients with appropriately selected metastatic renal cell carcinoma.

*J Clin Oncol* 13:688-696. © 1995 by American Society of Clinical Oncology.

# High Dose IL-2 In Metastatic Renal Cell Carcinoma: Summary of Efficacy

Response	Response Rate		Response Duration (months)	
	No.	%	Median	Range†
CR	12	5	NR	5+-62+
PR*	24	9	19	3-57+
PR + CR	36	14	20.3	3-62+

Abbreviation: NR, not yet reached.

\*Three partial responders had surgery while in PR and remain disease-free. Duration of response censored from date of surgery.

†Plus signs mean ongoing.

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# IL-2 Toxicity

Event by Body System	Grade (%)		
	All	3	4
<b>Cardiovascular</b>			
Hypotension	96	59	15
Arrhythmias	14	2	0
Supraventricular	5	2	1
Ventricular	1	0	0
Myocardial ischemia	2	1	<1
Myocardial infarction	2	0	2
Cardiac arrest	2	<1	2
Myocarditis	1	1	0
<b>Gastrointestinal</b>			
Nausea and vomiting	89	24	1
Diarrhea	81	20	2
Stomatitis	32	4	0
Gastrointestinal bleeding	15	3	1
Intestinal perforation	1	0	<1
<b>Neurologic</b>			
Mental status changes	82	23	5
Coma	2	0	2
Seizure (grand mal)	2	1	1
<b>Pulmonary</b>			
Dyspnea	57	16	1
Adult respiratory distress syndrome	1	<1	<1
Respiratory failure	3	<1	2
<b>Hepatic</b>			
Elevated bilirubin level	85	13	8
Elevated transaminase level	72	7	3
Elevated alkaline phosphatase level	77	8	<1
<b>Renal</b>			
Acidosis	19	4	2
Elevated BUN level	85	12	2
Oliguria/anuria	81	40	6
Serum creatinine elevation	81	11	3
<b>General</b>			
Fever and/or chills	97	19	5
Asthenia	39	4	0
Edema	55	2	0
Sepsis	8	4	2
<b>Hematologic</b>			
Thrombocytopenia	83	16	5
Anemia	99	15	3
<b>Other</b>			
Pruritus	53	4	0
Rash	25	1	0
Arthralgia	7	1	0
Myalgia	7	1	0

Fyfe et al. *Journal of Clinical Oncology*,  
Vol 13, No 3 (March), 1995: pp 688-696

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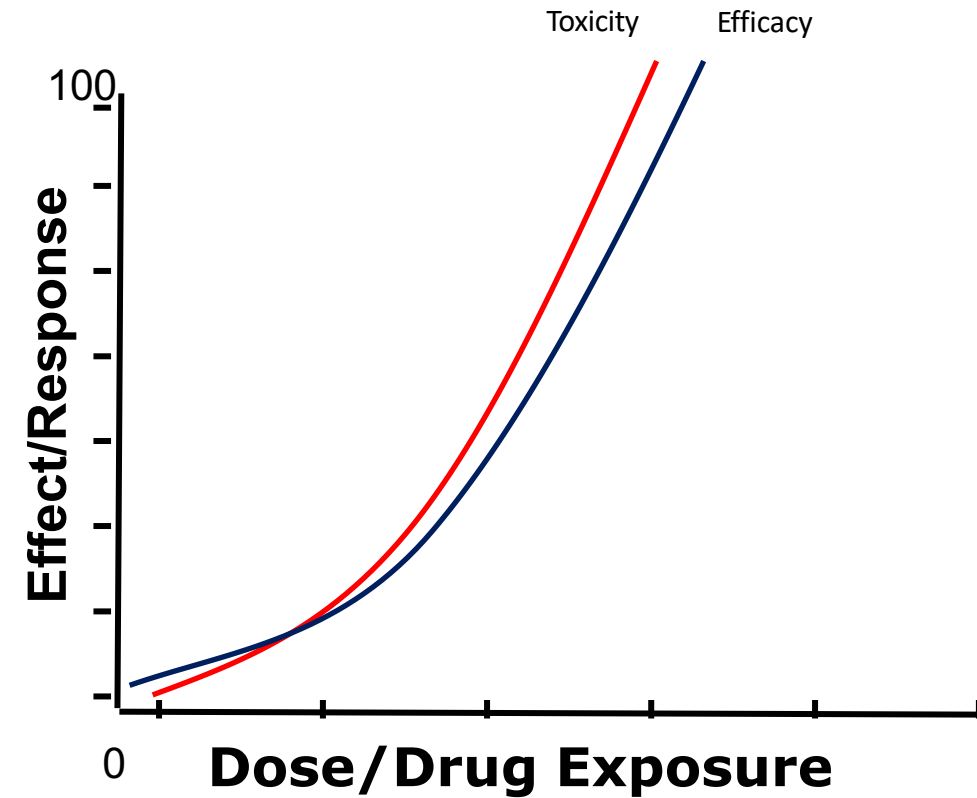
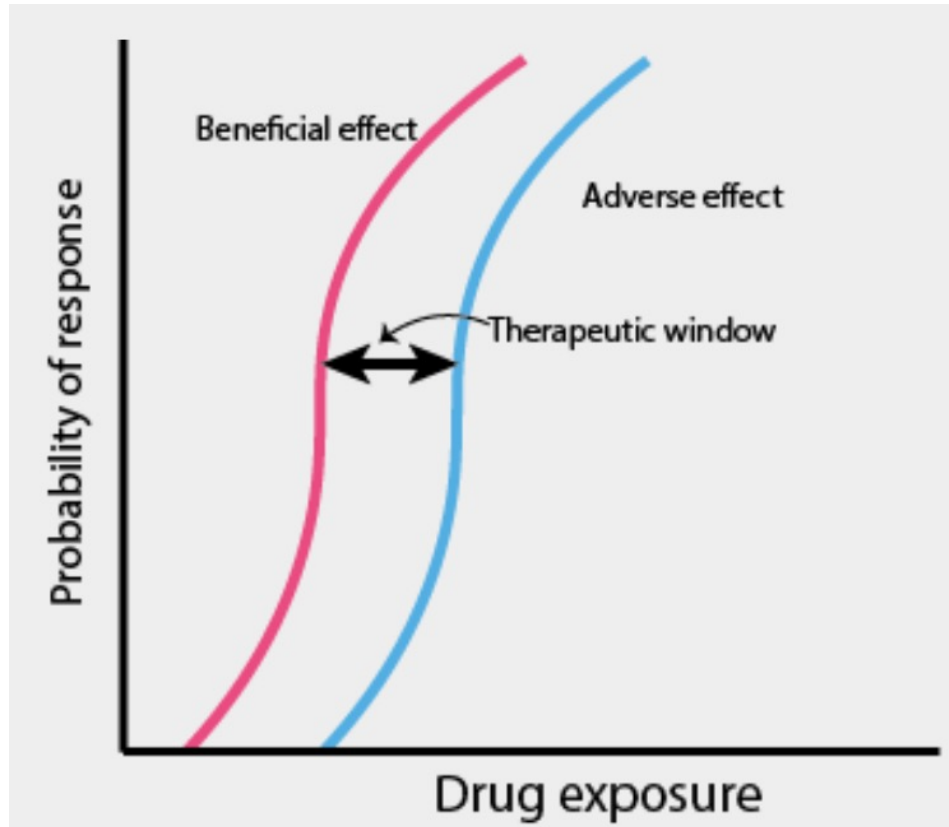


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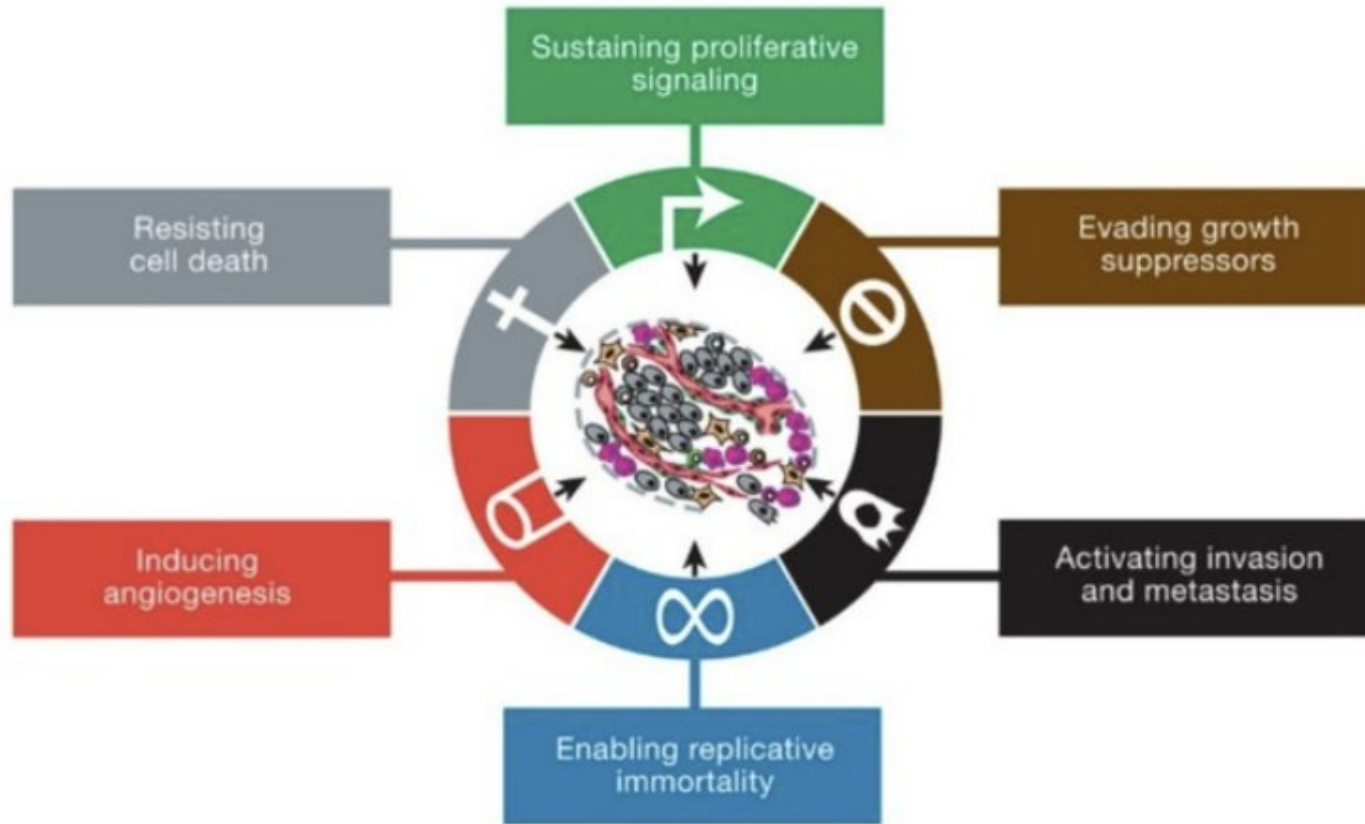
# Narrow Therapeutic Window



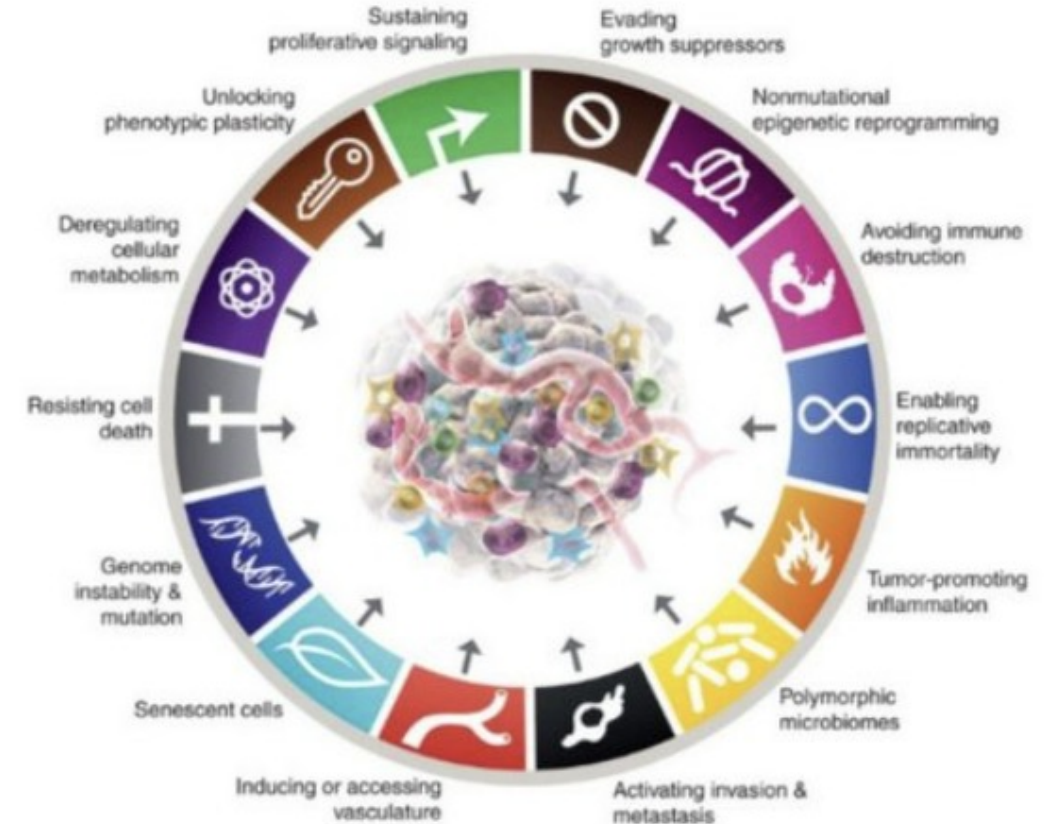
<https://www.fda.gov/advisory-committees/advisory-committee-calendar/updated-information-april-21-22-2022-meeting-oncologic-drugs-advisory-committee-meeting-announcement>

# Biology

# Hallmarks of Cancer



2000



2022

Hannahan & Weinberg, 2000, 2022



15 February 2001

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## the human genome

### Nuclear fission

Five-dimensional  
energy landscapes

### Seafloor spreading

The view from under  
the Arctic ice

### Career prospects

Sequence creates new  
opportunities



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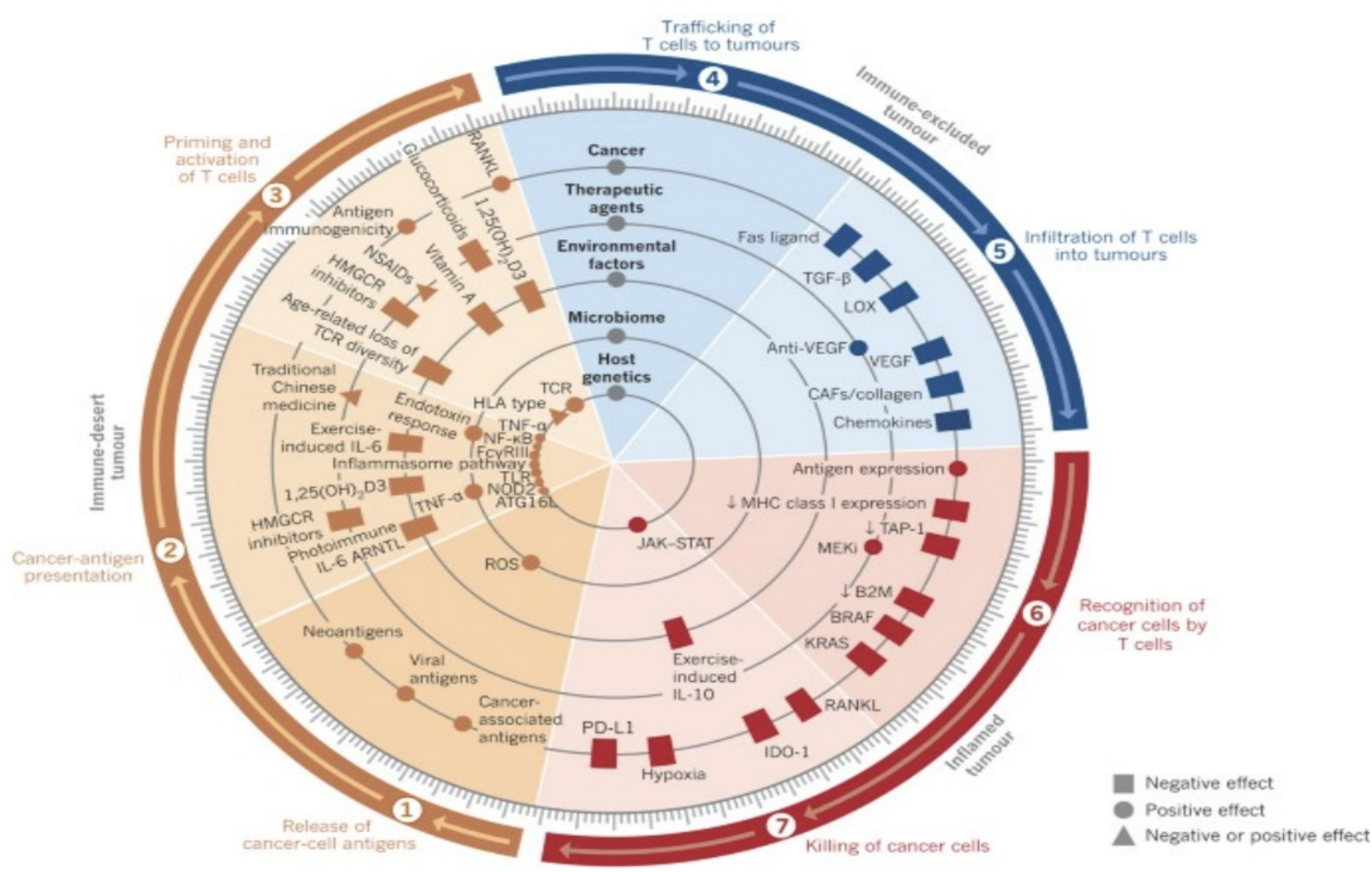


# The Human Genome Project

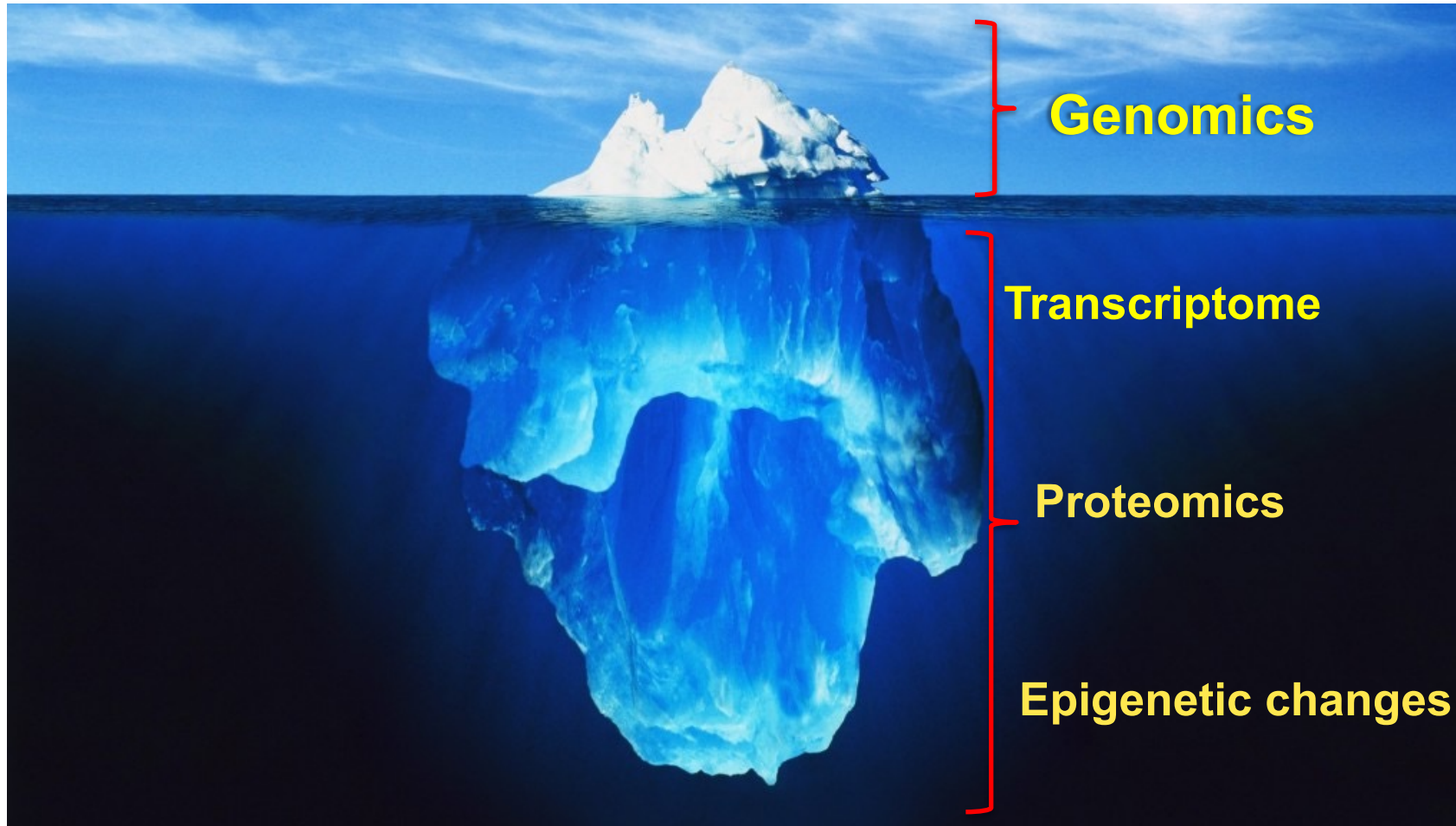
# Cancer-immune phenotypes.



