

# ***Combining mechanism and clinical data to explain chemotherapy efficacy***

Eugene F Douglass Jr

October 15, 2021

Assistant Professor

Pharmaceutical and Biomedical Sciences

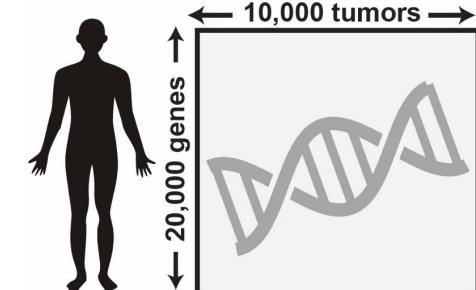
College of Pharmacy

University of Georgia

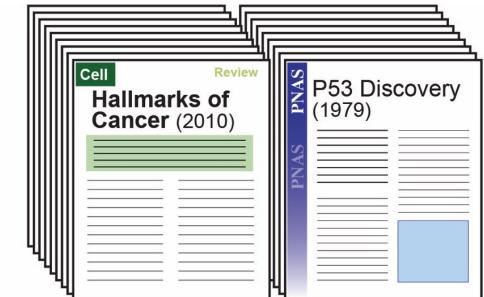
# Clinical “Big Data” + Experiment = Precision Medicine

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**Genomic vs Experimental “Big Data:”** *challenges and opportunities*



Sequencing Tech



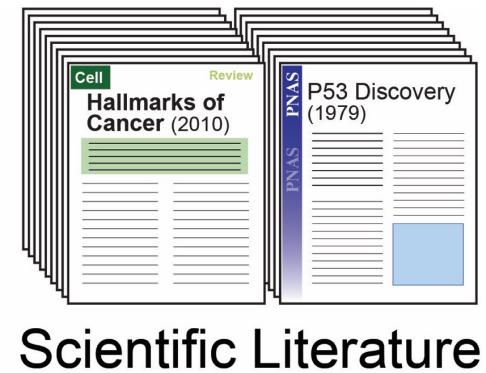
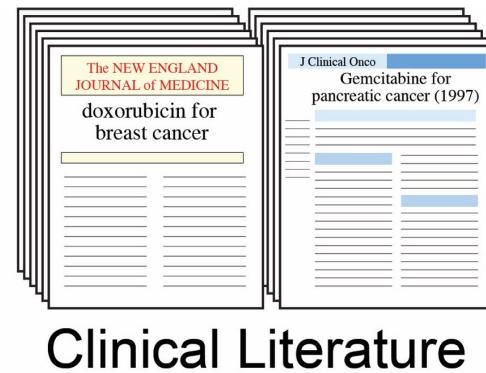
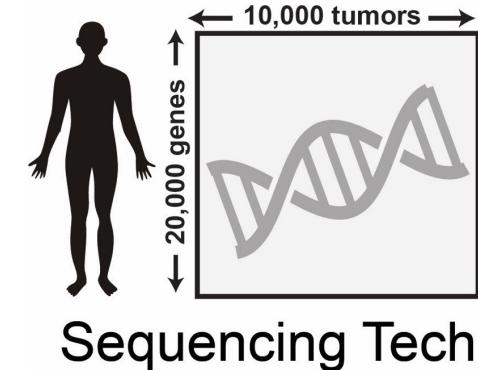
Scientific Literature

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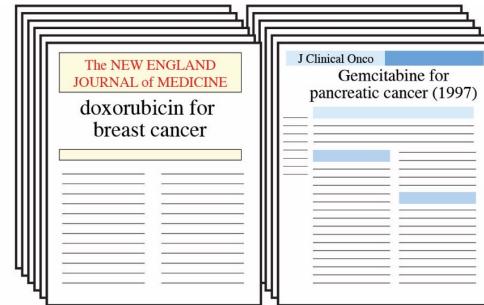
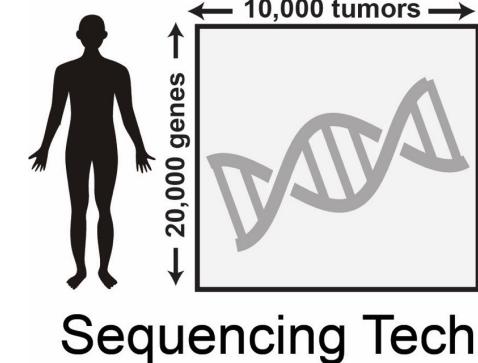
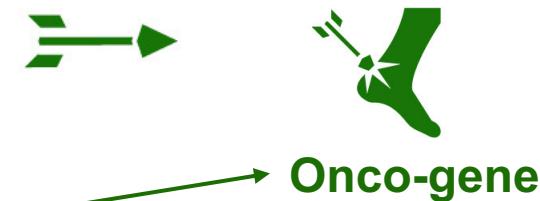
**History & Current Status of Cancer Therapy:** *1 drug / 1 target paradigm*



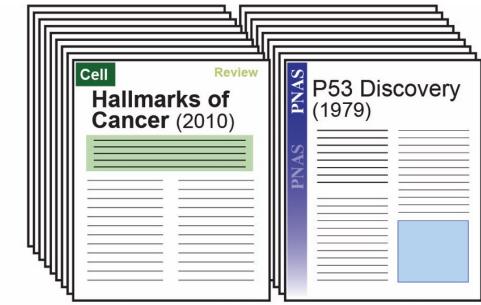
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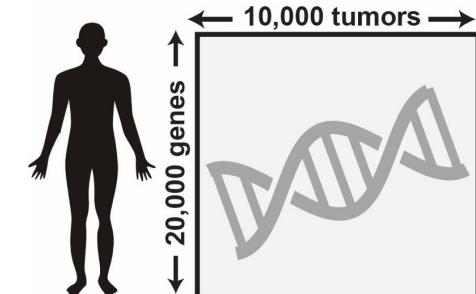
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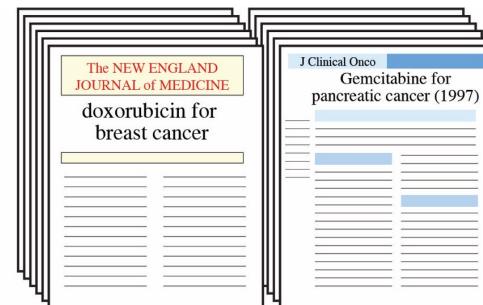
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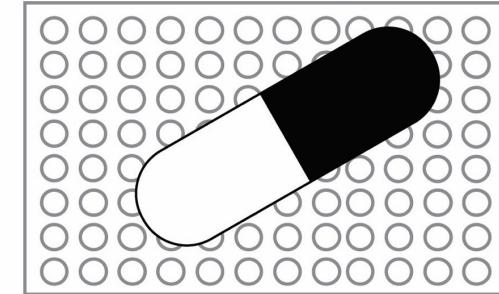
**Reconciling Clinical & Laboratory “Big Data”:** *1 drug / N-biomarkers*



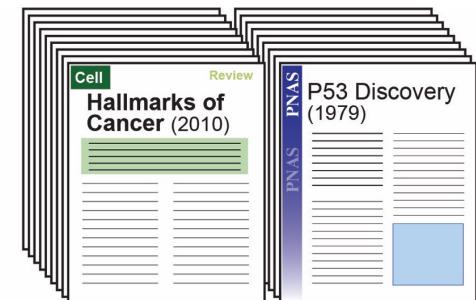
Sequencing Tech



Clinical Literature



Screening Technology



Scientific Literature

# Clinical “Big Data” + Experiment = Precision Medicine

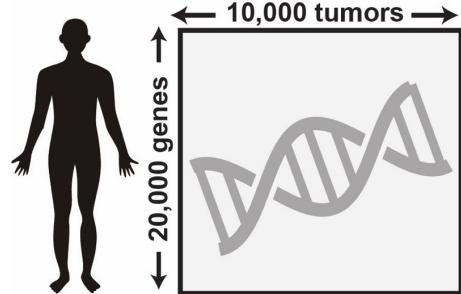
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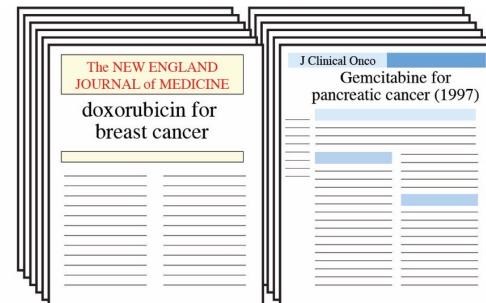
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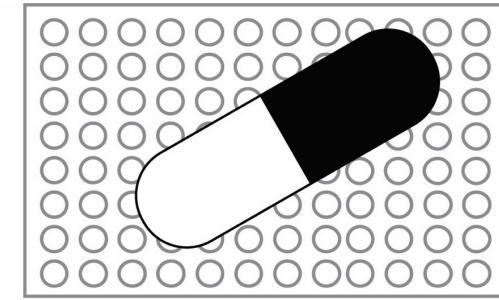
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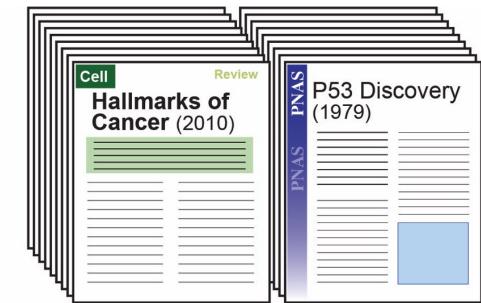
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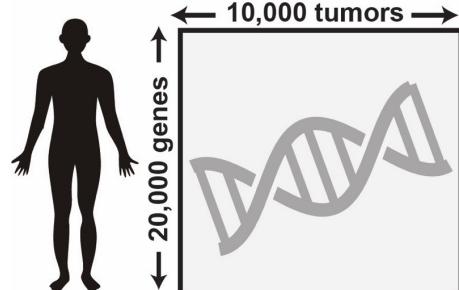
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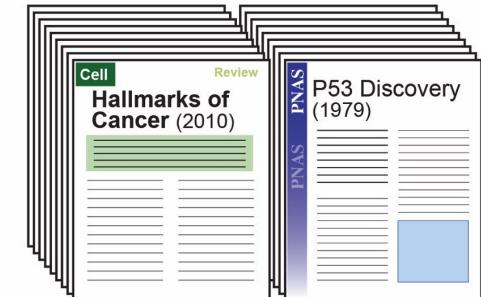
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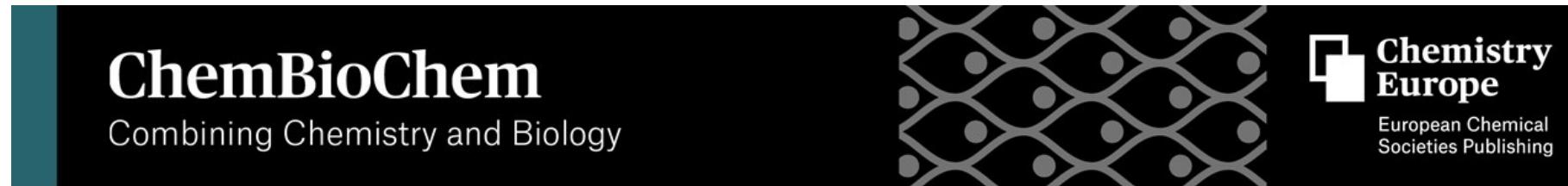


Sequencing Tech



Scientific Literature

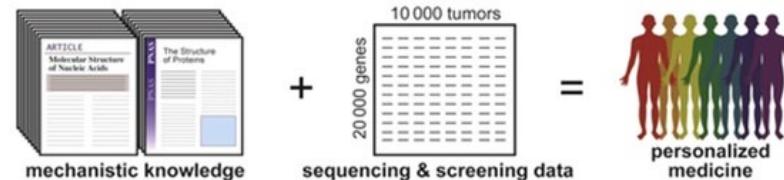
# Sequencing/Screening vs Experimental “Big Data”



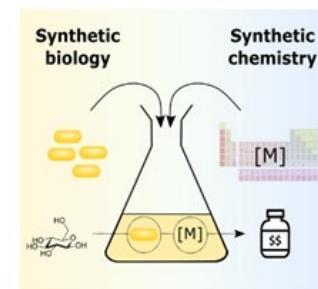
## Chemical Translational Biology – Early-Career Award Winners



Eugene Douglass  
University of Georgia  
Bridging “Big Data” and Mechanistic  
Insight To Enable Precision Medicine



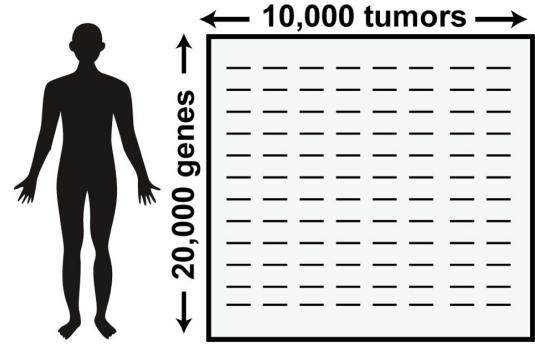
Joanna Sadler  
University of Edinburgh  
The Bipartisan Future of Synthetic  
Chemistry and Synthetic Biology



# Sequencing/Screening vs Experimental “Big Data”

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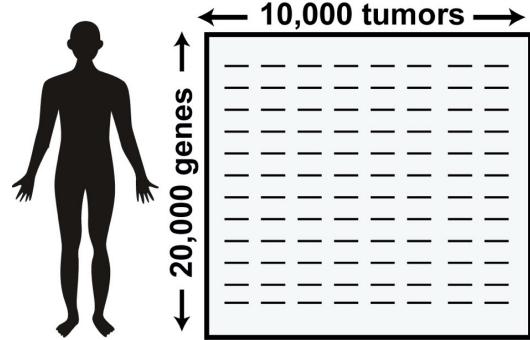
## Clinical Tumor Sequencing “Big Data”



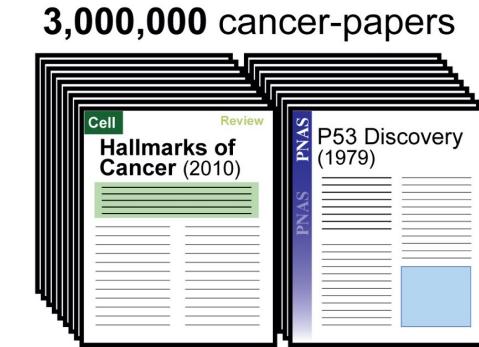
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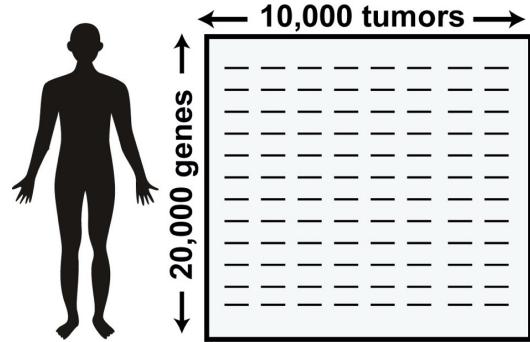


## Experimental Literature “Big Data”

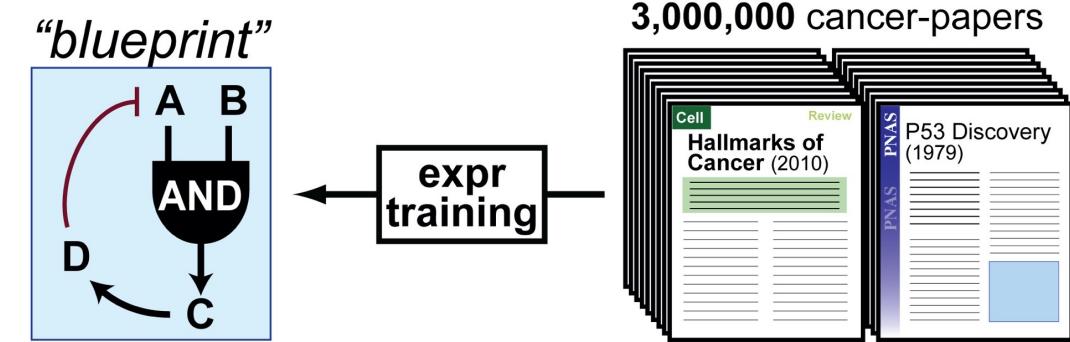


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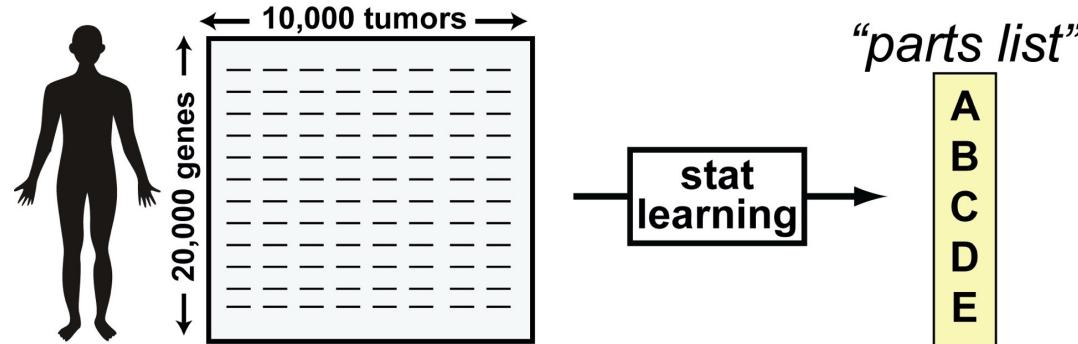


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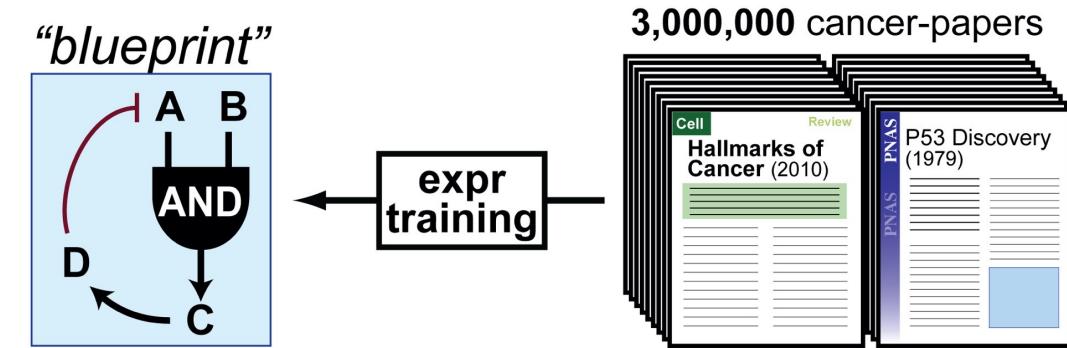


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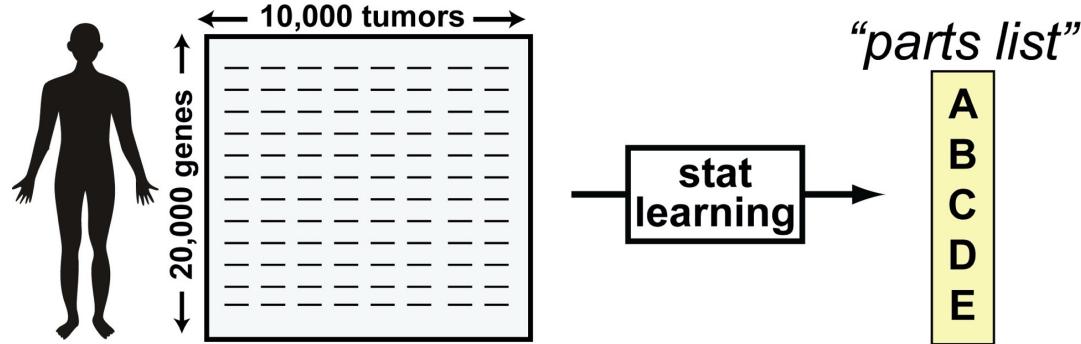


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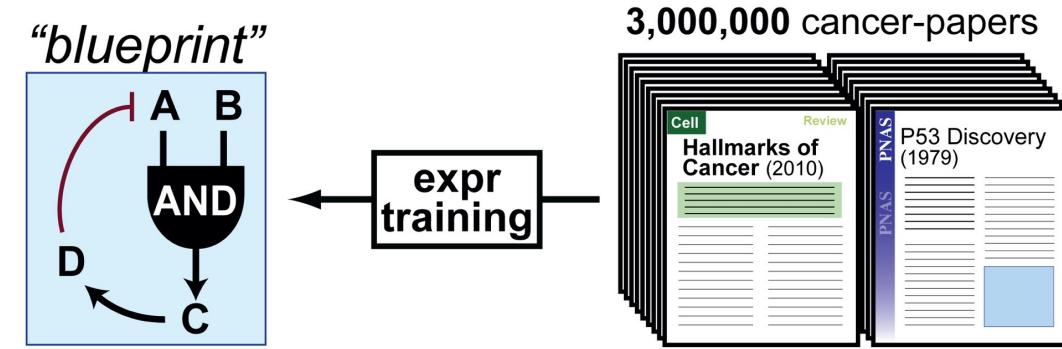


# Communication Barriers prevent Big Data Synergy

## Clinical Tumor Sequencing “Big Data”

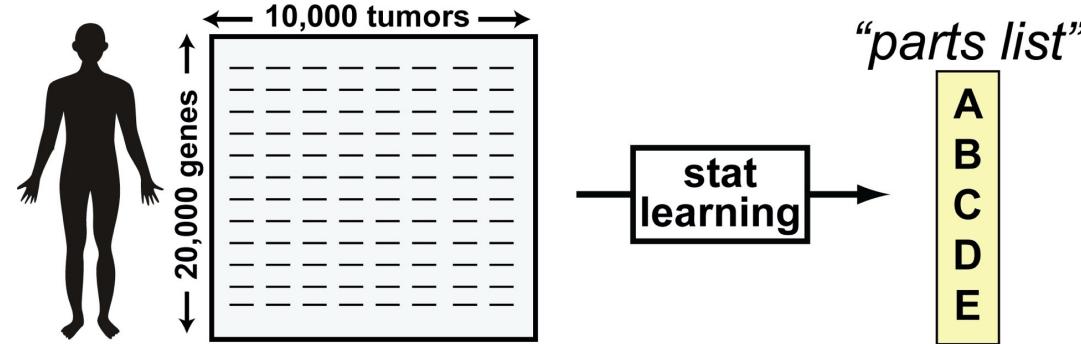


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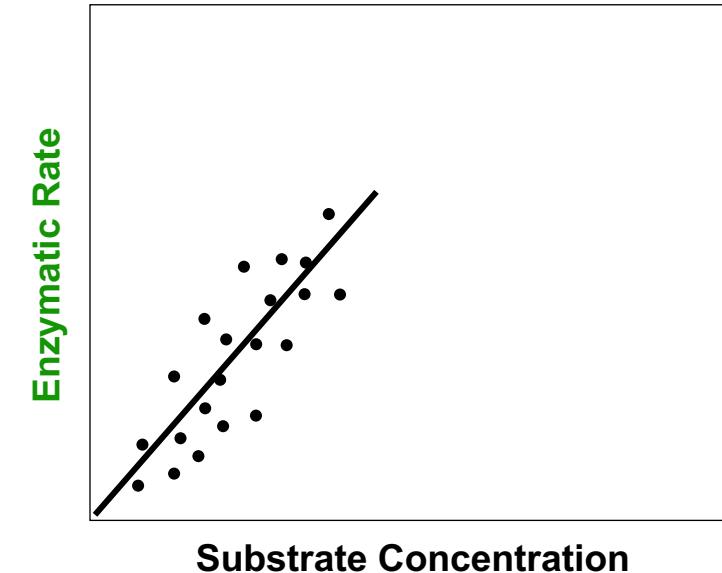
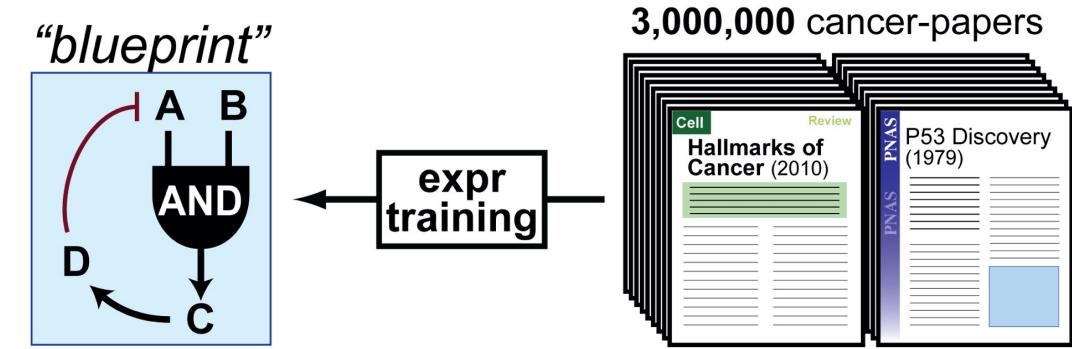
# Different Definitions of Proof: Correlation vs Causation

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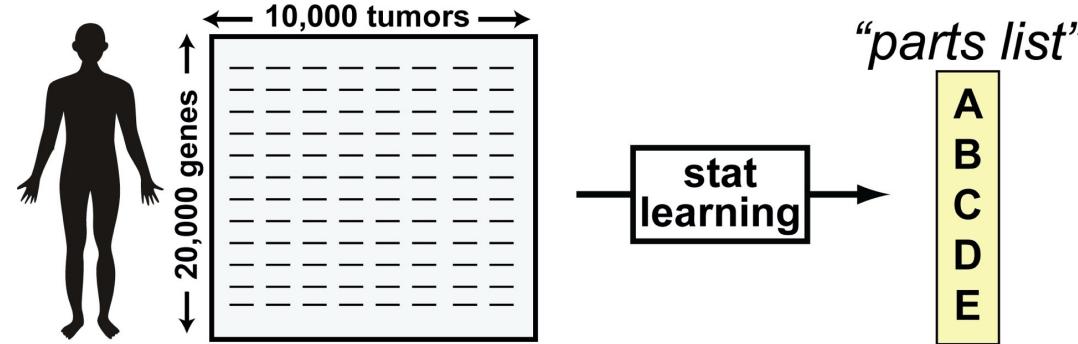
jargon