# Factors Influencing the Cancer Immune Set Point



# Major Scientific Tools

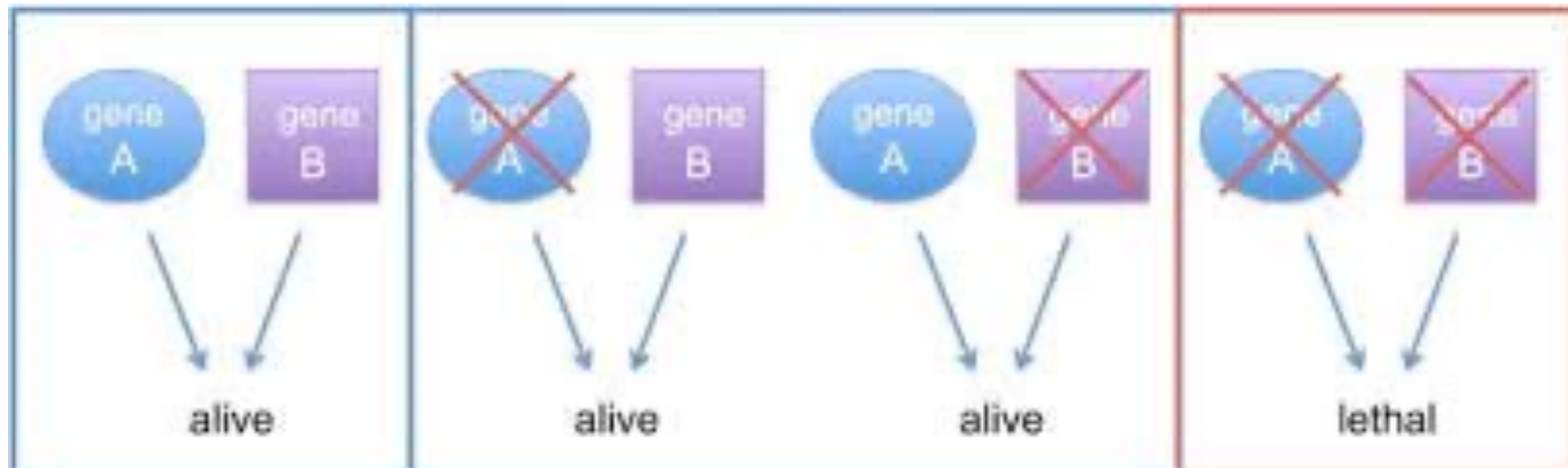




# Integrating Genetic Approaches into the Discovery of Anticancer Drugs

Leland H. Hartwell, Philippe Szankasi, Christopher J. Roberts, Andrew W. Murray, Stephen H. Friend\*

Science (1997), 278: 1064 - 1068



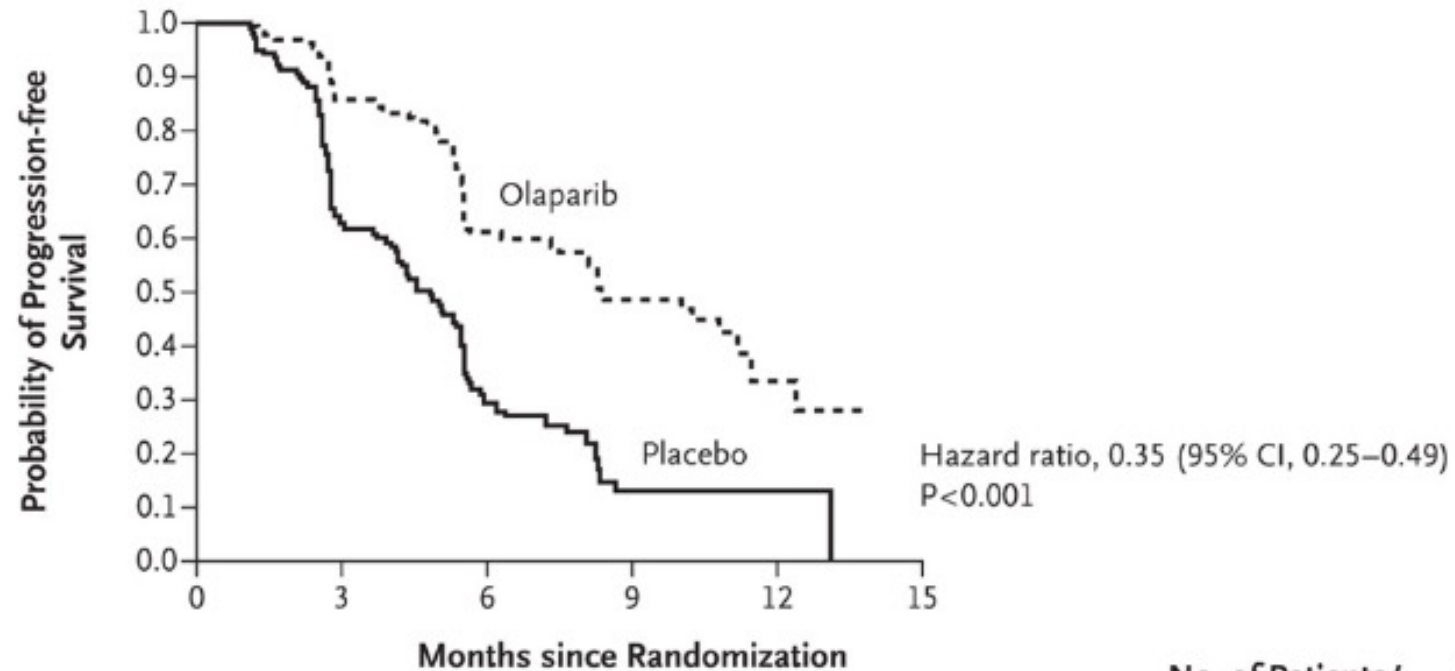
Tuesday, 20 December 2011

# **Company** *updates on olaparib development program*

**Company** today announced that its investigational compound olaparib will not progress into Phase III development for the maintenance treatment of serous ovarian cancer.

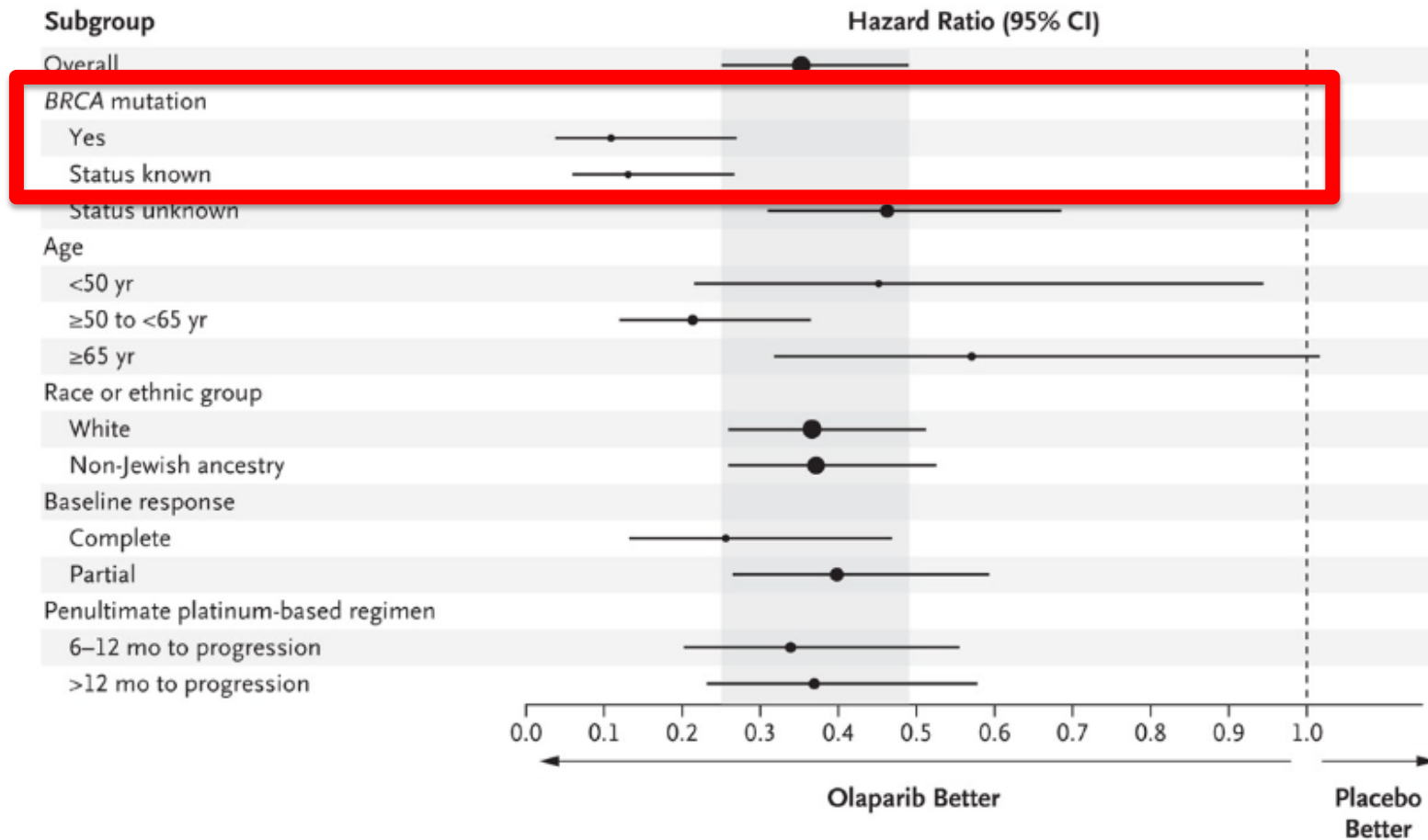
The decision to discontinue olaparib's development in serous ovarian cancer was made following a review of an interim analysis of a Phase II study (study 19) which indicated that the previously reported progression free survival benefit is unlikely to translate into an overall survival benefit, the definitive measure of patient benefit in ovarian cancer. In addition, attempts to identify a suitable tablet dose for use in Phase III studies have not been successful.

# Olaparib Maintenance in Platinum-sensitive Relapsed Ovarian Cancer: PFS



	No. at Risk						No. of Patients/ Total No. (%)	Median Progression-free Survival (mo)
	0	3	6	9	12	15		
Olaparib	136	104	51	23	6	0	60/136 (44.1)	8.4
Placebo	129	72	23	7	1	0	93/129 (72.1)	4.8

# Olaparib Maintenance in Platinum-sensitive Relapsed Ovarian Cancer: Subgroup Analysis of PFS





Contents lists available at ScienceDirect

## Gynecologic Oncology

journal homepage: [www.elsevier.com/locate/ygyno](http://www.elsevier.com/locate/ygyno)



### Efficacy and safety of olaparib monotherapy in germline *BRCA1/2* mutation carriers with advanced ovarian cancer and three or more lines of prior therapy



Susan M. Domchek<sup>a,\*</sup>, Carol Aghajanian<sup>b</sup>, Ronnie Shapira-Frommer<sup>c</sup>, Rita K. Schmutzler<sup>d</sup>, M. William Audeh<sup>e</sup>, Michael Friedlander<sup>f</sup>, Judith Balmaña<sup>g</sup>, Gillian Mitchell<sup>h,i</sup>, Georgeta Fried<sup>j</sup>, Salomon M. Stemmer<sup>k</sup>, Ayala Hubert<sup>l,m</sup>, Ora Rosengarten<sup>n</sup>, Niklas Loman<sup>o</sup>, Jane D. Robertson<sup>p,1</sup>, Helen Mann<sup>p</sup>, Bella Kaufman<sup>c</sup>

<sup>a</sup> *Basser Research Center and Abramson Cancer Center, Philadelphia, PA, USA*

<sup>b</sup> *Memorial Sloan Kettering Cancer Center and Weill Cornell Medical College, New York, NY, USA*

<sup>c</sup> *Sheba Medical Center, Tel Hashomer, Israel*

<sup>d</sup> *Center for Familial Breast and Ovarian Cancer and Center of Integrated Oncology, Cologne, Germany*

<sup>e</sup> *Samuel Oschin Cancer Institute, Los Angeles, CA, USA*

<sup>f</sup> *Prince of Wales Clinical School, University of New South Wales, Sydney, Australia*

<sup>g</sup> *Vall d'Hebron University Hospital and Vall d'Hebron Institute of Oncology, Barcelona, Spain*

<sup>h</sup> *Peter MacCallum Cancer Centre, Melbourne, Australia*

<sup>i</sup> *Sir Peter MacCallum Department of Oncology, University of Melbourne, Melbourne, Australia*

<sup>j</sup> *Institute of Oncology, Rambam Health Care Campus, Haifa, Israel*

<sup>k</sup> *Rabin Medical Center, Petah Tikva, Israel*

<sup>l</sup> *Hadassah-Hebrew University Hospital Sharett Institute of Oncology, Jerusalem, Israel*

<sup>m</sup> *Sharett Institute of Oncology, Jerusalem, Israel*

<sup>n</sup> *Shaare Zedek Medical Centre, Jerusalem, Israel*

<sup>o</sup> *Skånes universitetssjuk Lund, Lund, Sweden*

<sup>p</sup> *AstraZeneca, Macclesfield, UK*

Objective response rate and duration of response stratified by platinum sensitivity status in patients with gBRCA1/2m ovarian cancer and measurable disease at baseline who had received  $\geq 3$  prior chemotherapy regimens (n = 137).

Platinum sensitivity status <sup>a</sup> (N = total patients with measurable disease)	Confirmed responders <sup>b</sup> n	ORR, % (95% CI)	Median DoR, months (95% CI)
Total (N = 137)	46	34 (26–42)	7.9 (5.6–9.6)
Platinum sensitive (N = 39) <sup>c</sup>	18	46 (30–63)	8.2 (5.6–13.5)
Platinum resistant (N = 81)	24	30 (20–41)	8.0 (4.8–14.8)
Platinum refractory (N = 14)	2	14 (2–43)	6.4 (5.4–7.4)
Platinum status unknown (N = 3)	2	67 (9–99)	6.3 (4.7–7.9)

Domchek, et al  
GYN Onc 2016

# SOLO1: Phase III trial of maintenance olaparib following platinum-based chemotherapy in newly diagnosed patients with advanced ovarian cancer and a *BRCA1/2* mutation

Kathleen Moore,<sup>1</sup> Nicoletta Colombo,<sup>2</sup> Giovanni Scambia,<sup>3</sup> Byoung-Gie Kim,<sup>4</sup> Ana Oaknin,<sup>5</sup> Michael Friedlander,<sup>6</sup> Alla Lisyanskaya,<sup>7</sup> Anne Floquet,<sup>8</sup> Alexandra Leary,<sup>9</sup> Gabe S. Sonke,<sup>10</sup> Charlie Gourley,<sup>11</sup> Susana Banerjee,<sup>12</sup> Amit Oza,<sup>13</sup> Antonio González-Martín,<sup>14</sup> Carol Aghajanian,<sup>15</sup> William Bradley,<sup>16</sup> Elizabeth S. Lowe,<sup>17</sup> Ralph Bloomfield,<sup>18</sup> Paul DiSilvestro<sup>19</sup>

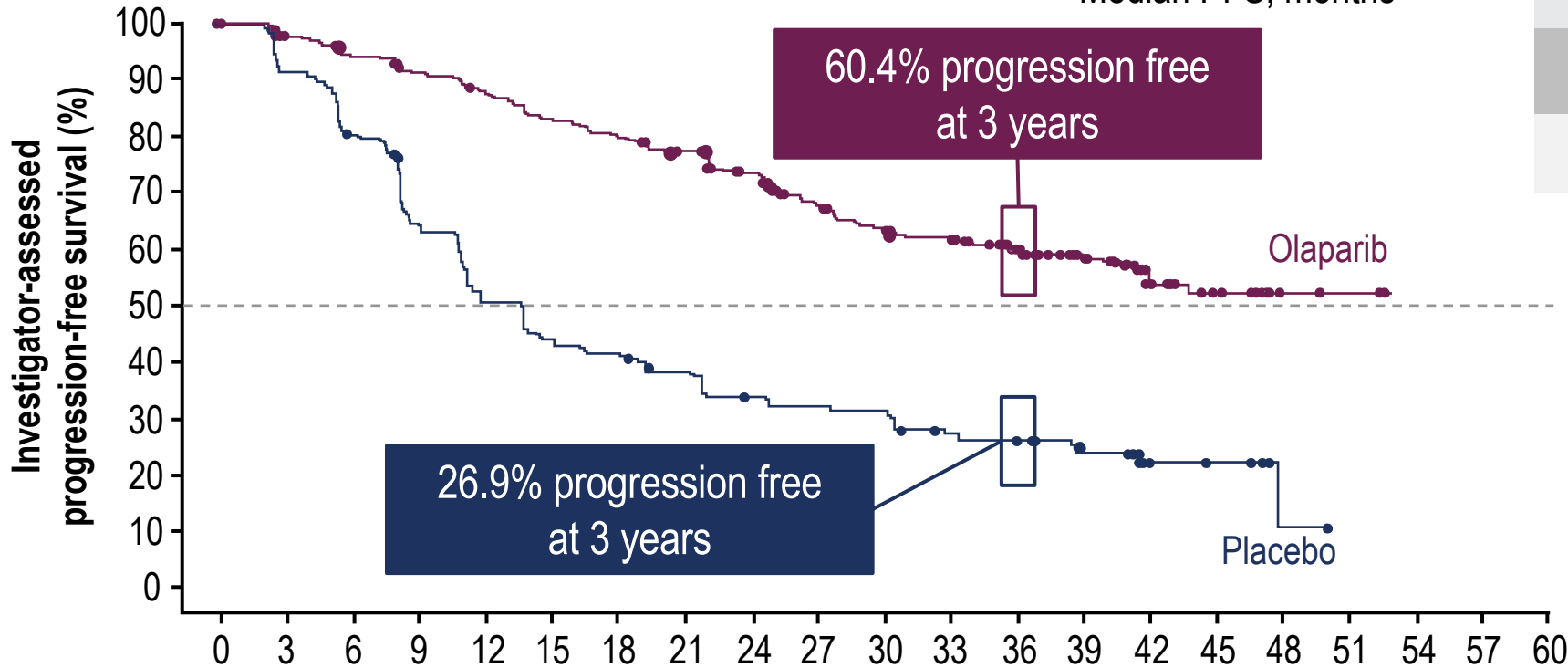
<sup>1</sup>Stephenson Cancer Center at the University of Oklahoma, Oklahoma City, OK, USA; <sup>2</sup>University of Milan-Bicocca and IEO, European Institute of Oncology IRCCS, Milan, Italy; <sup>3</sup>Fondazione Policlinico Universitario A. Gemelli IRCCS Università Cattolica, Rome, Italy; <sup>4</sup>Samsung Medical Center, Sungkyunkwan University School of Medicine, Seoul, Korea; <sup>5</sup>Vall d'Hebron University Hospital, Vall d'Hebron Institute of Oncology (VHIO), Barcelona, Spain; <sup>6</sup>University of New South Wales Clinical School, Prince of Wales Hospital, Randwick, Australia; <sup>7</sup>St Petersburg City Oncology Dispensary, St Petersburg, Russia; <sup>8</sup>Institut Bergonié, Comprehensive Cancer Centre, Bordeaux, France; <sup>9</sup>Gustave-Roussy Cancer Campus, Villejuif, France; <sup>10</sup>The Netherlands Cancer Institute, Amsterdam, The Netherlands; <sup>11</sup>Cancer Research UK Edinburgh Centre, Institute of Genetics and Molecular Medicine, University of Edinburgh, Edinburgh, UK; <sup>12</sup>The Royal Marsden NHS Foundation Trust and Institute of Cancer Research, London, UK; <sup>13</sup>Princess Margaret Cancer Centre, Toronto, ON, Canada; <sup>14</sup>Clinica Universidad de Navarra, Madrid, Spain; <sup>15</sup>Memorial Sloan Kettering Cancer Center, New York, NY, USA; <sup>16</sup>Froedtert and the Medical College of Wisconsin, Milwaukee, WI, USA; <sup>17</sup>AstraZeneca, Gaithersburg, MD, USA; <sup>18</sup>AstraZeneca, Cambridge, UK; <sup>19</sup>Women & Infants Hospital, Providence, RI, USA

esmo.org

# PFS by investigator assessment

Events (%) [50.6% maturity]

Median PFS, months



Olaparib (N=260)	Placebo (N=131)
102 (39.2)	96 (73.3)
NR	13.8
<b>HR 0.30</b>	
95% CI 0.23, 0.41; <i>P</i> <0.0001	

No. at risk

	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51	54	57	60
Olaparib	260	240	229	221	212	201	194	184	172	149	138	133	111	88	45	36	4	3	0	0	0
Placebo	131	118	103	82	65	56	53	47	41	39	38	31	28	22	6	5	1	0	0	0	0



The AACR logo features the letters 'AACR' in a bold, black, sans-serif font, with a green 'R' that has a white cross-like shape inside it.

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A colorful banner for the AACR 2022 Annual Meeting. It features a collage of images including people in lab coats, a microscope, and various scientific diagrams, all set against a background of colorful, abstract shapes. A green banner at the bottom of the collage contains the text 'APRIL 8-13, 2022 • #AACR22' in white.

APRIL 8-13, 2022 • #AACR22

**PETRA: A first-in-class, first-in-human trial of the** **CT007**  
**next-generation PARP1-selective inhibitor AZD5305 in**  
**patients with *BRCA1/2*, *PALB2* or *RAD51C/D* mutations**

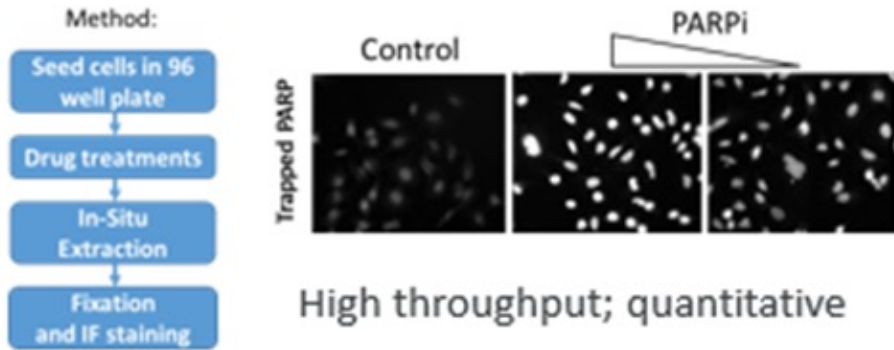
**Presenter: Timothy Yap, M.D.**

**Discussant: Patricia Mucci LoRusso, D.O., Ph.D.(h)**

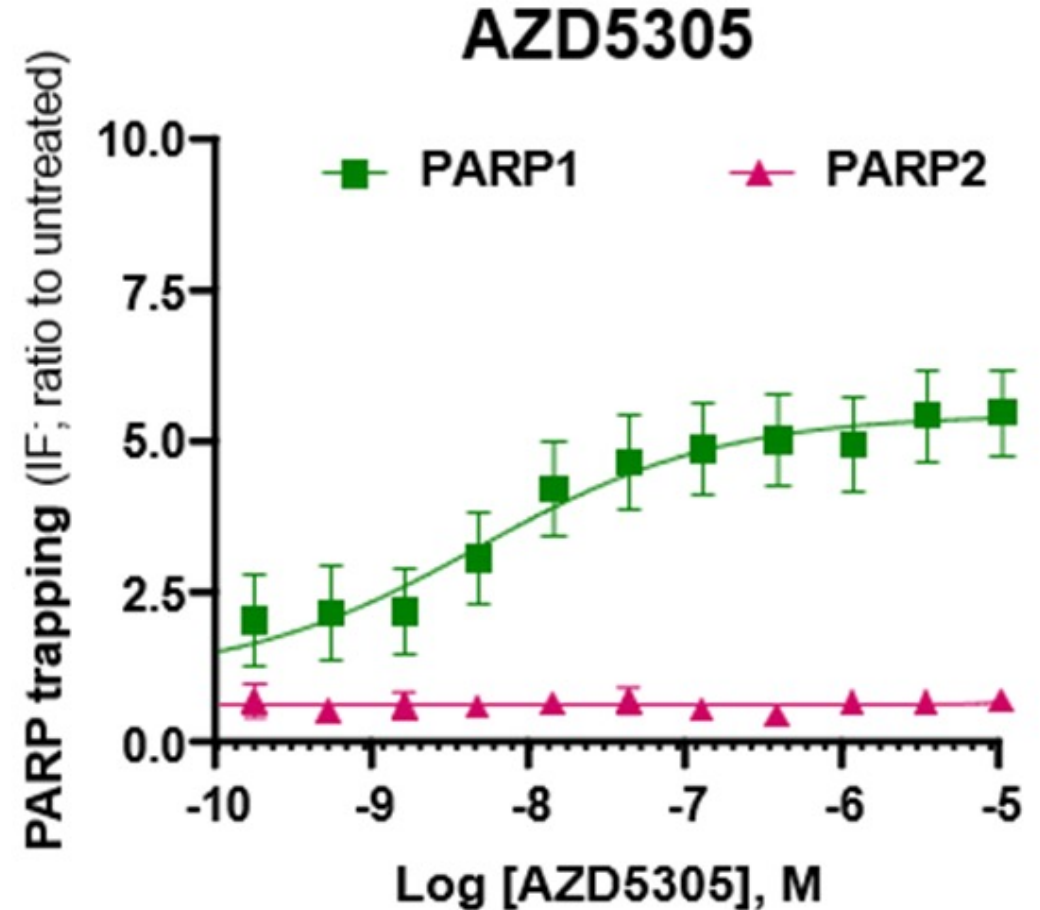
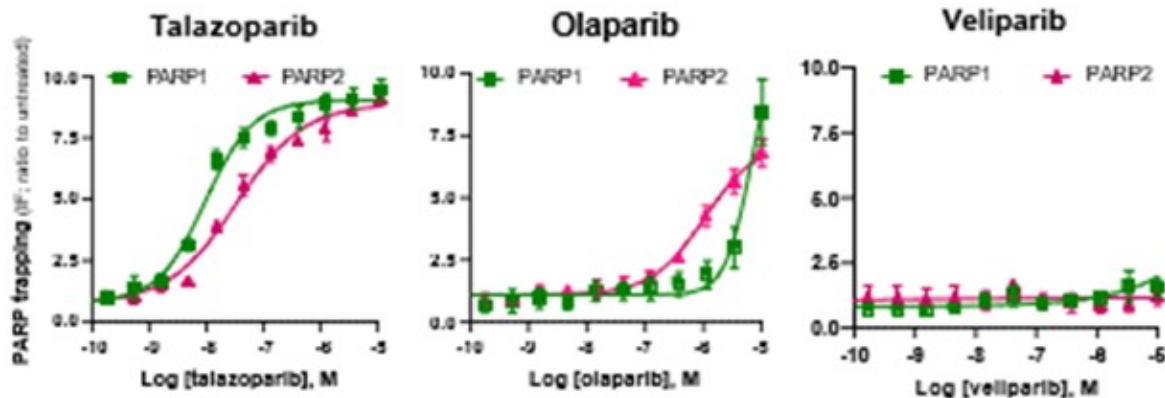
Professor of Medicine  
Associate Center Director: Experimental Therapeutics  
Yale University/Yale Cancer Center

# Trapping: AZD5305 selectively traps PARP1 onto the chromatin

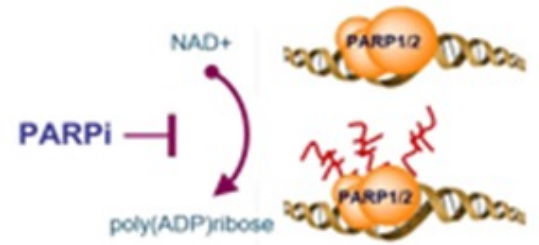
Novel PARP-trapping assay developed



Trapping profiles of clinical PARPi

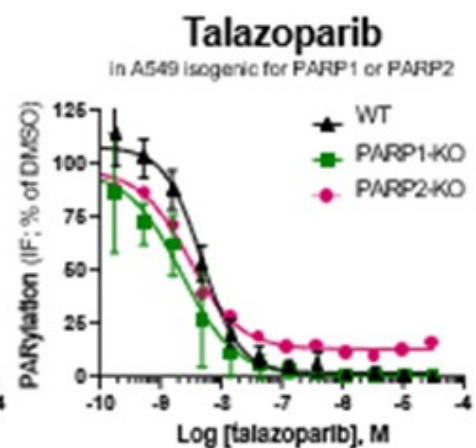
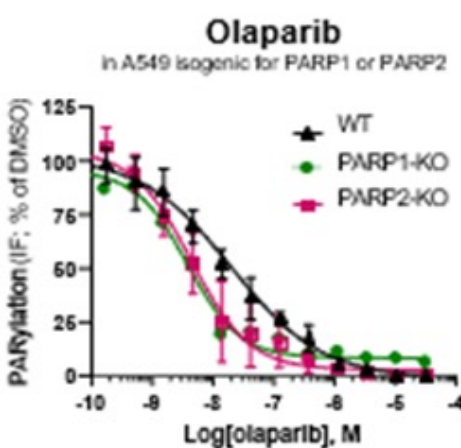
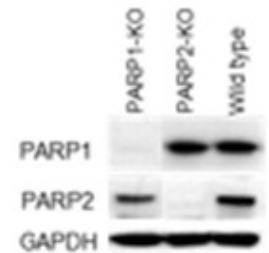


# PARylation: AZD5305 selectively inhibits PARP1 in cells

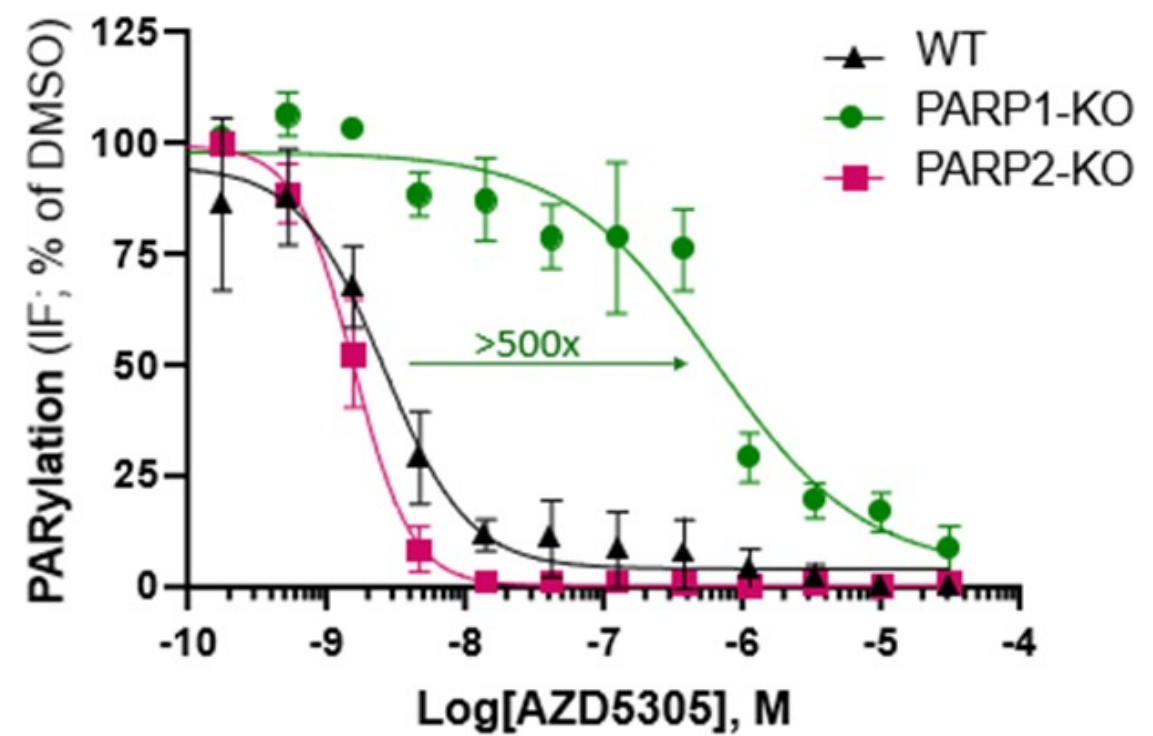


PARylation inhibition is most proximal biomarker for PARPi

PARylation contribution		
WT	PARP1	PARP2
PARP1-KO	/	PARP2
PARP2-KO	PARP1	/

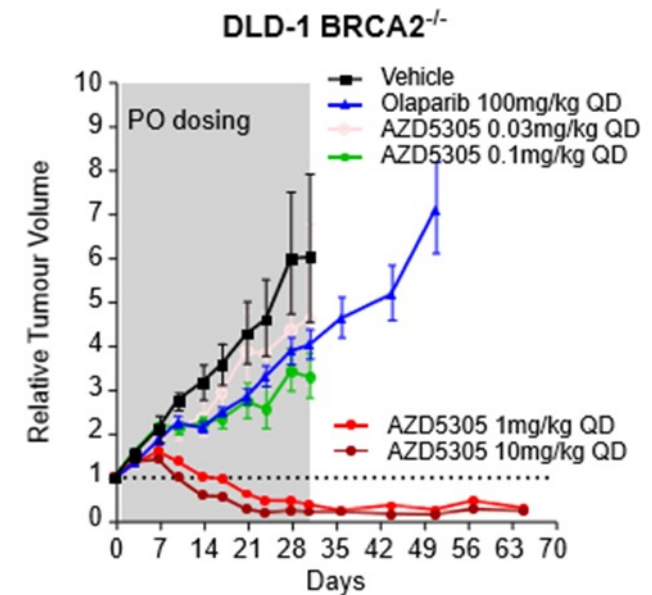
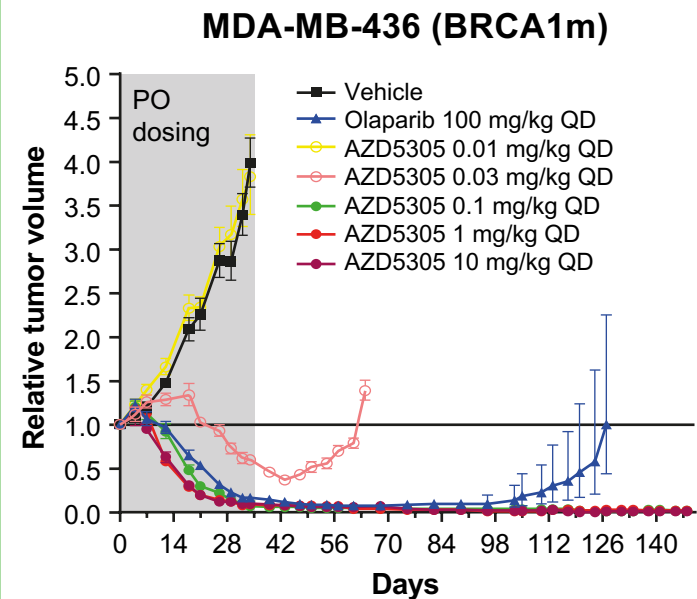


## AZD5305 PARylation in A549 isogenic for PARP1 or PARP2



# AZD5305 - more efficacious than first generation PARPi

- Durable regression observed following
- cessation of AZD5305 dosing<sup>4</sup>

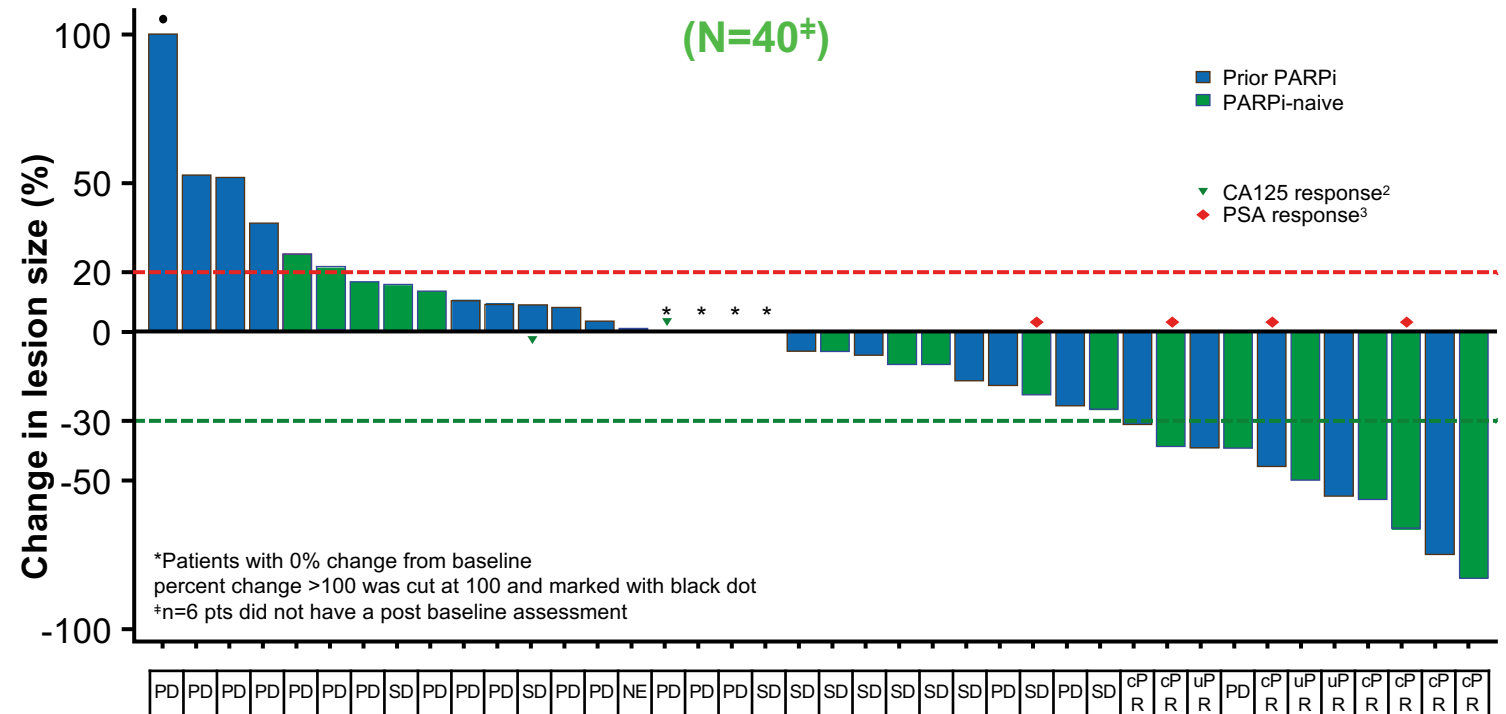


# Response vs Prior PARP Inhibitor Exposure

APRIL 8-13 • #AACR22

- 46 patients were included in the interim response analysis set<sup>1</sup>
- 6 patients were not evaluable<sup>2</sup> for RECIST v1.1 assessment
- Responses seen regardless of prior PARPi
- Few CRs at time of data cutoff
- Should we focus on response duration rather than response type?
- Were those PARPi exposed patients truly resistant to 1<sup>st</sup> generation inhibitors or was it drug exposure/pharmacologic resistance?
- Will combination strategies lead to improvement

Best % change in target lesions size by prior PARPi

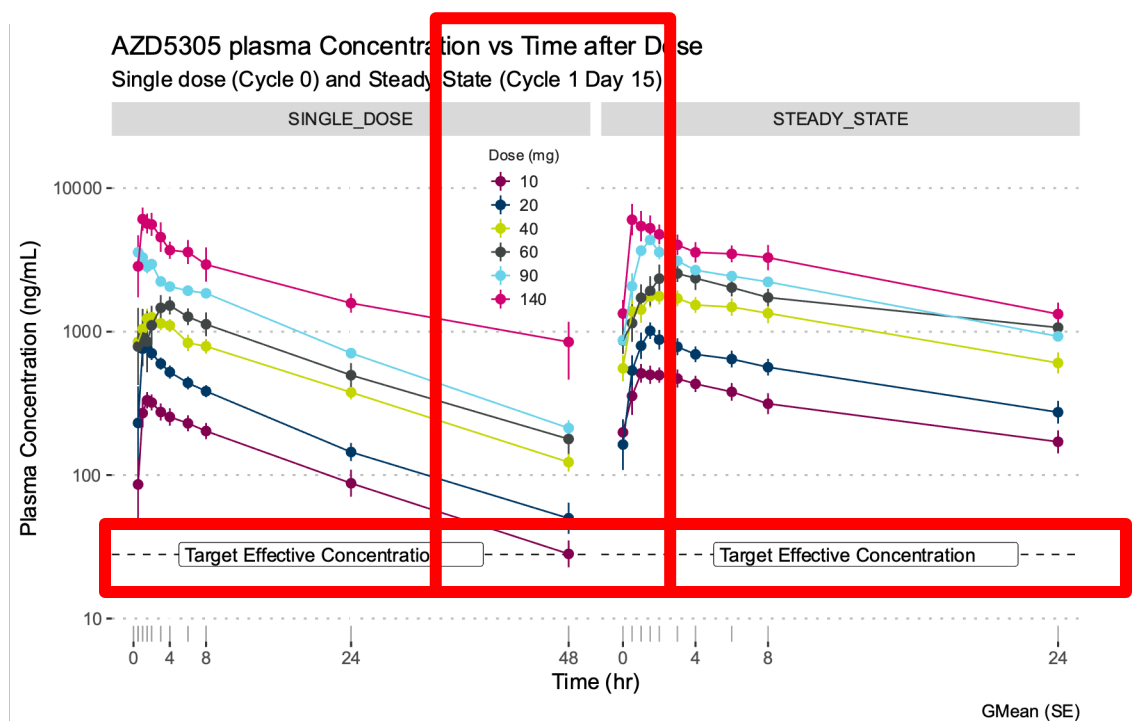


<sup>1</sup>All dosed subjects who had measurable disease at baseline and who received their first dose at least 13 weeks prior to data extract <sup>2</sup>n=5 did not have a follow up scan and n=1 had SD<7 weeks <sup>3</sup>GCIG CA125 response <sup>4</sup>PCWG3 PSA<sub>50</sub> response <sup>‡</sup>n=2 can be confirmed  
 cPR, confirmed Partial Response; uPR, unconfirmed Partial Response; SD, Stable disease; PD, progressive disease

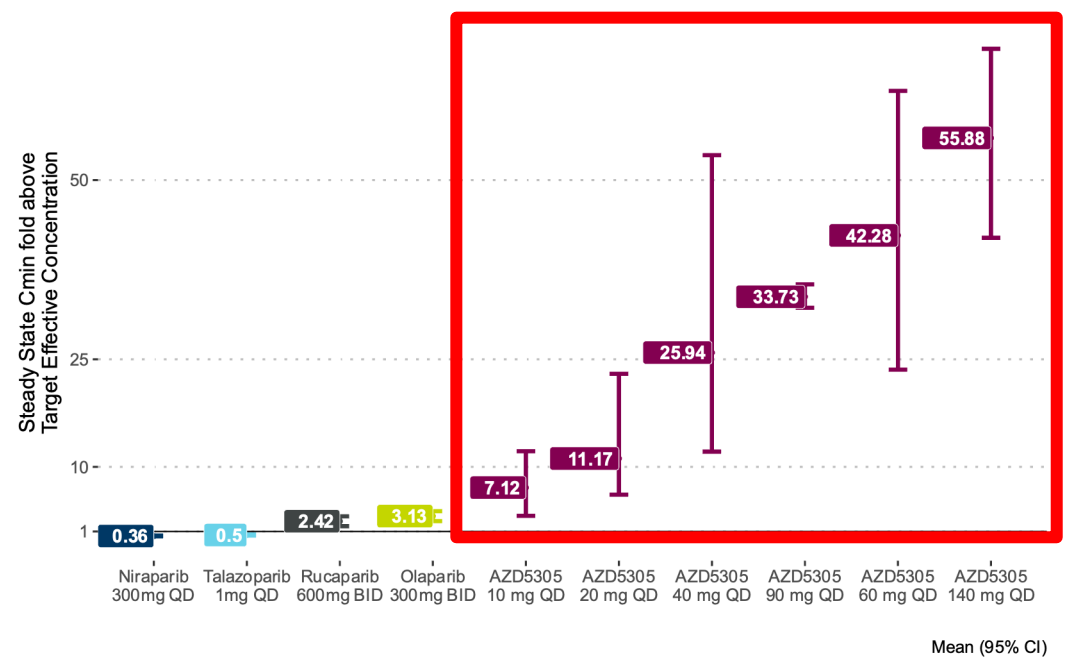
# AZD5305 Achieves Higher Fold Coverage over the TEC Compared to First Generation PARPi

APRIL 8-13 • #AACR22

## AZD5305 plasma concentration vs time after dose



## First-generation PARPi and AZD5305 C<sub>min</sub> fold above TEC



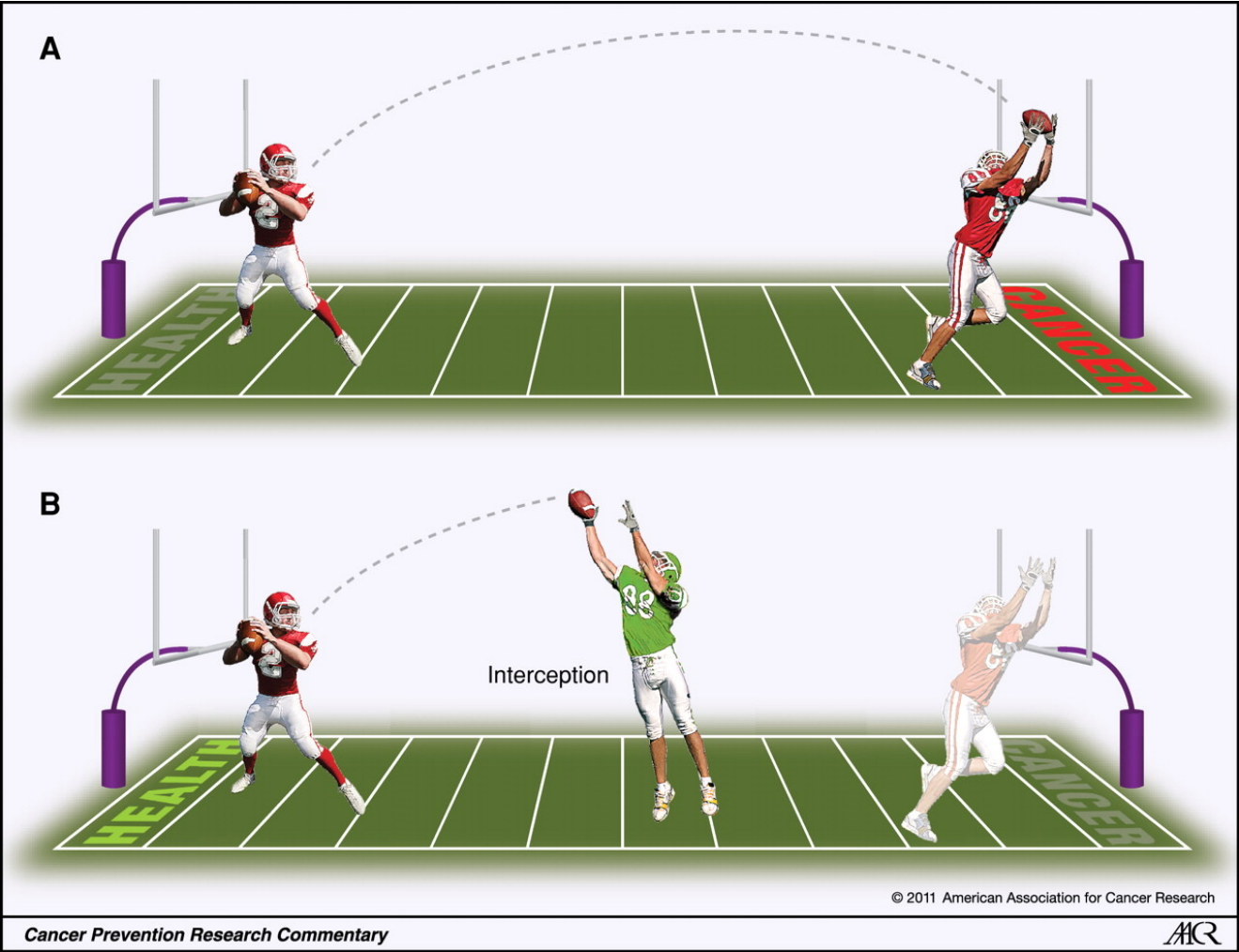
- Dose proportional increase in exposure (C<sub>max</sub> & AUC) observed with increase in dose (10 – 140 mg QD)
- Quick onset (T<sub>max</sub> 0.5-3 h) with mean terminal elimination half-life 13.1–16.4 h across cohorts
- Steady state C<sub>min</sub> above target effective concentration (TEC) in all patients with mean fold C<sub>min</sub>/TEC 7.12 and 55.88 at 10 mg and 140 mg QD, respectively