## Experimental Literature “Big Data”



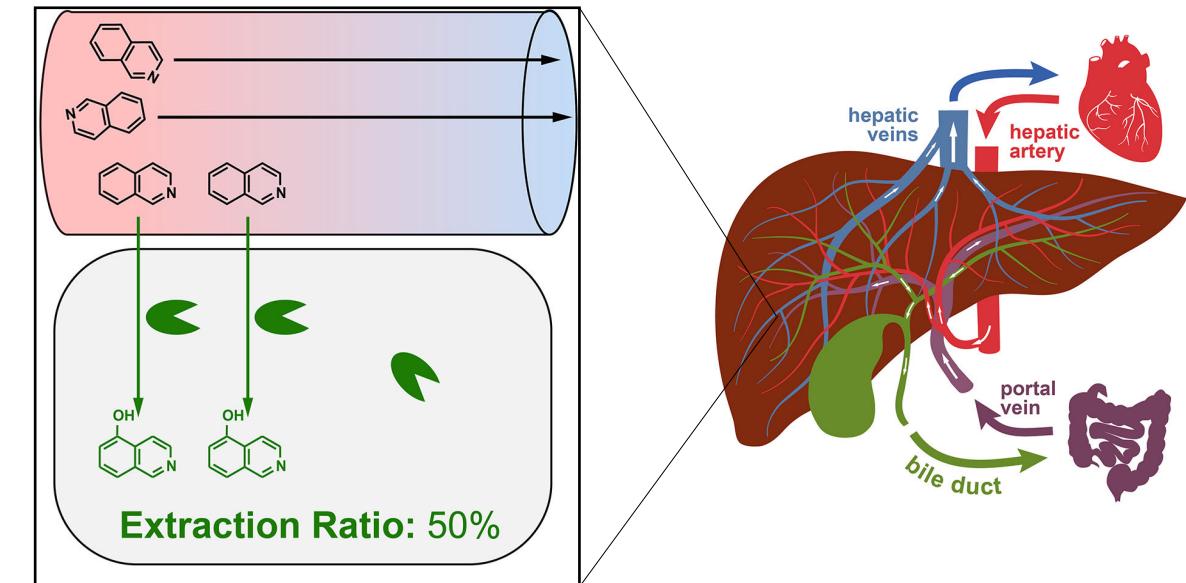
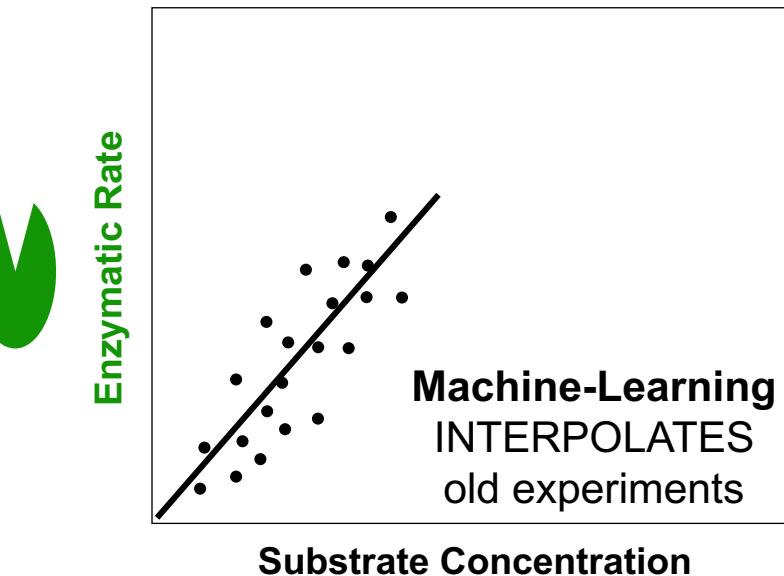
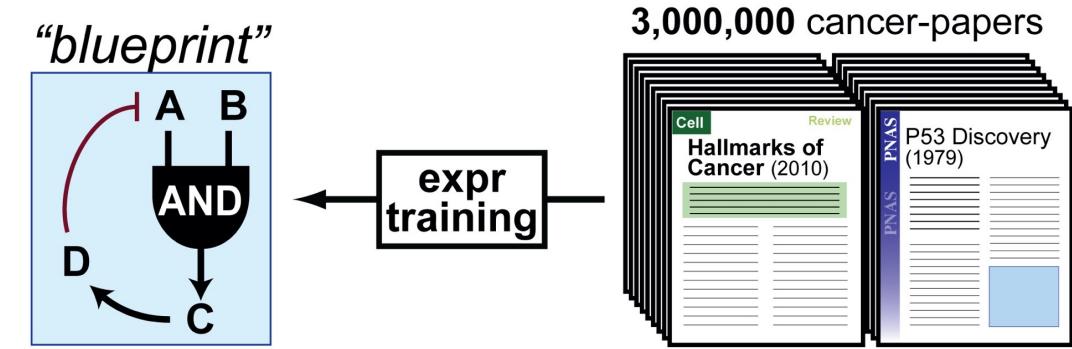
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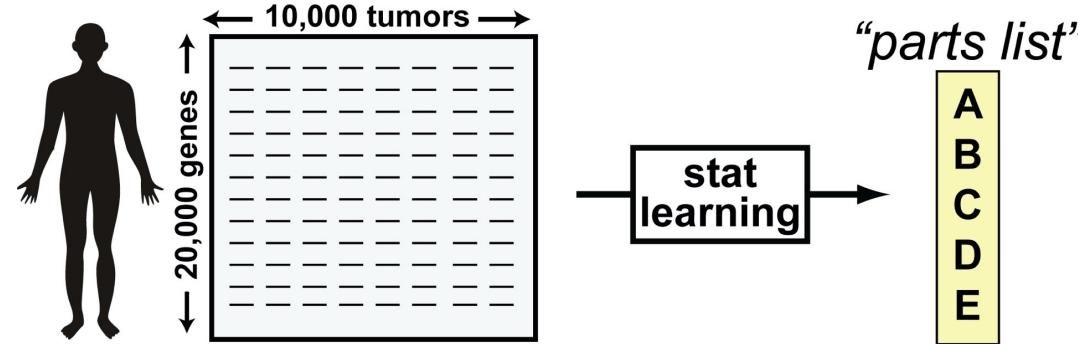
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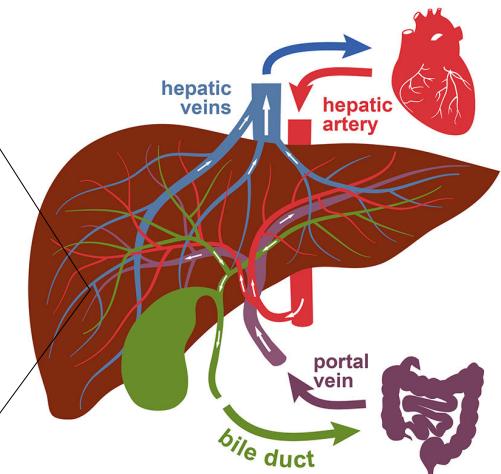
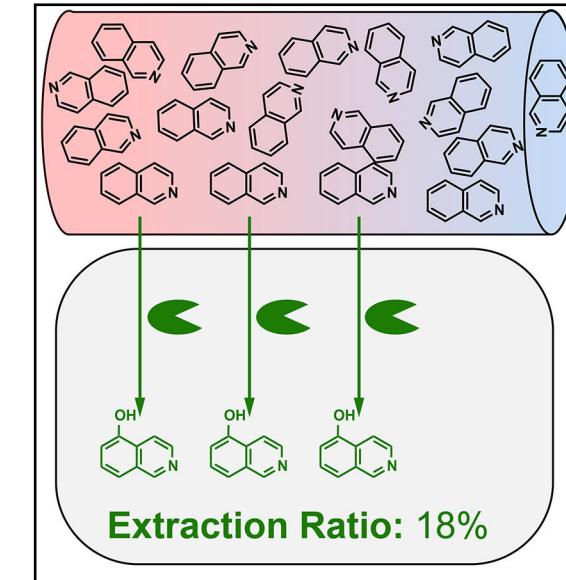
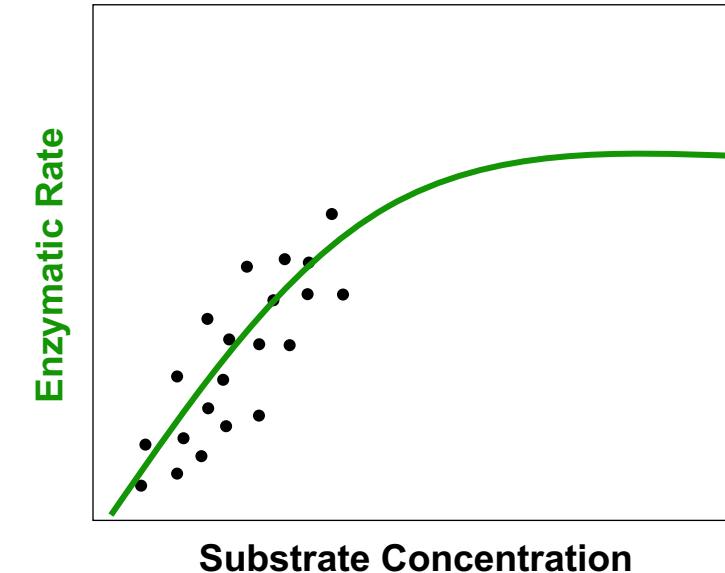
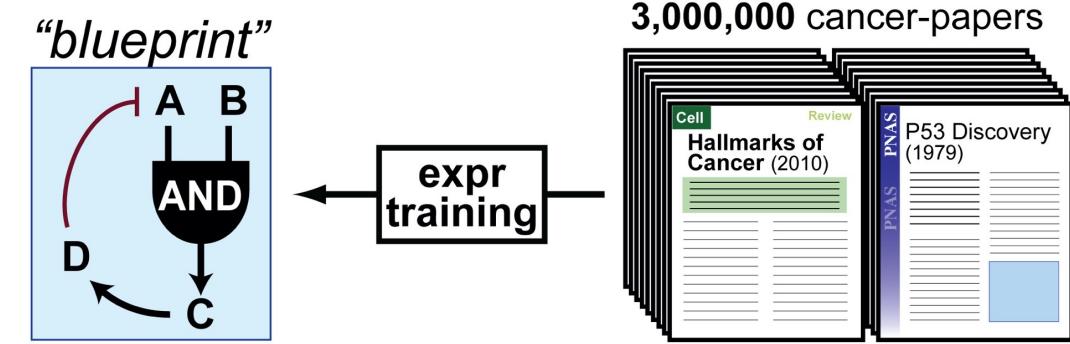
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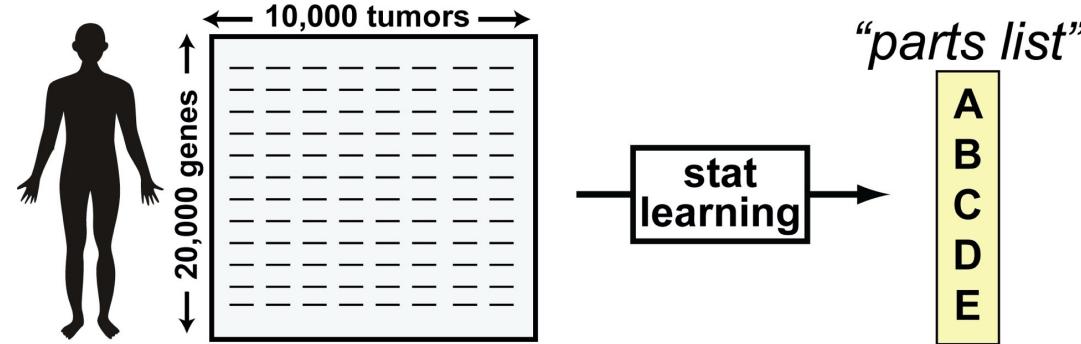
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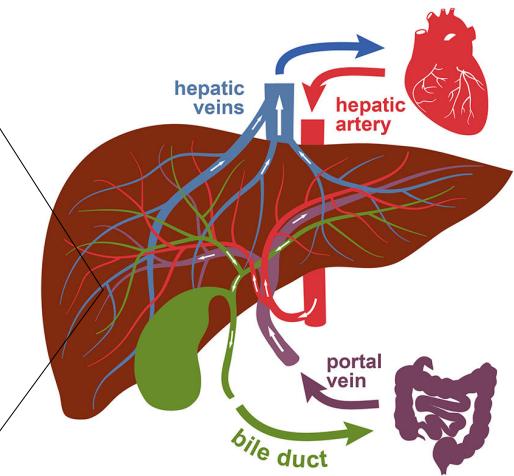
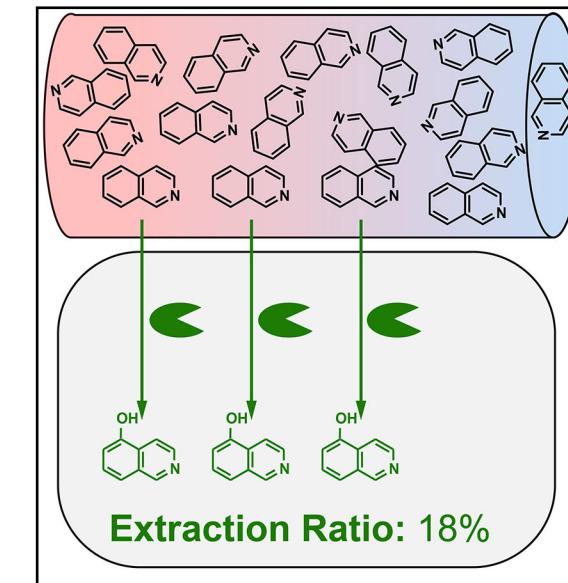
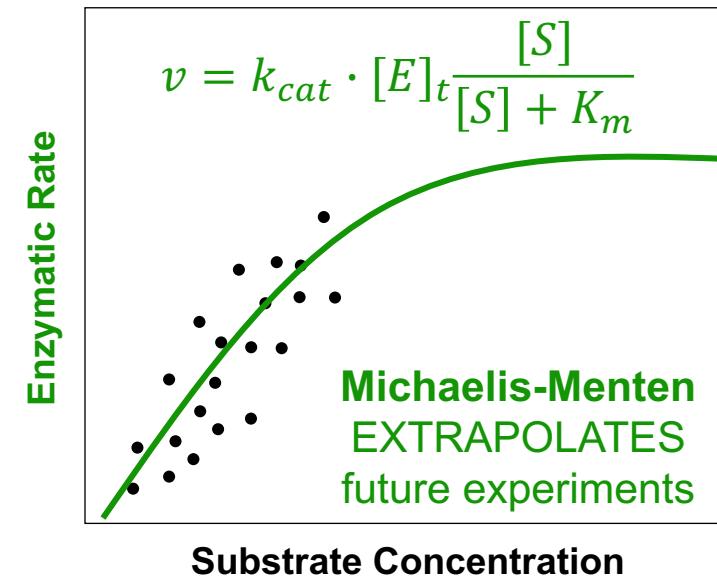
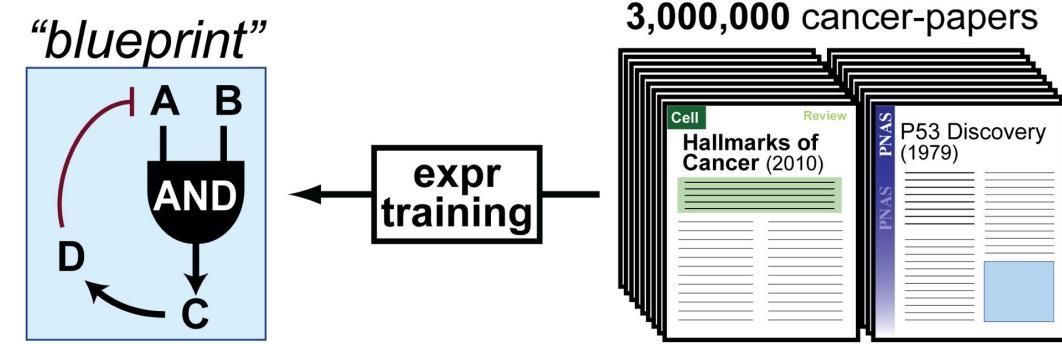
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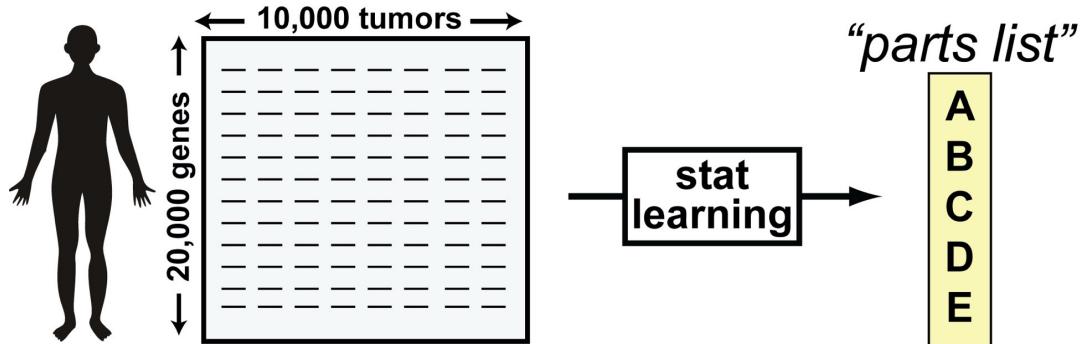
jargon

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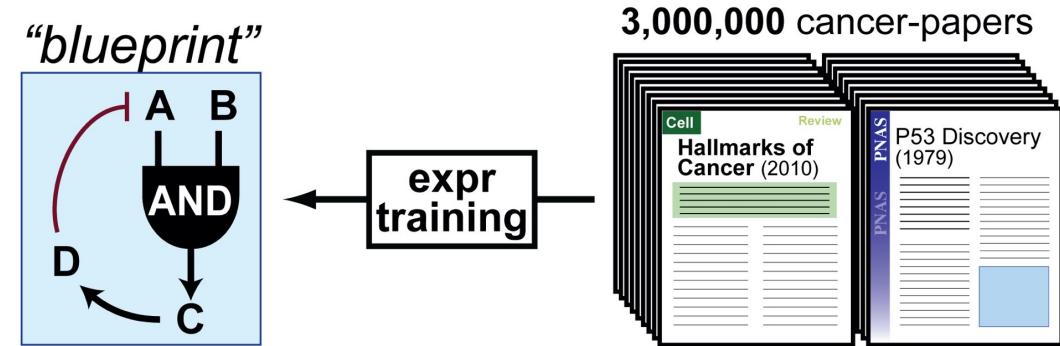


# Precedents for “Big Data” driving paradigm shifts

## Clinical Tumor Sequencing “Big Data”



## Experimental Literature “Big Data”



**Spectroscopy:**  
Black Body Radiation

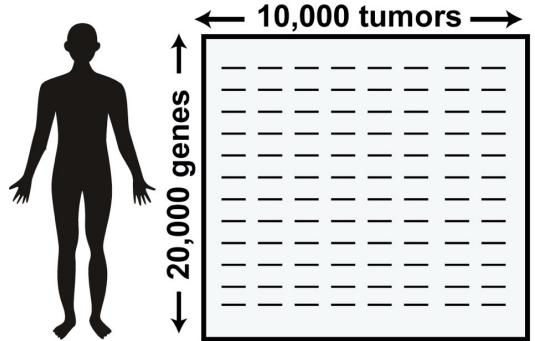
**Empirical Models:**  
Wien Approximation (1896)

**First Principles:**  
Plank's Law (1901)

**New Paradigm:**  
Quantum Mechanics

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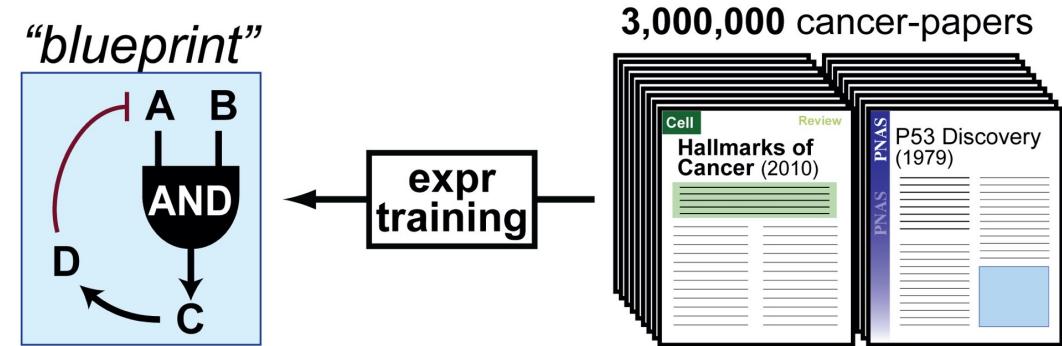


“parts list”

A  
B  
C  
D  
E

jargon

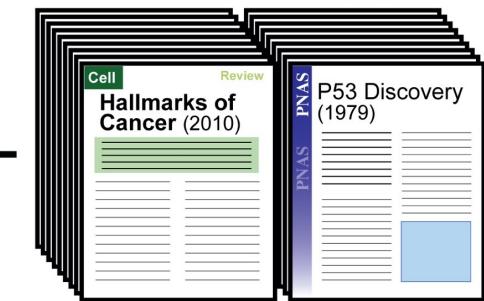
## Experimental Literature “Big Data”



“blueprint”

A  
B  
AND  
D  
C  
expr  
training

3,000,000 cancer-papers



**New Big Data:**  
Black Body Radiation

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Wien Approximation (1896)

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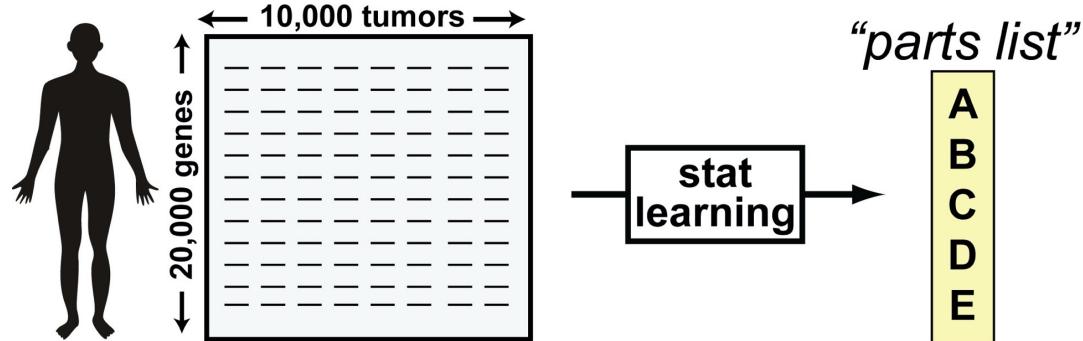
**New Paradigm:**  
Quantum Mechanics

*“The year was 1903...It was an ideal moment for an aspiring young man to enter the field. Half a century of laboratory research had generated an unparalleled backlog of data that demanded understanding.”*

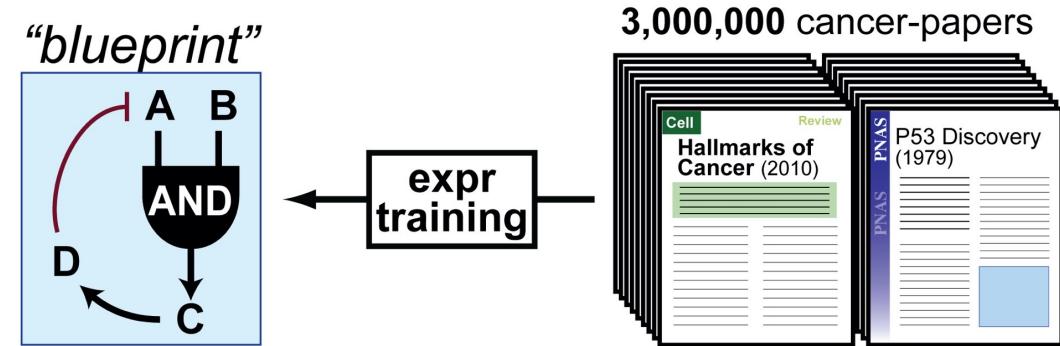
- Niels Bohr's Times, Clarendon Press 1991

# Precedents for “Big Data” driving paradigm shifts

## Clinical Tumor Sequencing “Big Data”



## Experimental Literature “Big Data”



**New Big Data:**  
Spectroscopy

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Wien Approximation (1896)

**First Principles:**  
Plank's Law (1901)

**New Paradigm:**  
Quantum Mechanics

**New Big Data:**  
Tumor genomics

**Empirical Models:**  
Machine learning

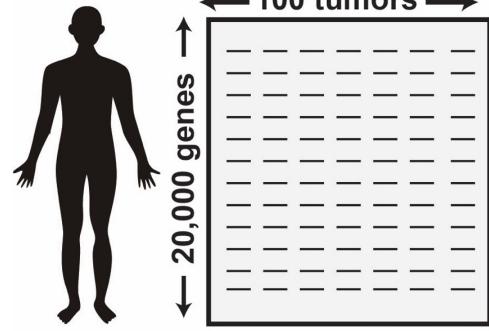
**First Principles:**  
??

**New Paradigm:**  
Precision Medicine

***Doctoral Student Educational Approach:  
computational scientists***

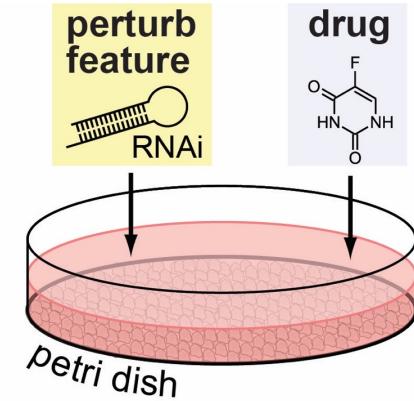
# *Student Training Experience: Columbia University*

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Computational Laboratory:  
***skeleton crew***

Experimental Laboratory:  
***fully staffed***



# *Student Training Experience: Columbia University*

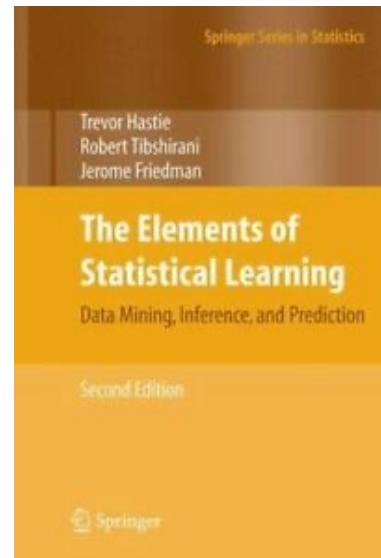
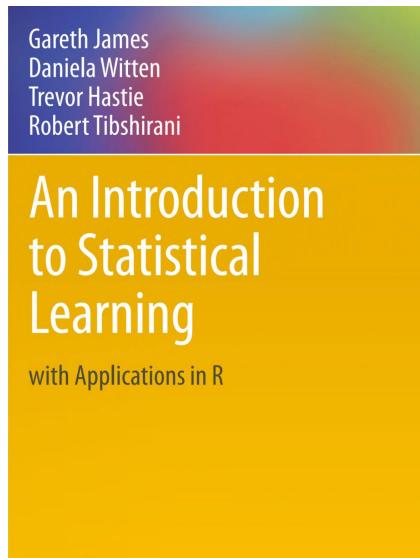
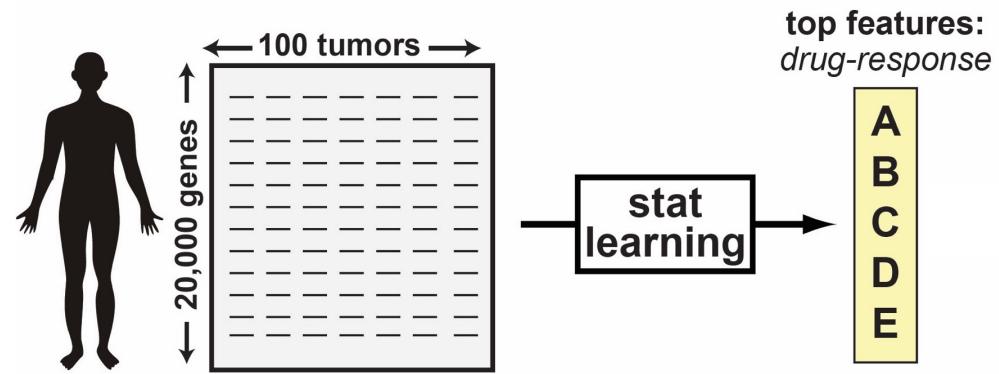


Computational Laboratory:  
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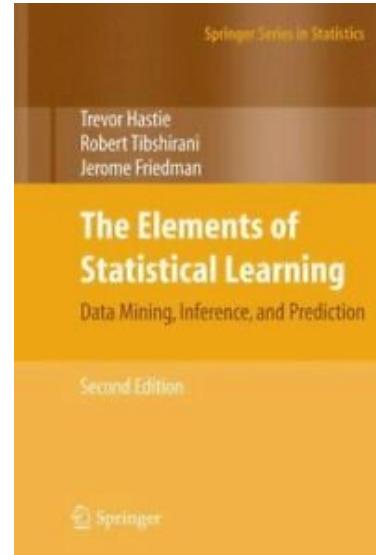
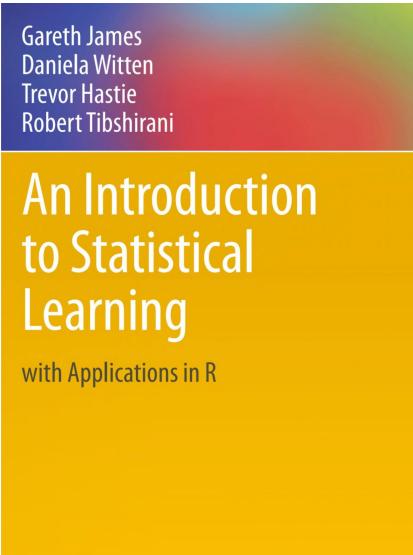
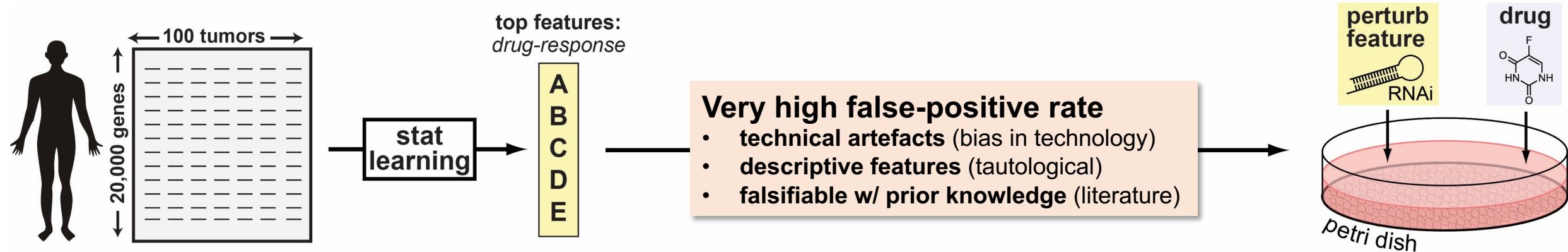
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# *Doctoral Training Philosophy: Columbia University*

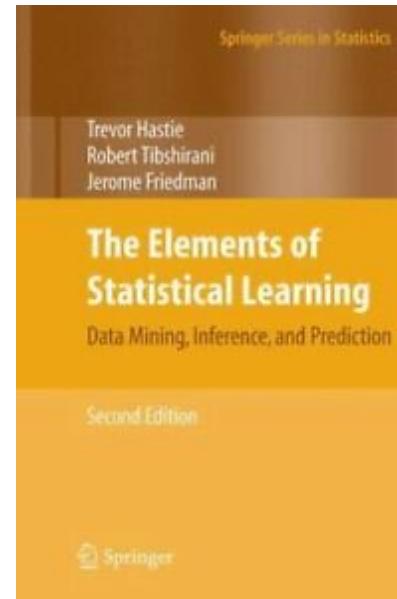
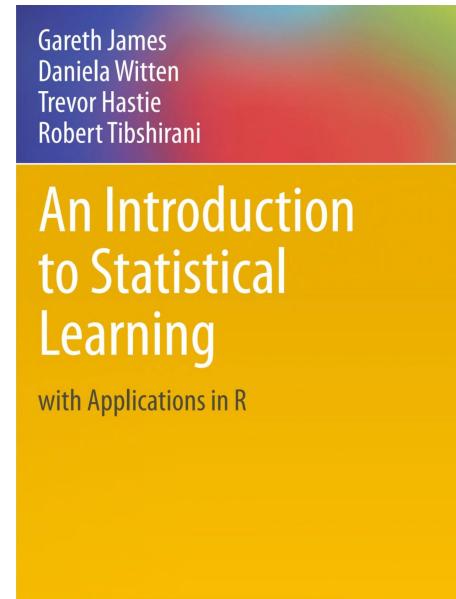
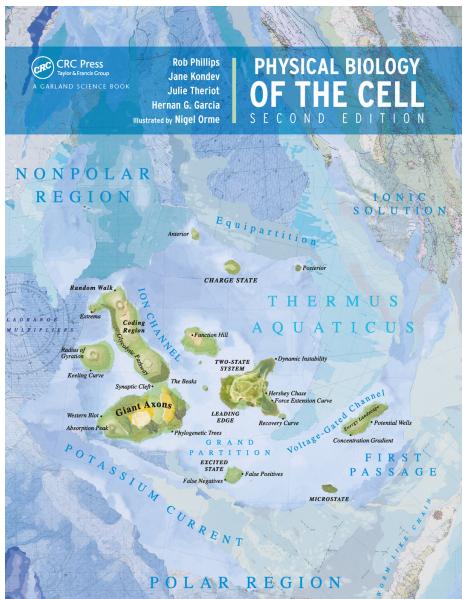
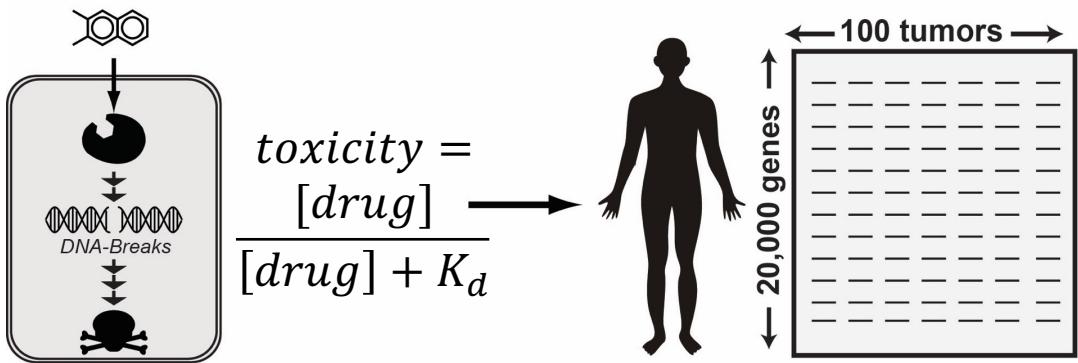
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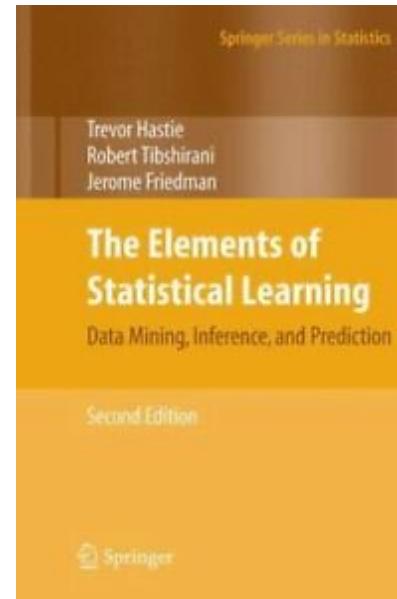
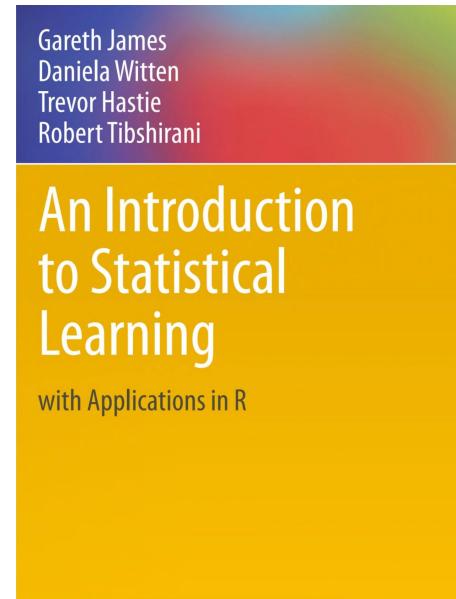
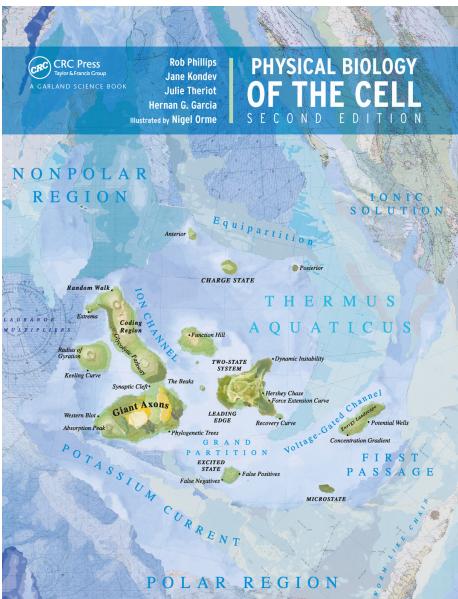
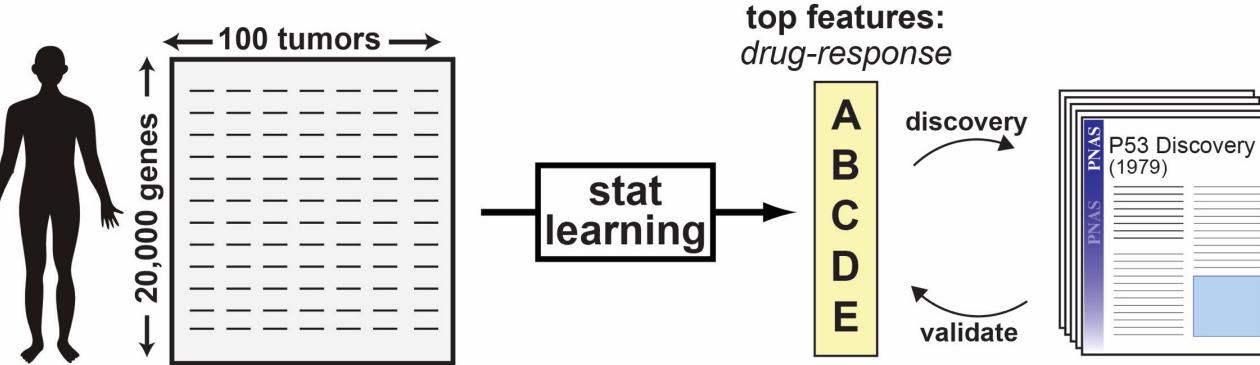
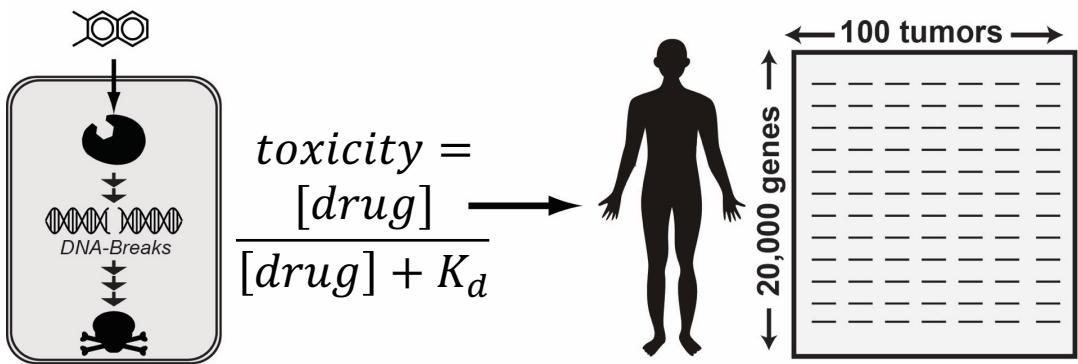
# Doctoral Training Philosophy: Columbia University



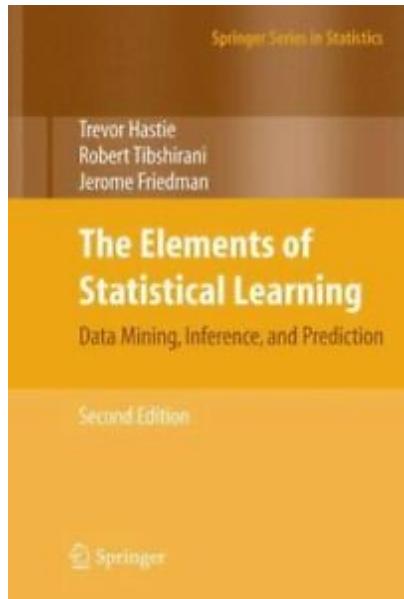
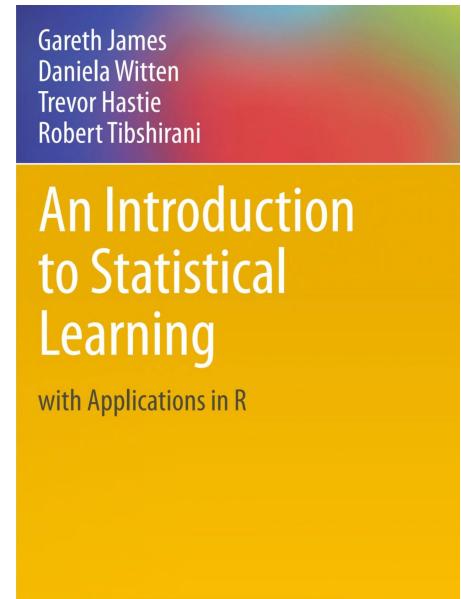
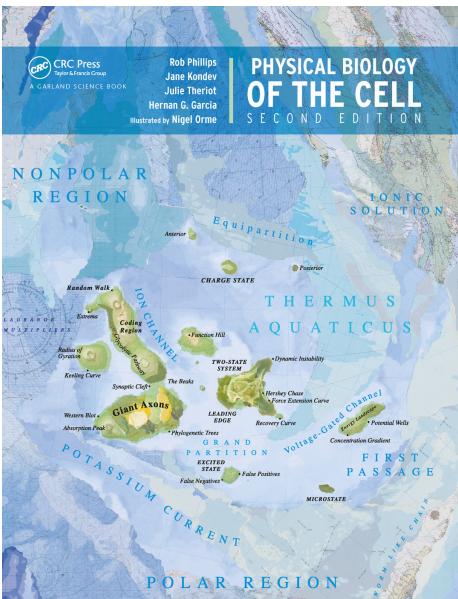
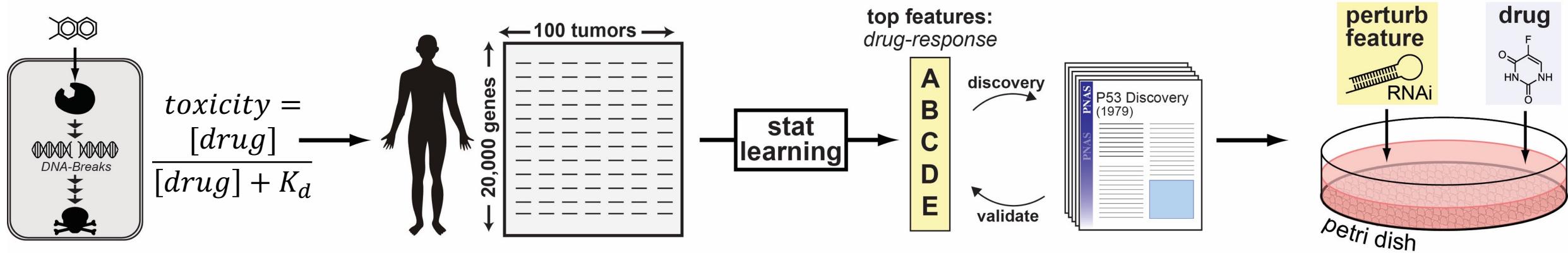
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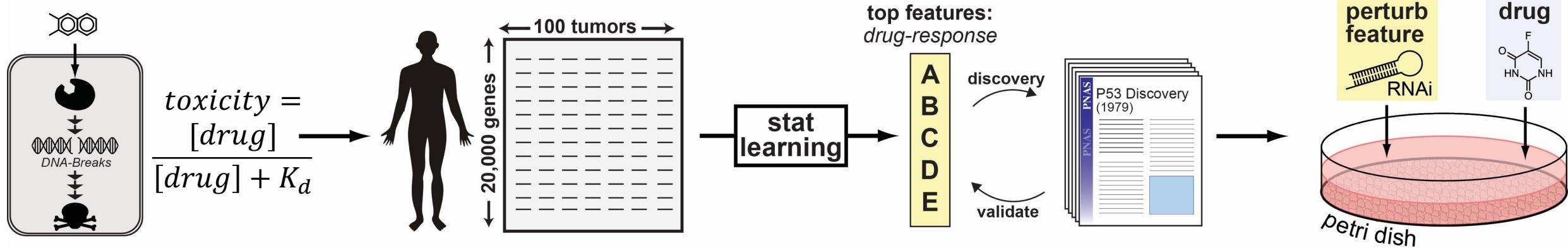
# Doctoral Training Philosophy: Columbia University



# *Doctoral Training Philosophy: Columbia University*



# Doctoral Training Philosophy: Columbia University



50 years of Data Science

David Donoho

Sept. 18, 2015  
Version 1.00

## Abstract

More than 50 years ago, John Tukey called for a reformation of academic statistics. In 'The Future of Data Analysis', he pointed to the existence of an as-yet unrecognized *science*, whose subject of interest was learning from data, or 'data analysis'. Ten to twenty years ago, John Chambers, Bill Cleveland and Leo Breiman independently once again urged academic statistics to expand its boundaries beyond the classical domain of theoretical statistics; Chambers called for more emphasis on data preparation and presentation rather than statistical modeling; and Breiman called for emphasis on prediction rather than inference. Cleveland even suggested the catchy name "Data Science" for his envisioned field.