# Results: Overall Safety Summary

AE category, n (%)	AZD5305 10 mg QD (n=8)	AZD5305 20 mg QD (n=19)	AZD5305 40 mg QD (n=17)	AZD5305 60 mg QD (n=10)	AZD5305 90 mg QD (n=3)	AZD5305 140 mg QD (n=4)	Total (N=61)
<b>Duration of therapy (months), median (range)</b>	4.8 (1.8-14.9)	2.8 (0.9-11.5)	2.0 (0.4-9.6)	1.95 (0-3.8)	4.8 (4.6-4.9)	0.7 (0.6-1.6)	2.1 (0-14.9)
<b>All TEAEs</b>	8 (100.0)	19 (100.0)	17 (100.0)	8 (80.0)	3 (100.0)	0	55 (90.2)
Grade ≥3 AEs	3 (37.5)	10 (52.6)	6 (35.3)	3 (30.0)	2 (66.7)	0	24 (39.3)
Serious TEAEs	0	9 (47.4)	5 (29.4)	2 (20.0)	0	0	16 (26.2)
Discontinuations	0	0	0	1 (10.0)	0	0	1 (1.6)
Dose reductions	1 (12.5)	0	1 (5.9)	0	0	0	2 (3.3)
<b>AZD5305-related TEAEs</b>	5 (62.5)	11 (57.9)	9 (52.9)	4 (40.0)	3 (100.0)	0	32 (52.5)
Grade ≥3 AEs	3 (37.5)	3 (15.8)	1 (5.9)	0	2 (66.7)	0	9 (14.8)
Serious TEAEs	0	0	0	0	0	0	0
Discontinuations	0	0	0	0	0	0	0
Dose reductions	1 (12.5)	0	1 (5.9)	0	0	0	2 (3.3)

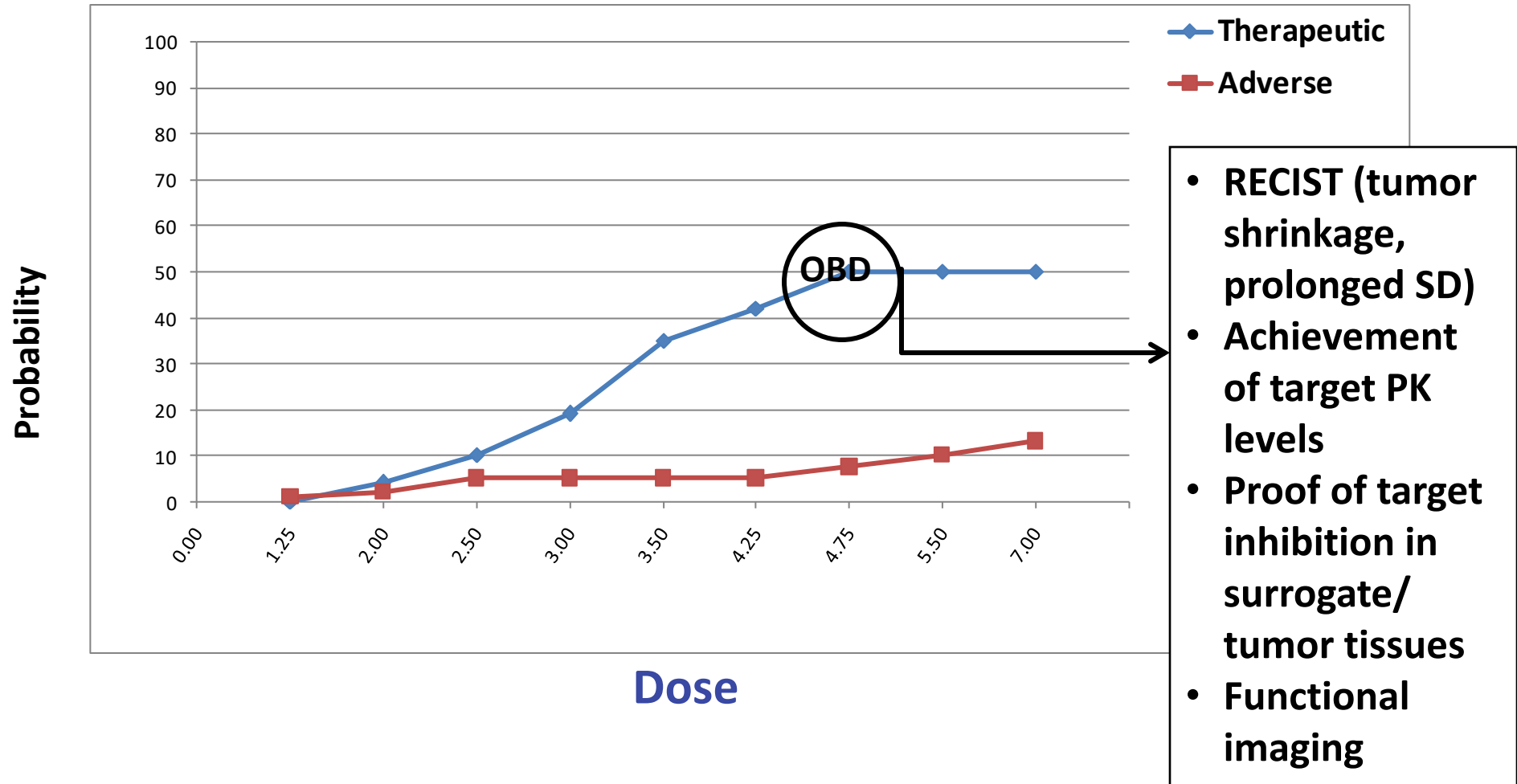
- AZD5305 was well tolerated across doses
- Only 2 patients (3.3%) had dose reductions (for grade 3 neutropenia and grade 1 thrombocytopenia related to AZD5305)
- At DCO there have been no DLTs and no AZD5305-related serious AEs or treatment discontinuations

# Cancer Interception





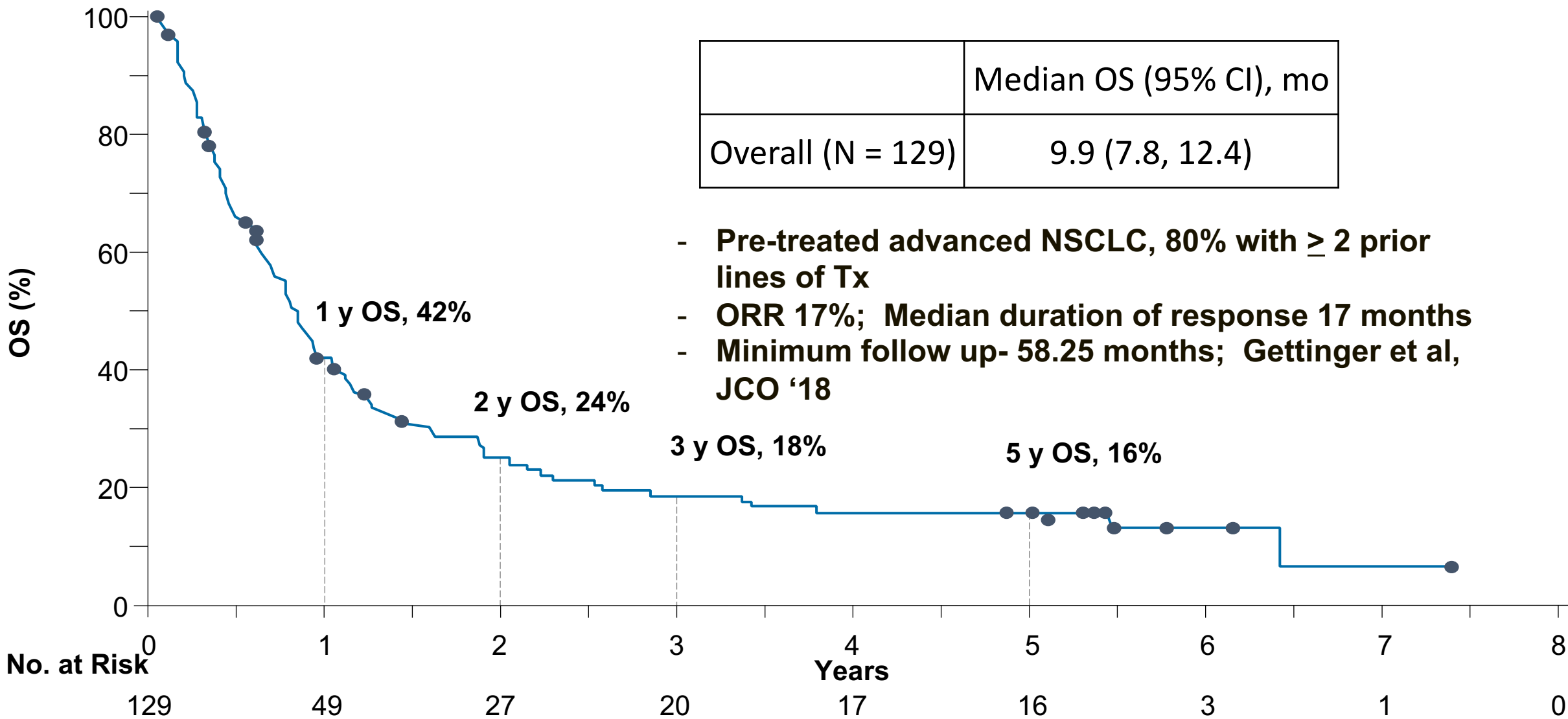
# Dose-Response: Efficacy and Toxicity



---

# Earlier Intervention

# CA209-003, Salvage Nivolumab for Adv NSCLC



	Median OS (95% CI), mo
Overall (N = 129)	9.9 (7.8, 12.4)

- Pre-treated advanced NSCLC, 80% with  $\geq 2$  prior lines of Tx
- ORR 17%; Median duration of response 17 months
- Minimum follow up- 58.25 months; Gettinger et al, JCO '18



# Keynote 24- Front Line

(biomarker driven)

## 5 year OS:

Median

26.3 mo (18.3-40.4 mo) 31.9% vs.

13.4 mo (9.4-18.3 mo) 16.3%

HR: 0.62 (0.48-0.81)

## 3 Year PFS:

Median

7.7 mo (6.1-10.2 mos) 22.8% vs.

5.5 mo (4.2-6.2 mo) 4.1%

HR: 0.50 (0.39-0.65)

## Overall Response Rate:

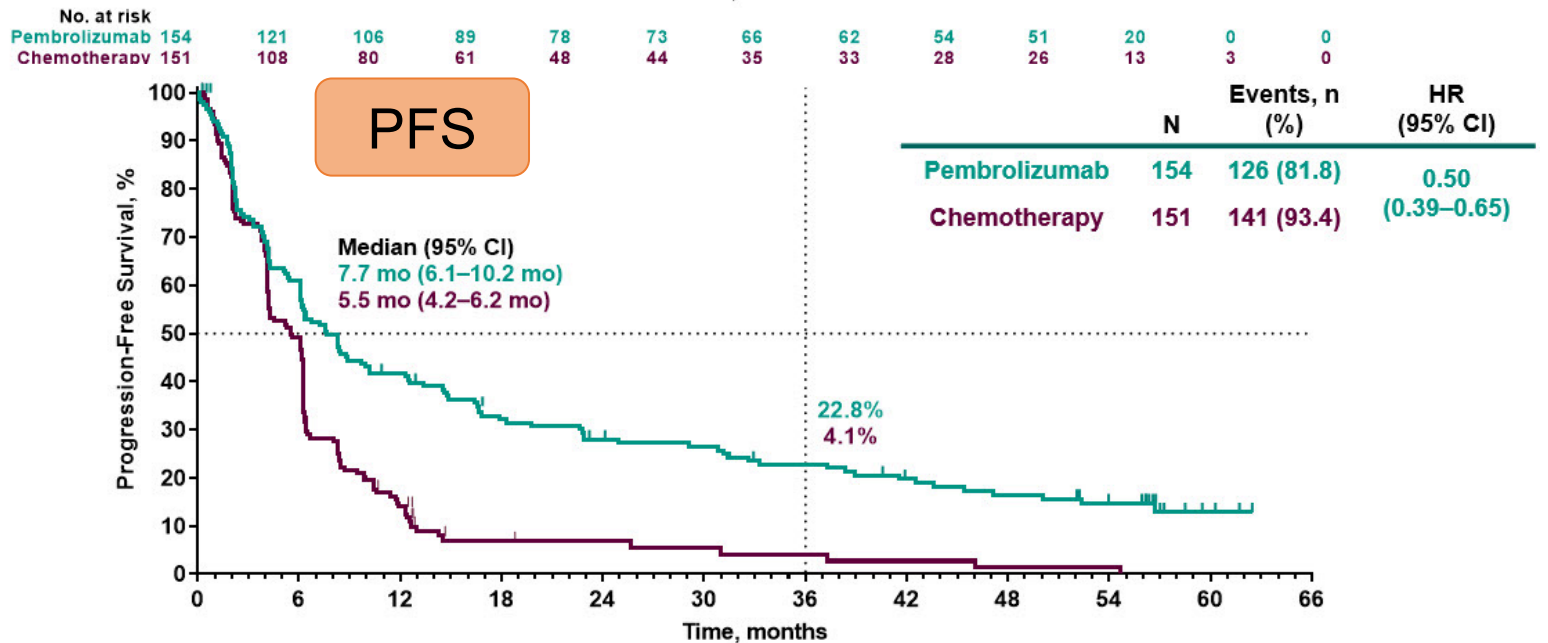
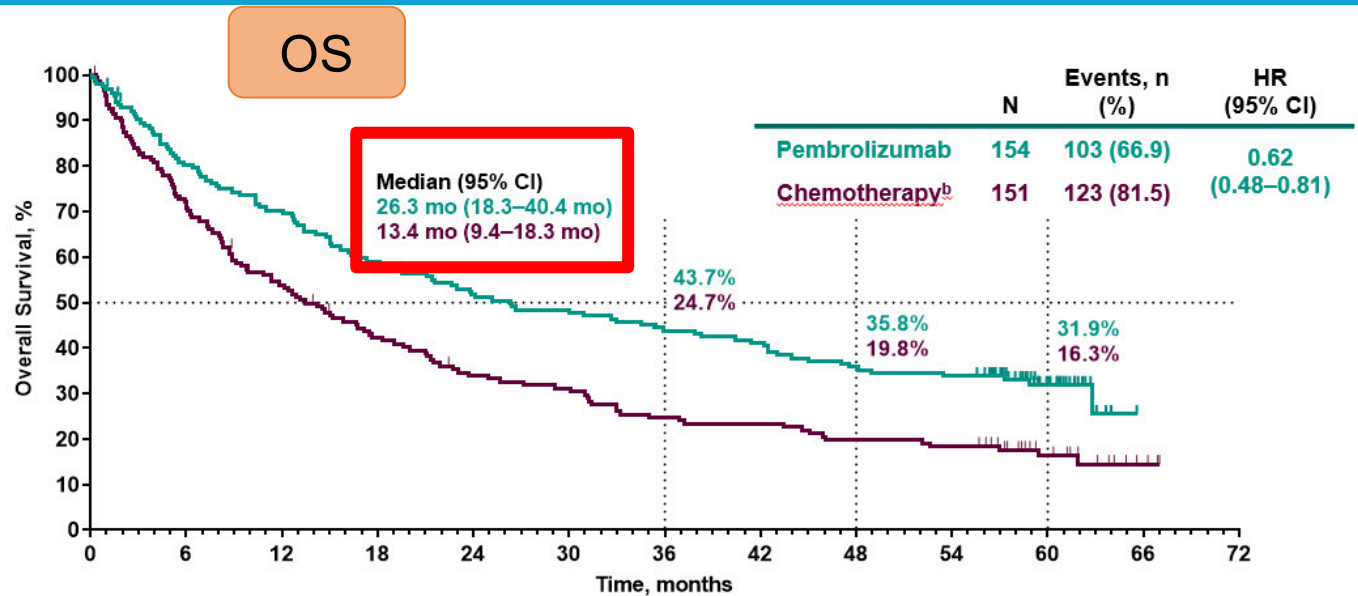
46.1% vs. 31.1%

## Partial Response:

41.6% vs. 31.1%

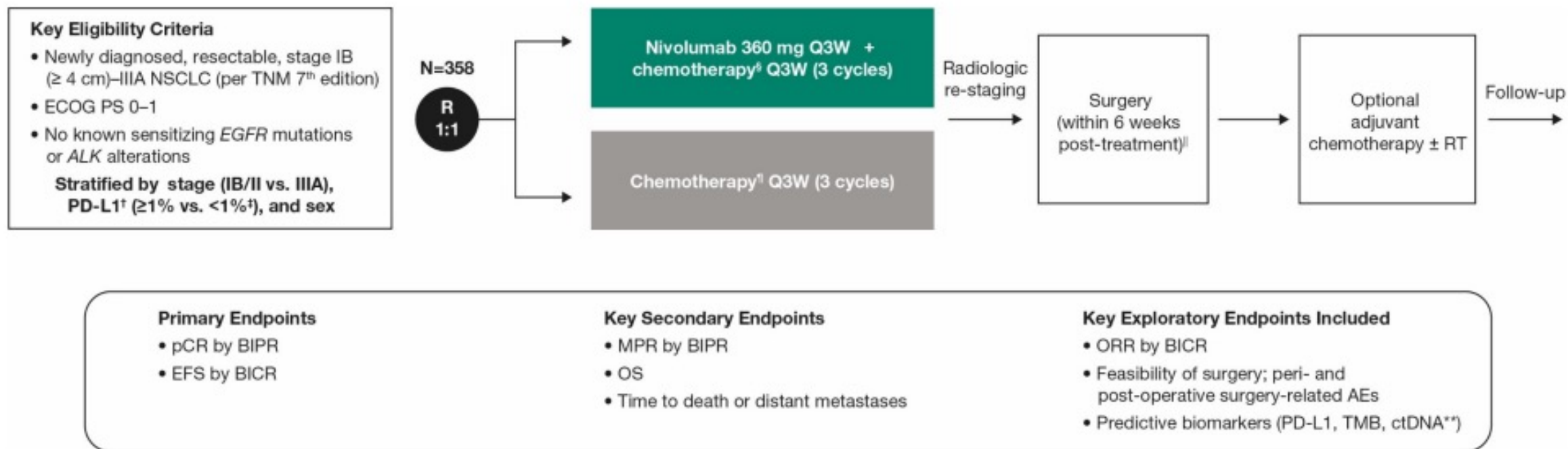
## Complete Response:

4.5% vs. 0

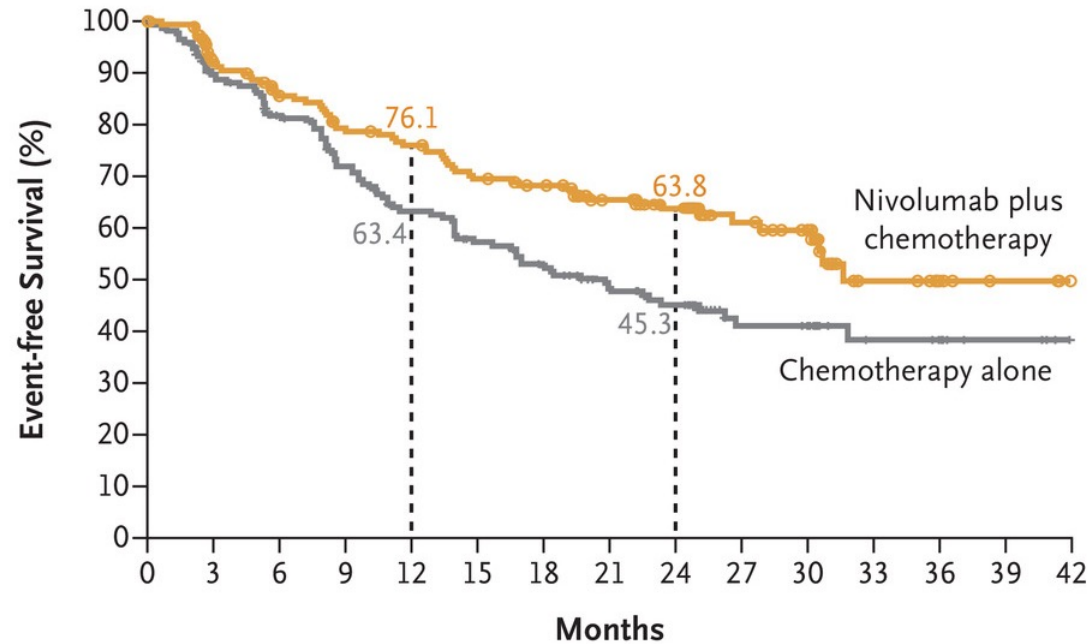


No. at risk	0	6	12	18	24	30	36	42	48	54	60	66
Pembrolizumab	154	92	62	46	38	36	30	24	20	15	3	0
Chemotherapy	151	73	20	6	5	4	3	2	1	1	0	0

# CHECKPOINT 816 STUDY DESIGN



# EVENT-FREE SURVIVAL ACCORDING TO BLINDED INDEPENDENT CENTRAL REVIEW



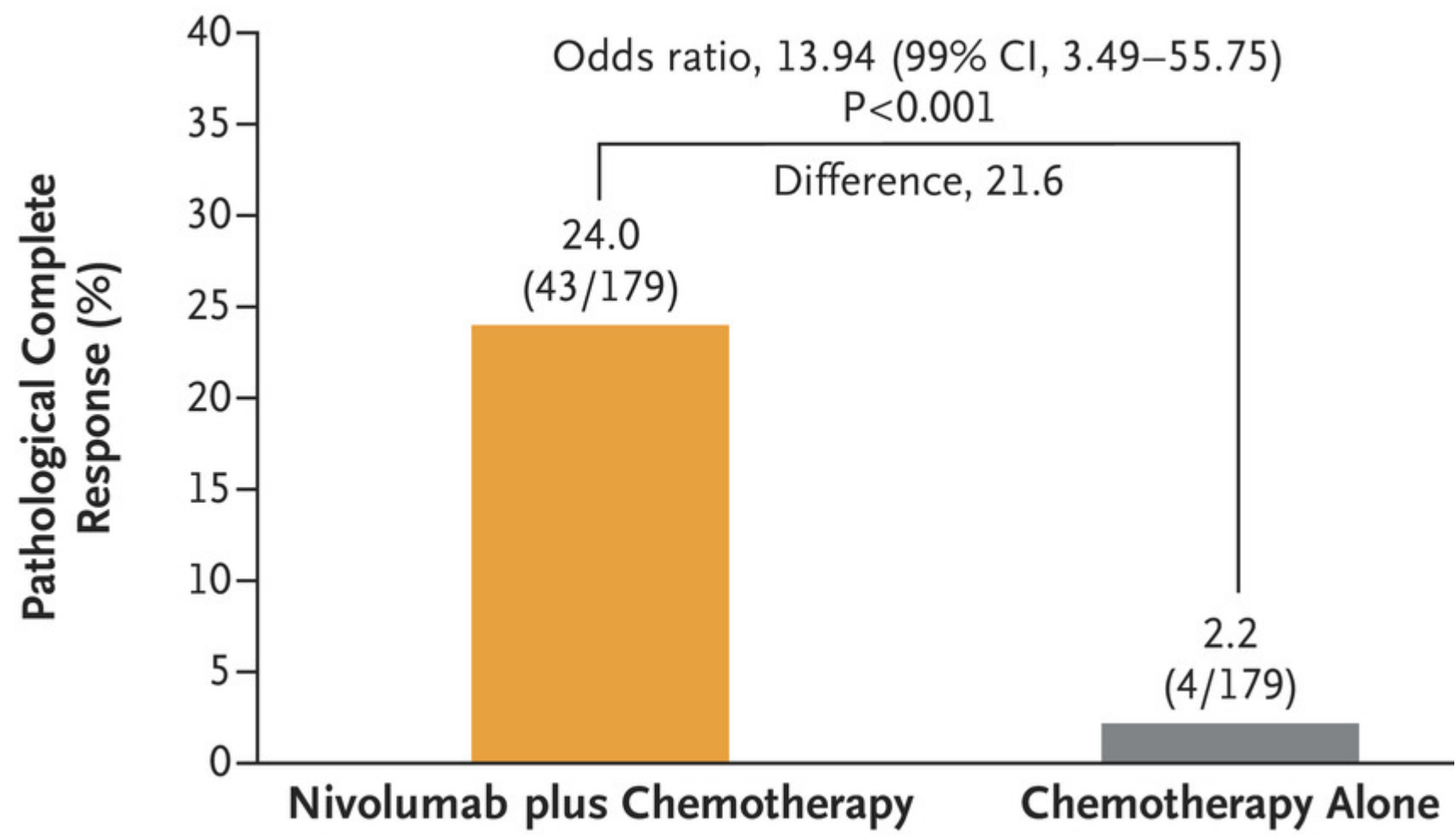
	No. of Patients	Median Event-free Survival (95% CI) mo
<b>Nivolumab plus Chemotherapy</b>	179	31.6 (30.2–NR)
<b>Chemotherapy Alone</b>	179	20.8 (14.0–26.7)

Hazard ratio for disease progression, disease recurrence, or death, 0.63 (97.38% CI, 0.43–0.91)  
P=0.005

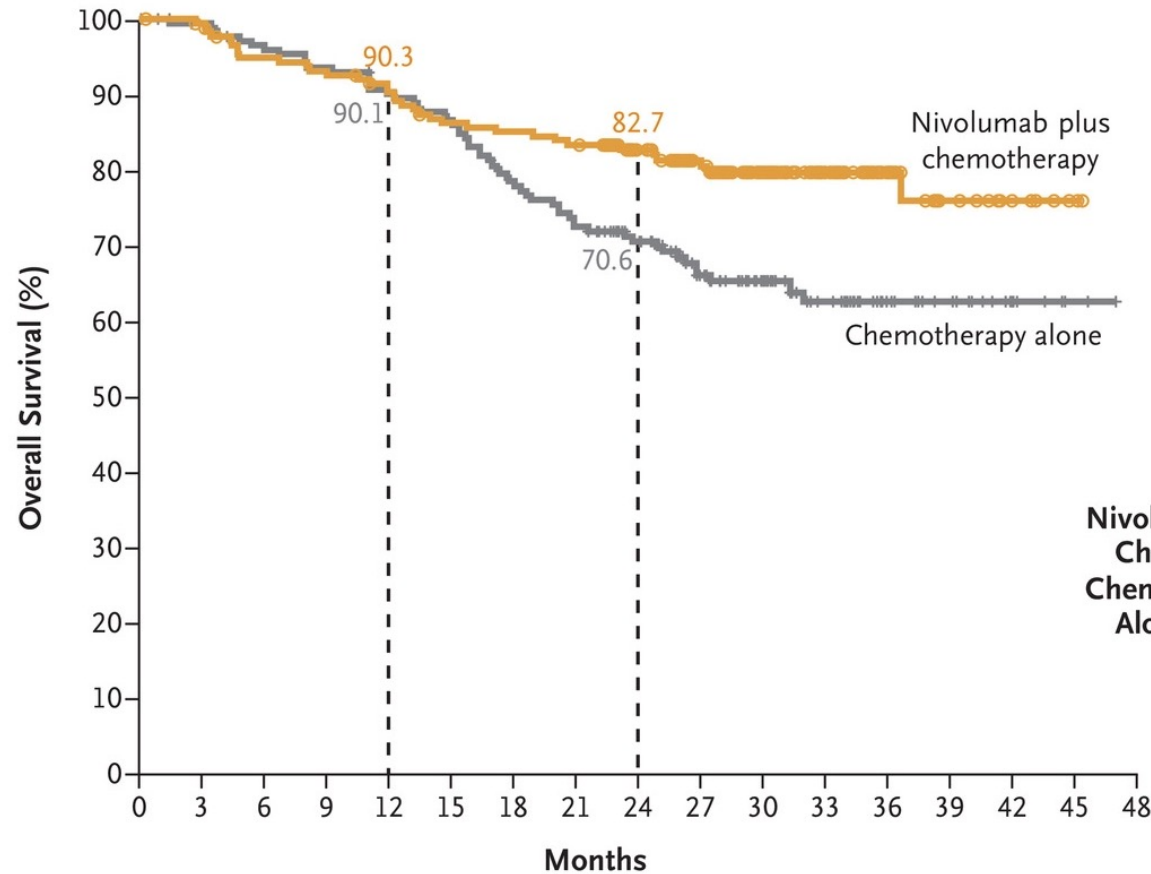
## No. at Risk

Nivolumab plus chemotherapy	179	151	136	124	118	107	102	87	74	41	34	13	6	3	0
Chemotherapy alone	179	144	126	109	94	83	75	61	52	26	24	13	11	4	0

# PATHOLOGICAL COMPLETE RESPONSE ACCORDING TO BLINDED INDEPENDENT PATHOLOGICAL REVIEW



# OVERALL SURVIVAL



	No. of Patients	Median Overall Survival (95% CI) mo
Nivolumab plus Chemotherapy	179	NR (NR–NR)
Chemotherapy Alone	179	NR (NR–NR)

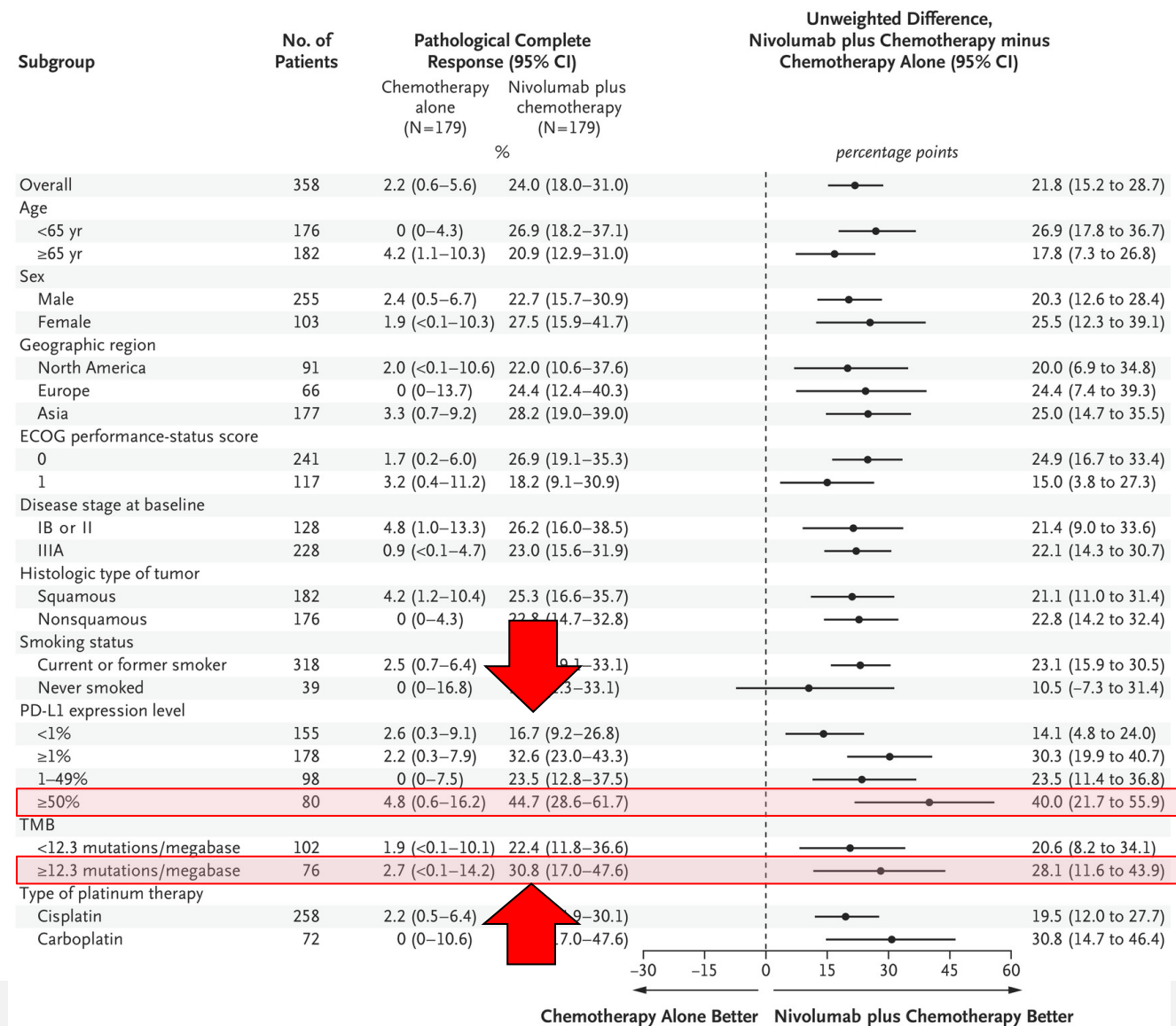
Hazard ratio for death, 0.57  
(99.67% CI, 0.30–1.07)  
P=0.008

## No. at Risk

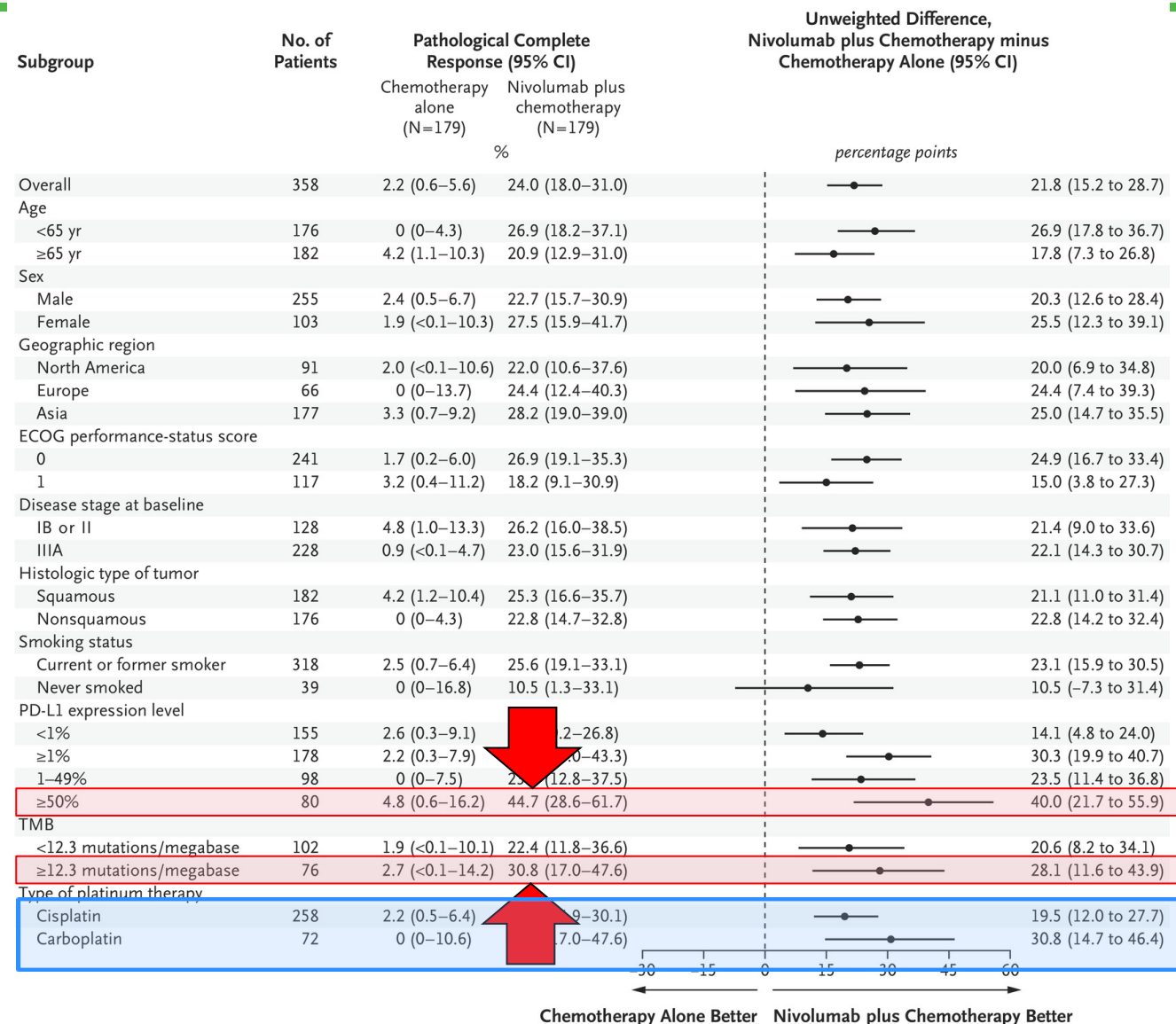
Nivolumab plus chemotherapy	179	176	166	163	156	148	146	143	122	101	72	48	26	16	7	3	0
Chemotherapy alone	179	172	165	161	154	148	133	123	108	80	59	41	24	16	7	2	0



# PATHOLOGICAL COMPLETE RESPONSE ACCORDING TO BLINDED INDEPENDENT PATHOLOGICAL REVIEW



# PATHOLOGICAL COMPLETE RESPONSE ACCORDING TO BLINDED INDEPENDENT PATHOLOGICAL REVIEW



• Newly diagnosed, resectable Stage Ib ( $\geq 4\text{cm}$ ) to IIIa (N2) NSCLC  
• PD-L1 all-comers  
• TMB all-comers  
• EGFR/ALK/ROS WT

Core tissue biopsy available at screening or amenable to re-biopsy  
Stratify by:  
- Stage (IB-II vs III)  
- PD-L1 Status  
- TMB Status  
- Gender

**Arm 1:**  
TMB  $\geq 10$   
and PDL1  $\geq 50\%$

**Arm 1A:**  
Nivolumab + Carboplatin Doublet

**Arm 1B:** Nivolumab + Carboplatin Doublet + Drug X

**Arm 2:**  
TMB  $< 10$   
or PDL1  $< 50\%$

**Arm 2A:**  
Nivolumab + Carboplatin Doublet

**Arm 2B:** Nivolumab + Carboplatin Doublet + Drug X

**Arm 3:**  
TMB  $< 10$   
and PDL1  $< 50\%$

**Arm 3A:**  
Nivolumab + Carboplatin Doublet

**Arm 3B:** Nivolumab + Carboplatin Doublet + Drug X

Surgery

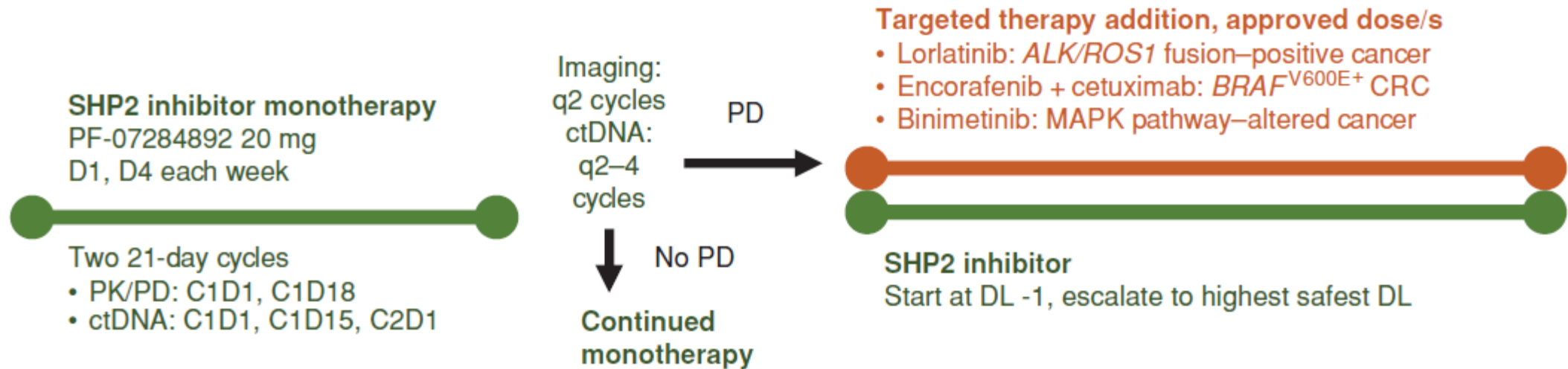
## RESEARCH BRIEF

# SHP2 Inhibition Sensitizes Diverse Oncogene-Addicted Solid Tumors to Re-treatment with Targeted Therapy

Alexander Drilon<sup>1</sup>, Manish R. Sharma<sup>2</sup>, Melissa L. Johnson<sup>3</sup>, Timothy A. Yap<sup>4</sup>, Shirish Gadgeel<sup>5</sup>, Dale Nepert<sup>6</sup>, Gang Feng<sup>7</sup>, Micaela B. Reddy<sup>6</sup>, Allison S. Harney<sup>6</sup>, Mohamed Elsayed<sup>6</sup>, Adam W. Cook<sup>6</sup>, Christina E. Wong<sup>6</sup>, Ronald J. Hinklin<sup>6</sup>, Yutong Jiang<sup>6</sup>, Eric N. Brown<sup>6</sup>, Nickolas A. Neitzel<sup>6</sup>, Ellen R. Laird<sup>6</sup>, Wen-I Wu<sup>6</sup>, Anurag Singh<sup>6</sup>, Ping Wei<sup>8</sup>, Keith A. Ching<sup>8</sup>, John J. Gaudino<sup>6</sup>, Patrice A. Lee<sup>6</sup>, Dylan P. Hartley<sup>6</sup>, and S. Michael Rothenberg<sup>6,8</sup>

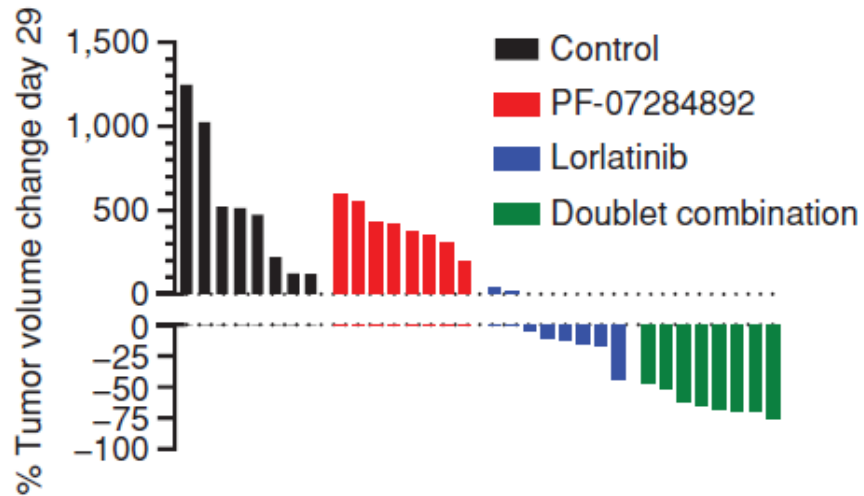
# SHP2 inhibitor PF-07284892 Phase I Study Schema

Oncogene-driven solid tumors with progression on matched targeted therapy  
*ALK/ROS1* fusion-positive cancer, *BRAF*<sup>V600E+</sup> CRC, or MAPK pathway-altered cancer

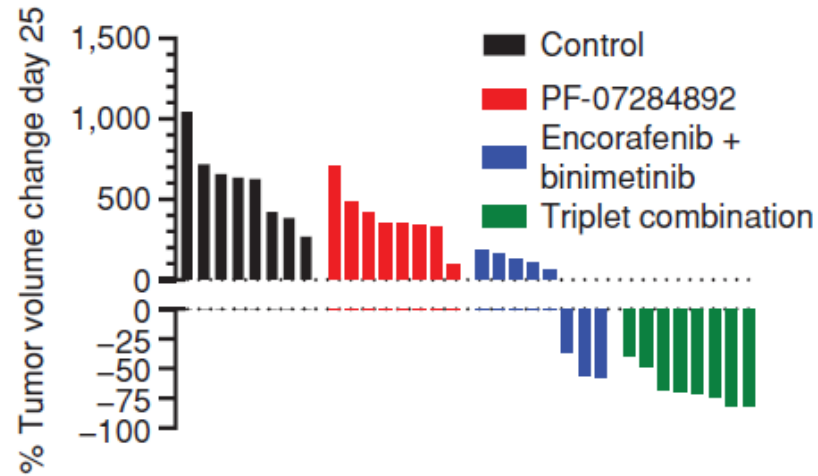


# PF-07284892 *In Vitro* Efficacy

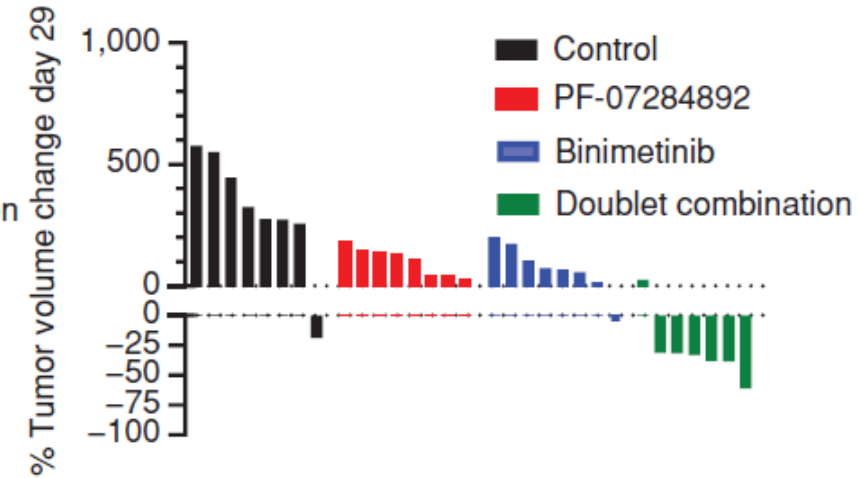
NCI-H3122 lorlatinib-resistant  
*ALK* fusion-positive NSCLC