## FIELD SPECIFIC:

- 1. Data Preparation:** data-QC and organizing/cleaning
- 2. Data Transformation:** emphasize meaningful info (Fourier)
- 3. Data Visualization:** effective communication

## FIELD AGNOSTIC:

- 4. Computing:** languages, cluster computing
- 5. Machine Learning:** "Adequate Stat Model"
- 6. Mathematics:** Deterministic Modeling or "True Model"

# *Original Teaching Material: practicallyscience.com*

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MATH

COMPUTING

PHYSICS

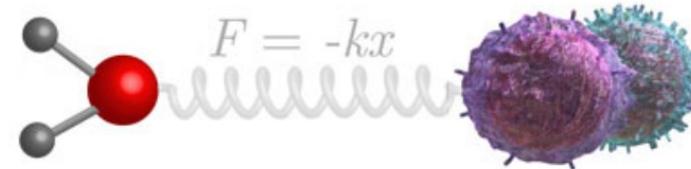
CHEMISTRY

BIOLOGY

TIMELINES

RUNNING

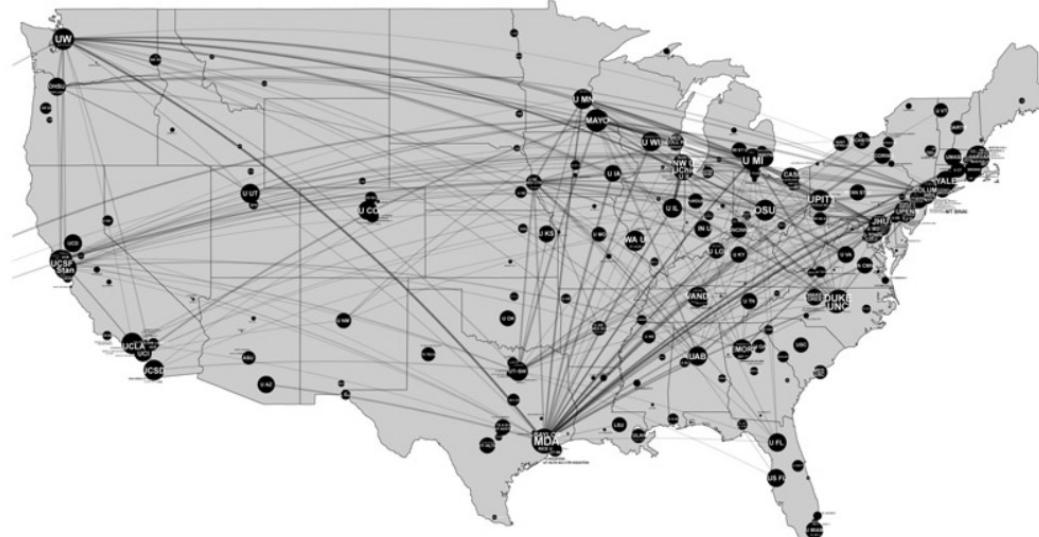
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*conceptual  
infographics for  
interdisciplinary  
scientists*

## Cancer Research in the USA

Posted on [October 28, 2018](#) | Comments Off



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### ABOUT US

Practically Science was started by two Yale PhD students in 2012. Its goal is to make single-sheet summaries of common interdisciplinary methods, ideas, etc.

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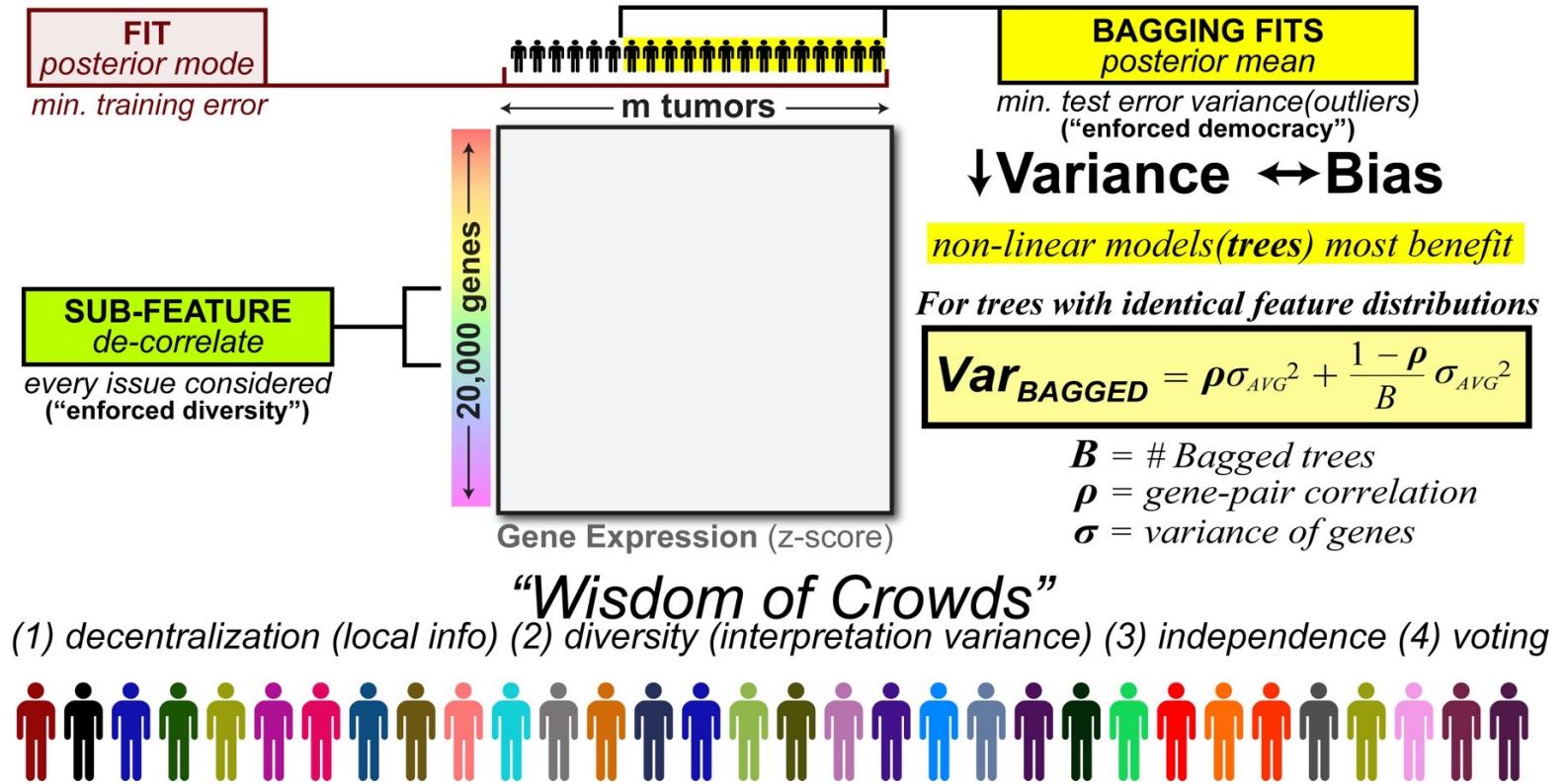
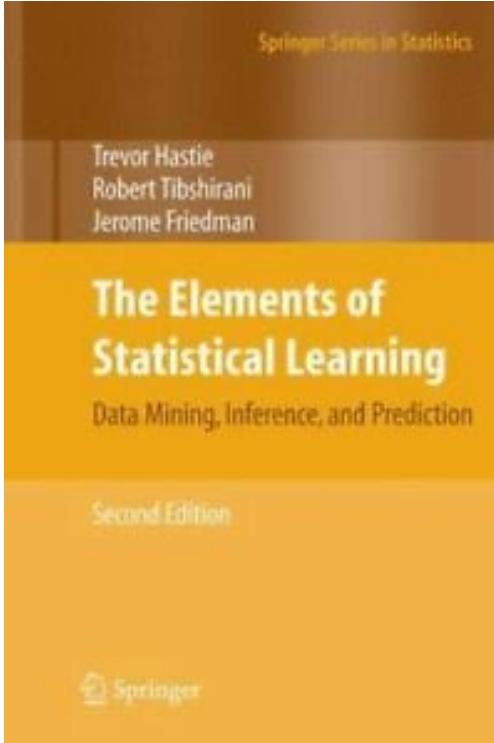
### PLEASE FOLLOW & LIKE US :)



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### BLOGROLL

# Original Teaching Material: graphical summaries



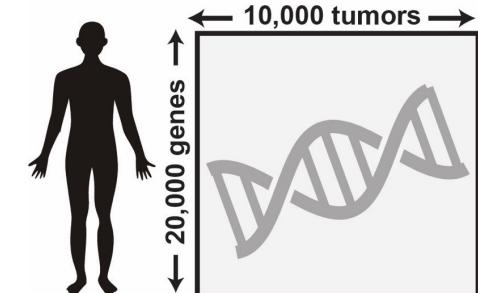
# Clinical “Big Data” + Experiment = Precision Medicine

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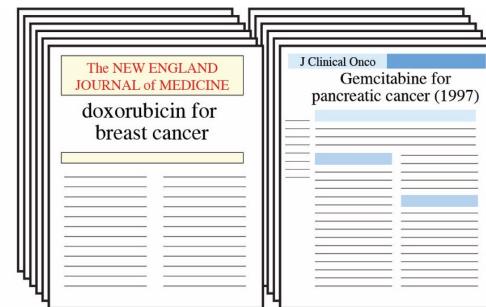
Genomic vs Experimental “Big Data:” *challenges and opportunities*

**History & Current Status of Cancer Therapy:** *1 drug / 1 target paradigm*

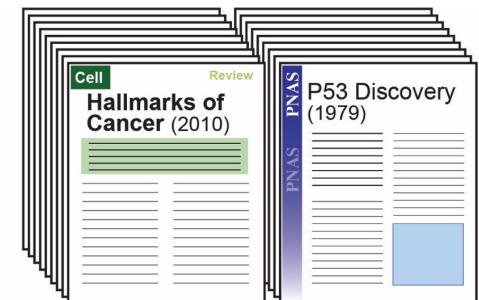
**Reconciling Clinical & Laboratory “Big Data”:** *1 drug / N-biomarkers*



Sequencing Tech

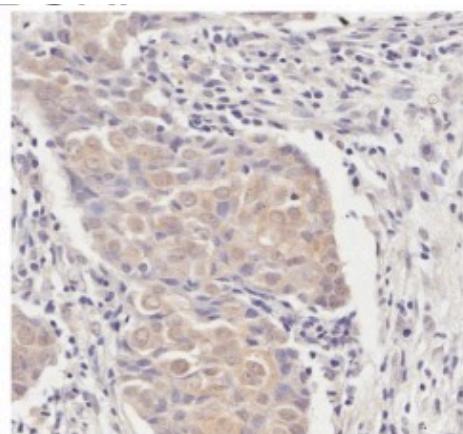
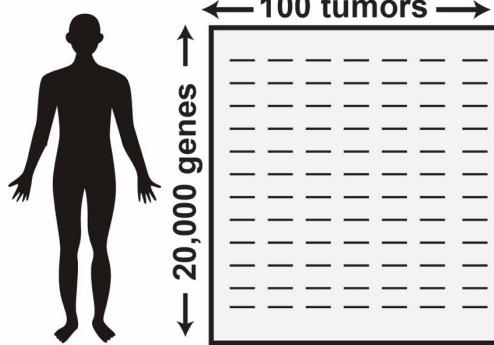


Clinical Literature

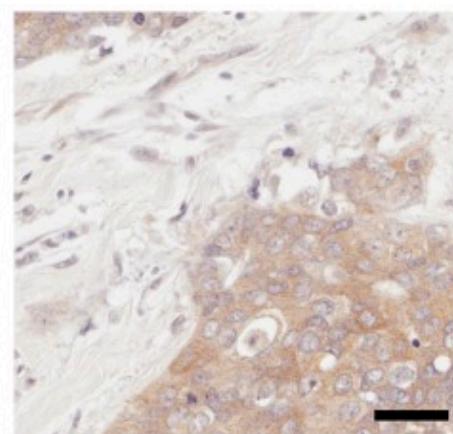


Scientific Literature

# Emory Collaboration: Kevin Kalinsky & Bill Li

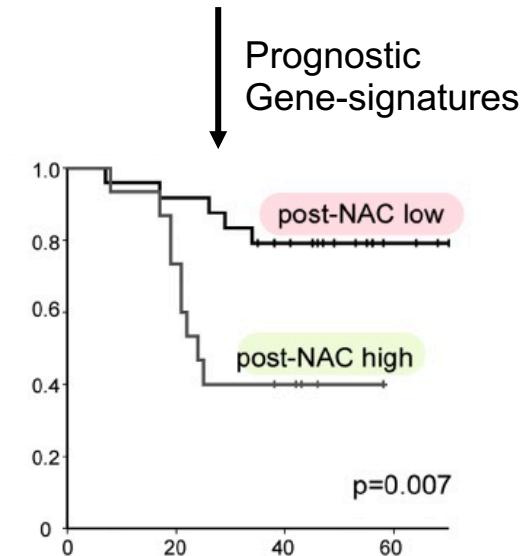
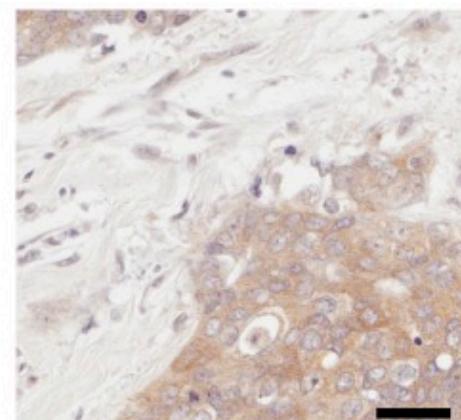
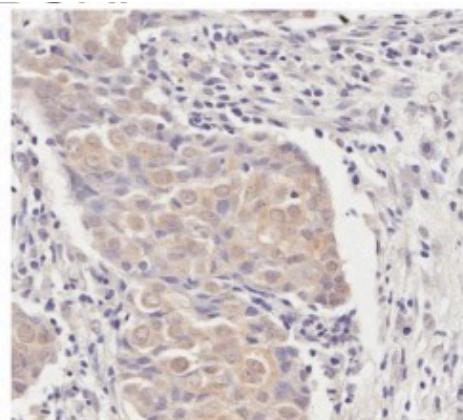
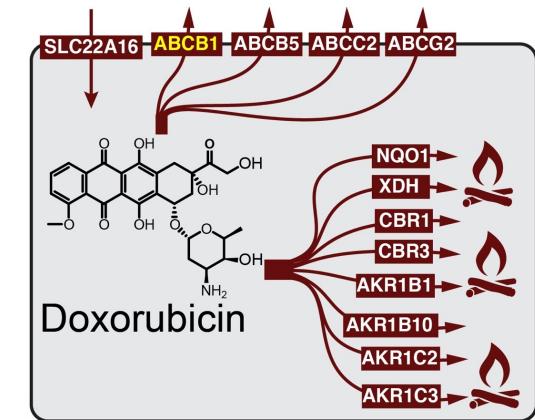
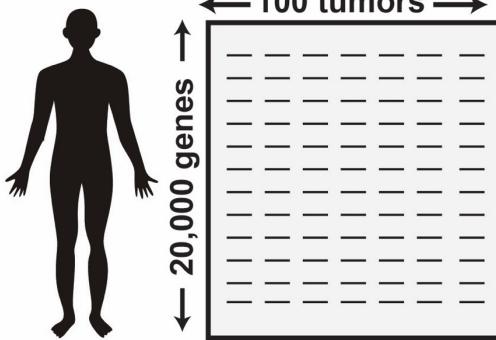


pre-NAC

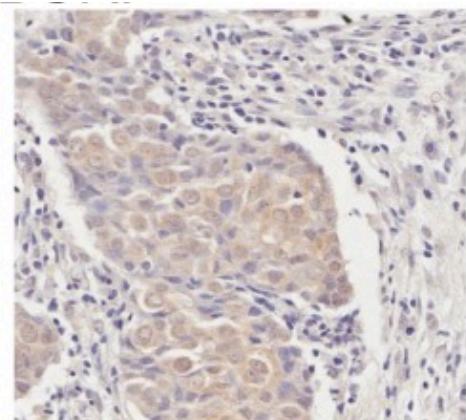
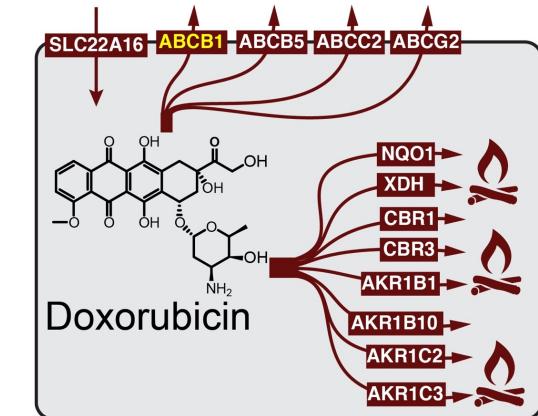
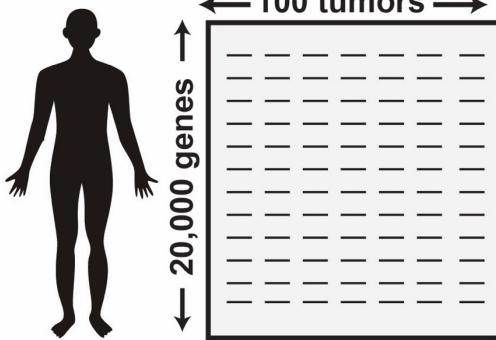


post-NAC

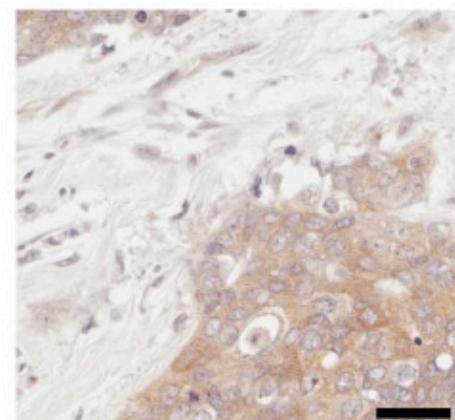
# Emory Collaboration: Kevin Kalinsky & Bill Li



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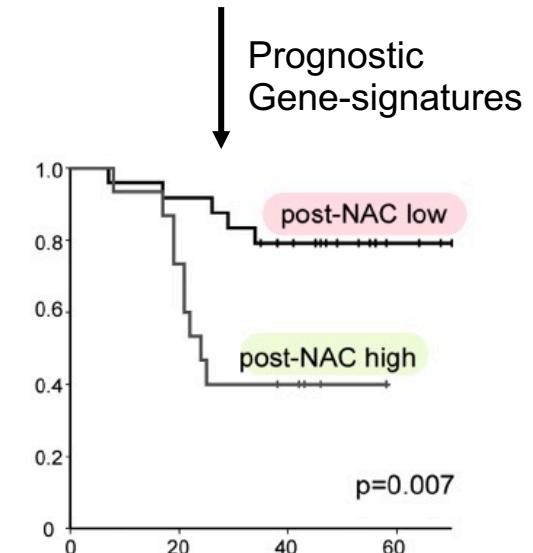


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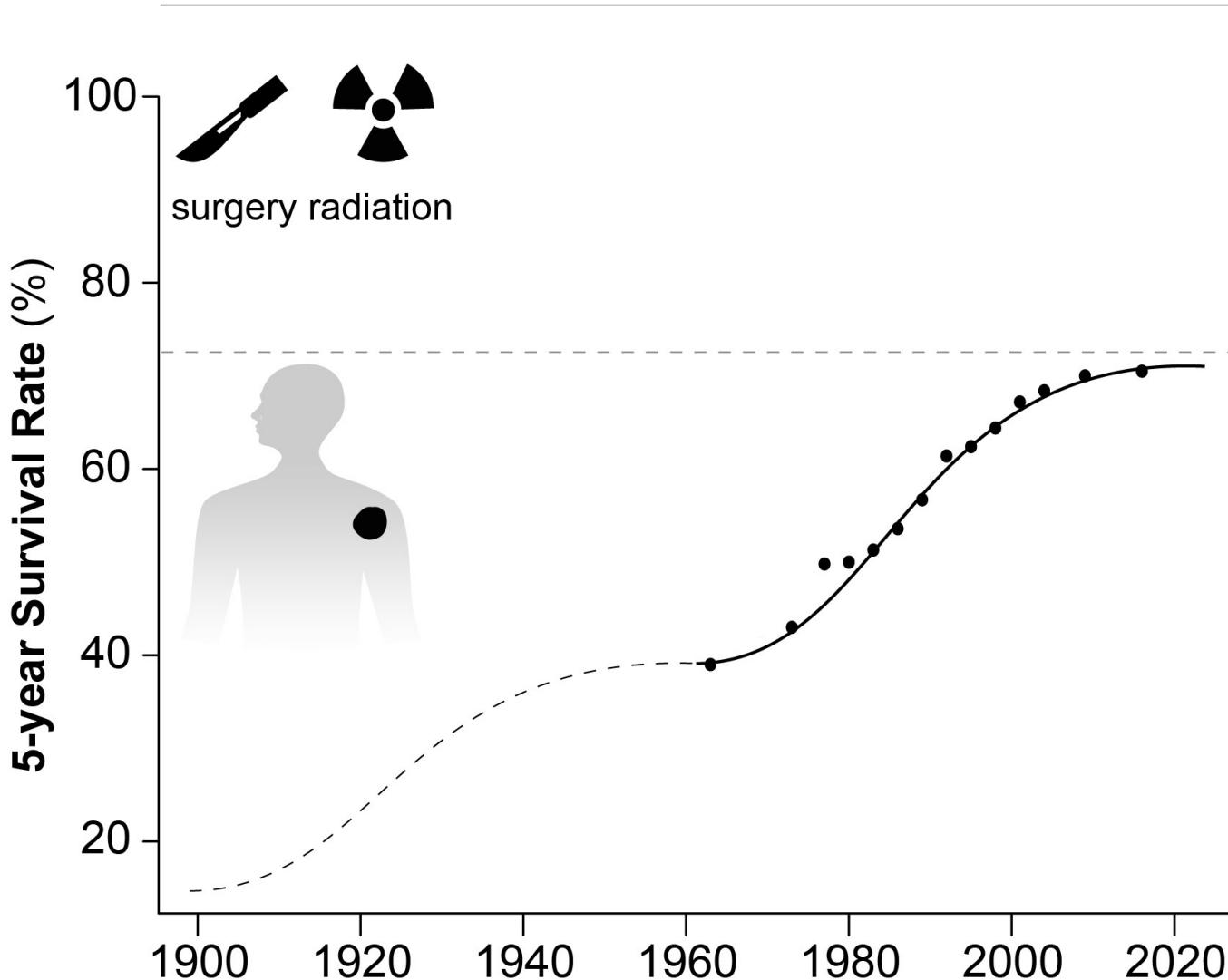


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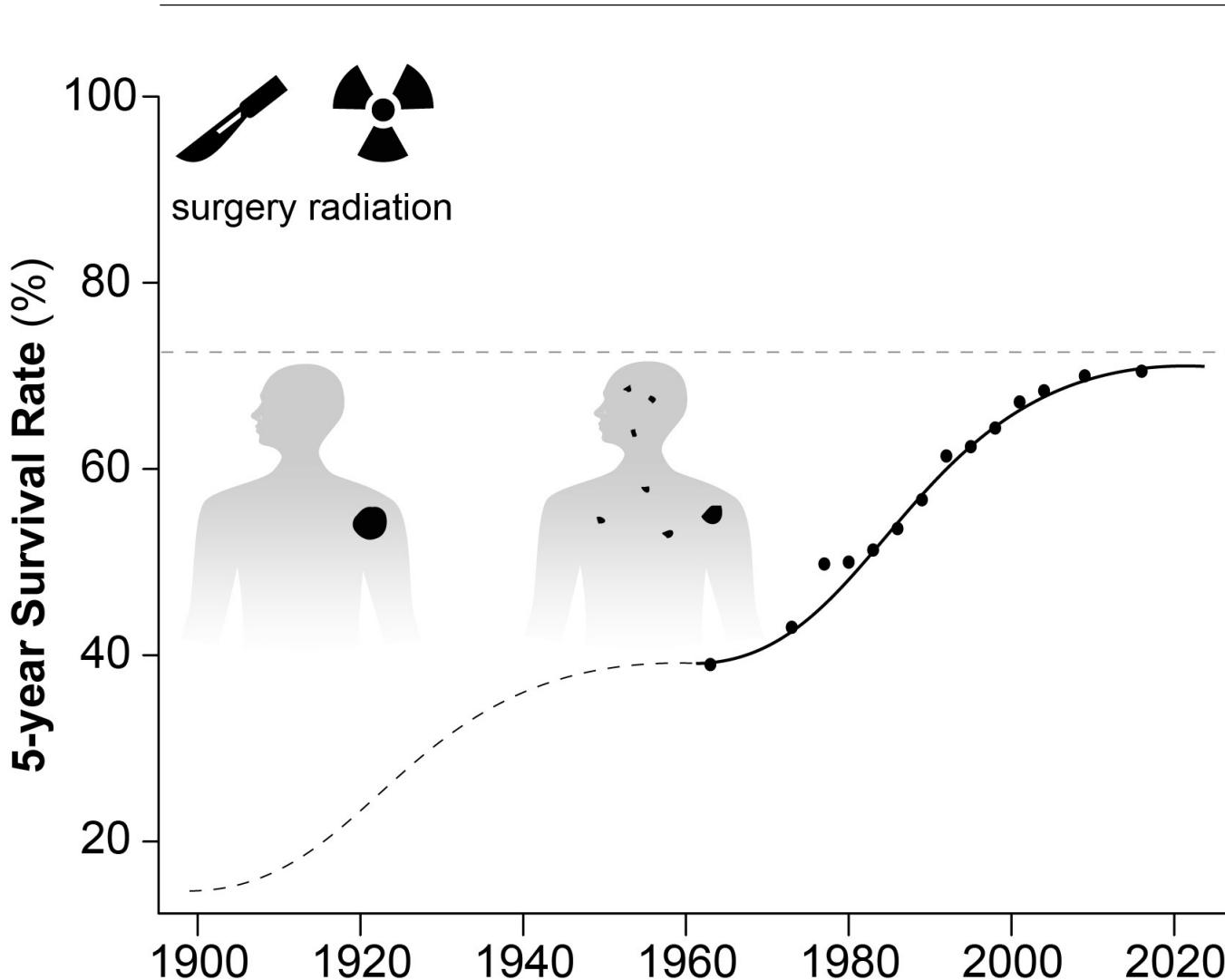
Predict drug-response  
Identify drugs to overcome resistance



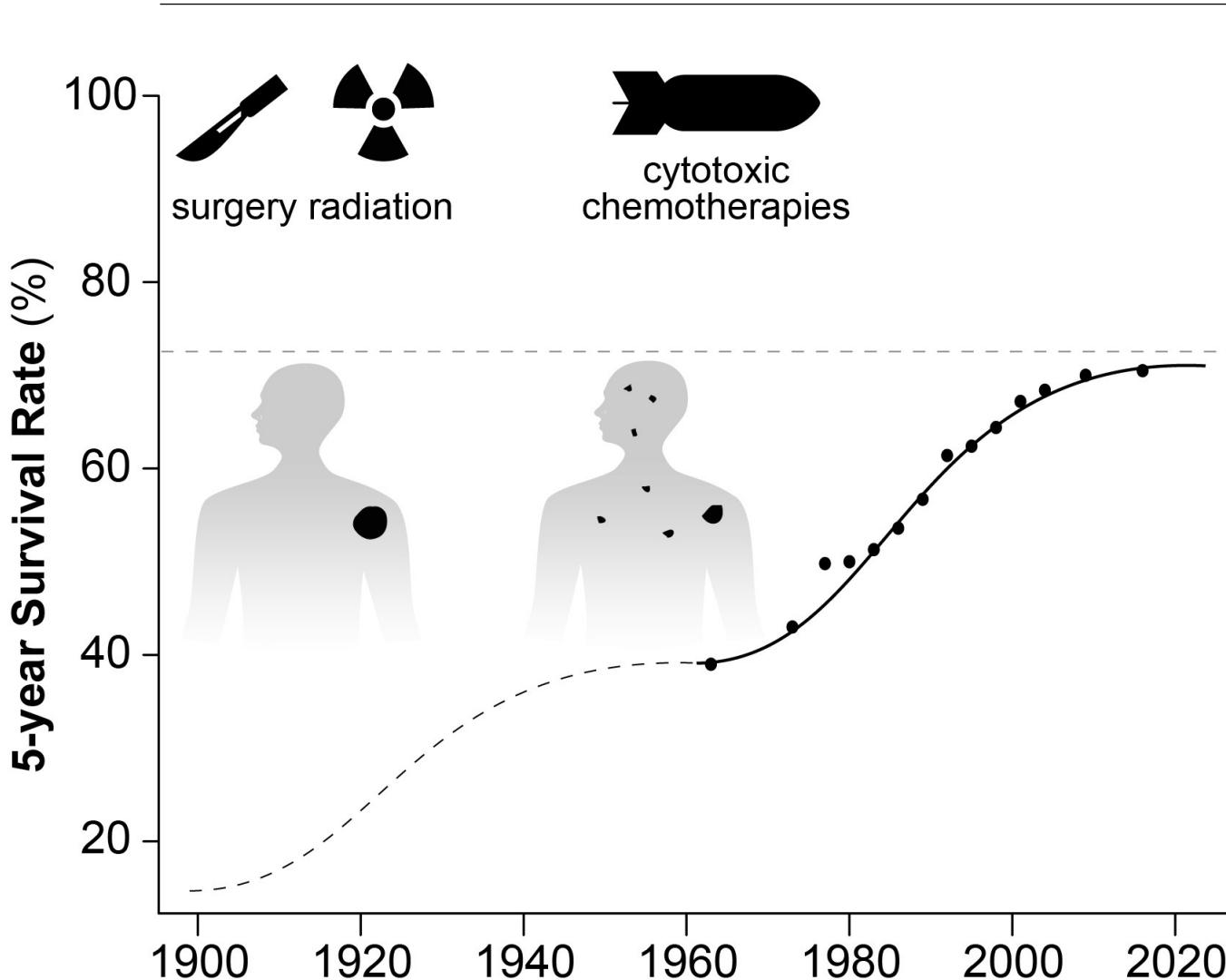
# History of Cancer Treatment: *localized cancers*



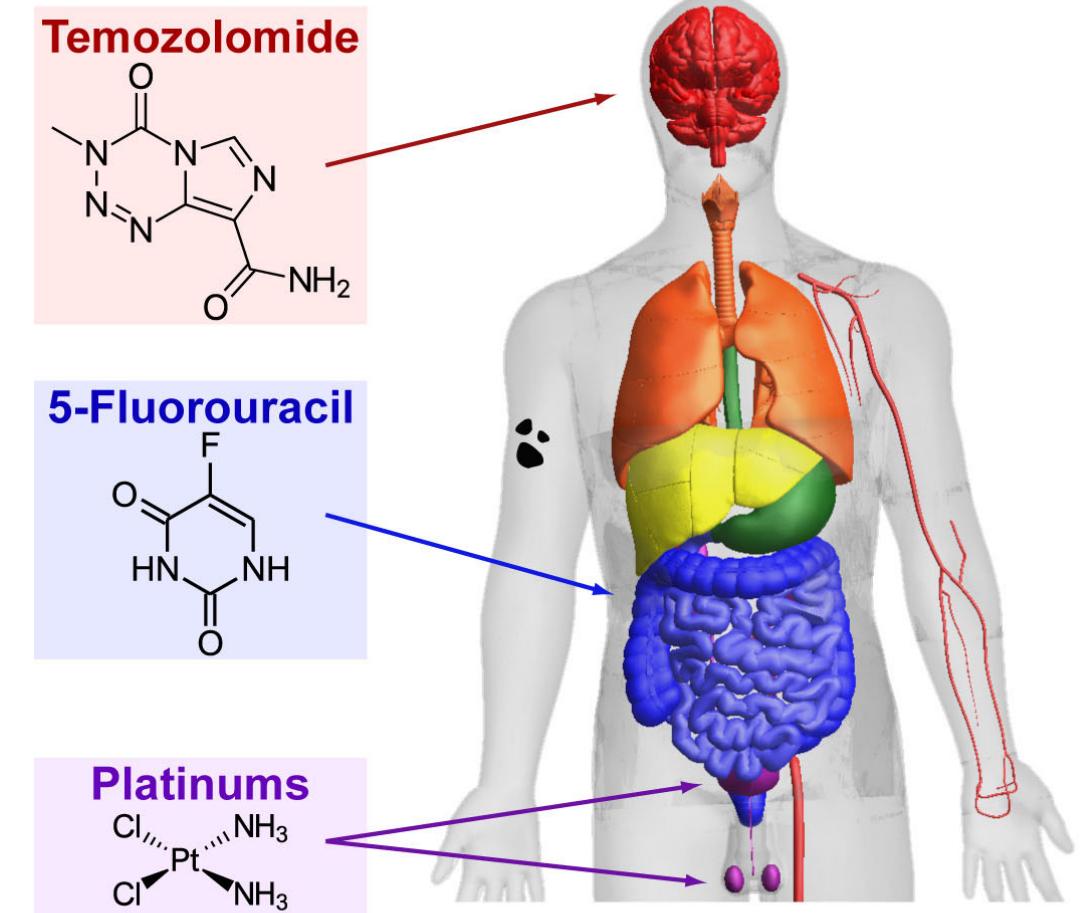
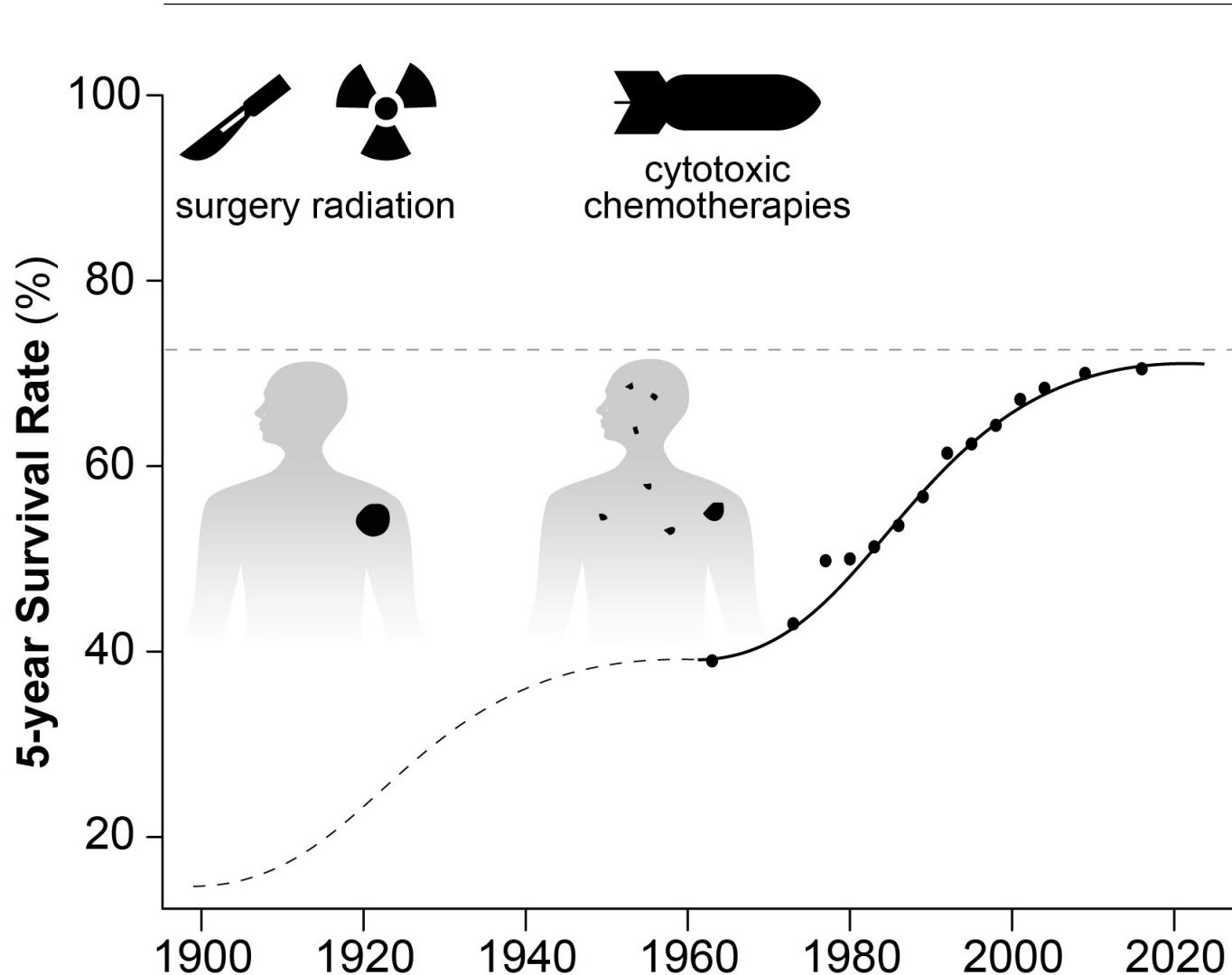
# History of Cancer Treatment: *metastatic cancer*



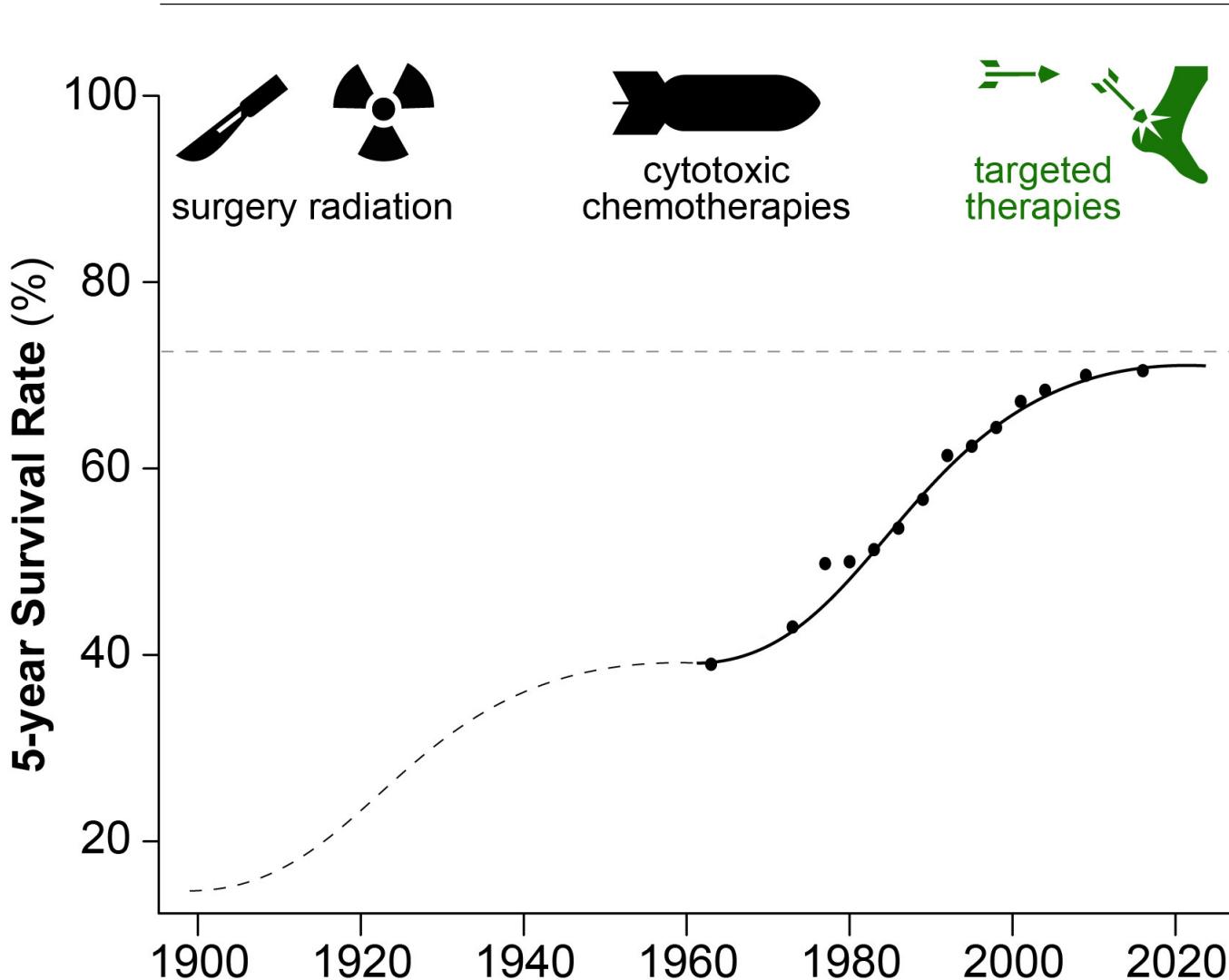
# History of Cancer Treatment: *metastatic cancer*



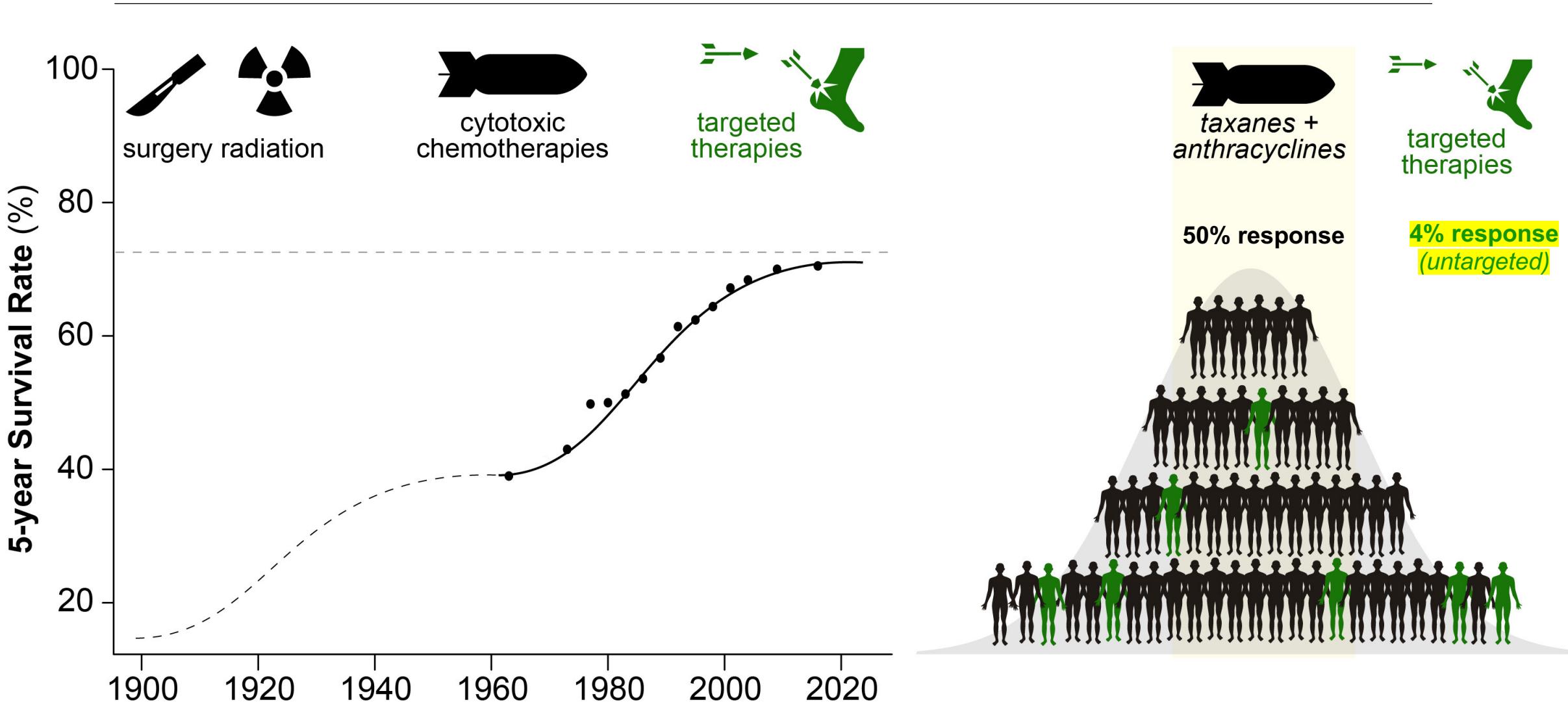
# History of Cancer Treatment: *metastatic cancer*



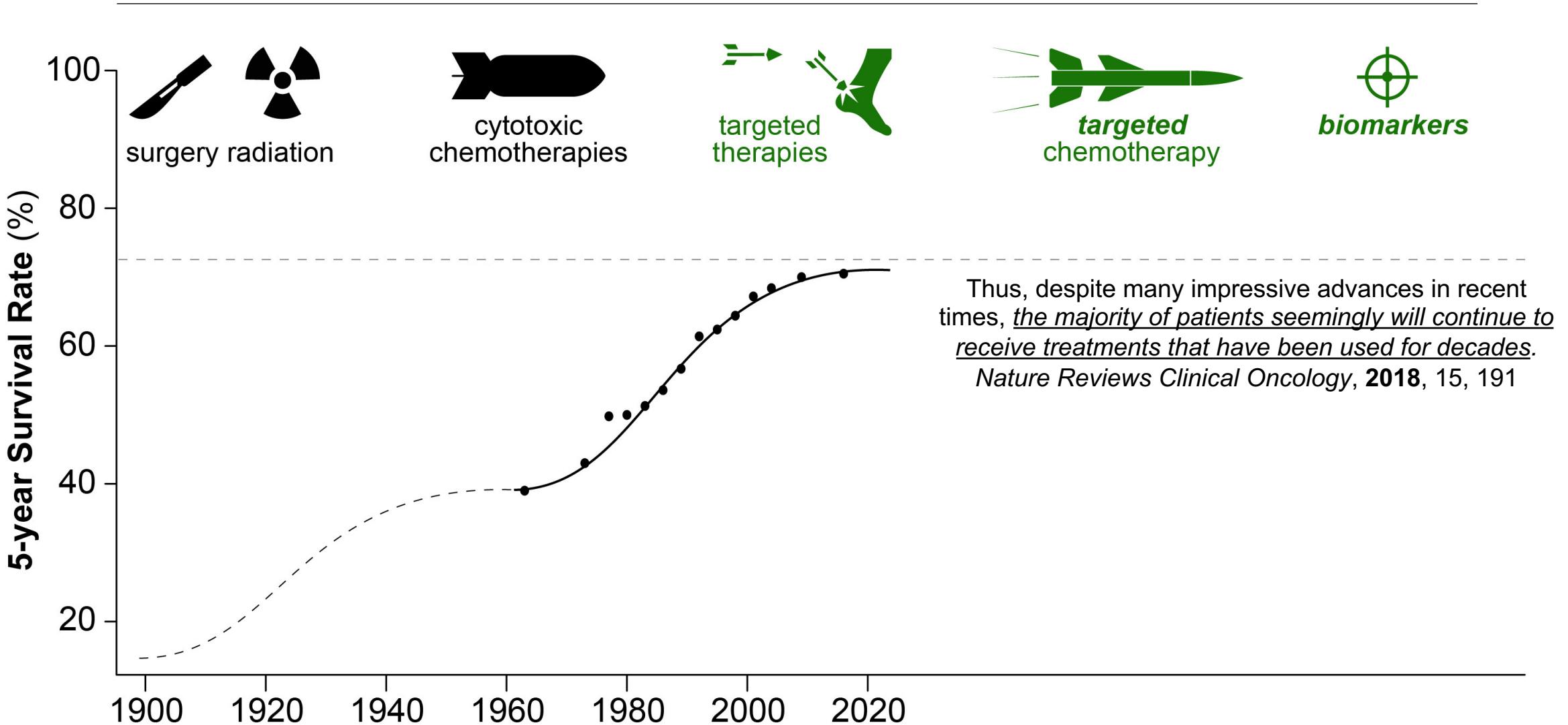
# History of Cancer Treatment: *targeted therapies?*



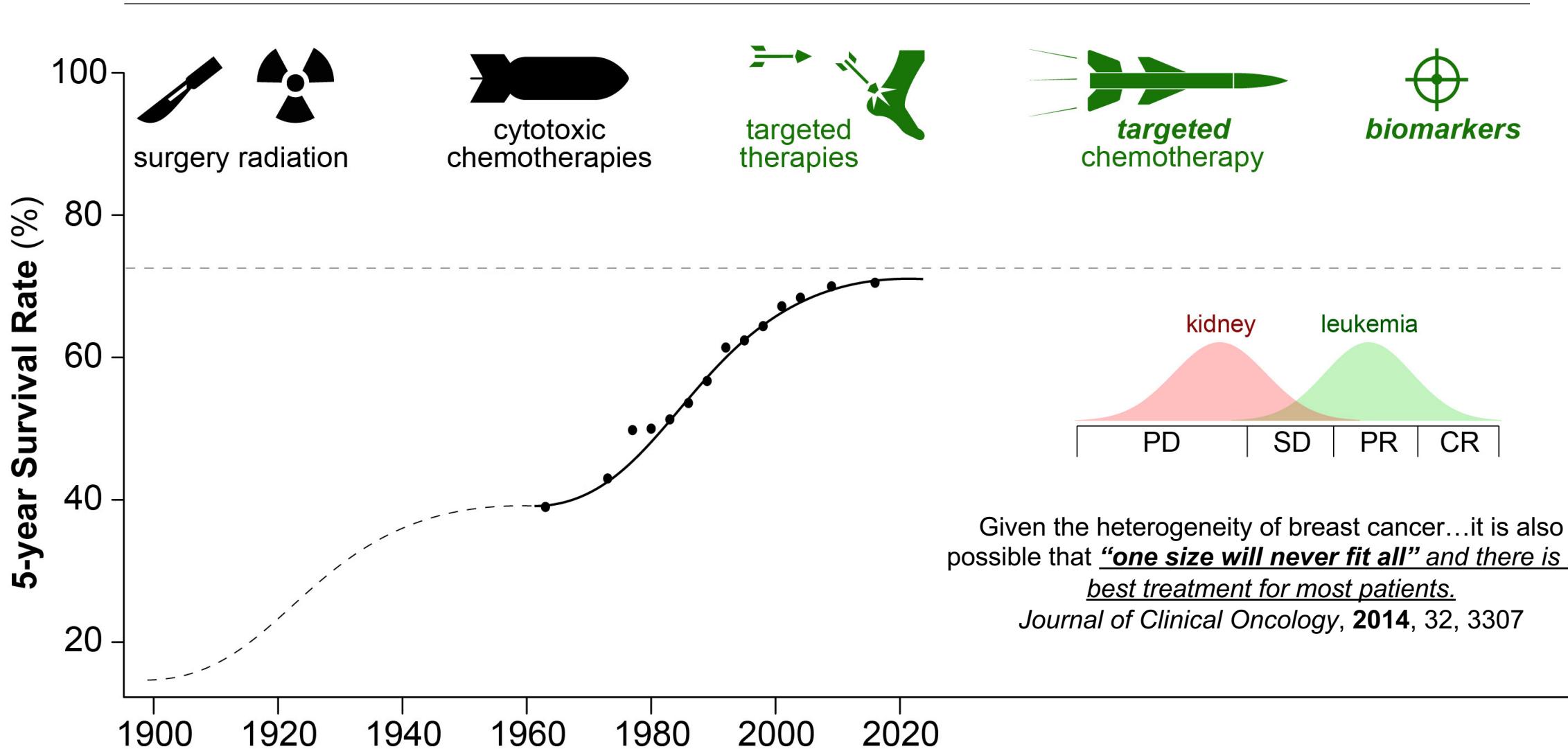
# History of Cancer Treatment: *targeted therapies?*



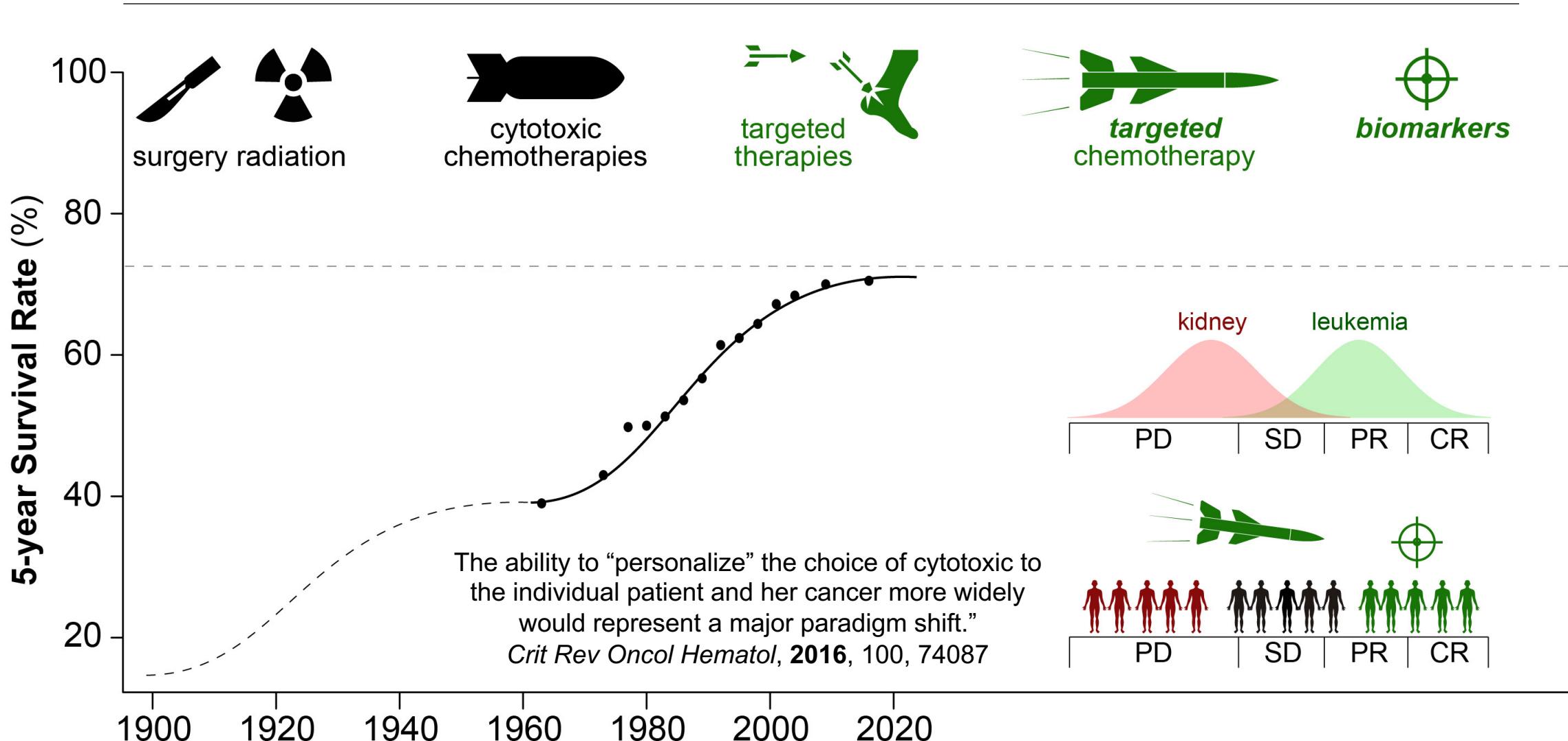
# History of Cancer Treatment: *targeted chemotherapies?*