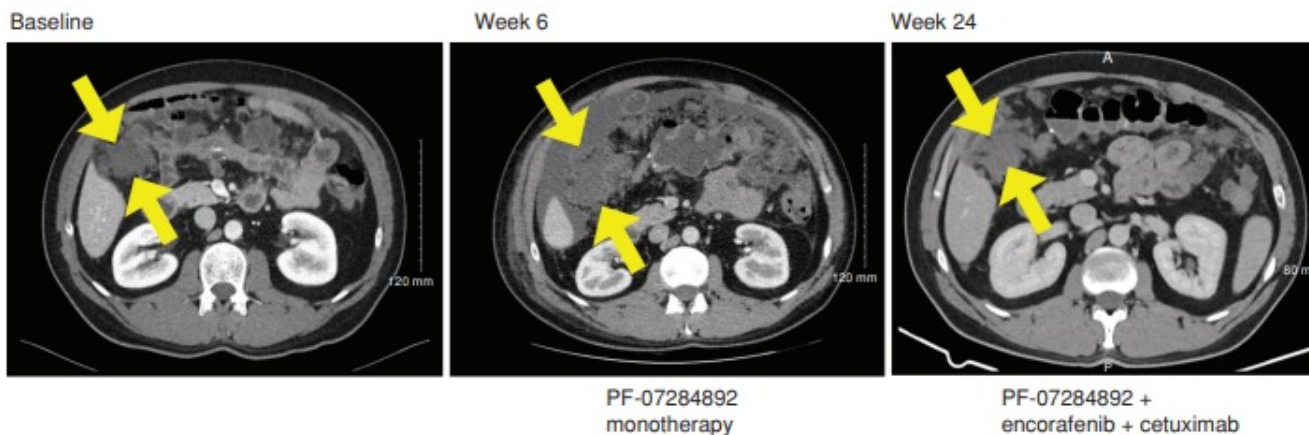
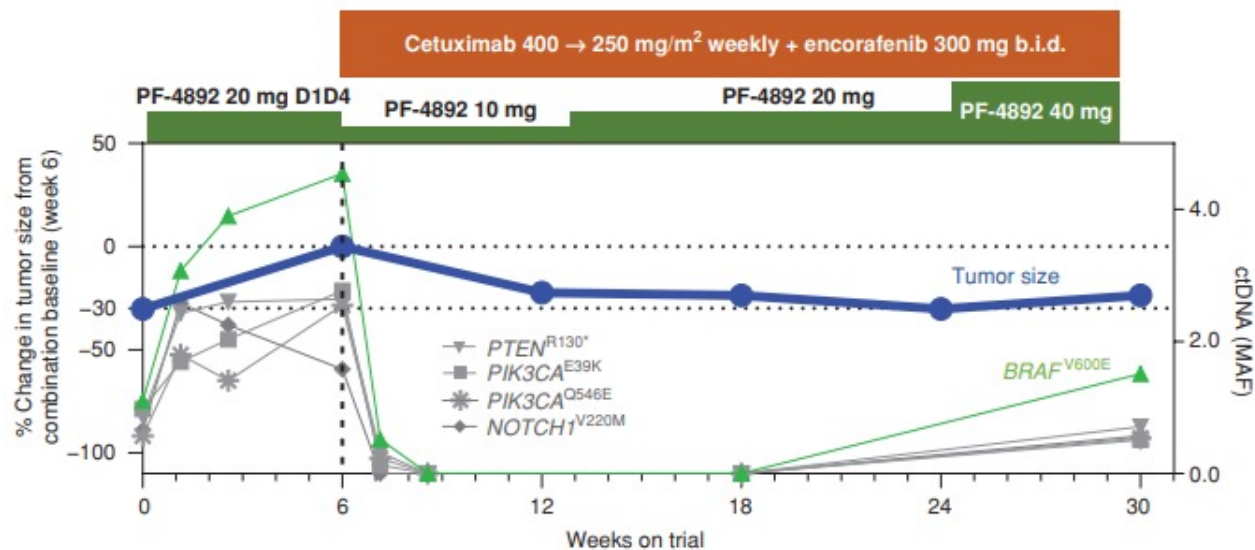
VACO-432  
*BRAF*<sup>V600E</sup>-mutant CRC



MIA PaCa-2  
*KRAS*<sup>G12C</sup>-mutant PDAC



# PF-07284892 overcomes intrinsic resistance to encorafenib + cetuximab in a BRAFV600E-mutant CRC patient



# Novel Technologies

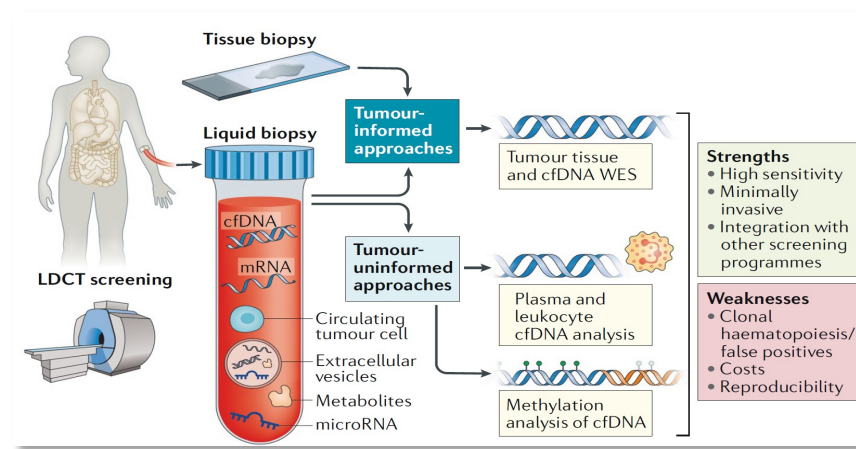
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YaleNewHaven**Health**  
Smilow Cancer Hospital

Yale **CANCER**  
CENTER  
A Comprehensive Cancer Center Designated  
by the National Cancer Institute

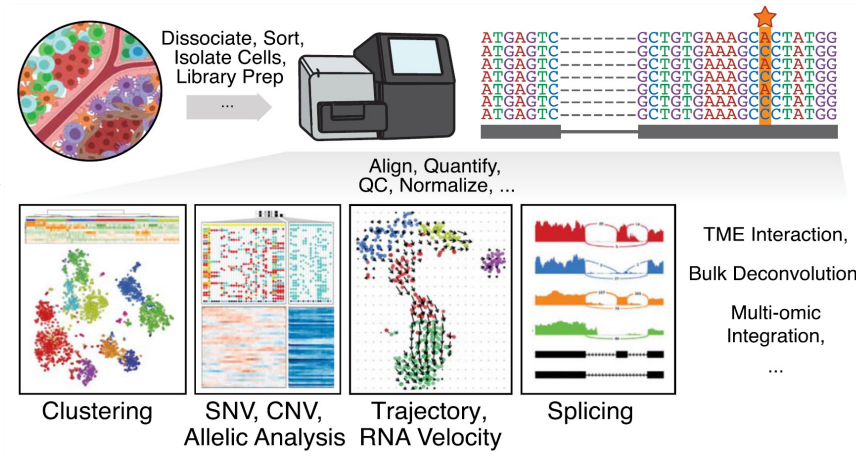
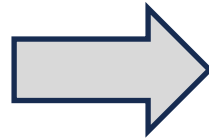


# Liquid biopsy



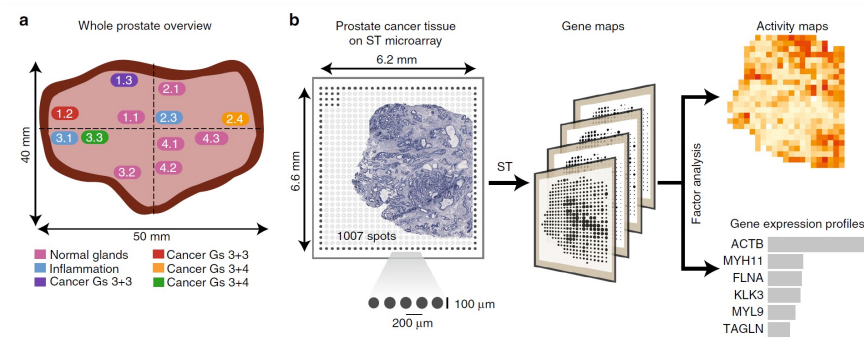
Rolfo and Russo., 2020. Nature Reviews Clinical Oncology

# Single-cell transcriptomics



Fan et al., 2020. Experimental & Molecular Medicine volume

# Spatial transcriptomics & proteomics



Berglund et al., 2020. Nature Communications

## Advanced integrative computational analysis & AI




# Camonsertib in DNA damage response-deficient advanced solid tumors: phase 1 trial results

Received: 14 October 2022

Accepted: 12 May 2023

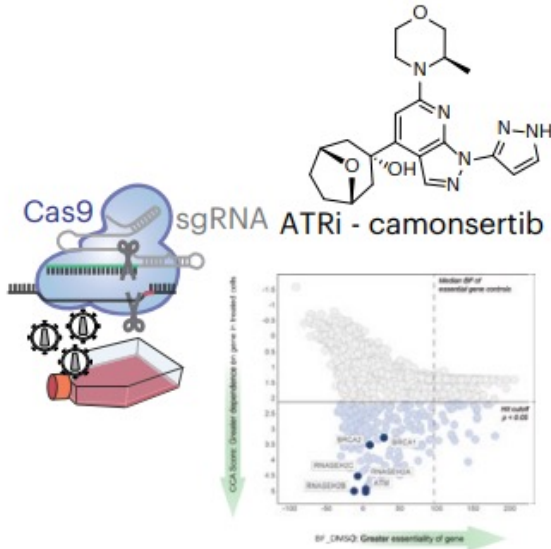
Published online: 5 June 2023

 Check for updates

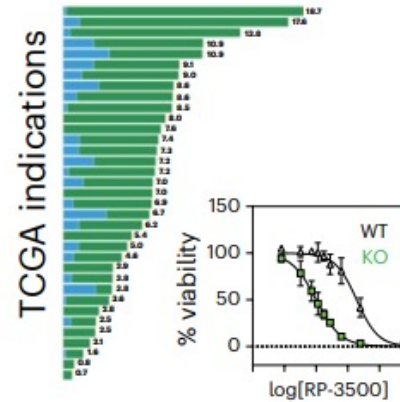
Timothy A. Yap <sup>1</sup>✉, Elisa Fontana<sup>2</sup>, Elizabeth K. Lee <sup>3</sup>, David R. Spigel<sup>4</sup>, Martin Højgaard <sup>5</sup>, Stephanie Lheureux<sup>6</sup>, Niharika B. Mettu<sup>7</sup>, Benedito A. Carneiro<sup>8</sup>, Louise Carter<sup>9</sup>, Ruth Plummer<sup>10</sup>, Gregory M. Cote<sup>11</sup>, Funda Meric-Bernstam <sup>1</sup>, Joseph O'Connell<sup>12</sup>, Joseph D. Schonhoft <sup>12</sup>, Marisa Wainszelbaum<sup>12</sup>, Adrian J. Fretland <sup>12</sup>, Peter Manley<sup>12,15</sup>, Yi Xu<sup>12</sup>, Danielle Ulanet<sup>12</sup>, Victoria Rimkunas<sup>12</sup>, Mike Zinda<sup>12</sup>, Maria Koehler<sup>12</sup>, Ian M. Silverman <sup>12</sup>, Jorge S. Reis-Filho <sup>13</sup> & Ezra Rosen<sup>14</sup>

# SNIPRx CRISPR–Cas9-enabled chemogenomic screen to identify ATRi-sensitizing and synthetic lethal alterations for patient selection

Chemogenomic CRISPR screen to identify synthetic lethal gene LoF



Cancer prevalence, experimental validation and biomarker feasibility

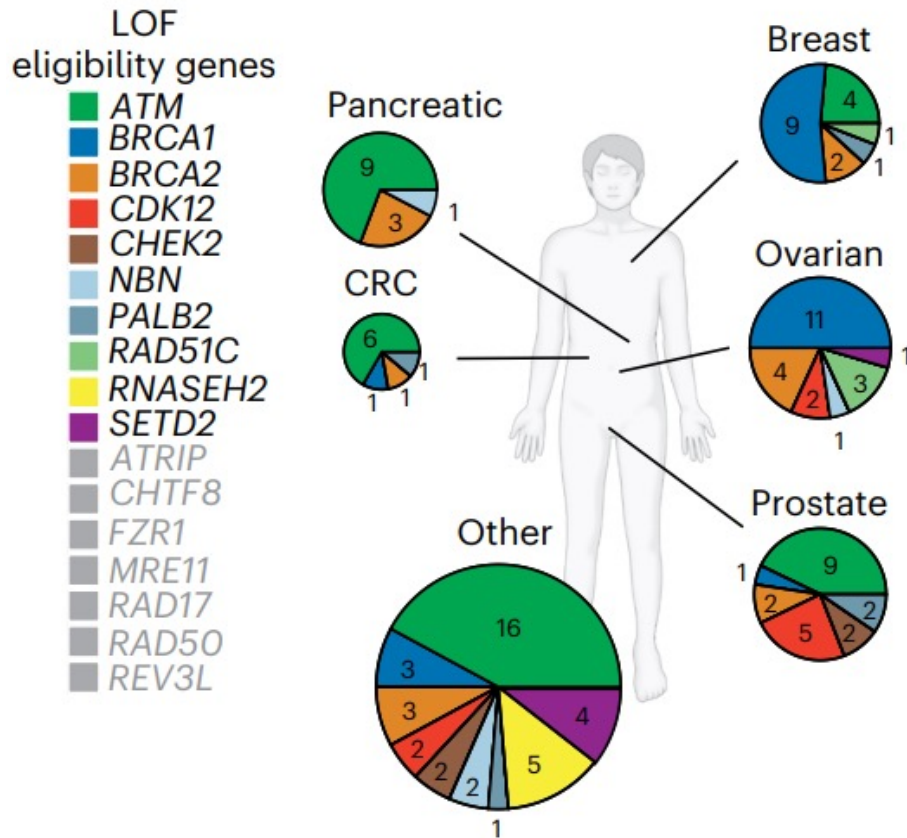


List of prioritized gene alterations

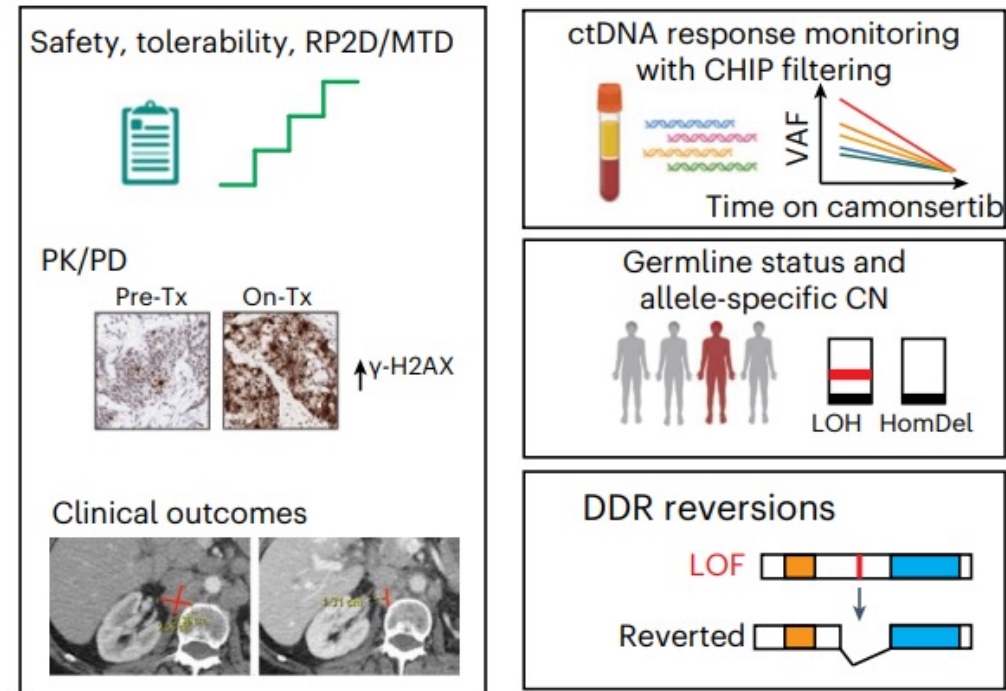
Patients with advanced or metastatic solid tumors with LoF in the following genes:  
*ATM, ATRIP, BRCA1, BRCA2, CDK12, CHTF8, FZR1, MRE11, NBN, PALB2, RAD17, RAD50, RAD51B/C/D, REV3L, RNASEH2A/B or SETD2*

# Patient enrollment by gene and tumor type and overview of pre-planned analyses

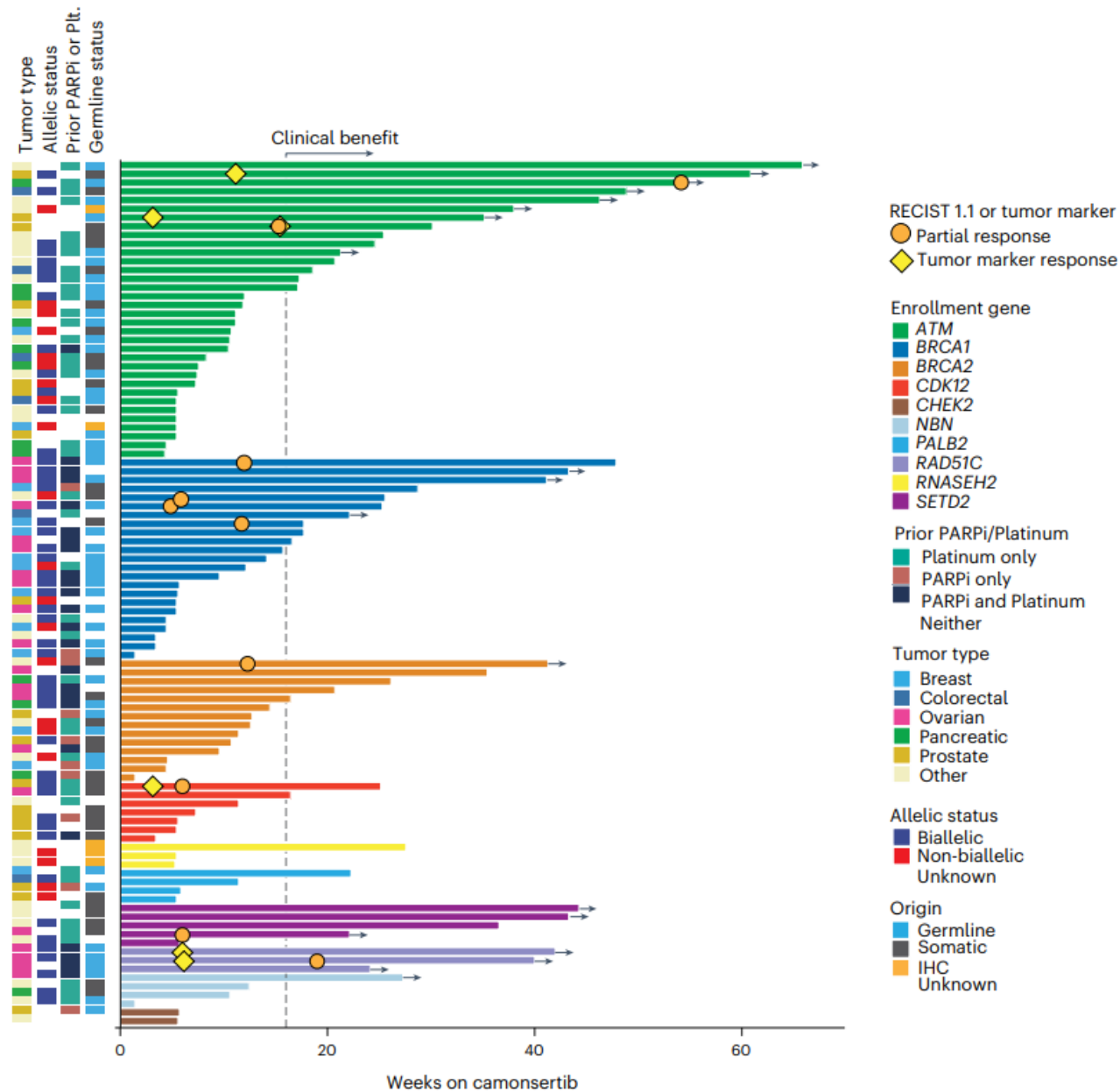
TRESR enrollment for camonsertib monotherapy based on CRISPR screen ( $n = 120$ )