# History of Cancer Treatment: *targeted chemotherapies?*

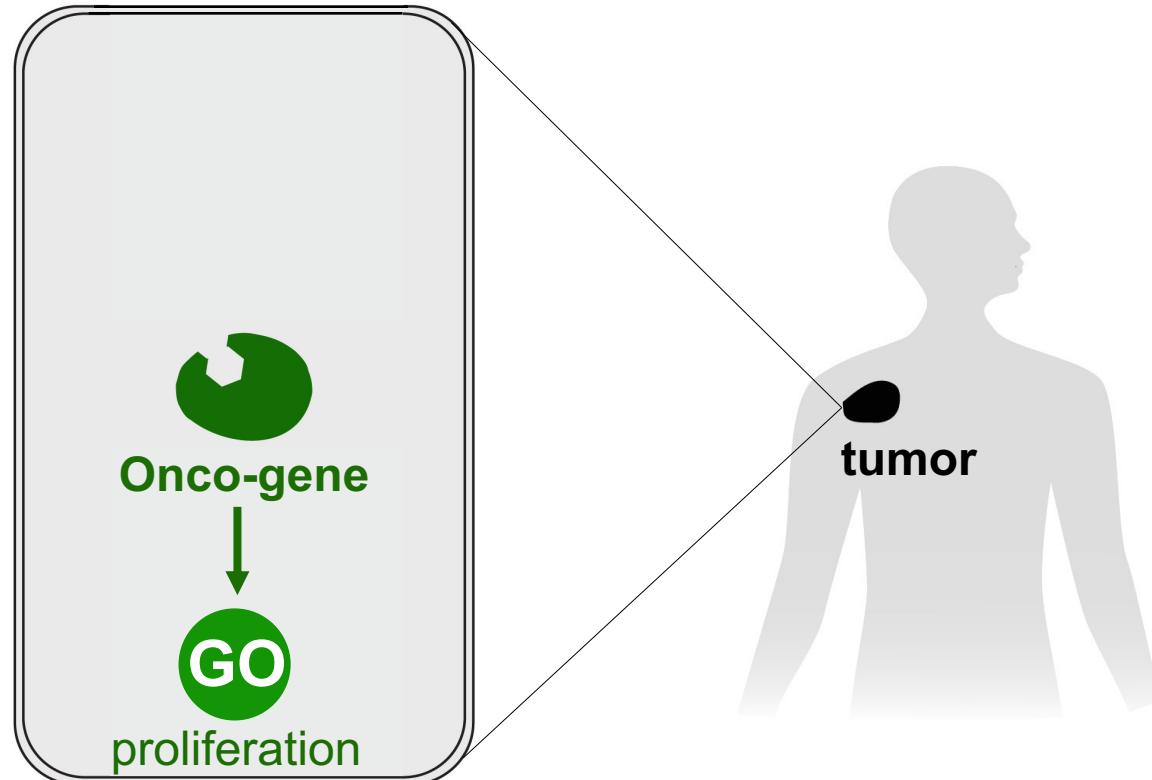


# History of Cancer Treatment: *targeted chemotherapies?*

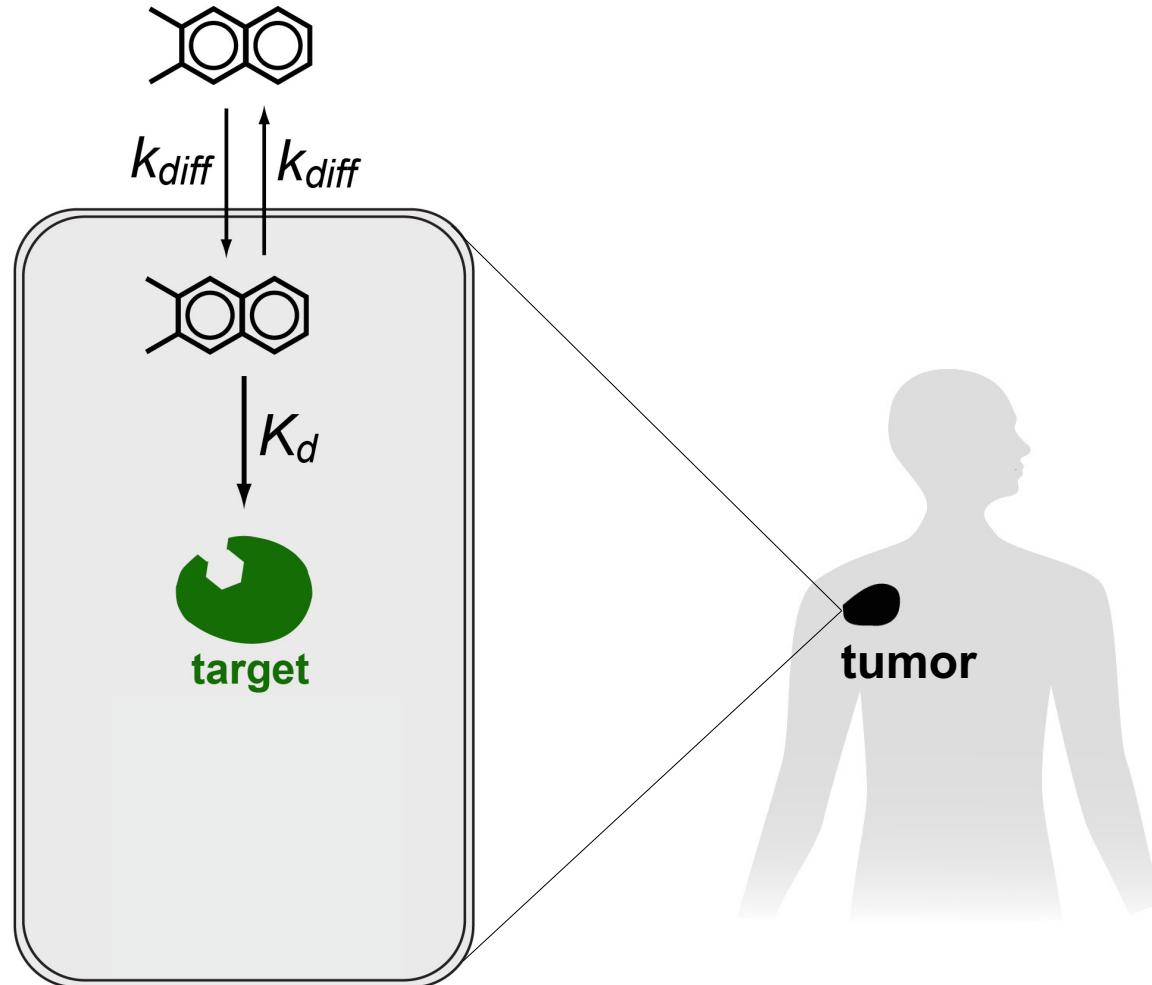


# Laboratory Drug-sensitivity: dose-response curves

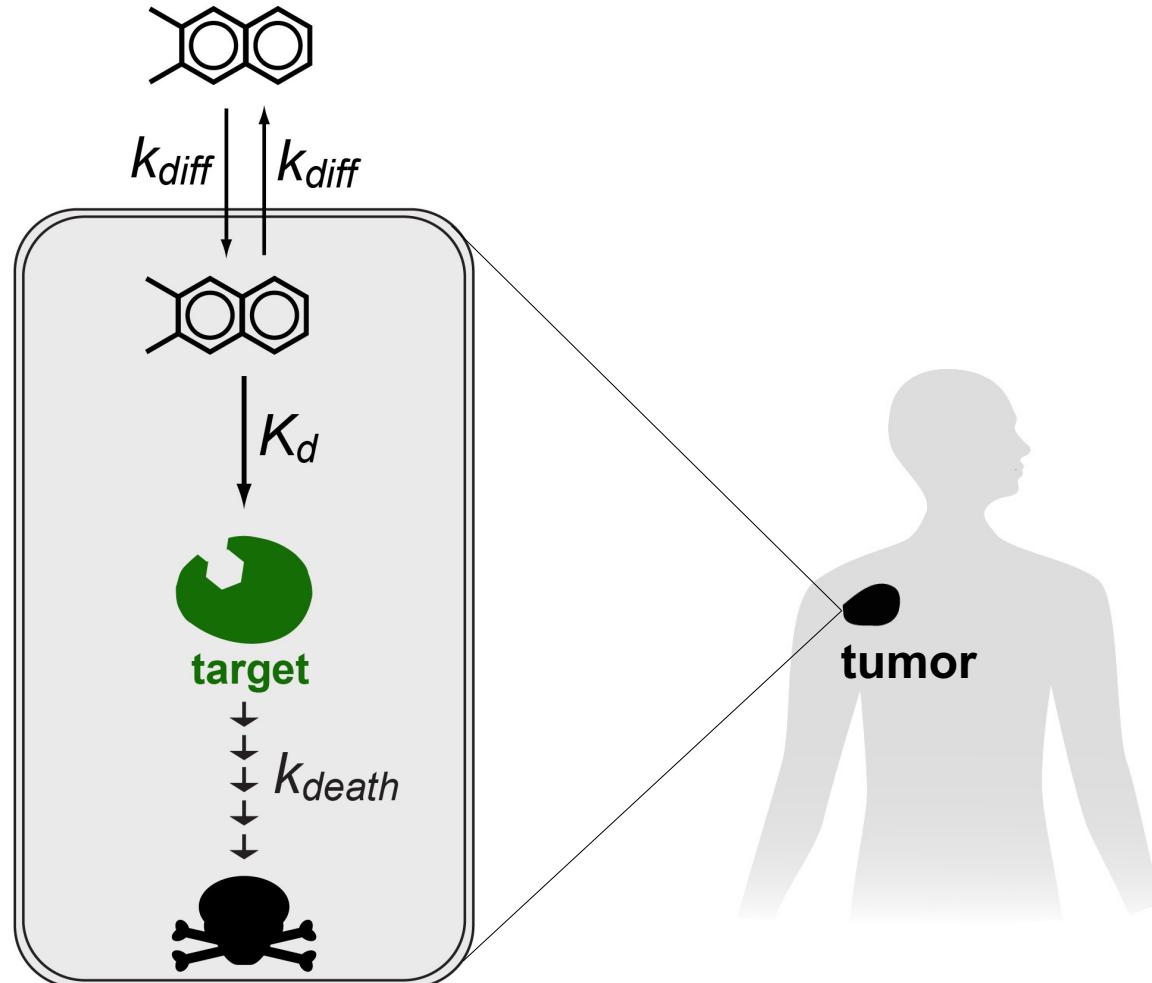
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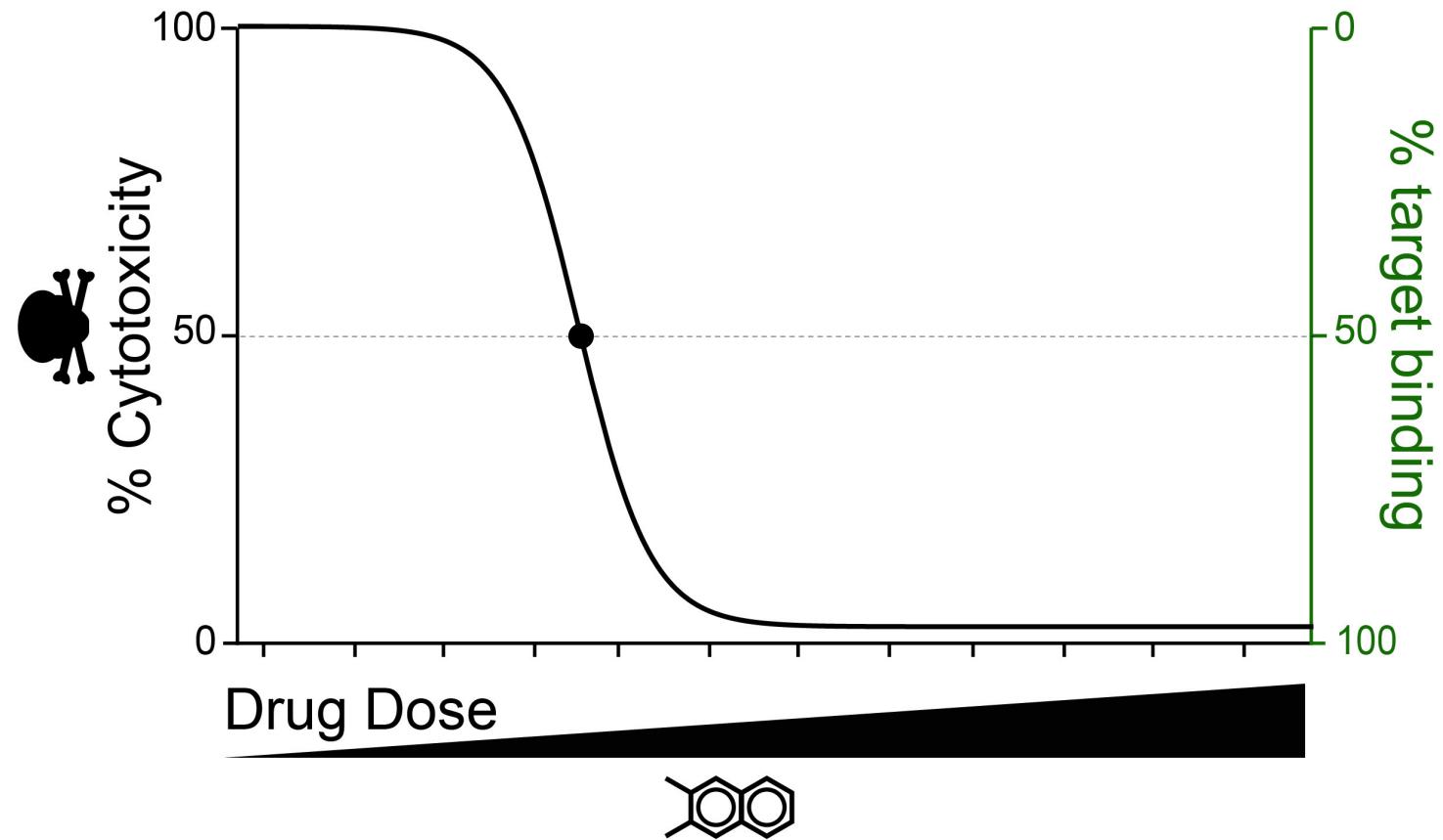
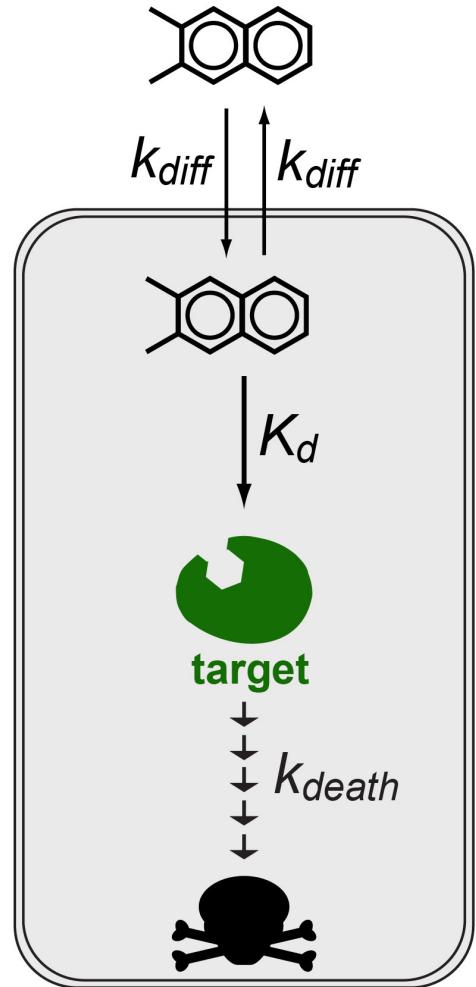
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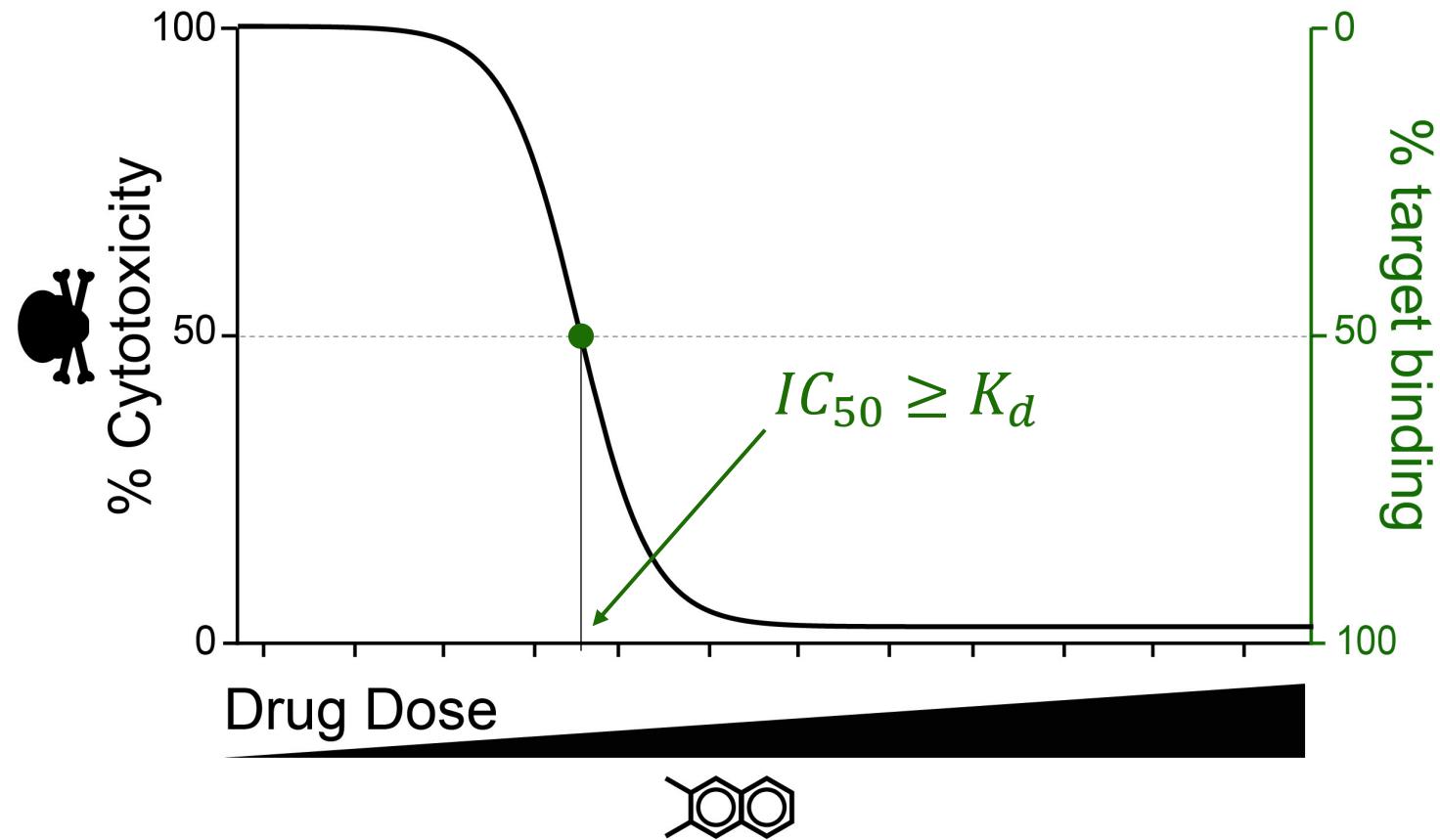
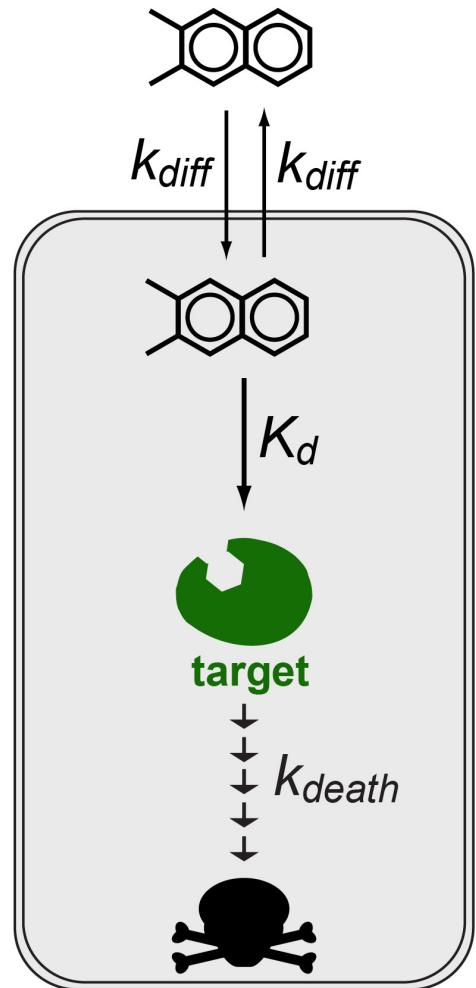
# Laboratory Drug-sensitivity: dose-response curves



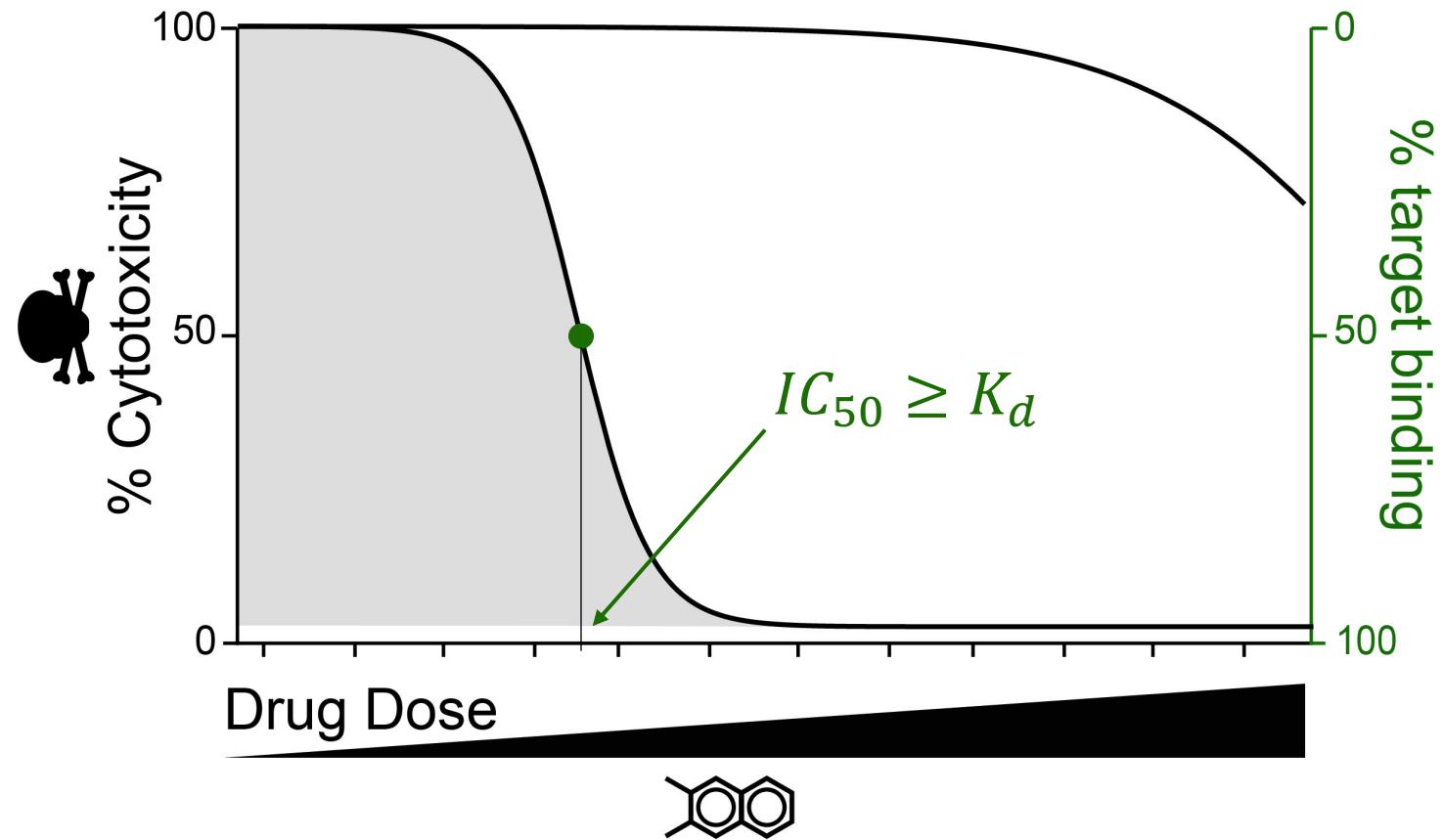
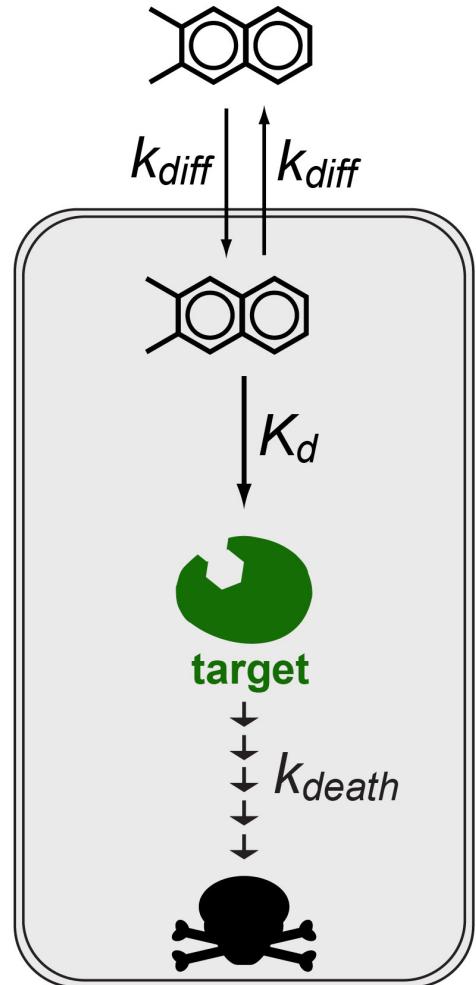
# Laboratory Drug-sensitivity: dose-response curves



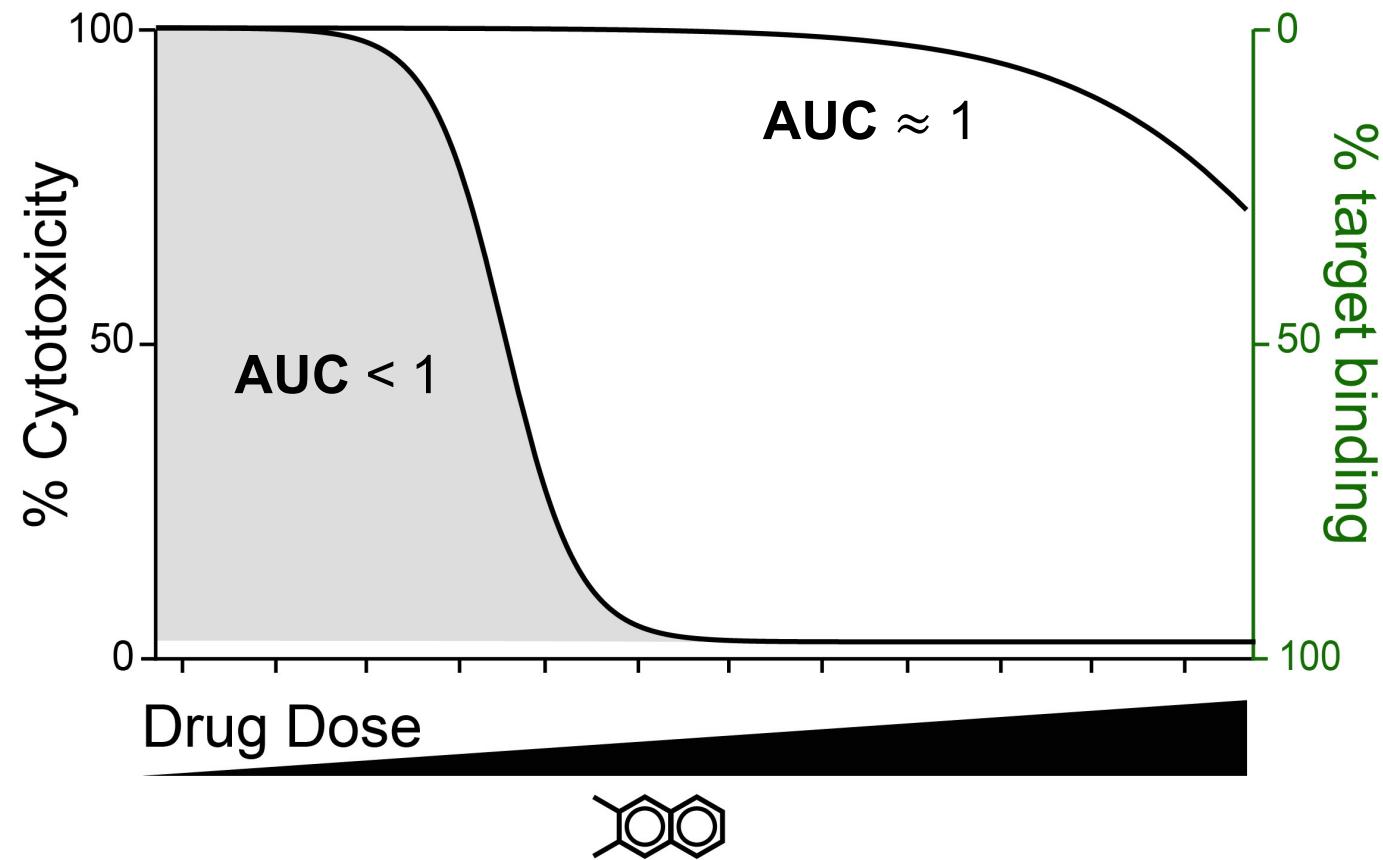
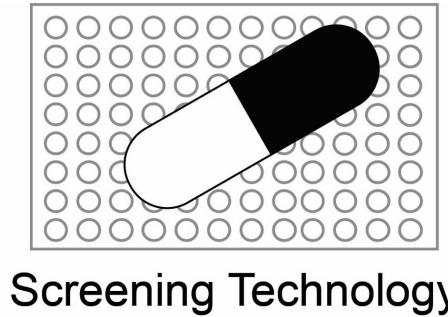
# Laboratory Drug-sensitivity: dose-response curves



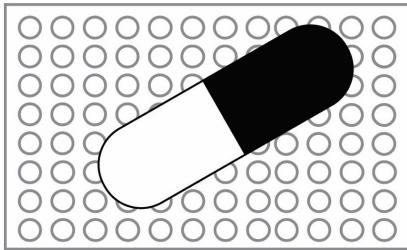
# Laboratory Drug-sensitivity: dose-response curves



# Laboratory Drug-sensitivity: dose-response curves



# Laboratory Drug-sensitivity: dose-response curves

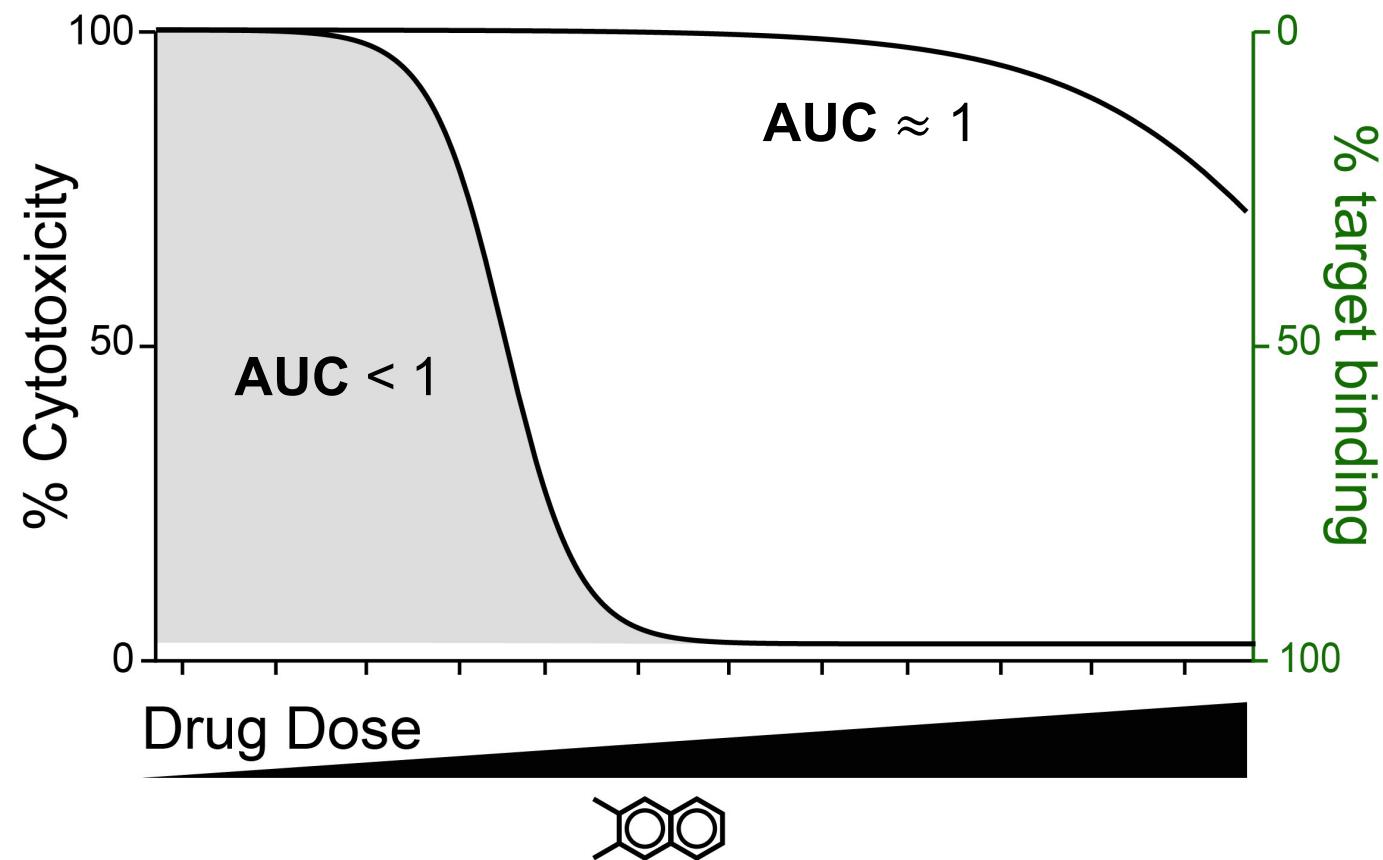


Screening Technology

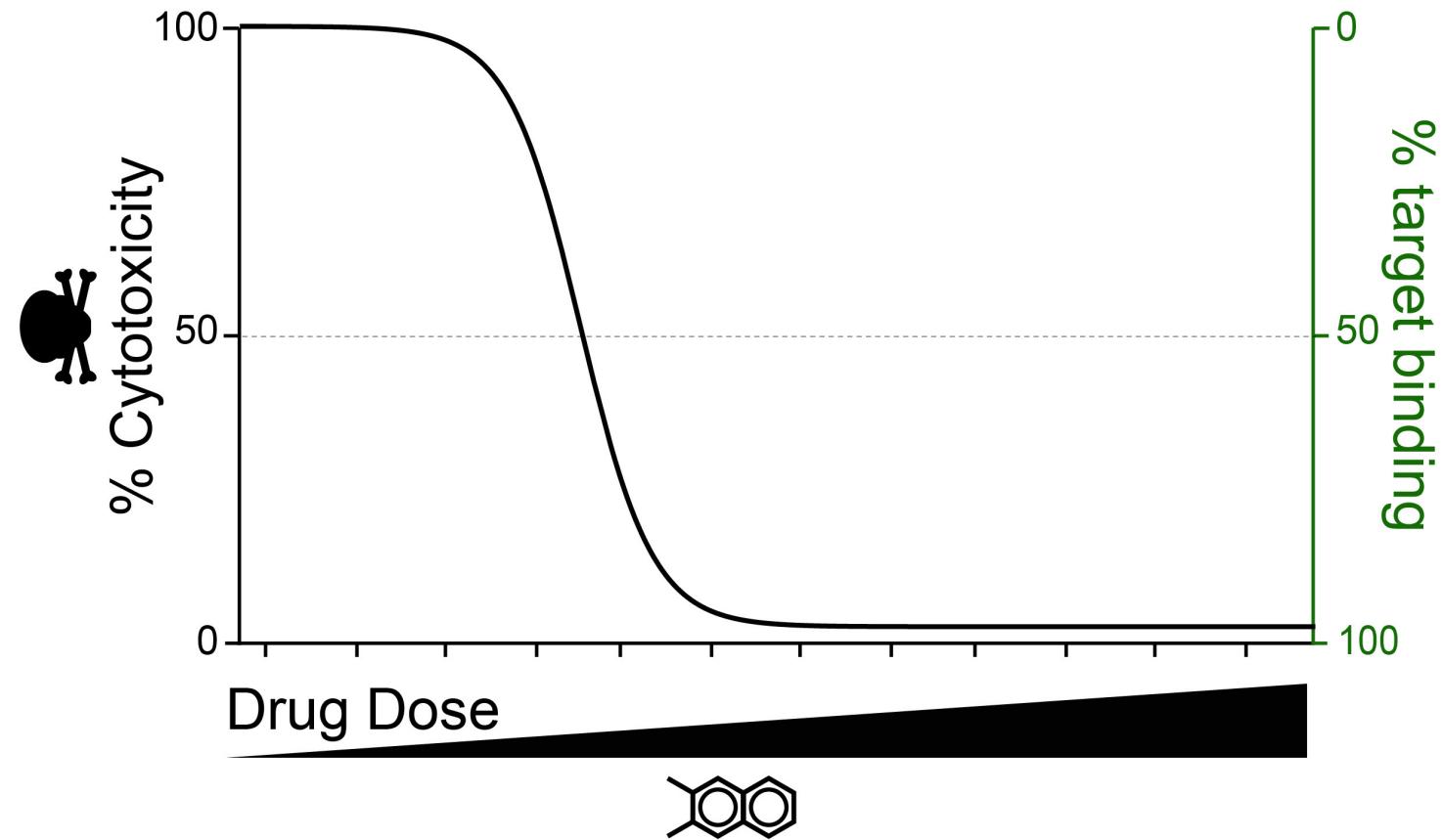
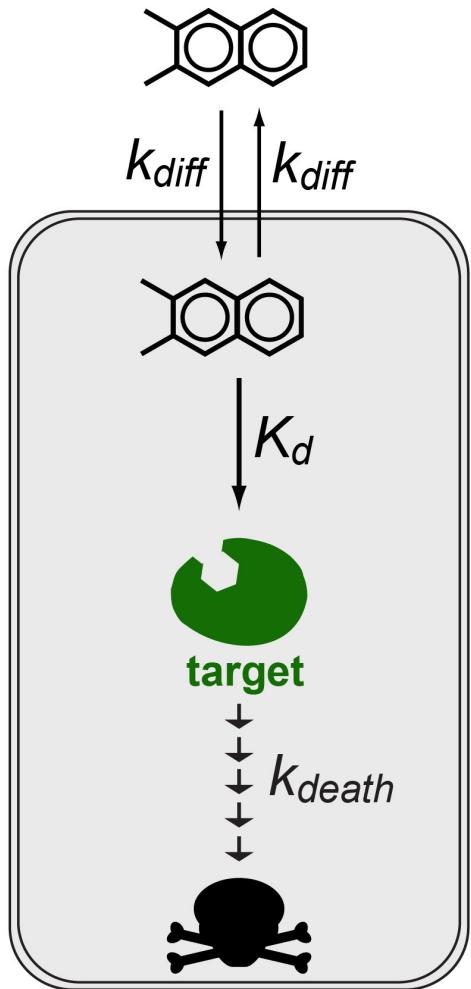
Cancer Therapeutic Response Portal (CTRP)  
**545 drugs x 887 cell-lines**  
*Cell*, 2013, 154, 1151

Genomics of Drug Sensitivity (GDSC)  
**473 drugs x 972 cell-lines**  
*Cell*, 2016, 166, 740

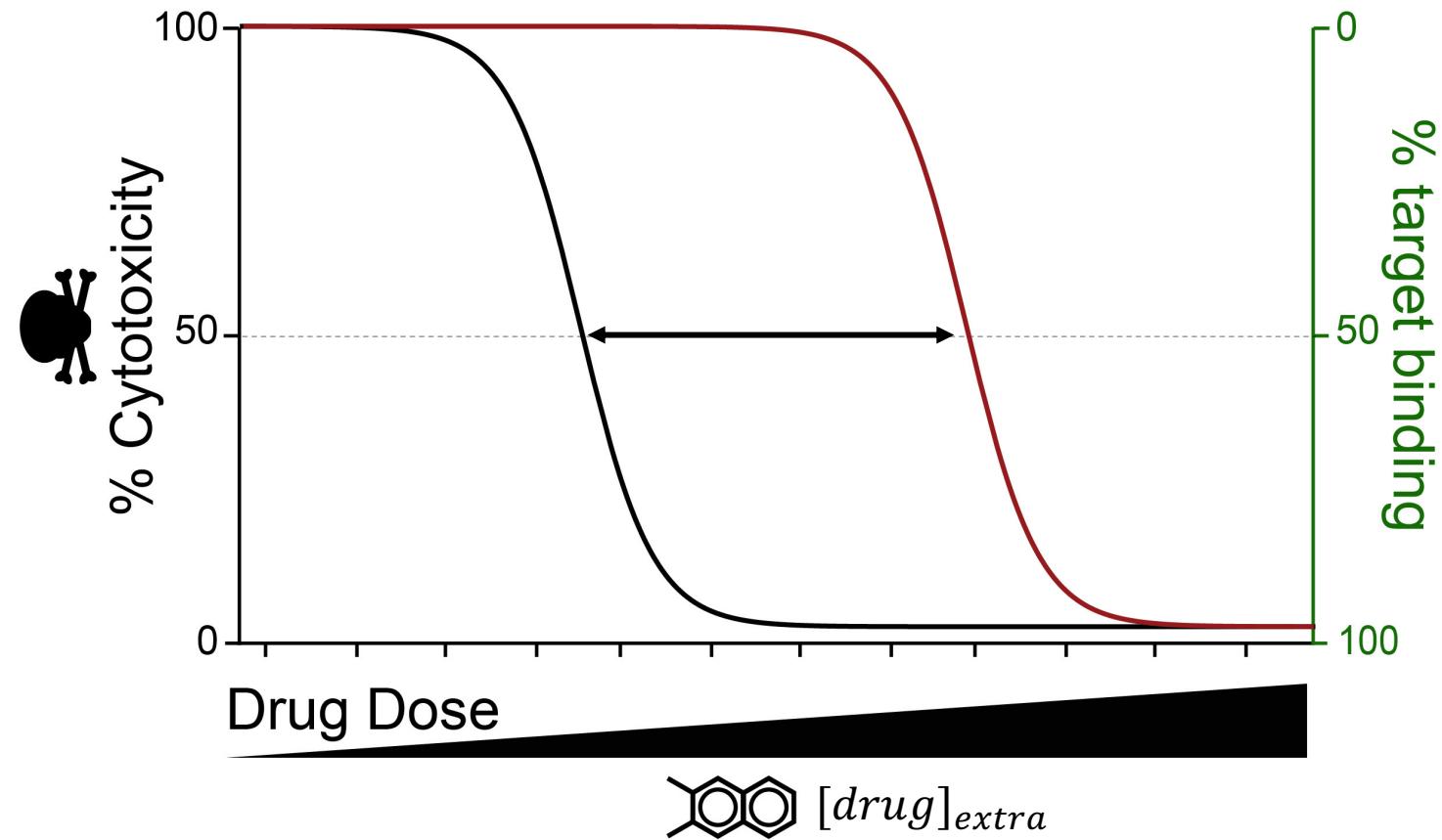
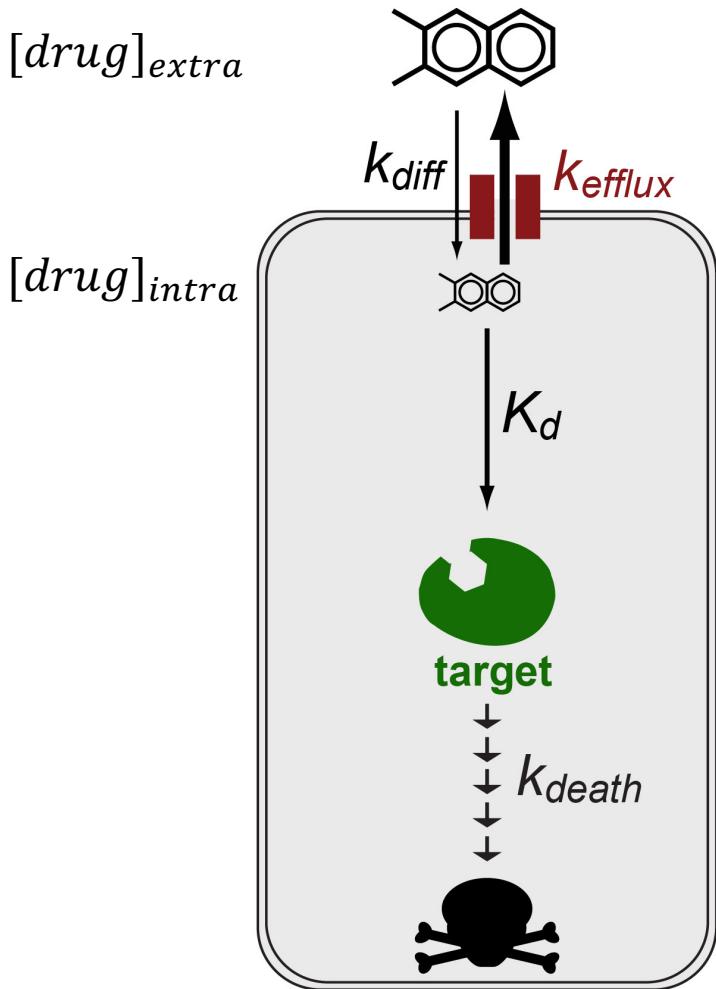
Profiling Relative Inhibition Simultaneously  
in Mixtures(PRISM)  
**1,448 drugs x 480 cell-lines**  
*Nature Cancer*, 2020, 1, 235



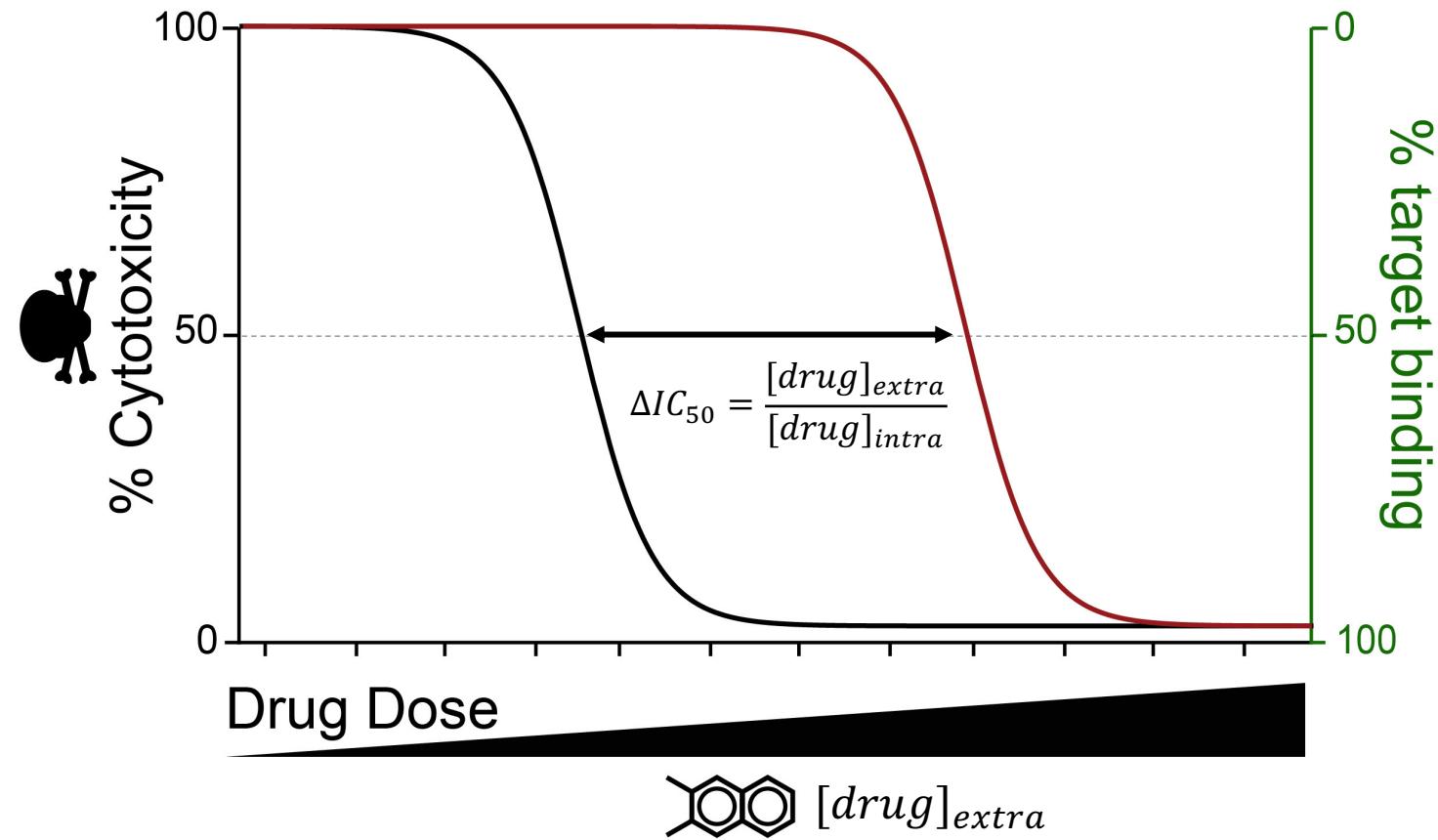
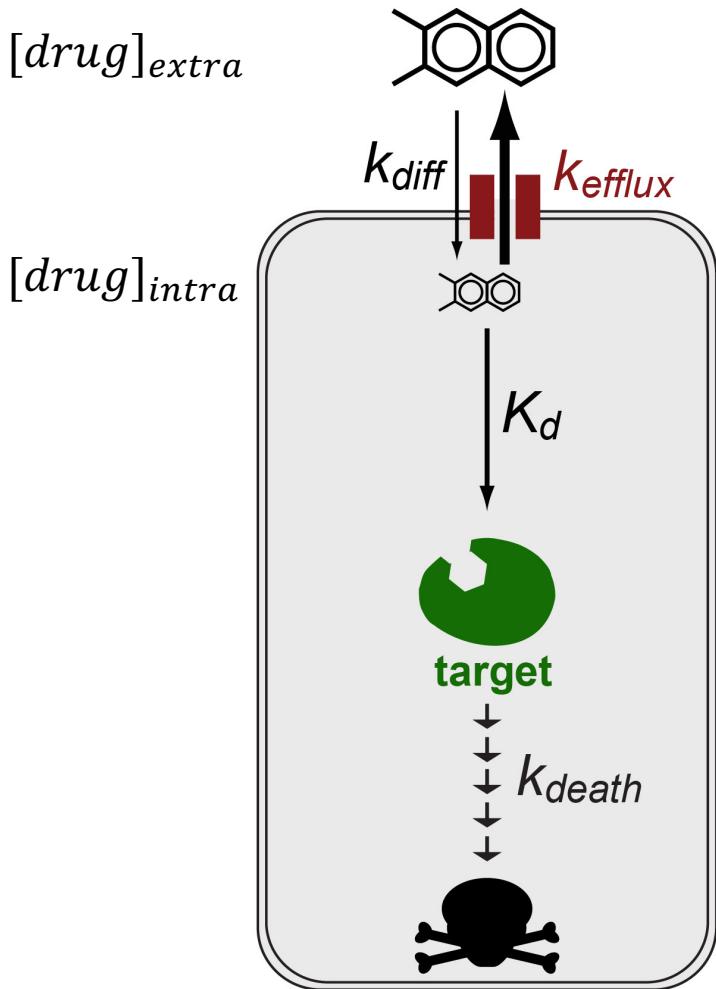
# Drug Resistance: Laboratory Data



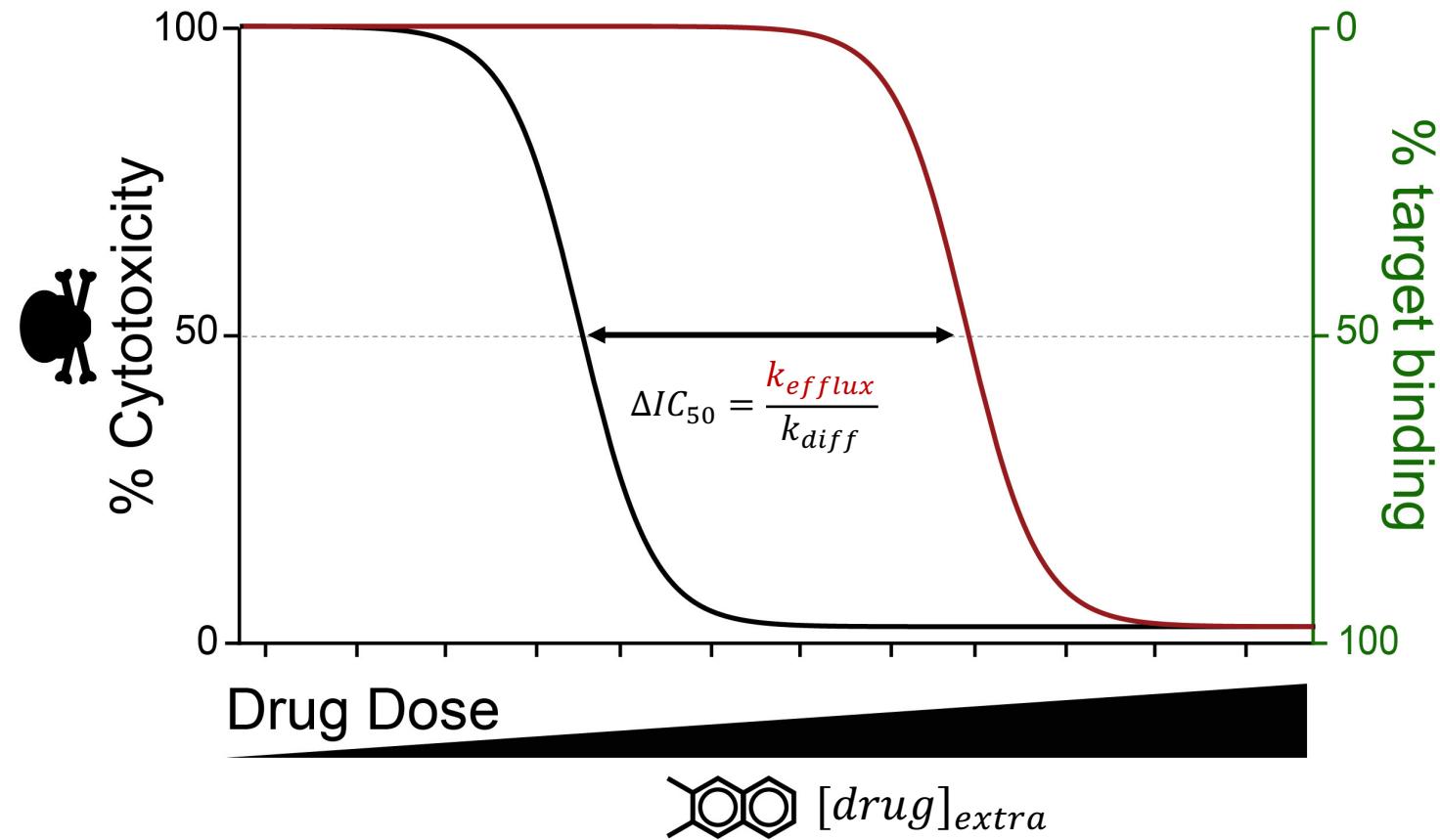
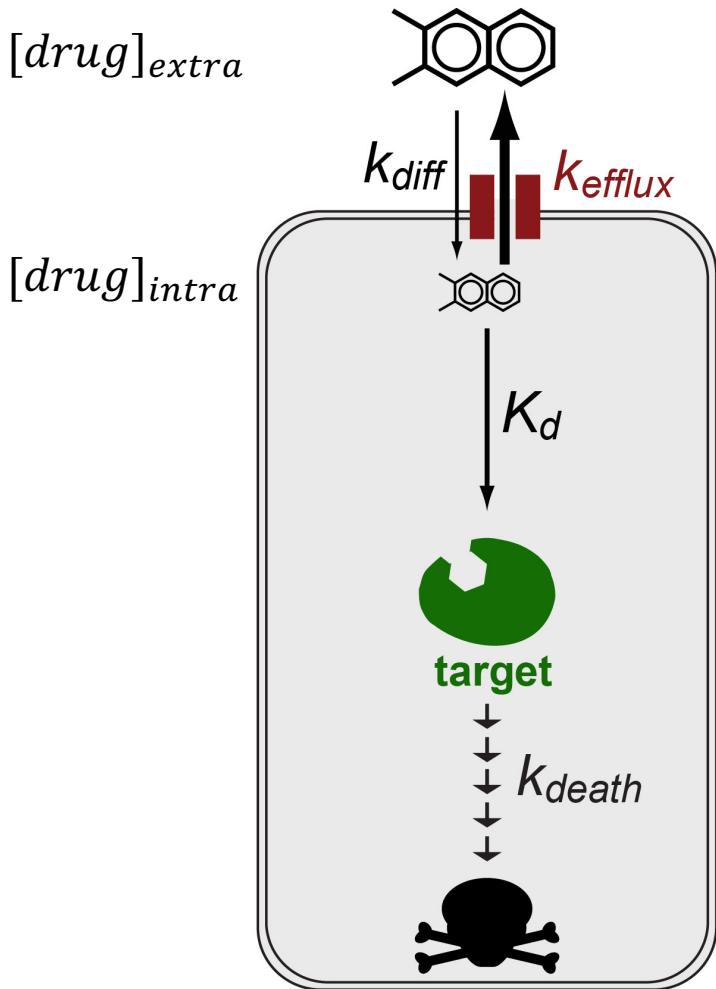
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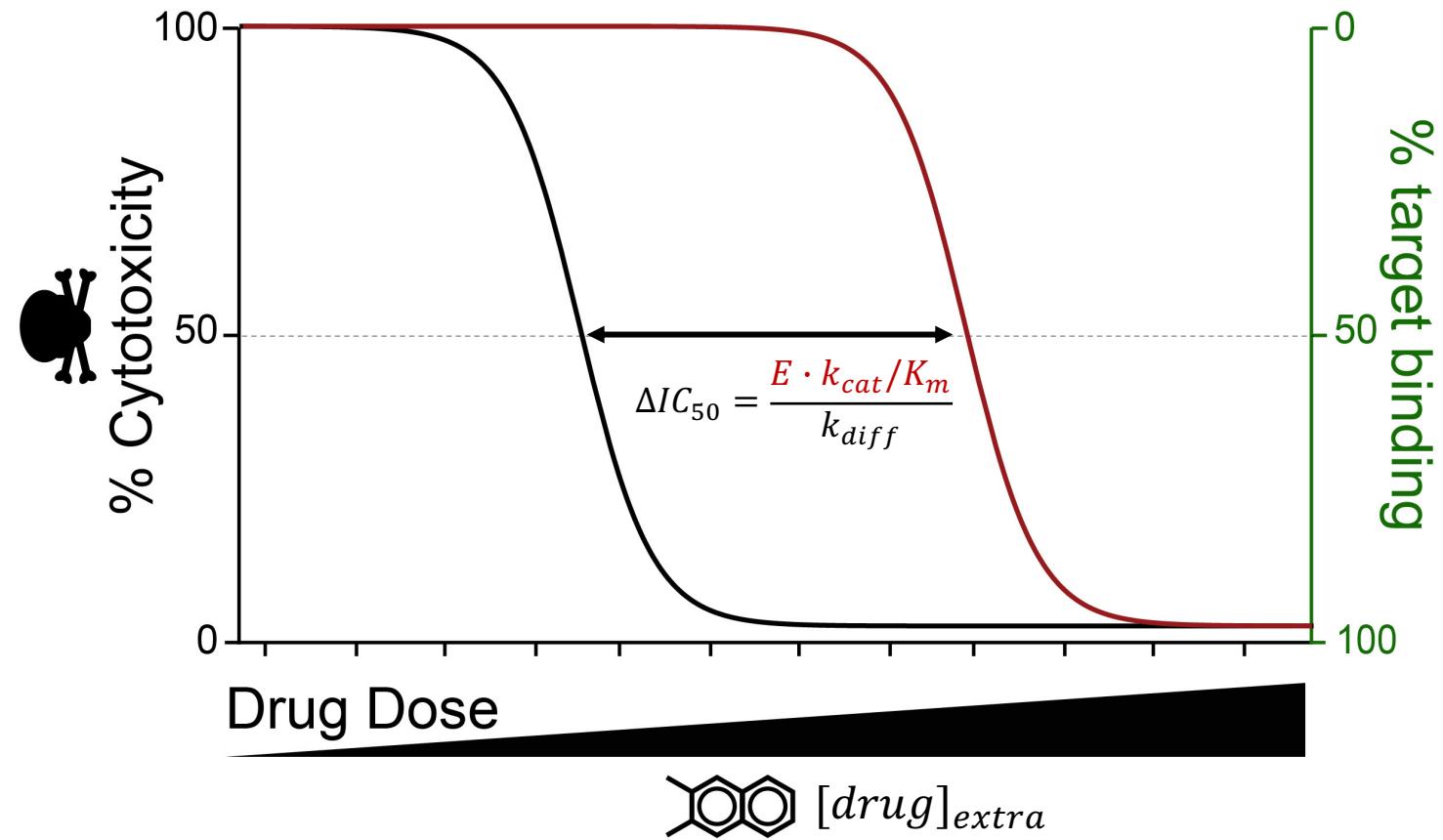
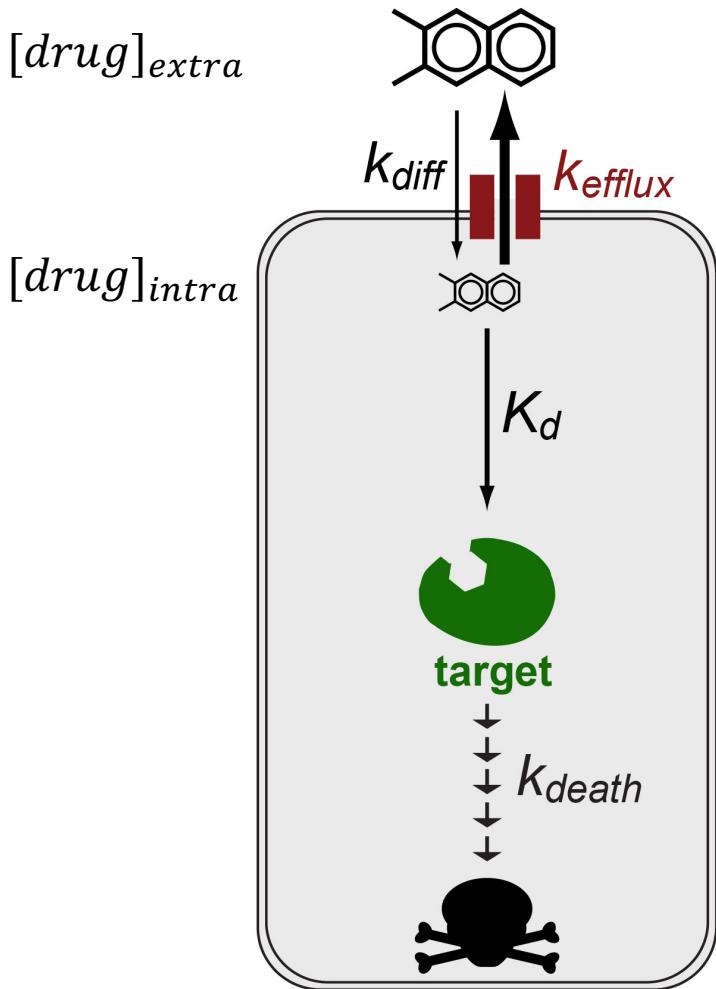
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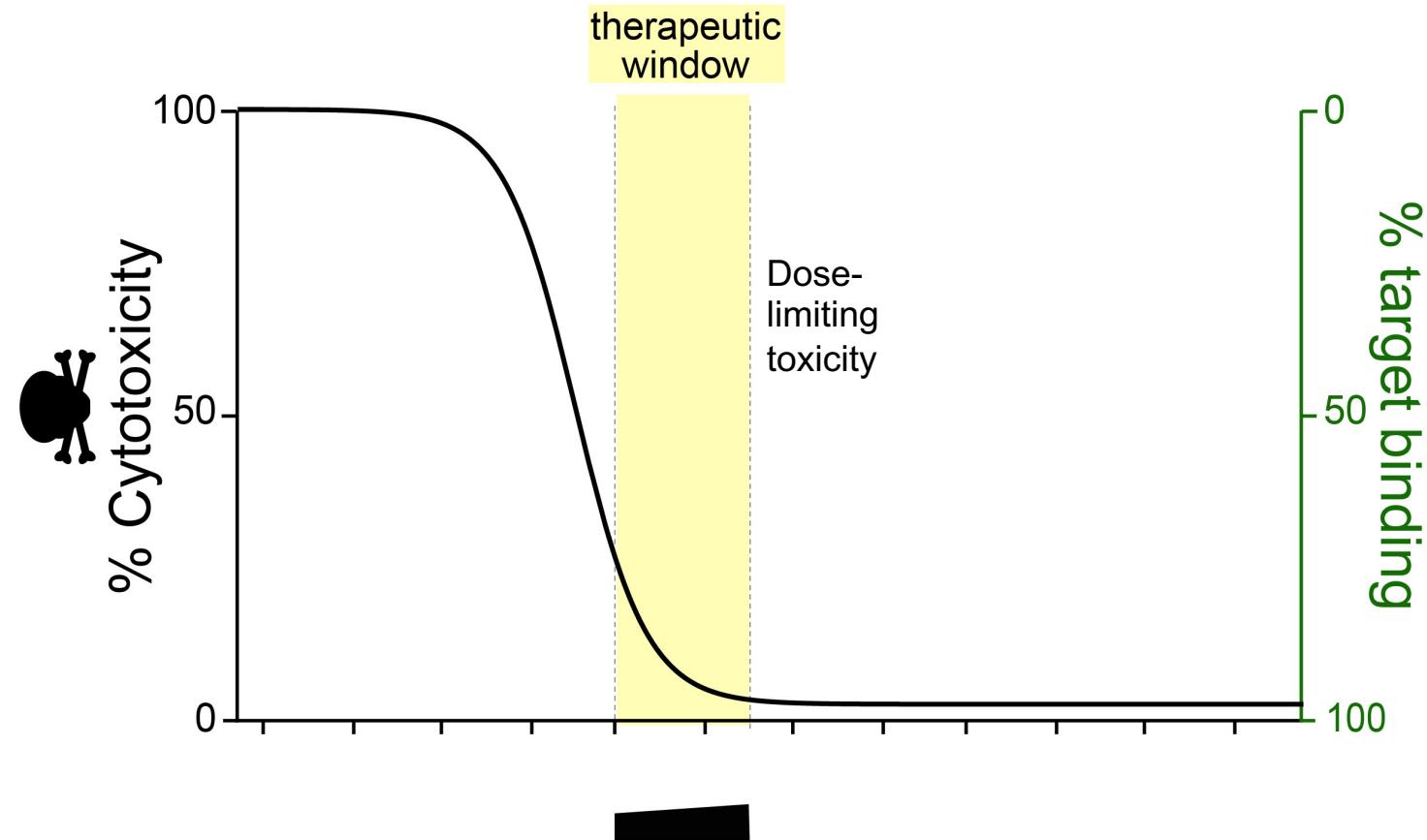
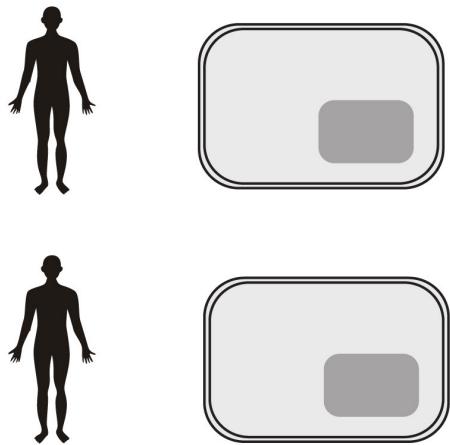
# Drug Resistance: Laboratory Data



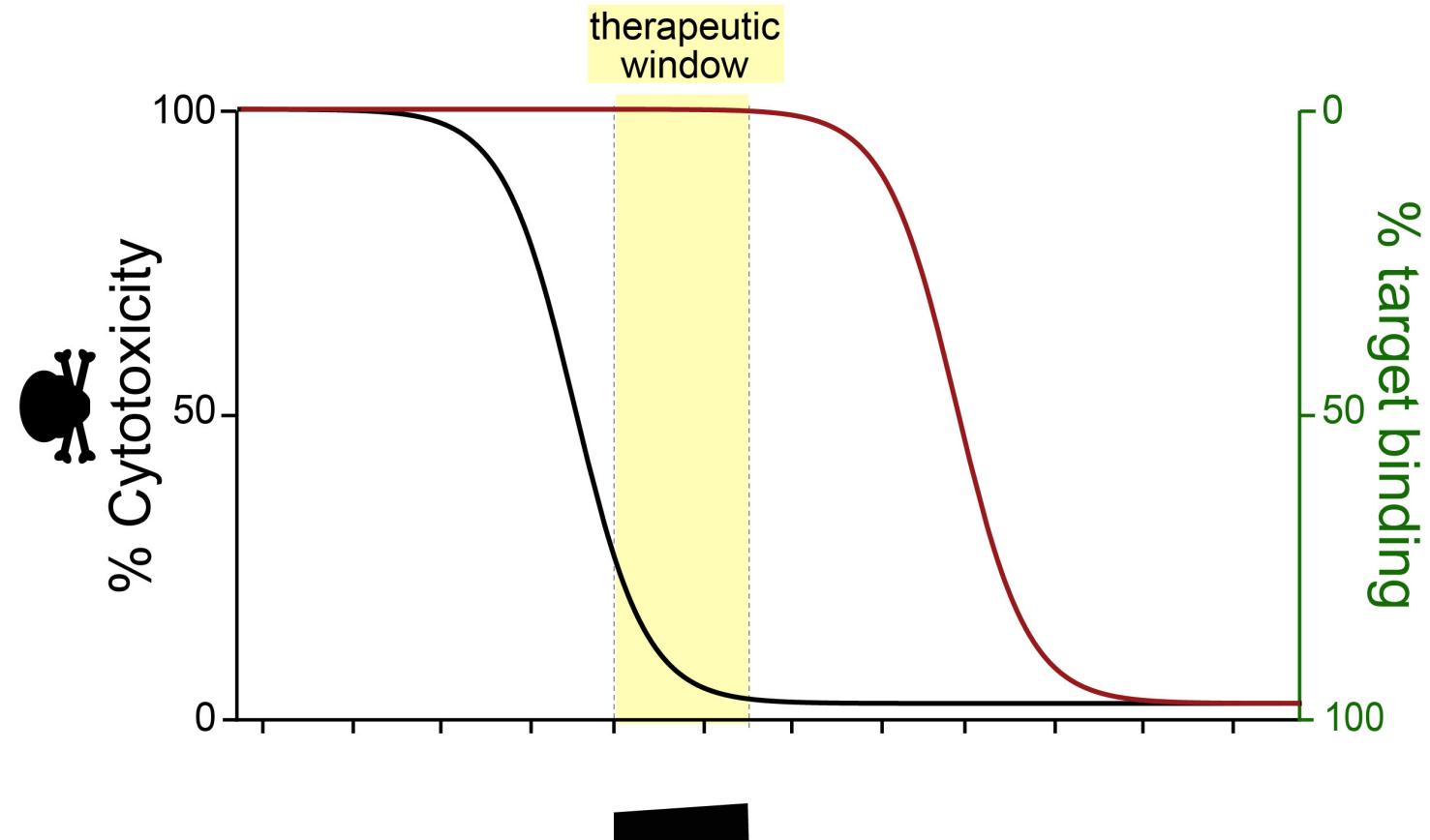
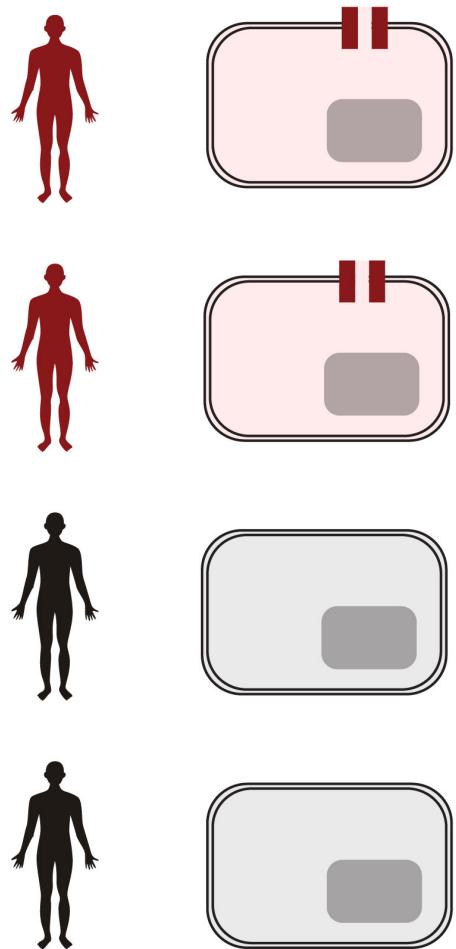
# Drug Resistance: Laboratory Data



# Drug Resistance: clinical response rates



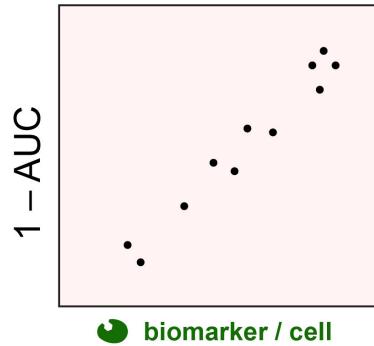
# Drug Resistance: clinical response rates



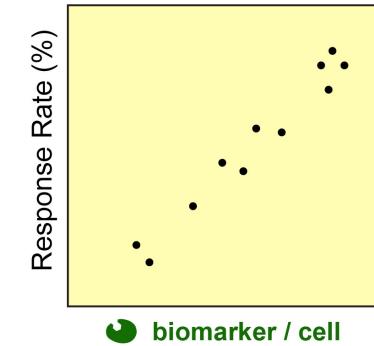
# Laboratory vs Clinical Biomarker analysis:



*in vitro* data sets



*clinical* data sets



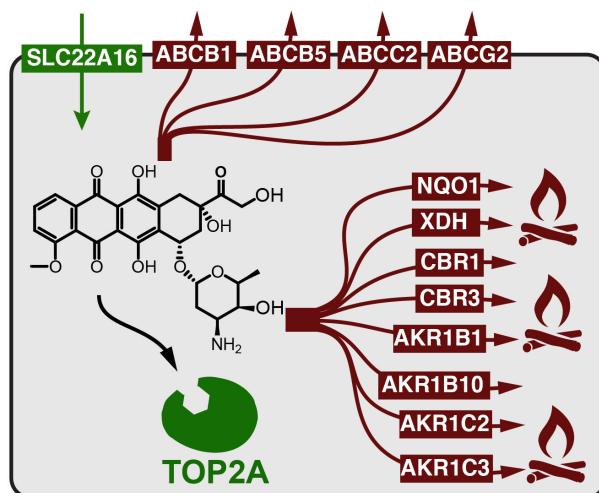
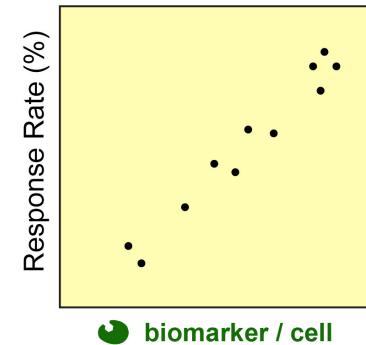
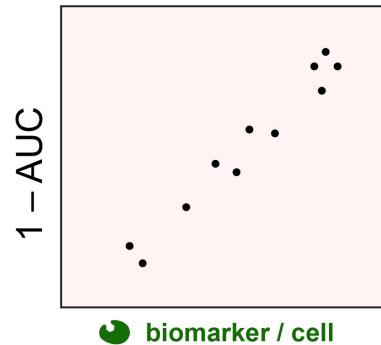
# Cytotoxic chemotherapies: 1-drug / N-biomarkers



*in vitro* data sets



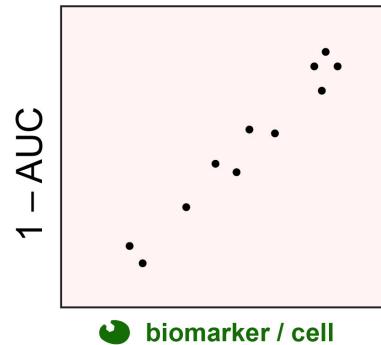
*clinical* data sets



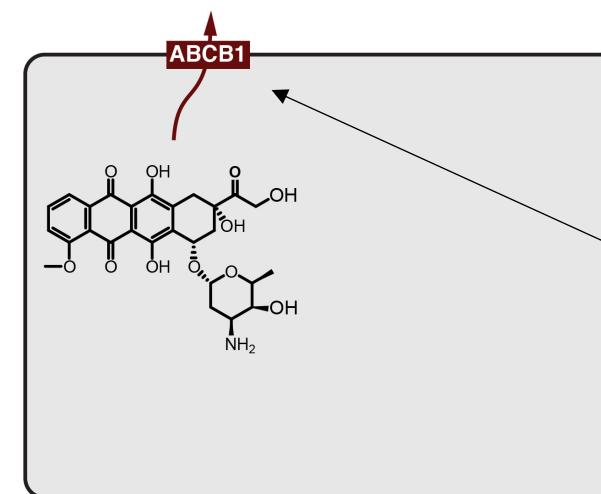
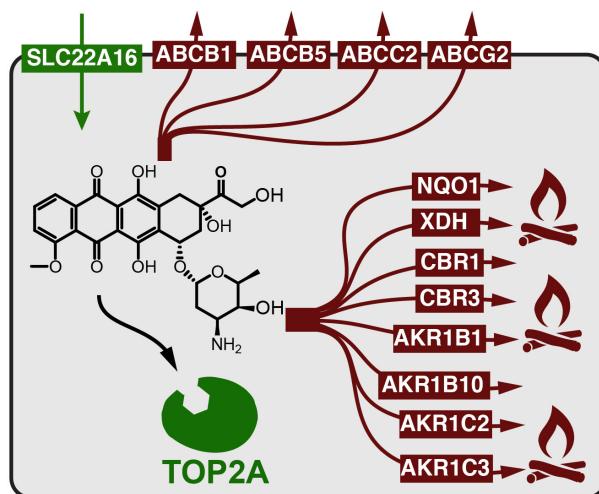
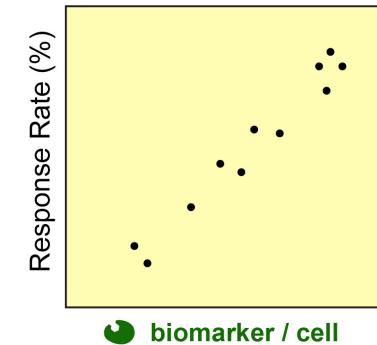
# Cytotoxic chemotherapies: 1-drug / N-biomarkers