Phase 1 objectives and pre-planned genomic analysis

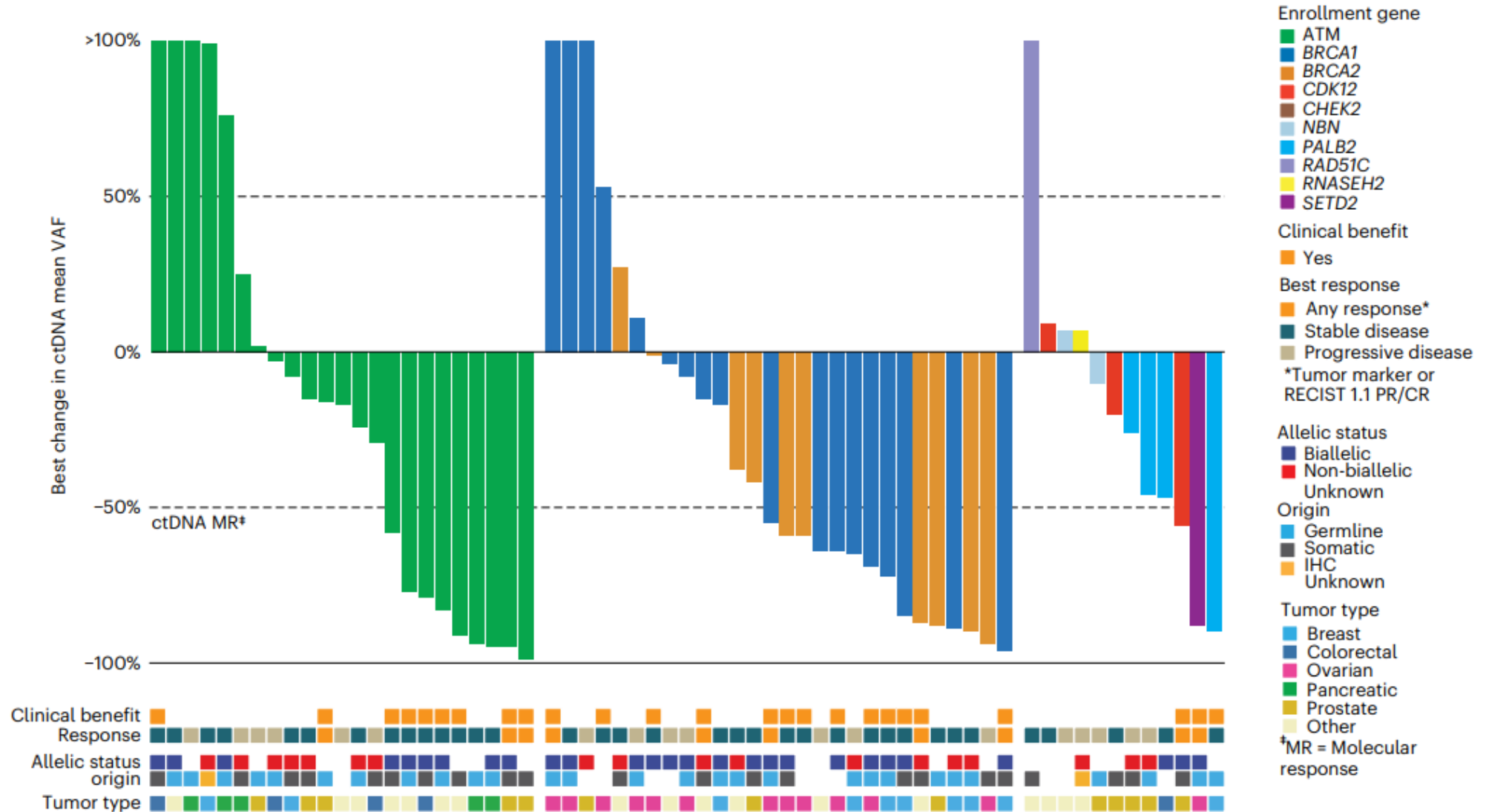


# Clinical outcomes in TRESR: Duration of treatment by genotype



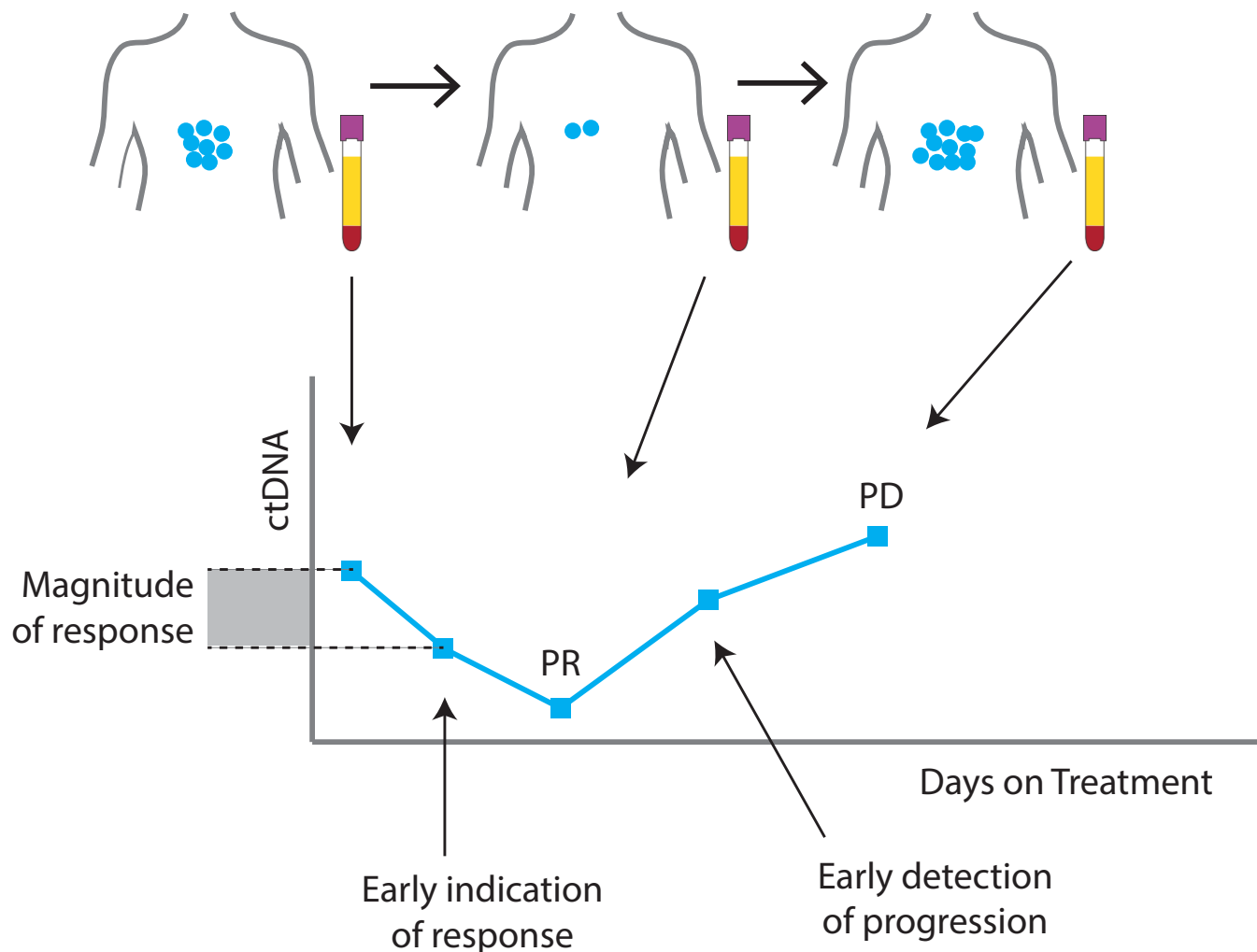
Yap et al. *Nature Medicine*. Vol 29. June 2023

# ctDNA MRs in TRESR: Best ctDNA response by enrollment gene





# Can we monitor metastatic cancer patients more frequently than every two months using a ctDNA blood test?



What would it need to be?

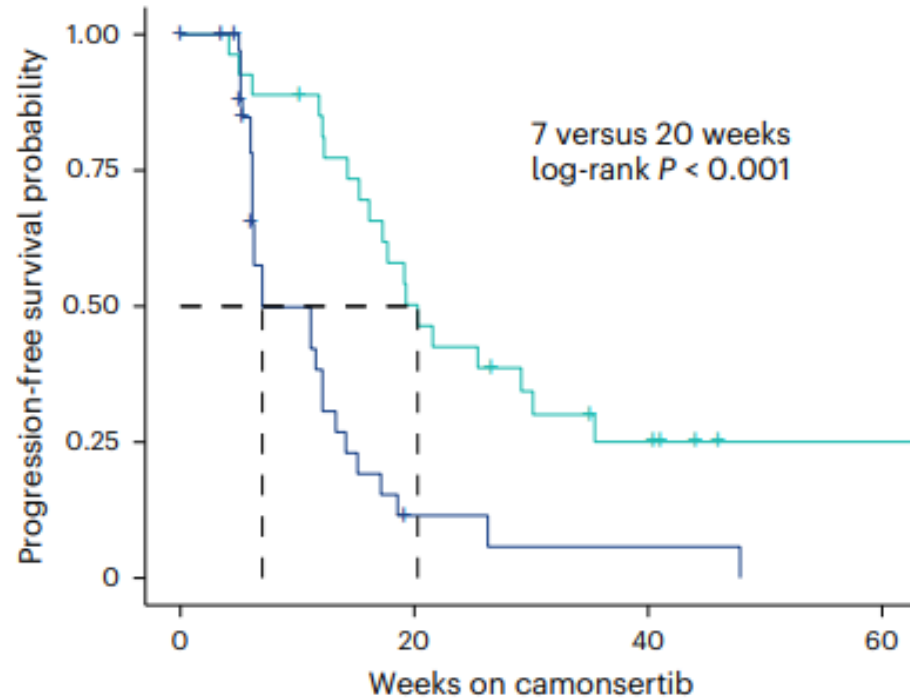
- A pan-cancer treatment monitoring test
- Weekly testing
- Low cost, fast turnaround
- Low sample requirement
- Logistically straightforward

Why do we need it?

- To individualize treatment decisions for each patient in real-time
- To improve clinical trial design and read-out, move away from average response across a group of patients

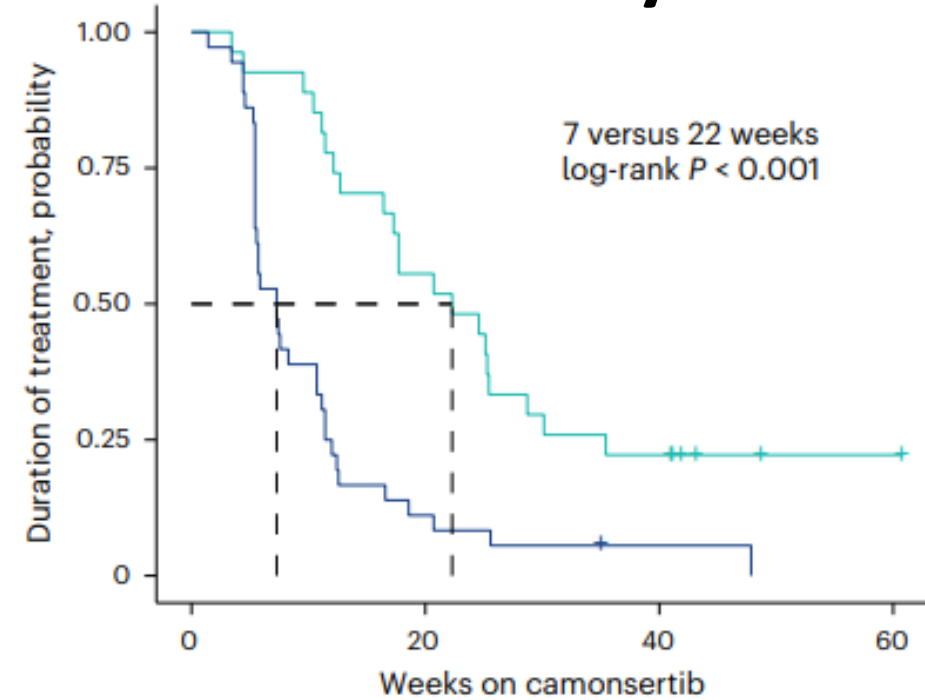
# ctDNA MRs in TRESR: PFS and DOT

## PFS



	0	20	40	60
Number at risk				
ctDNA MR	27	13	5	1
No MR	36	2	1	0

## DOT by MR



	0	20	40	60
Number at risk				
ctDNA MR	27	15	6	1
No MR	36	4	1	0



ORIGINAL ARTICLE

# Single-Agent Divarasil (GDC-6036) in Solid Tumors with a *KRAS* G12C Mutation

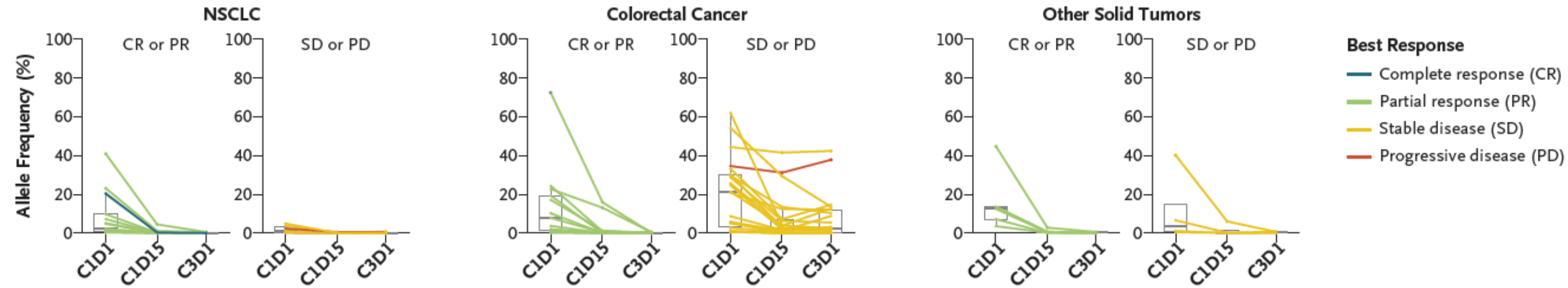
Adrian Sacher, M.D., Patricia LoRusso, D.O., Manish R. Patel, M.D.,  
Wilson H. Miller, Jr., M.D., Ph.D., Elena Garralda, M.D.,  
Martin D. Forster, M.D., Ph.D., Armando Santoro, M.D.,  
Alejandro Falcon, M.D., Tae Won Kim, M.D., Ph.D., Luis Paz-Ares, M.D.,  
Samantha Bowyer, M.B., B.Ch., M.P.H., Maria de Miguel, M.D.,  
Sae-Won Han, M.D., Ph.D., Matthew G. Krebs, M.B., Ch.B., Ph.D.,  
Jong-Seok Lee, M.D., Michael L. Cheng, M.D., Kathryn Arbour, M.D.,  
Erminia Massarelli, M.D., Ph.D., Yoonha Choi, Ph.D., Zhen Shi, Ph.D.,  
Sandhya Mandlekar, Ph.D., Mark T. Lin, M.D., Ph.D., Stephanie Royer-Joo, Engr.,  
Julie Chang, Ph.D., Neekesh V. Dharia, M.D., Ph.D., Jennifer L. Schutzman, M.D., Ph.D.,  
and Jayesh Desai, M.B., B.S., for the GO42144 Investigator and Study Group\*





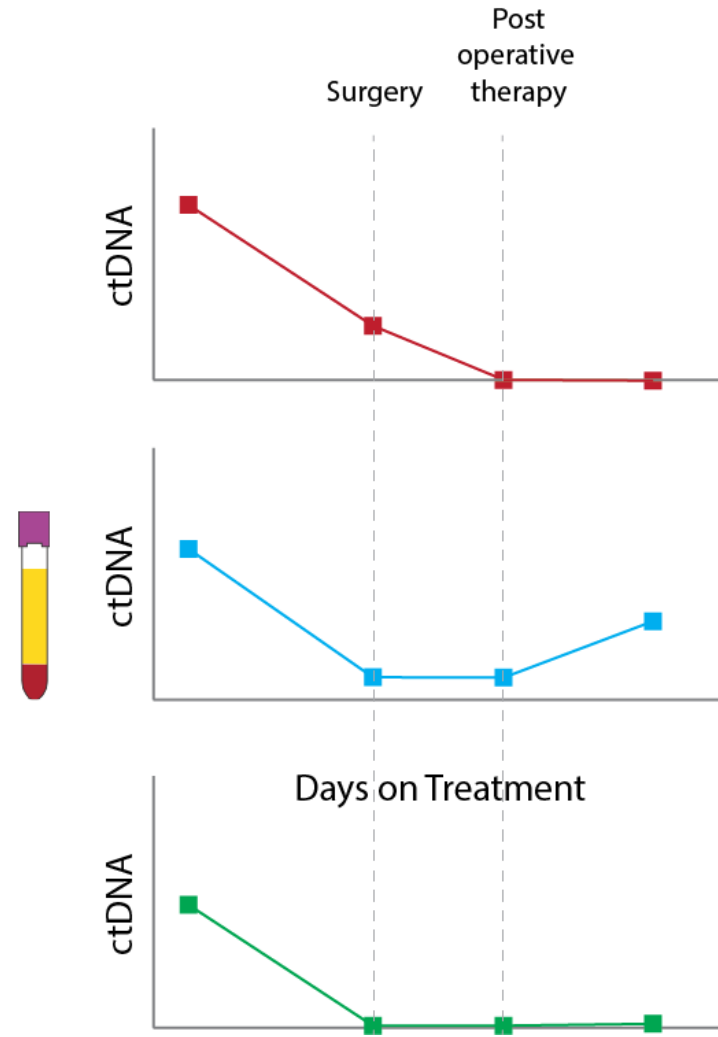
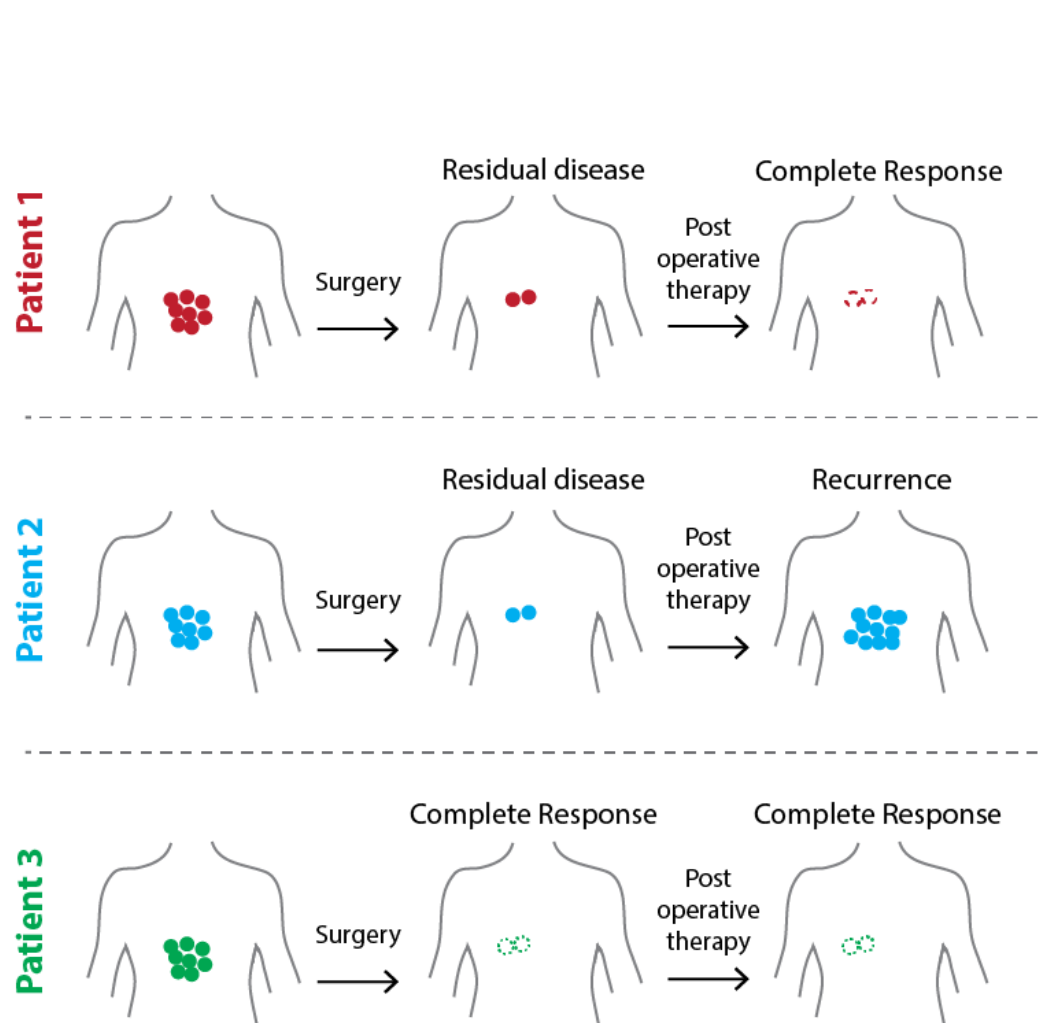
# Divarasilab: Biomarkers of Response and Resistance

## A KRAS G12C Variant Allele Frequency

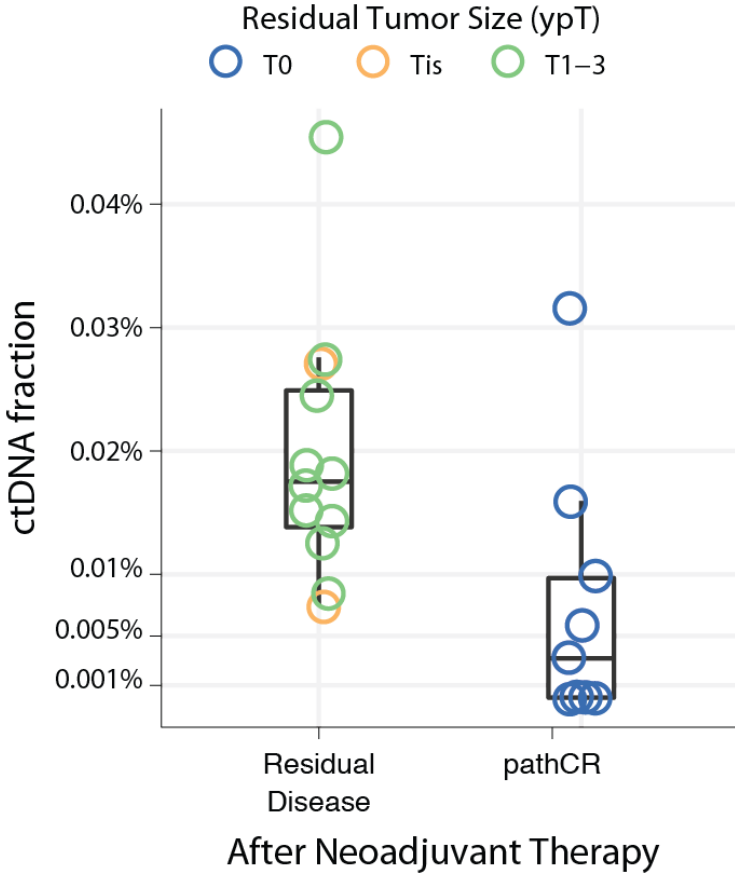
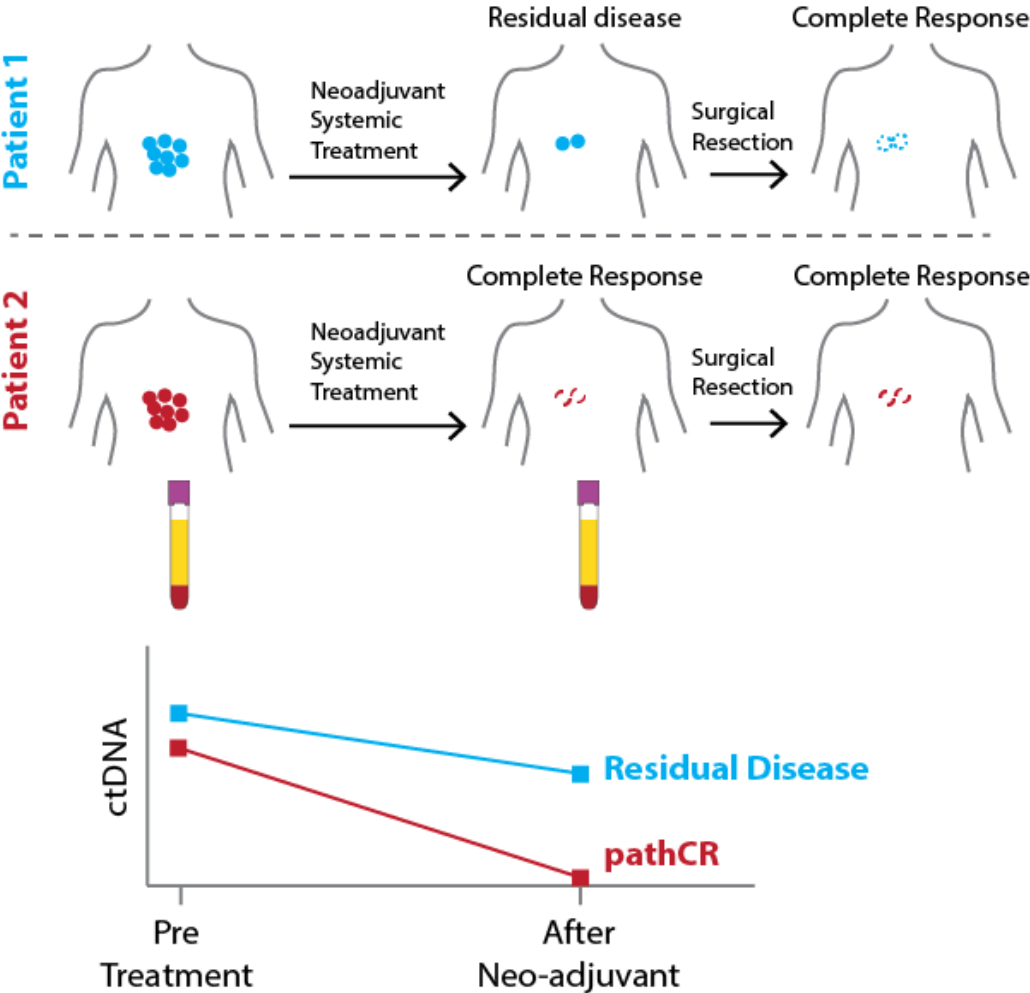