*in vitro* data sets



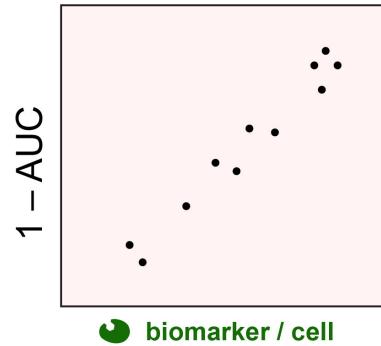
clinical data sets



# Cytotoxic chemotherapies: 1-drug / N-biomarkers



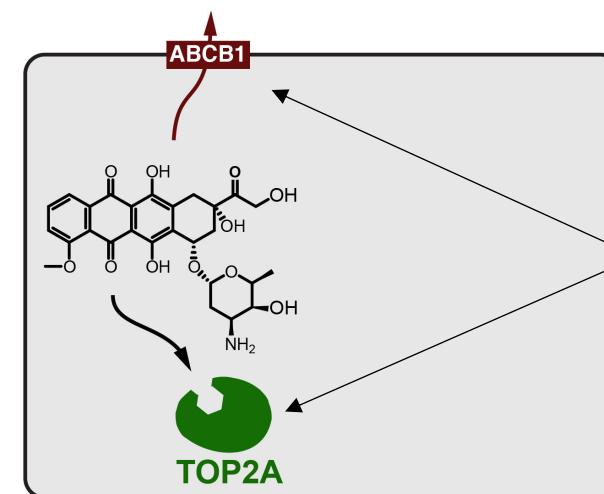
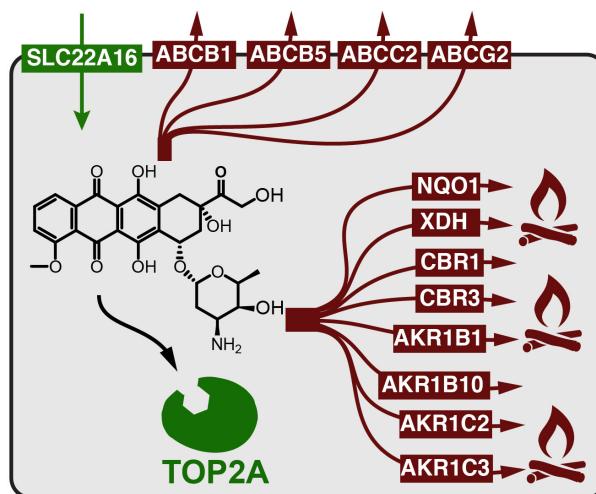
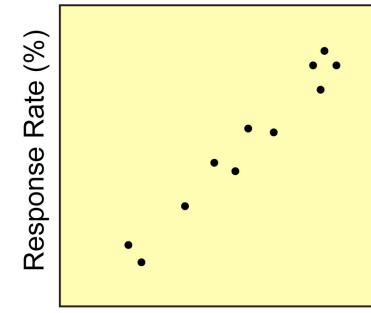
*in vitro* data sets



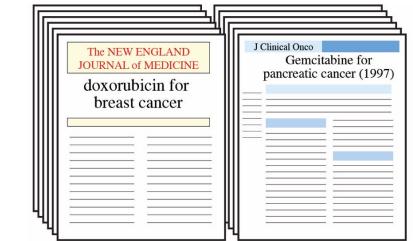
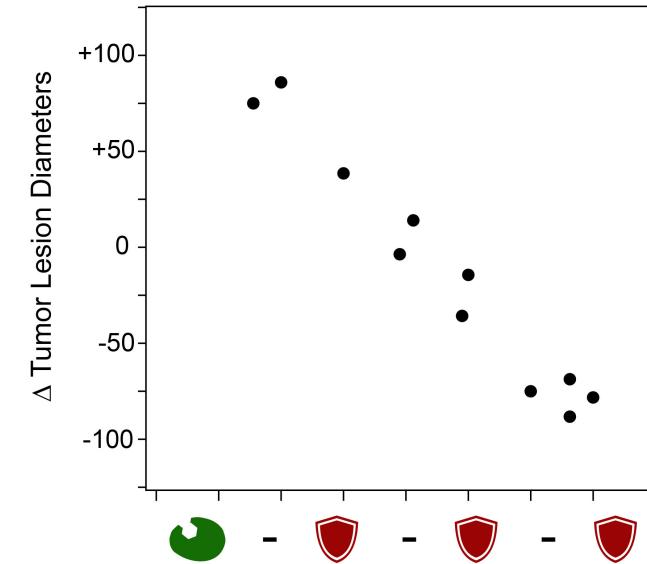
biomarker / cell



*clinical* data sets

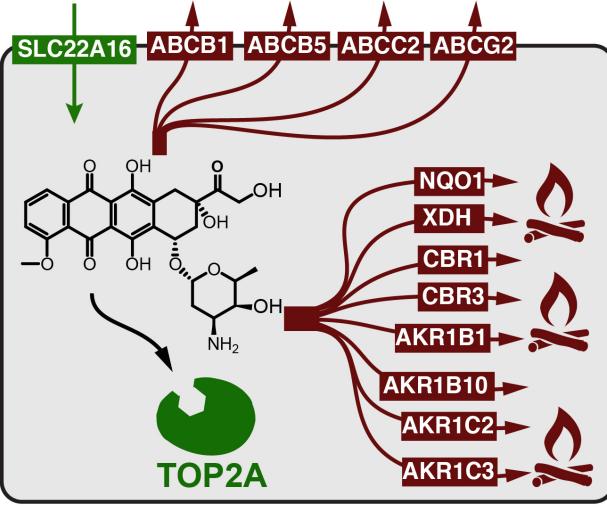
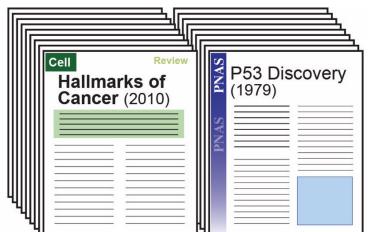


# How do you combine N-biomarkers?

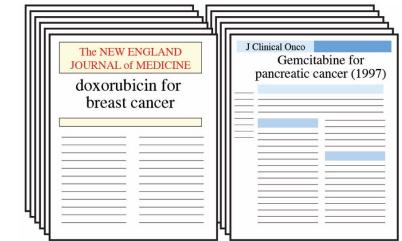
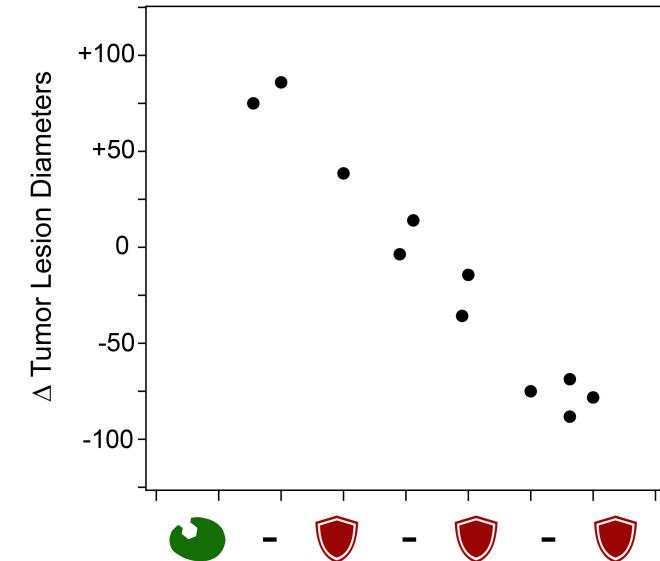


Clinical Literature

# How do you combine N-biomarkers?

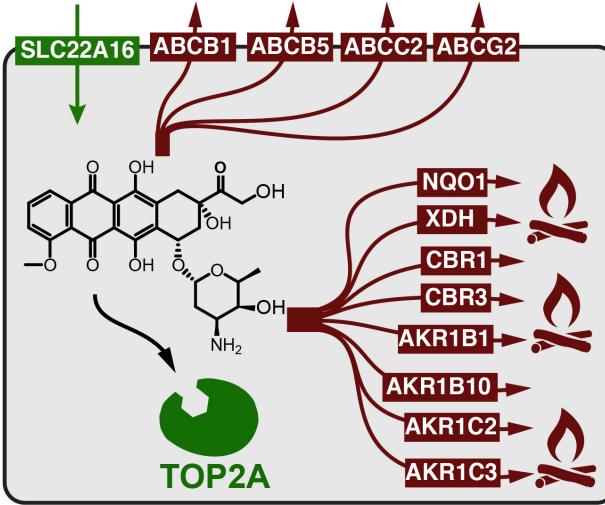
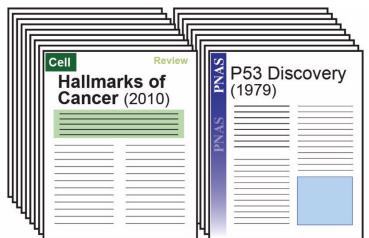


Scientific Literature

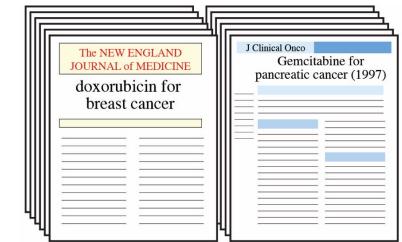
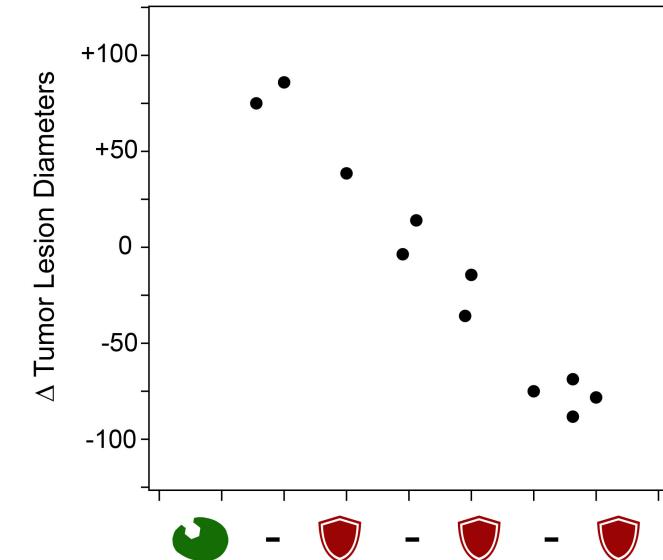


Clinical Literature

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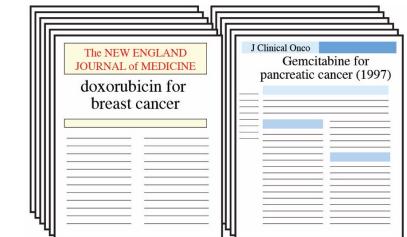
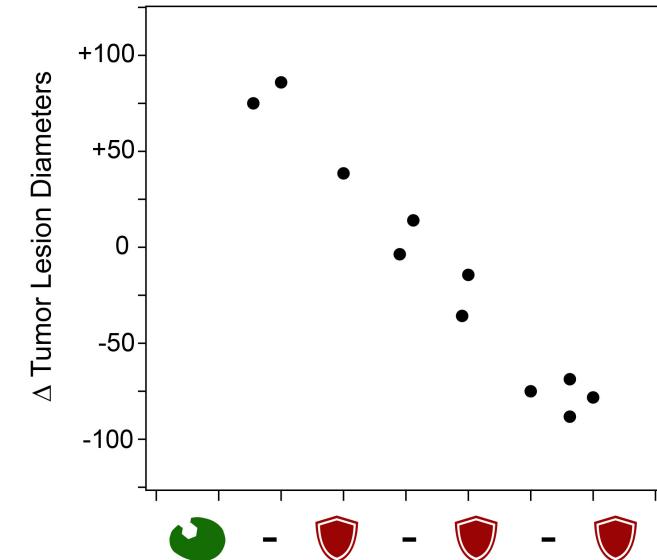
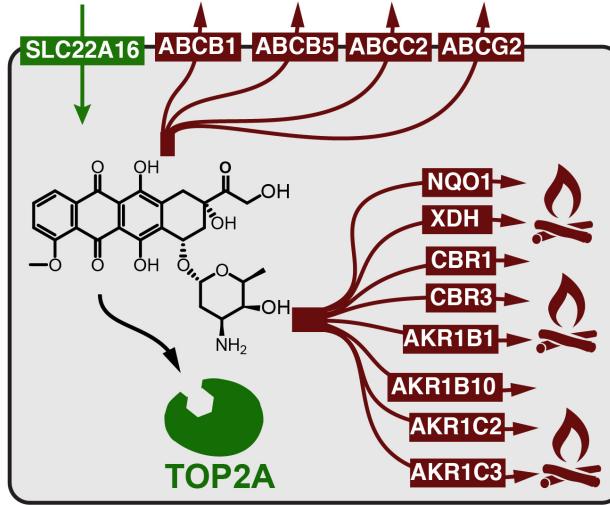
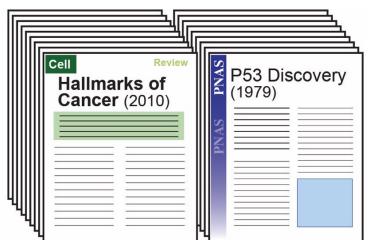
Scientific Literature



Clinical Literature

$$efflux = \sum_{i=1}^N k_{cat}[E]_t \frac{[drug]_{intra}}{[drug]_{intra} + K_m}$$

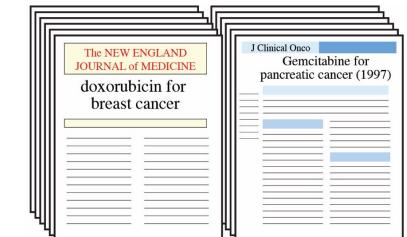
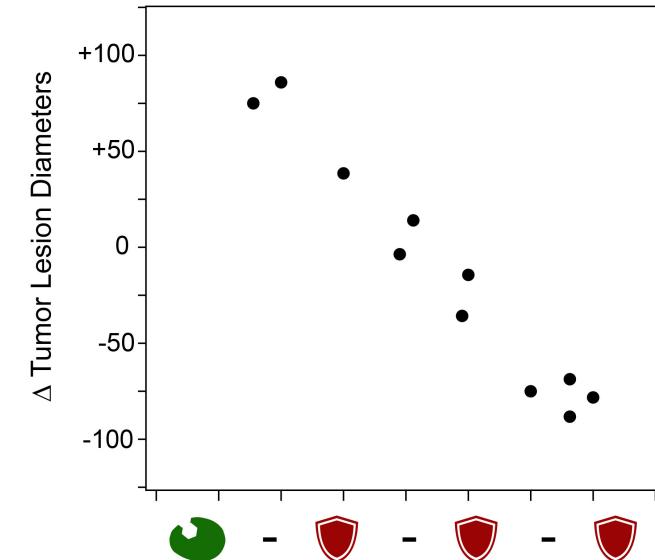
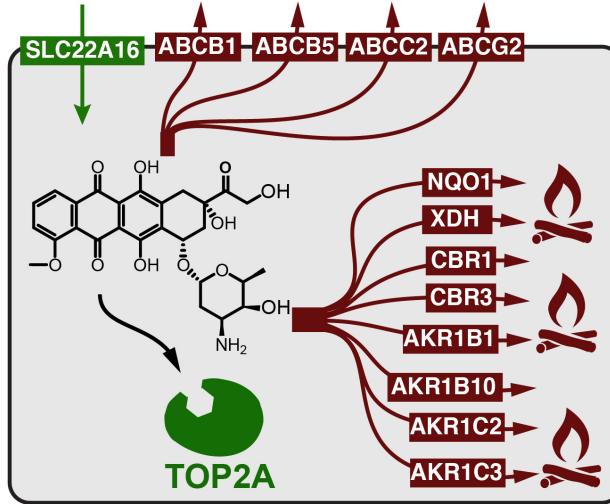
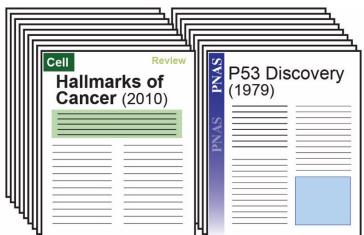
# How do you combine N-biomarkers?



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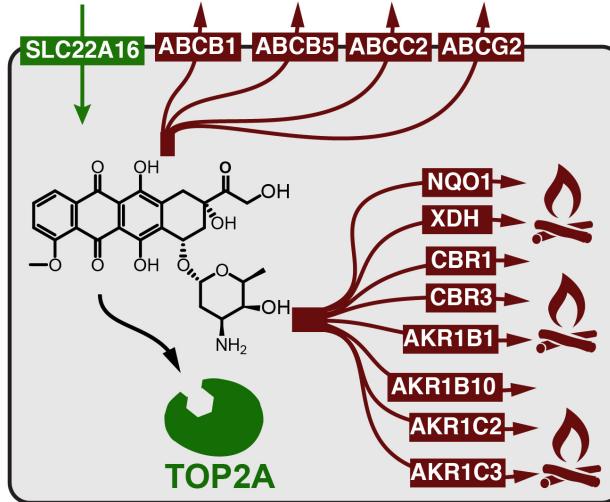
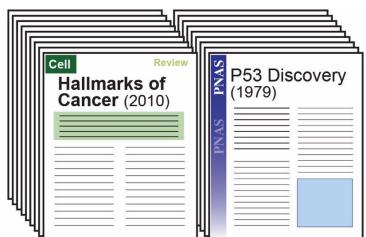


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$$efflux = k_{resist} [drug]_{intra}$$

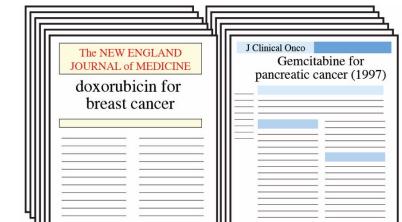
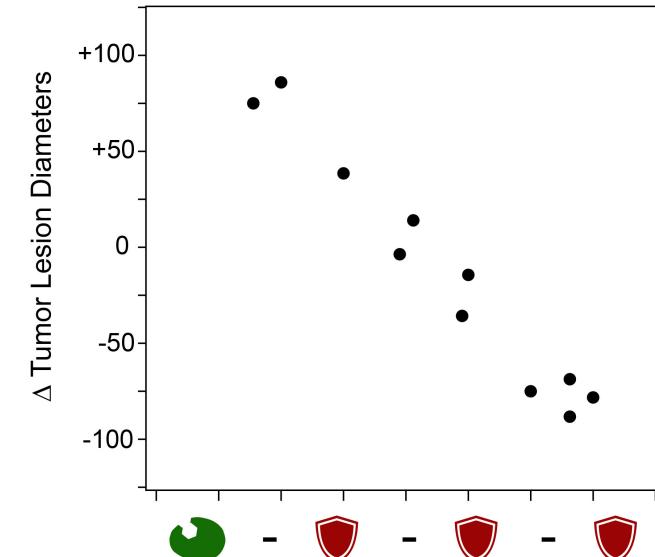
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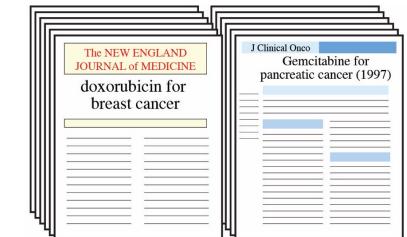
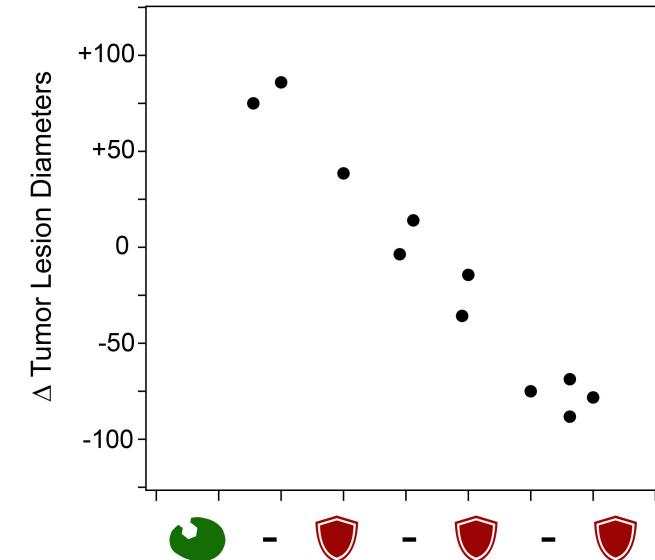
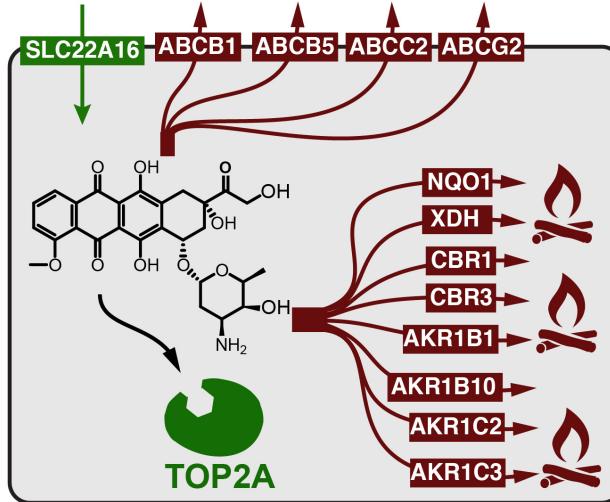
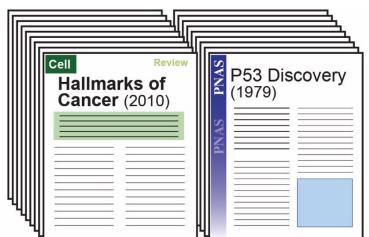
$$efflux = k_{resist}[drug]_{intra}$$



$$influx = efflux$$

$$k_{diff}[drug]_{extra} = k_{resist}[drug]_{intra}$$

# How do you combine N-biomarkers?



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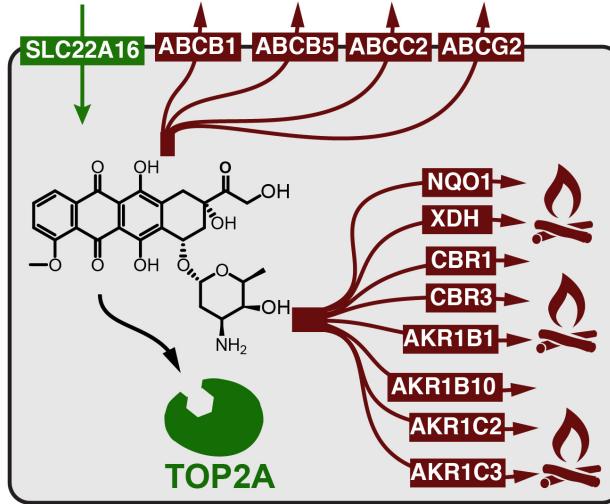
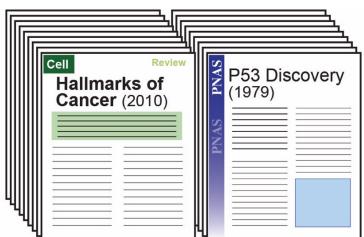
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$$\frac{[drug]_{extra}}{[drug]_{intra}} = \frac{k_{resist}}{k_{diff}}$$

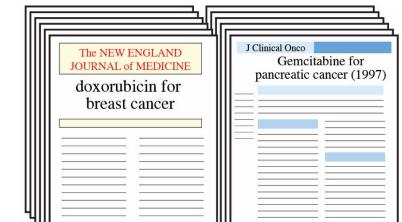
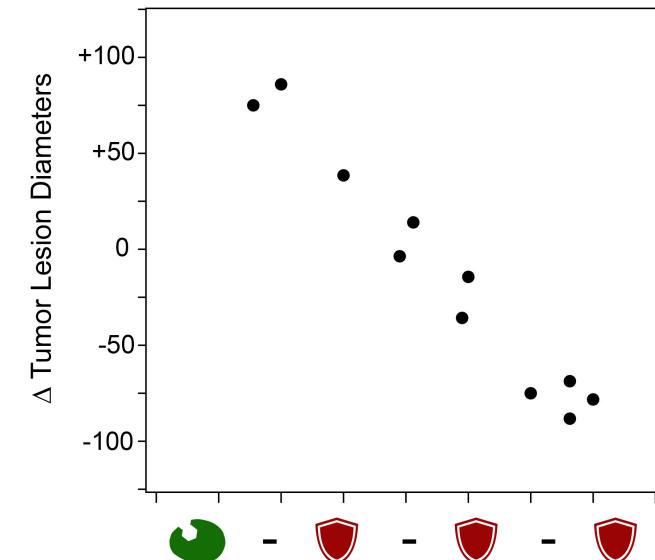
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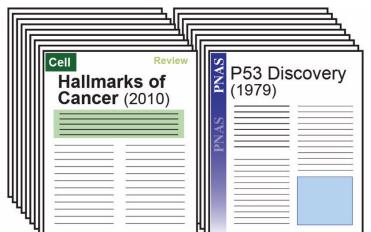


$$influx = efflux$$

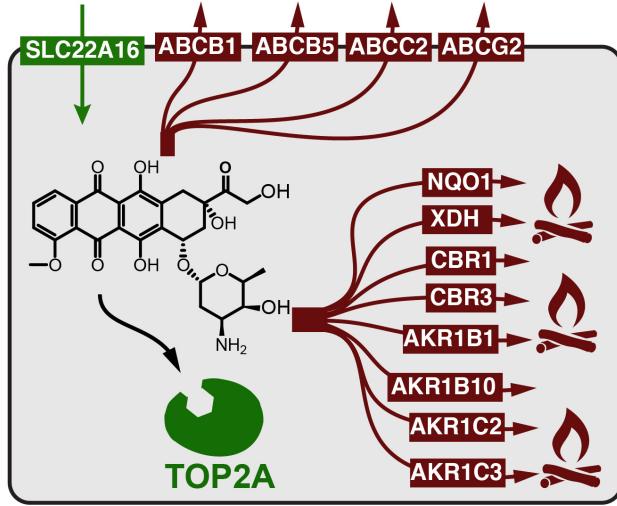
$$k_{diff} [drug]_{extra} = k_{resist} [drug]_{intra}$$

$$\Delta IC_{50} = \frac{[drug]_{extra}}{[drug]_{intra}} = \frac{k_{resist}}{k_{diff}}$$

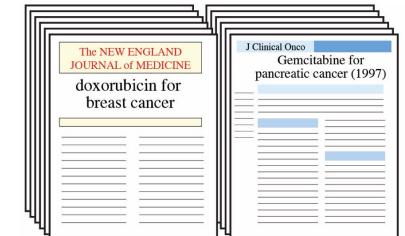
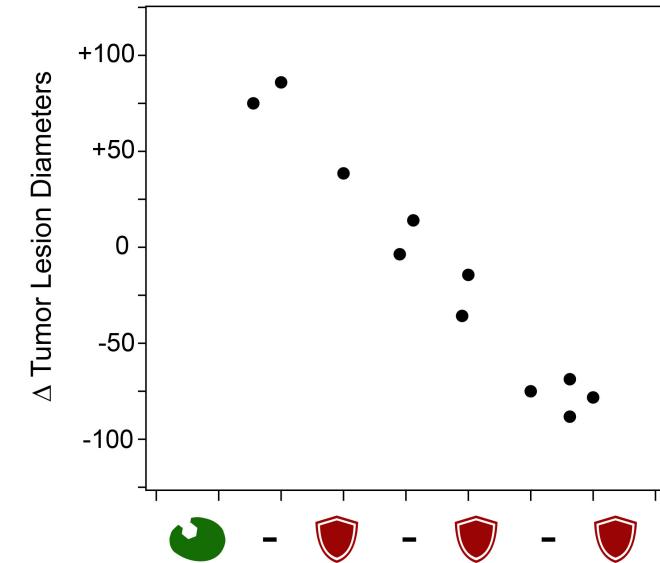
# How do you combine N-biomarkers?



Scientific Literature



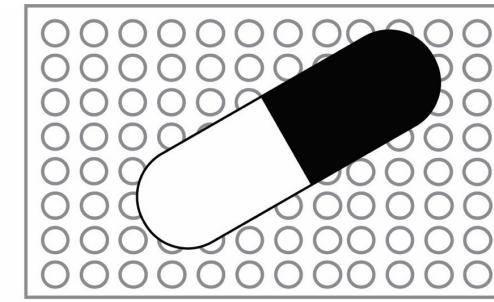
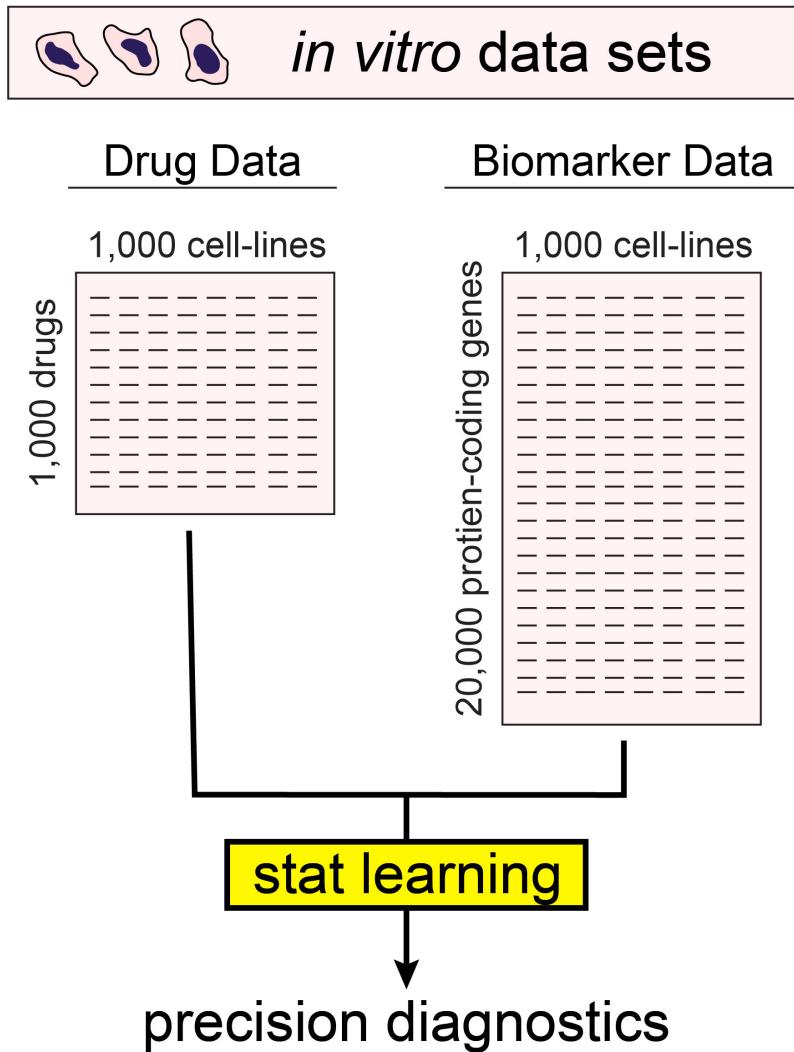
$$\Delta IC_{50} = \frac{k_{cat}/K_m}{k_{diff}} [ABCB1] + \frac{k_{cat}/K_m}{k_{diff}} [ABCG2] \dots$$



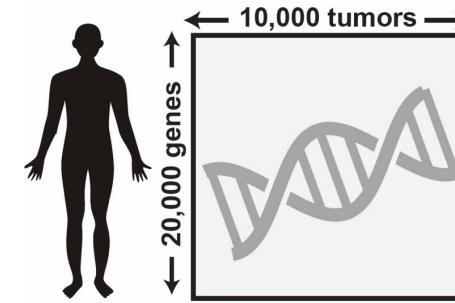
Clinical Literature

$$\Delta IC_{50} = \frac{k_{resist}}{k_{diff}} = \sum_{i=1}^N \frac{k_{cat}/K_m}{k_{diff}} [E]_t$$