# Can ctDNA analysis help individualize treatment of early-stage cancers by detecting minimal residual disease (MRD)?

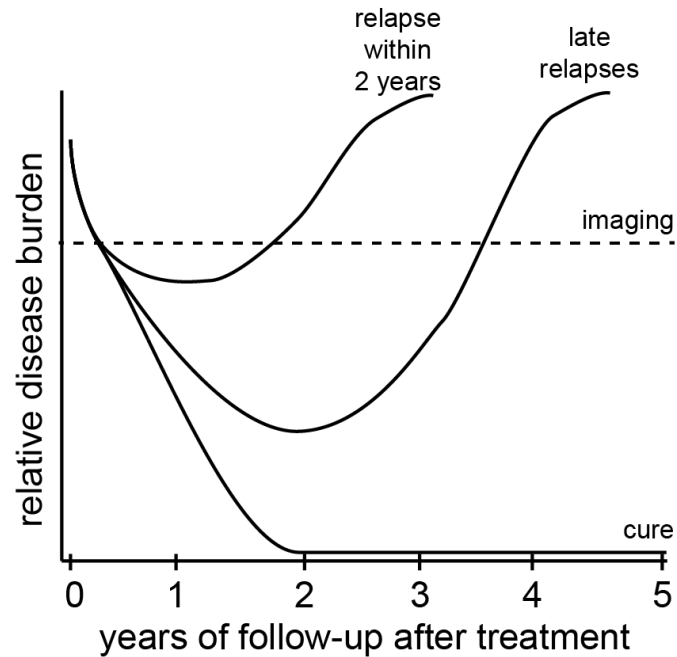
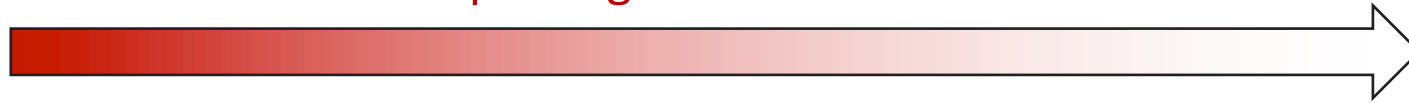


# After neoadjuvant therapy, ctDNA levels were significantly lower in patients with breast cancer who achieved pathCR

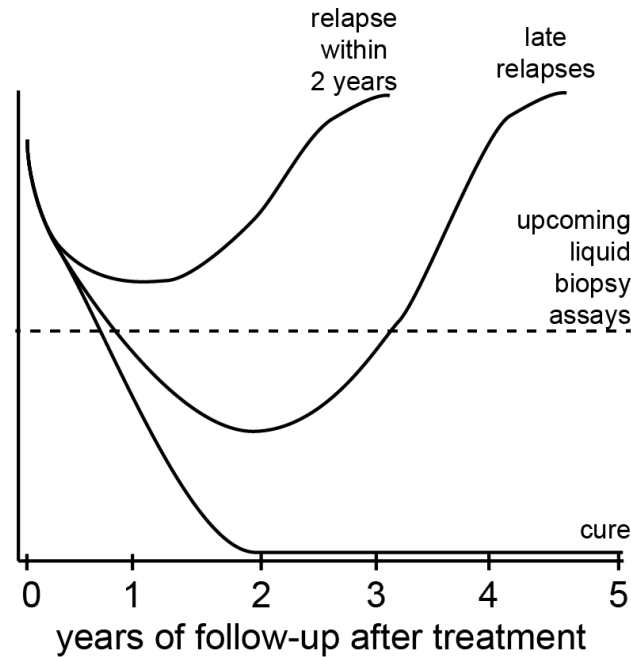


# How sensitive do we need to get for treatment de-escalation studies?

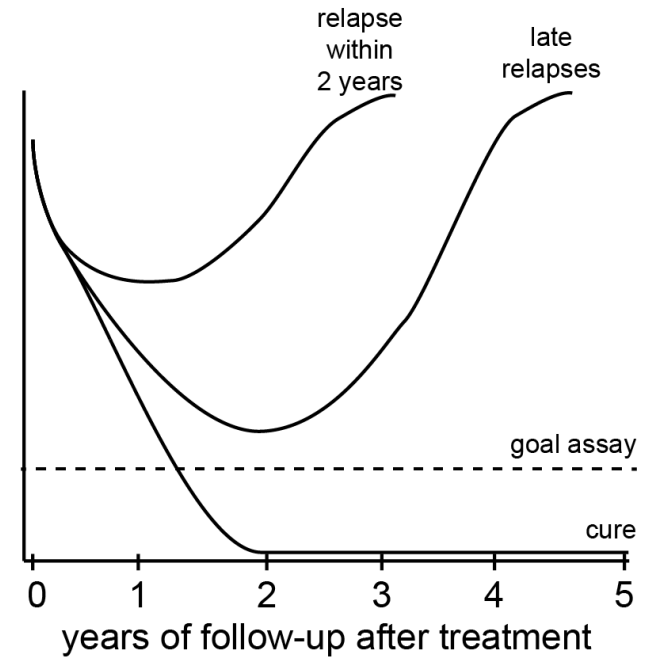
Improving limit of detection



Current clinical practice



Treatment escalation trials  
High Positive Predictive Value

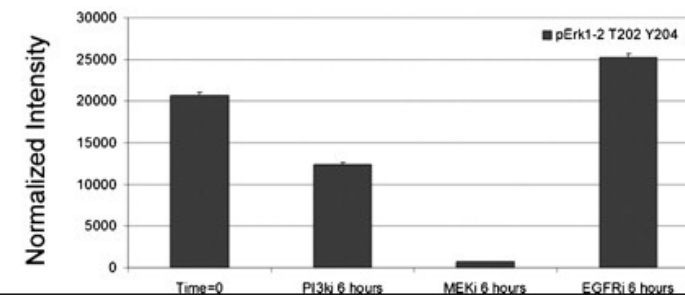
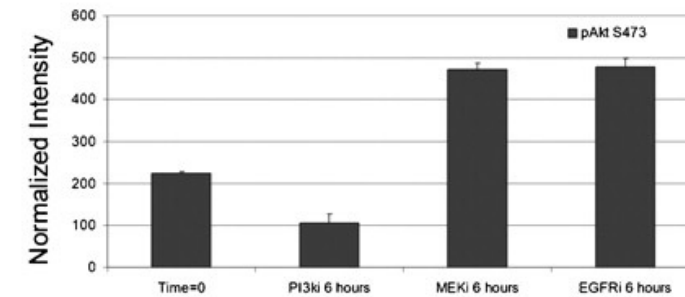
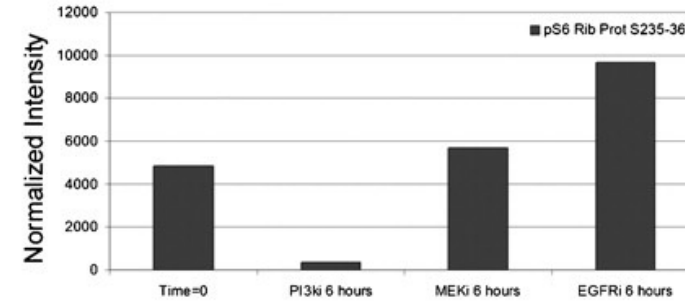
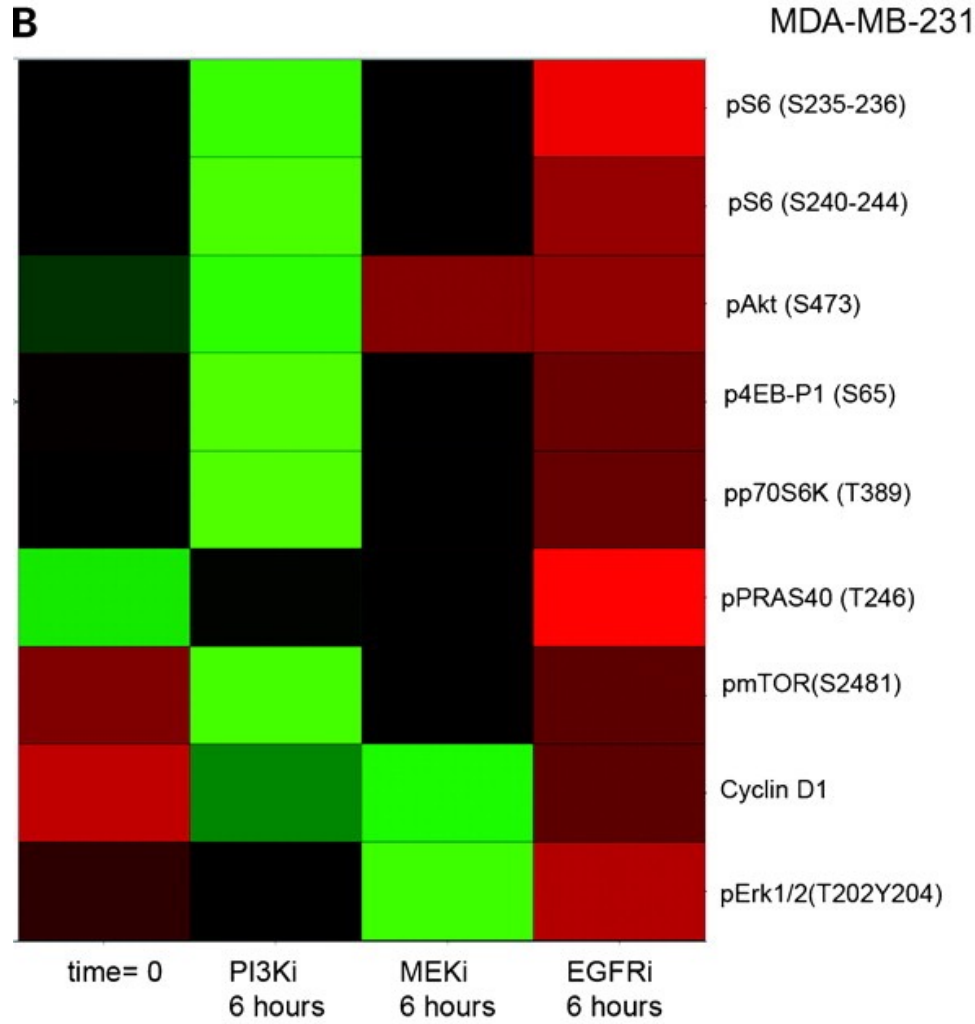


Treatment de-escalation trials  
Need to achieve high Negative Predictive Value



# PhosphoProteomics

# Pharmacodynamic modulation of signaling pathways by targeted kinase inhibitors



# Novel Quantitative HER2 Assay for Determining Dynamic HER2 Expression in the HER2 IHC 0 “Ultra-Low” Setting: Implications for Precision Therapy in HER2- Breast Cancer

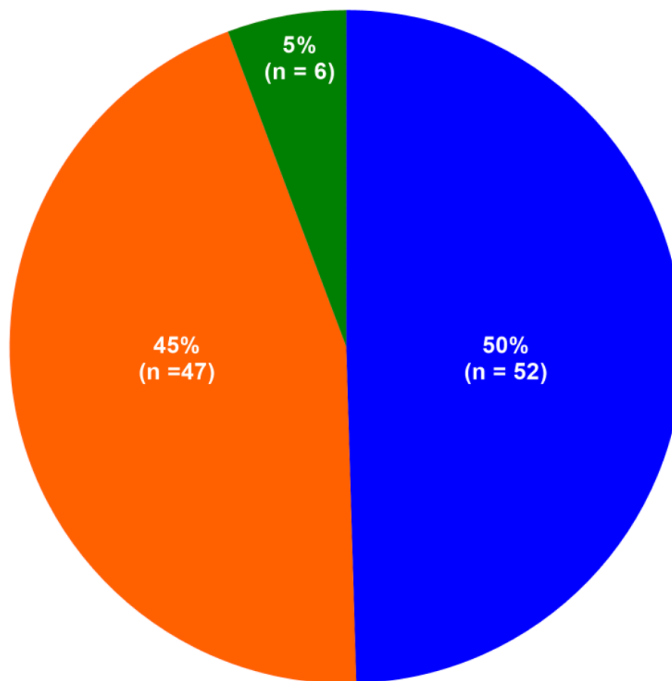
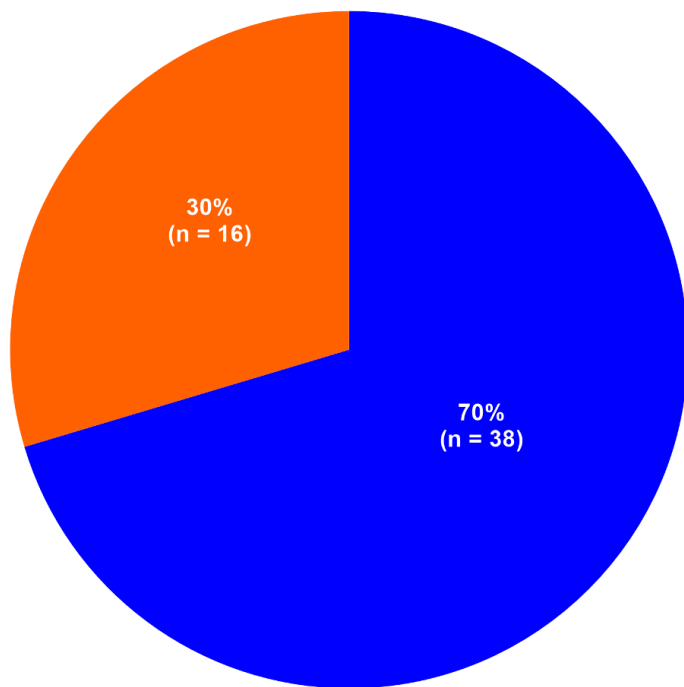
Brian Corgiat<sup>1</sup>, Joyce O’Shaughnessy<sup>2</sup>, Patricia LoRusso<sup>3</sup>, Kris Weinberg<sup>1</sup>, Justin Davis<sup>1</sup>, Chelsea Gawryletz<sup>3</sup>, Emanuel F Petricoin III<sup>4</sup>

Contact: Emanuel Petricoin  
epetrico@gmu.edu

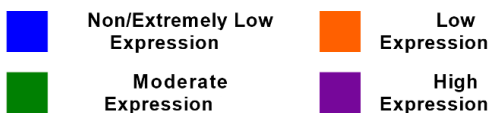
<sup>1</sup>Theralink Technologies, Inc.; <sup>2</sup>Baylor University Medical Center, Texas Oncology, US Oncology, Dallas, TX, <sup>3</sup>Yale Cancer Center, <sup>4</sup>UCHealth- Cancer Care and Hematology, <sup>5</sup>George Mason University

### ER- Cohort (n = 54)

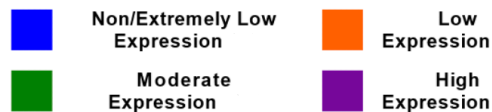
### ER+ Cohort (n = 105)



#### HER2 Score by RPPA



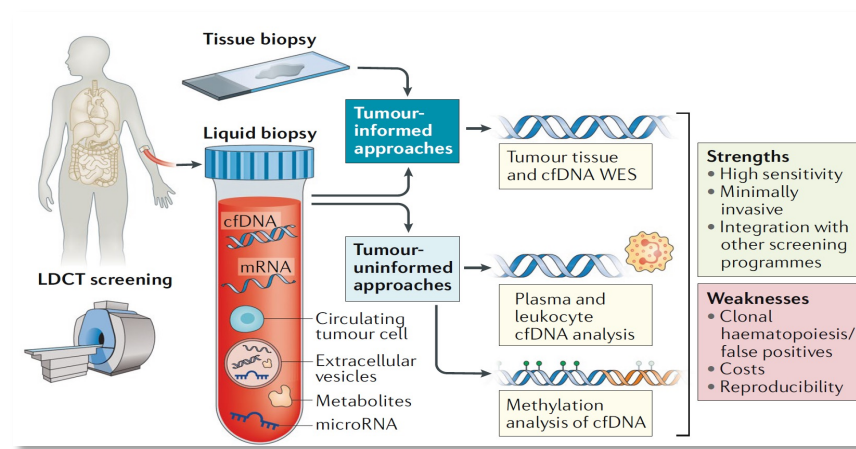
#### HER2 Score by RPPA



## RESULTS:

When evaluated by a quantitative HER2 assay, cases that were defined as HER2 LOW (1+ by IHC or 2+ by IHC and FISH-) actually had very little to no actual HER2 expression in 70% (ER-, N=38) to 50% (ER+, N=52) of time.

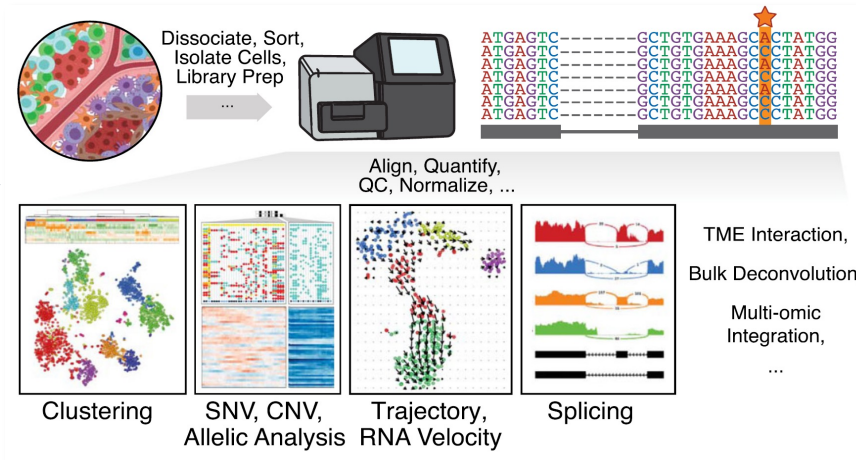
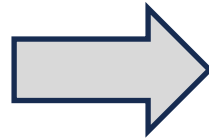
# Liquid biopsy



Rolfo and Russo., 2020. Nature Reviews Clinical Oncology

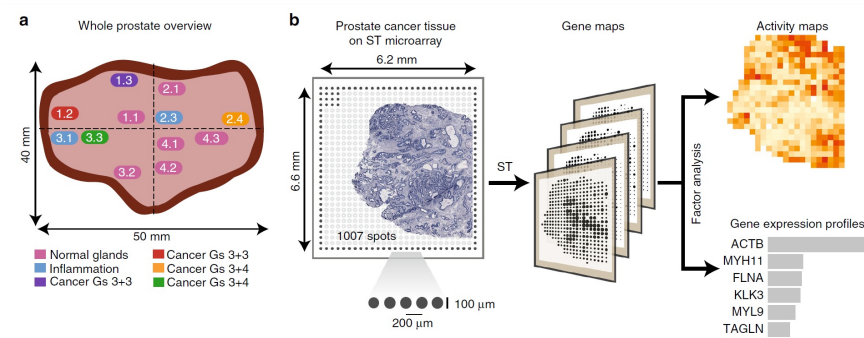
- Strengths**
- High sensitivity
  - Minimally invasive
  - Integration with other screening programmes
- Weaknesses**
- Clonal haematopoiesis/ false positives
  - Costs
  - Reproducibility

# Single-cell transcriptomics



Fan et al., 2020. Experimental & Molecular Medicine volume

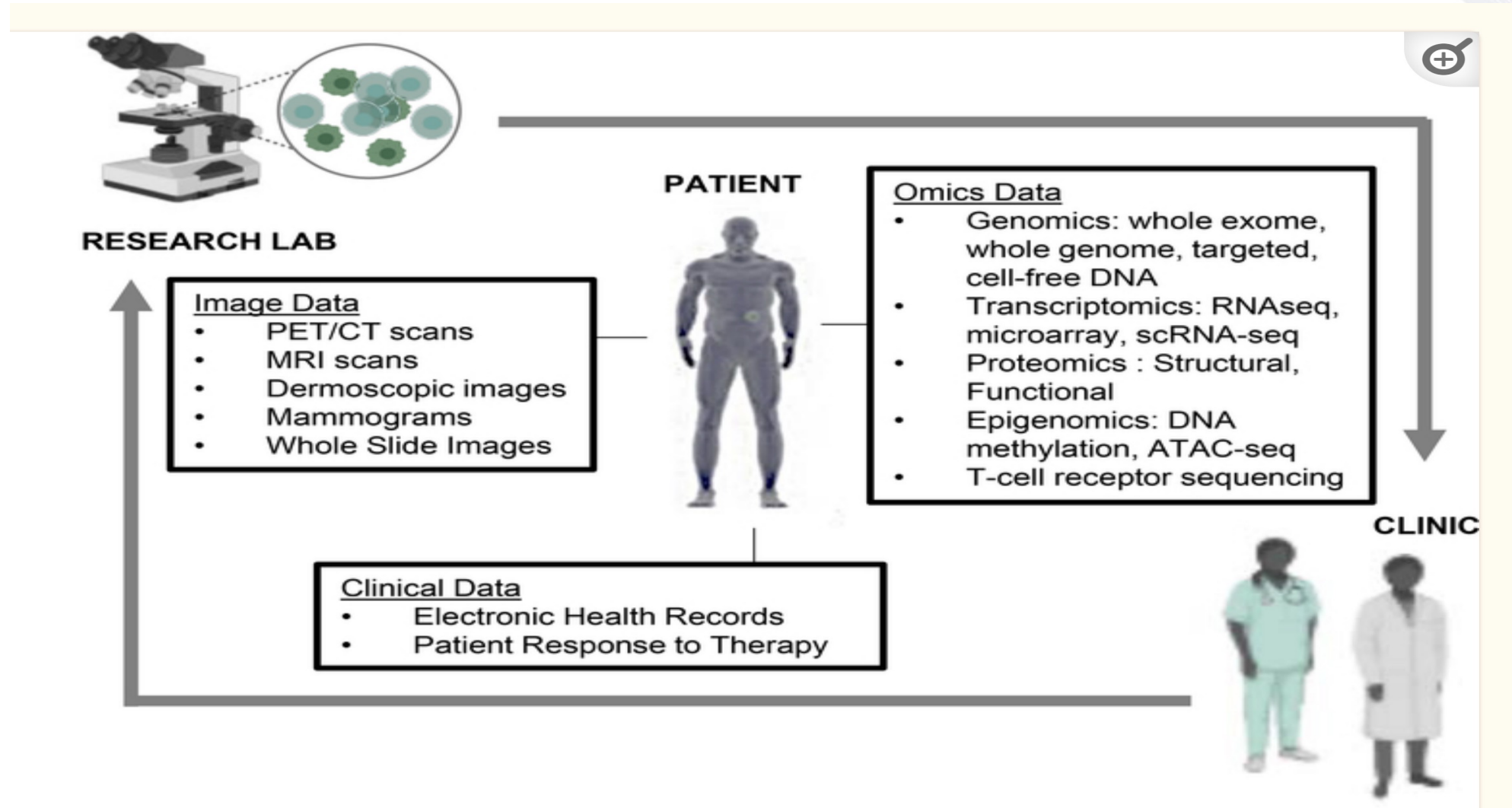
# Spatial transcriptomics & proteomics



Berglund et al., 2020. Nature Communications

## Advanced integrative computational analysis & AI

# Patient Derived Dataset AI models for Bench-to-Bedside Cancer Care



# Conclusions

- Drug discovery & development have come a long way since declaring the War on Cancer
- However, we realize more than ever we have a long way to go
- Exciting times integrating novel technologies into the development of new anticancer agents to unfold the mysteries of cancer biology so as to enhance therapeutic outcomes for patients

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# Thank You!!!