# Previous Pharmacogenomic Learning:

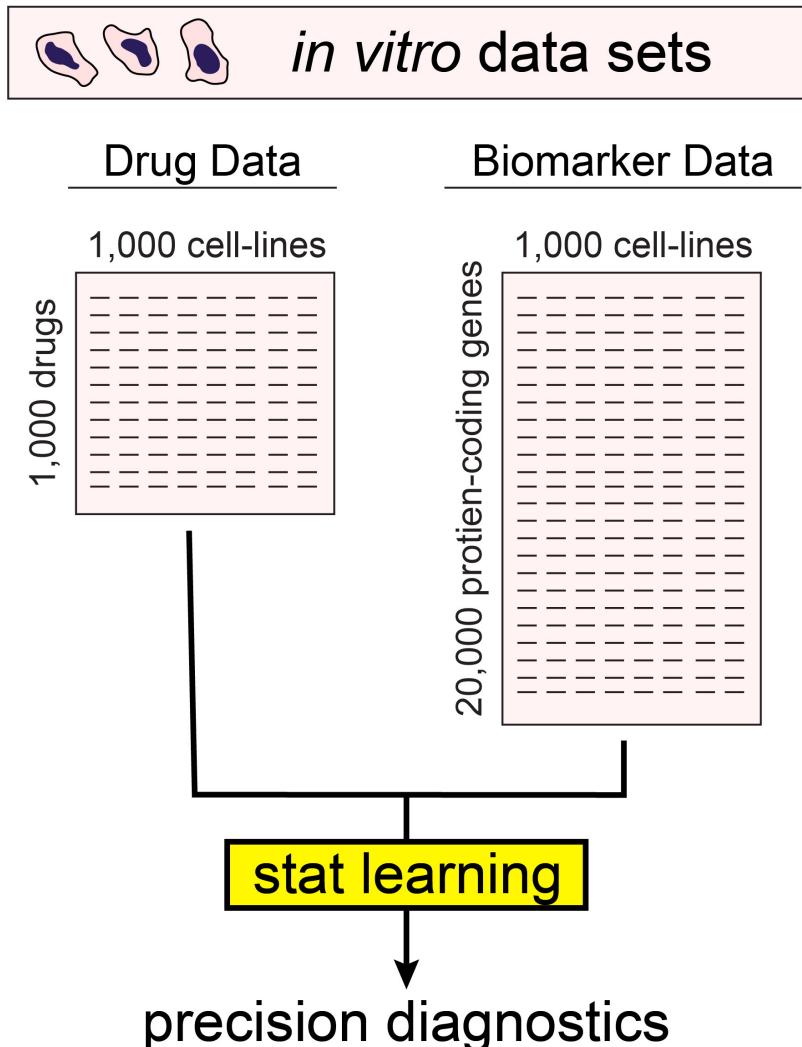


Screening Technology



Sequencing Tech

# Previous Pharmacogenomic Learning:

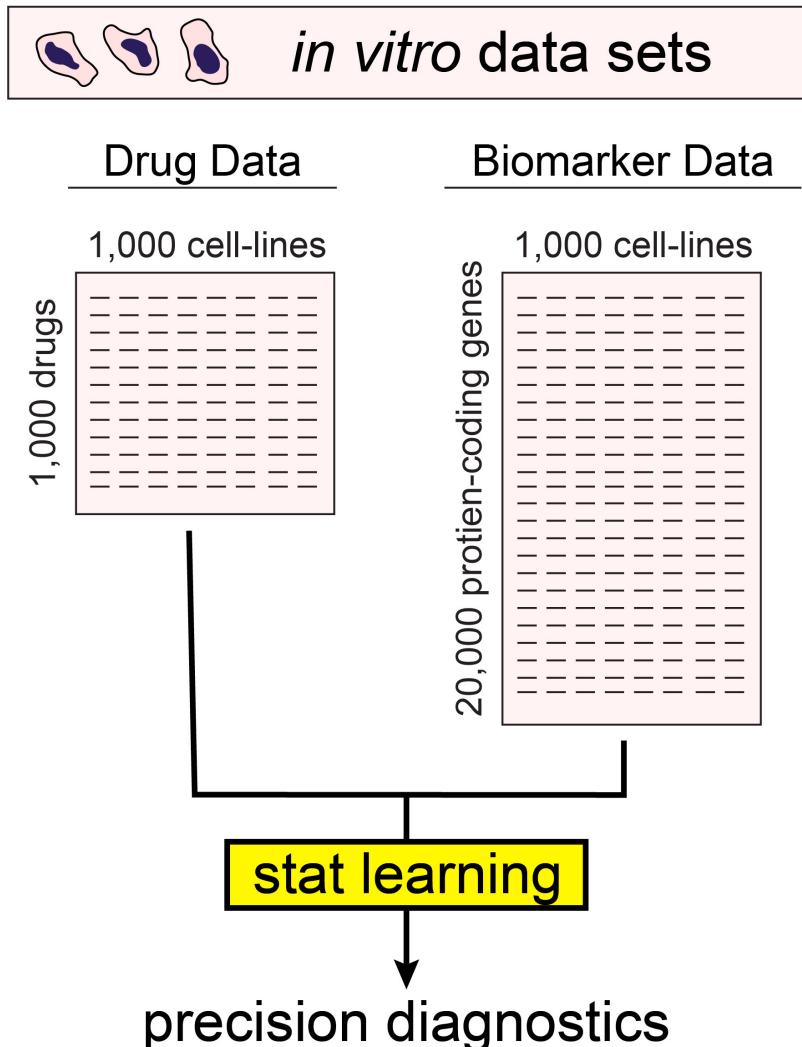


*In a pan-cancer analysis, the most predictive data type was gene expression, closely followed by the tissue of origin of the cell lines. By comparison, genomic features performed poorly.*

*Cell*, 2016, 166, 740–754

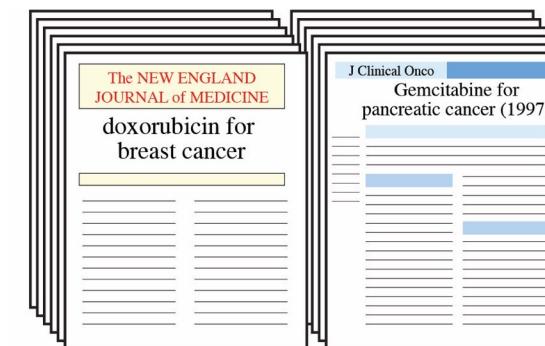
*Cell*, 2016, 166, 740  
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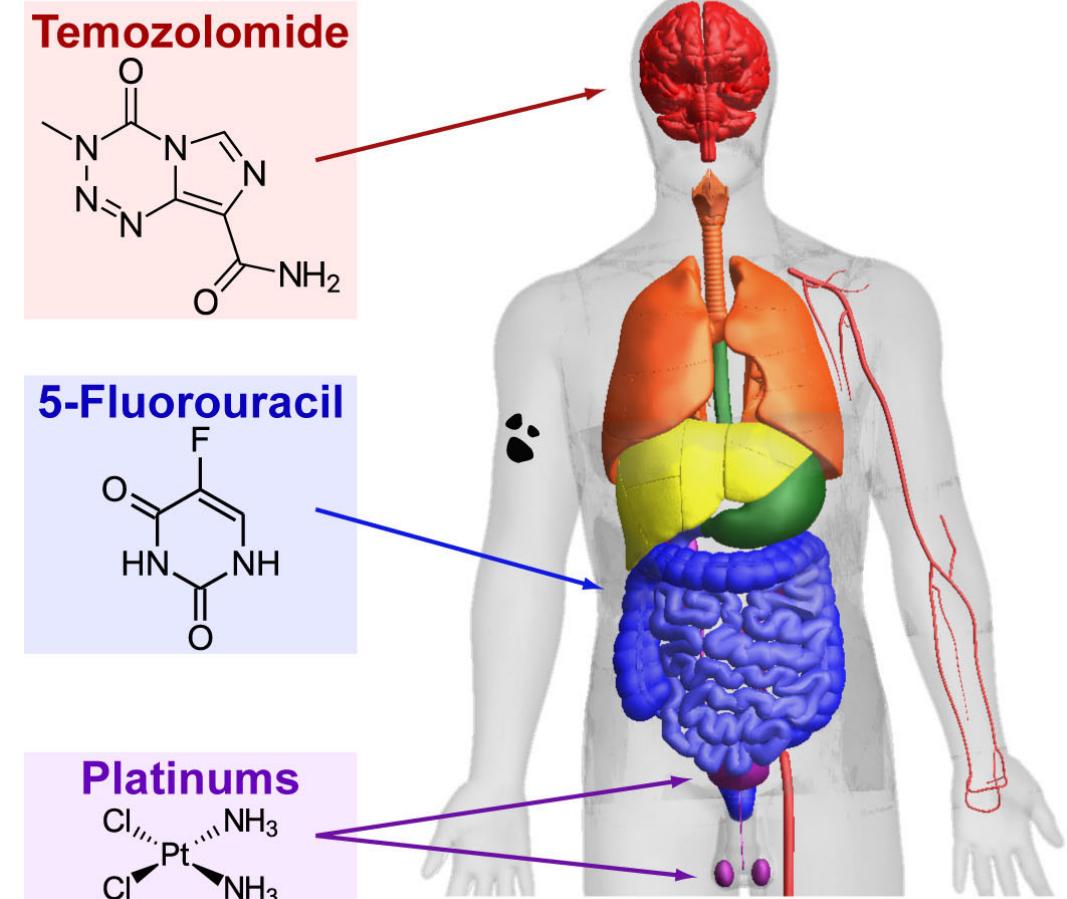
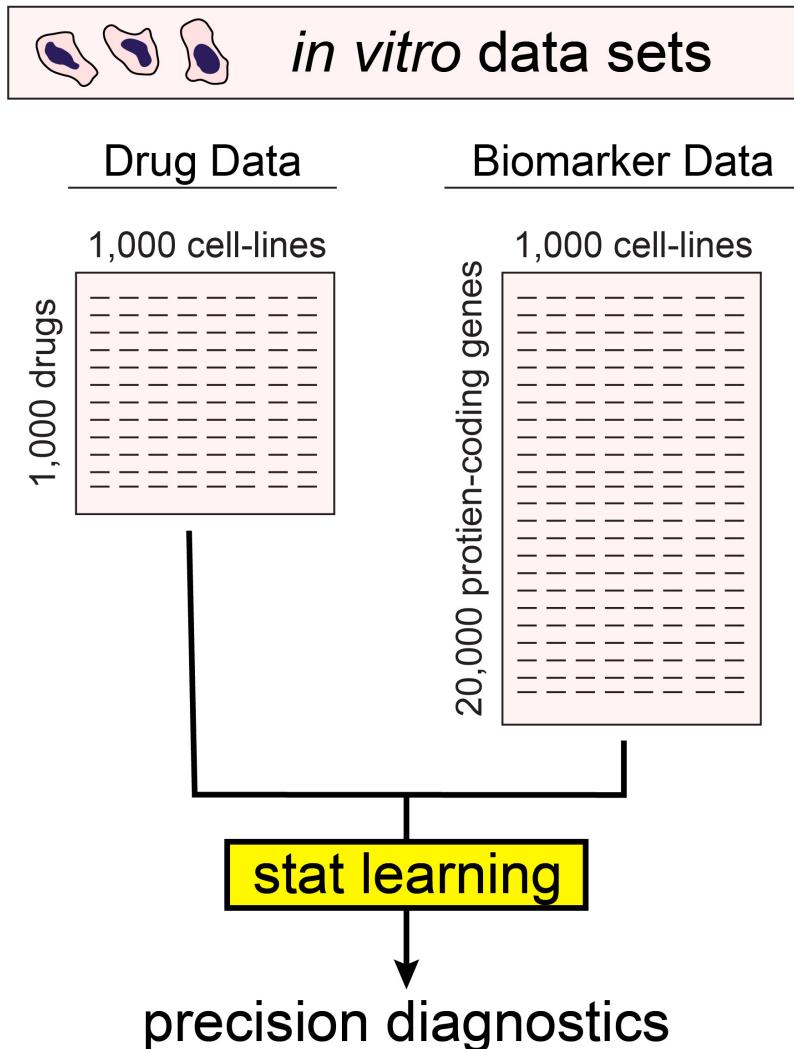
*Cell*, 2016, 166, 740–754, 79



Clinical Literature

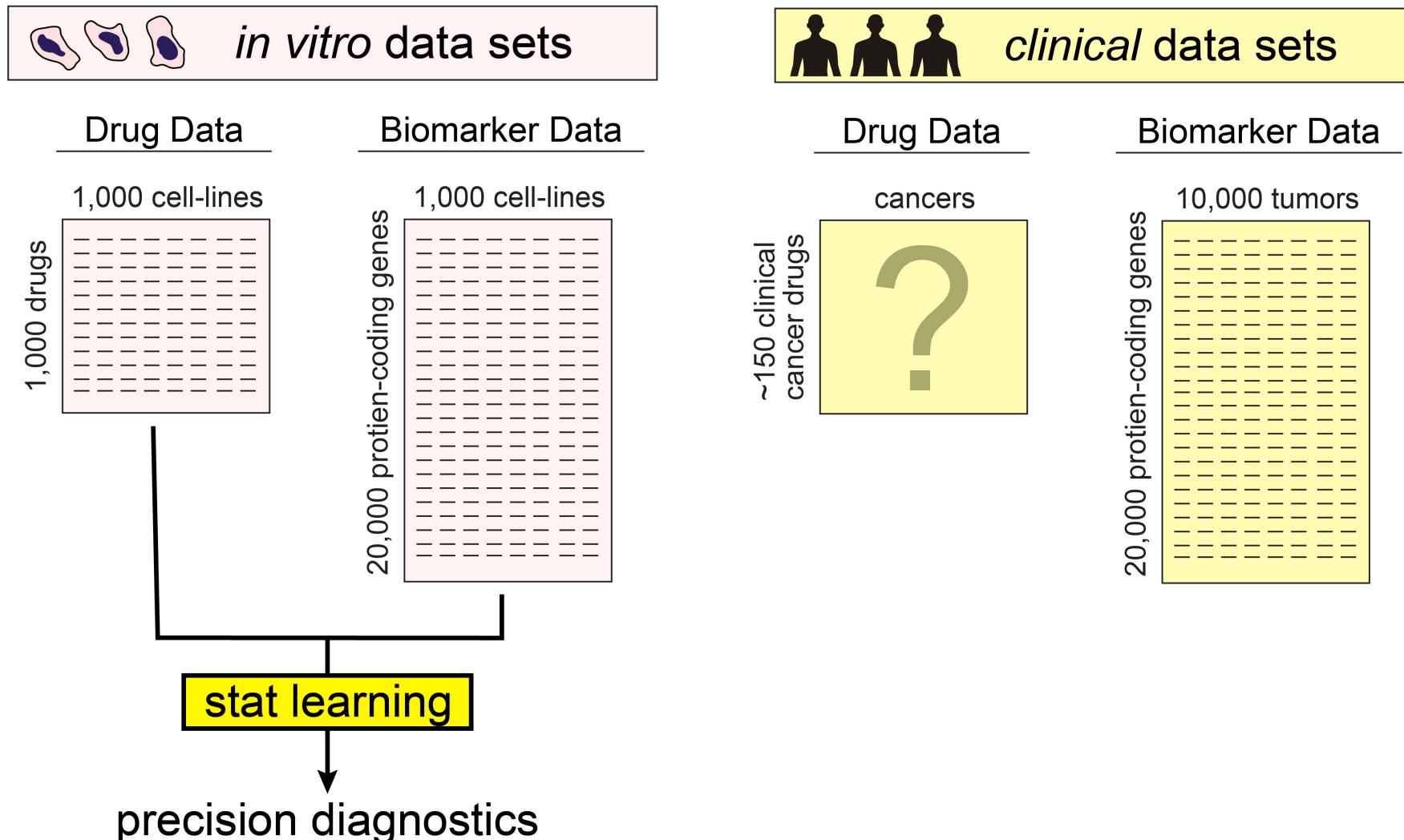
Drug Sensitivity(GDSC): *Cell*, 2016, 166, 740  
Drug Sensitivity(CTRP): *Cell*, 2013, 154, 1151

# Previous Pharmacogenomic Learning:

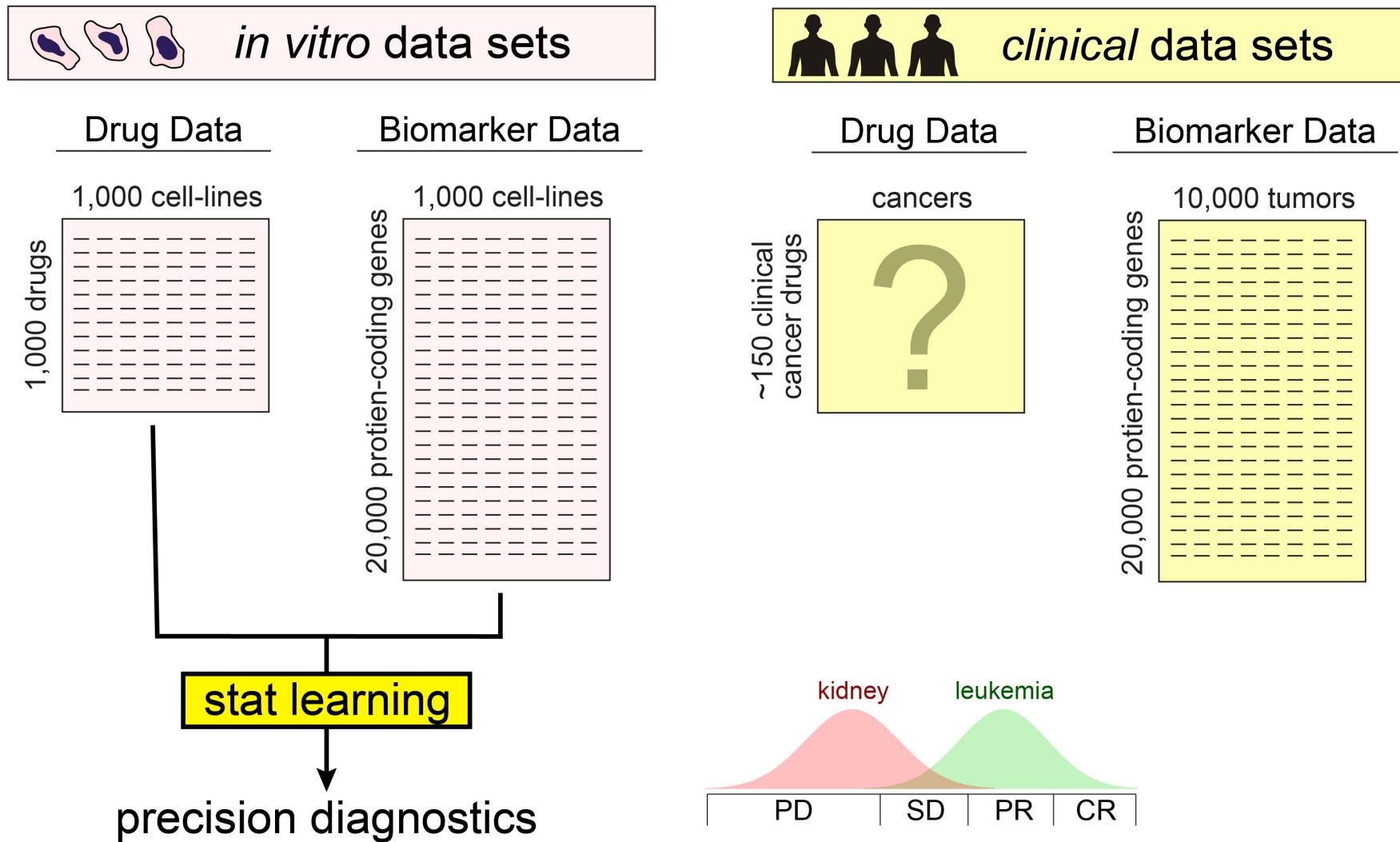


Drug Sensitivity(GDSC): *Cell*, 2016, 166, 740  
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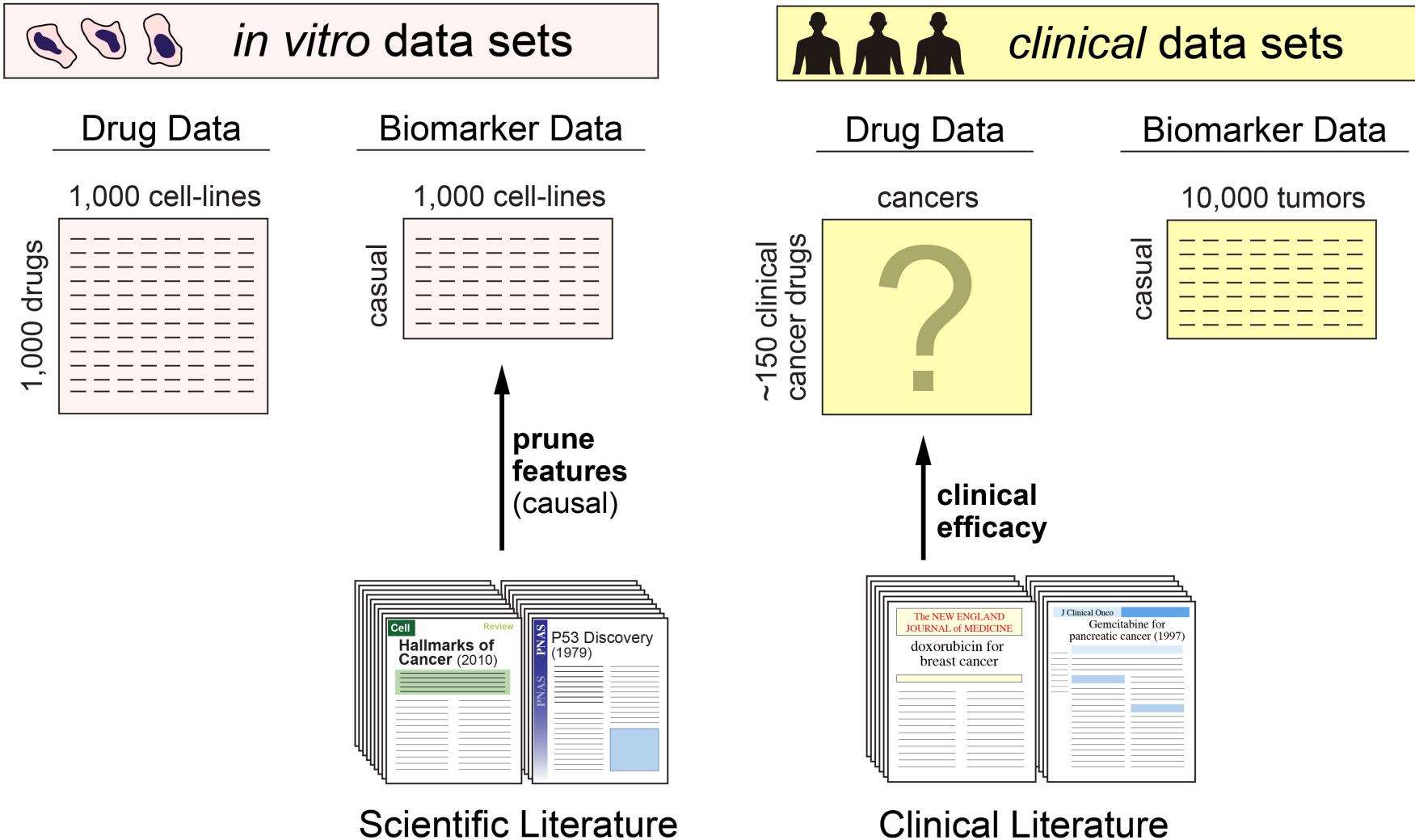
# Data-gap within clinical literature:



# Data-gap within clinical literature:



# Douglass Lab Approach: prune features & build database



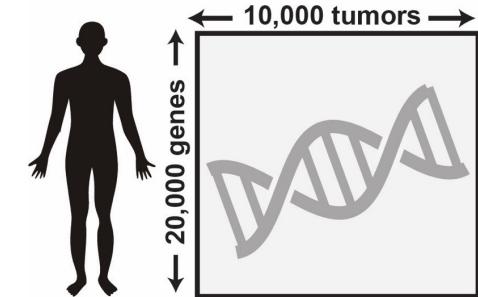
# Clinical “Big Data” + Experiment = Precision Medicine

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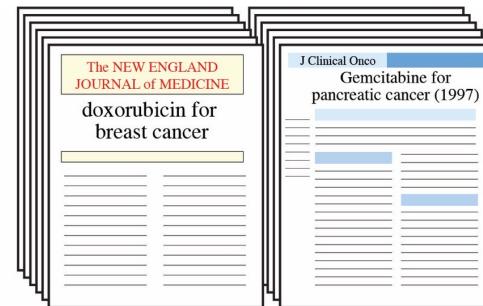
Genomic vs Experimental “Big Data:” *challenges and opportunities*

History & Current Status of Cancer Therapy: *1 drug / 1 target paradigm*

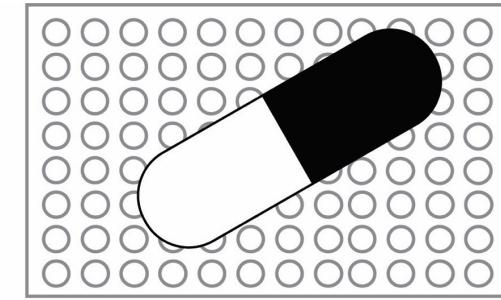
Reconciling Clinical & Laboratory “Big Data”: *1 drug / N-biomarkers*



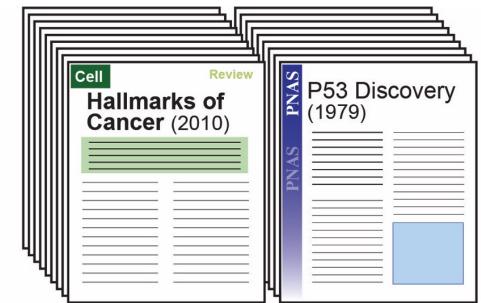
Sequencing Tech



Clinical Literature



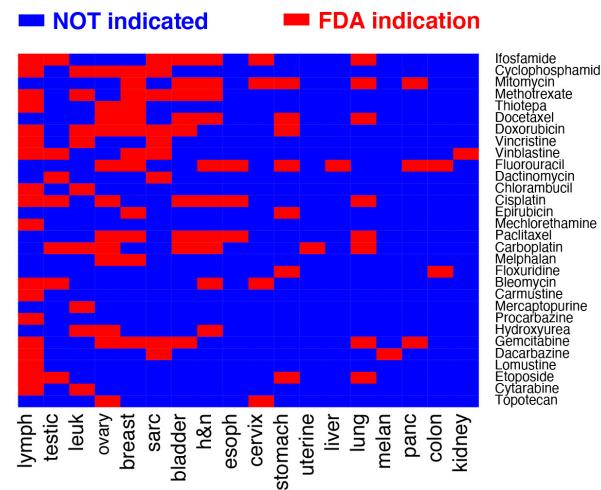
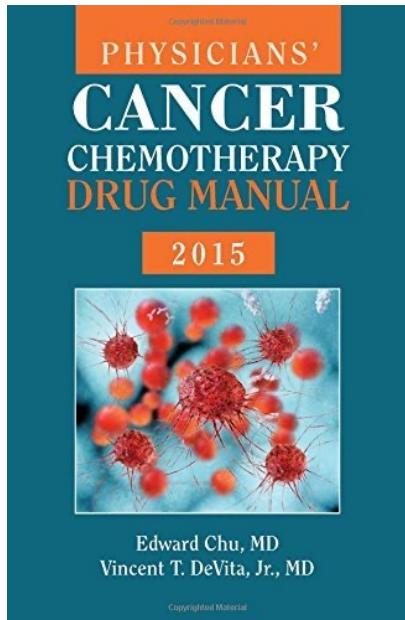
Screening Technology



Scientific Literature

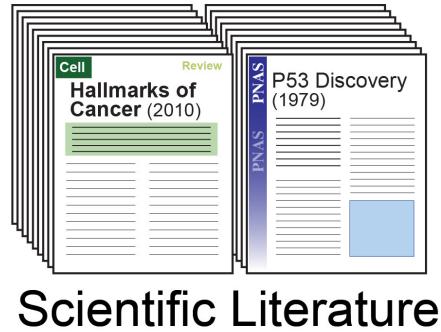
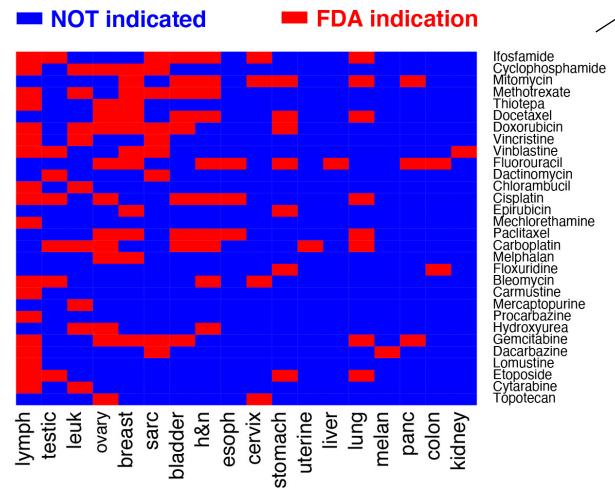
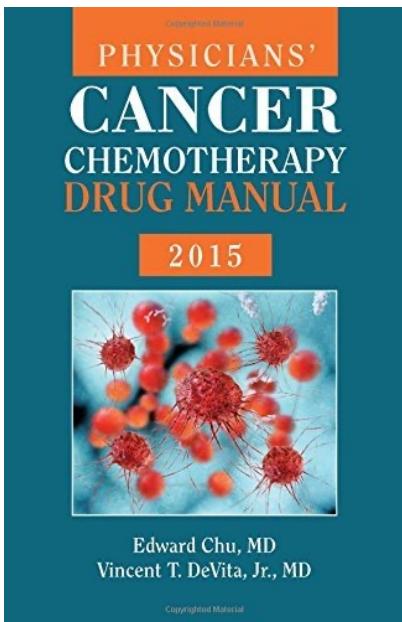
# Douglass Lab Approach: prune features & build database

## Define Top Clinical Chemotherapies



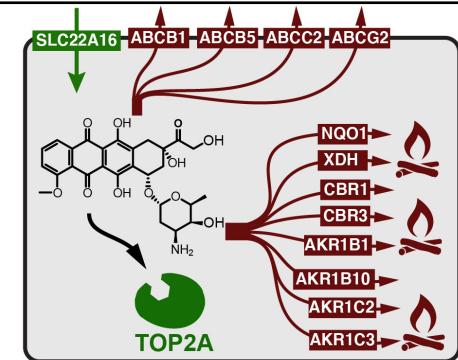
# Douglass Lab Approach: prune features & build database

## Define Top Clinical Chemotherapies



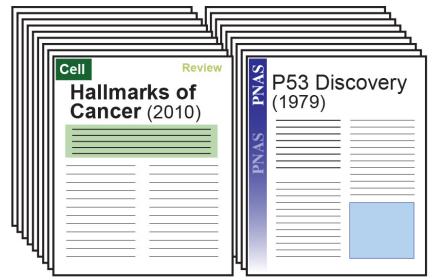
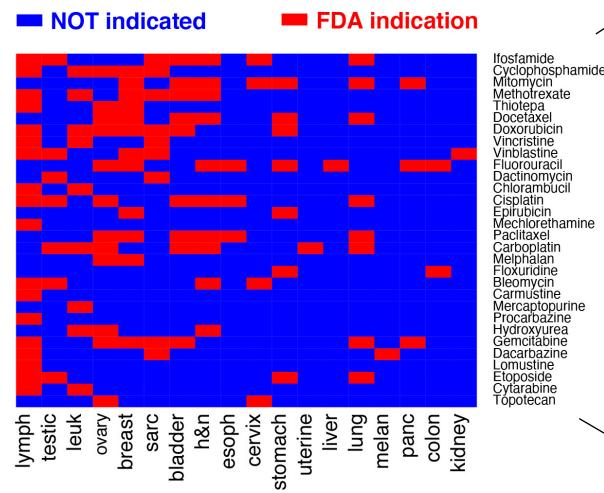
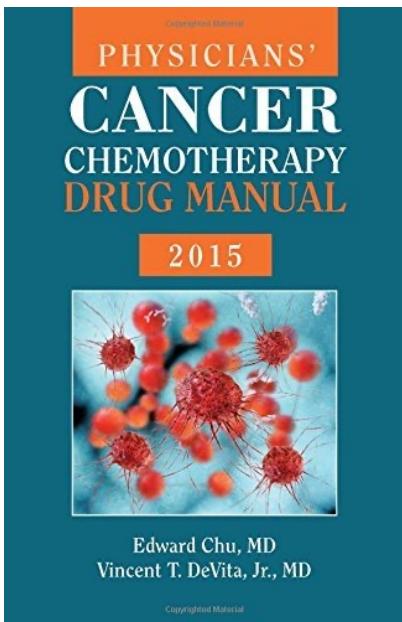
Scientific Literature

## Causal Biomarker Lists

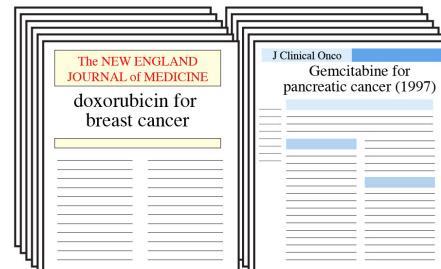


# Douglass Lab Approach: prune features & build database

## Define Top Clinical Chemotherapies

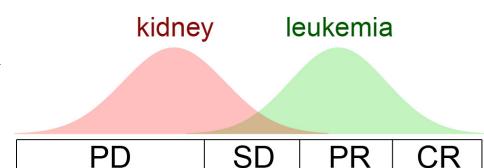
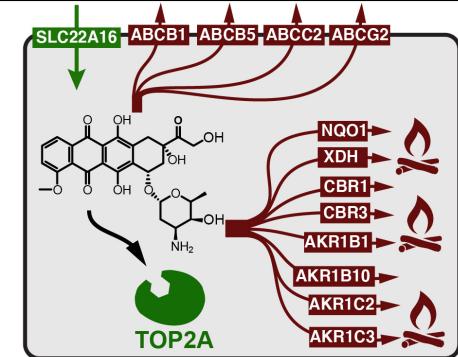


Scientific Literature



Clinical Literature

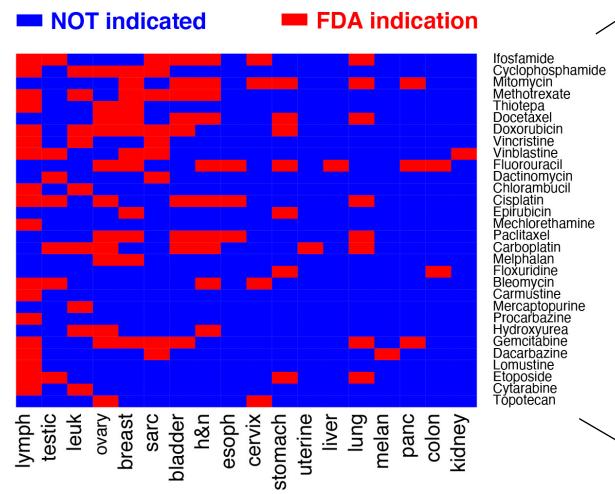
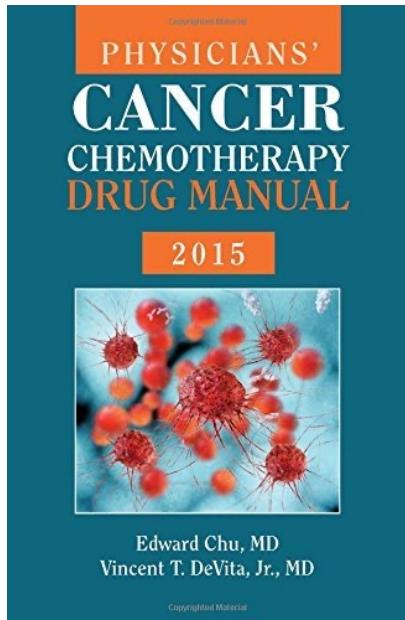
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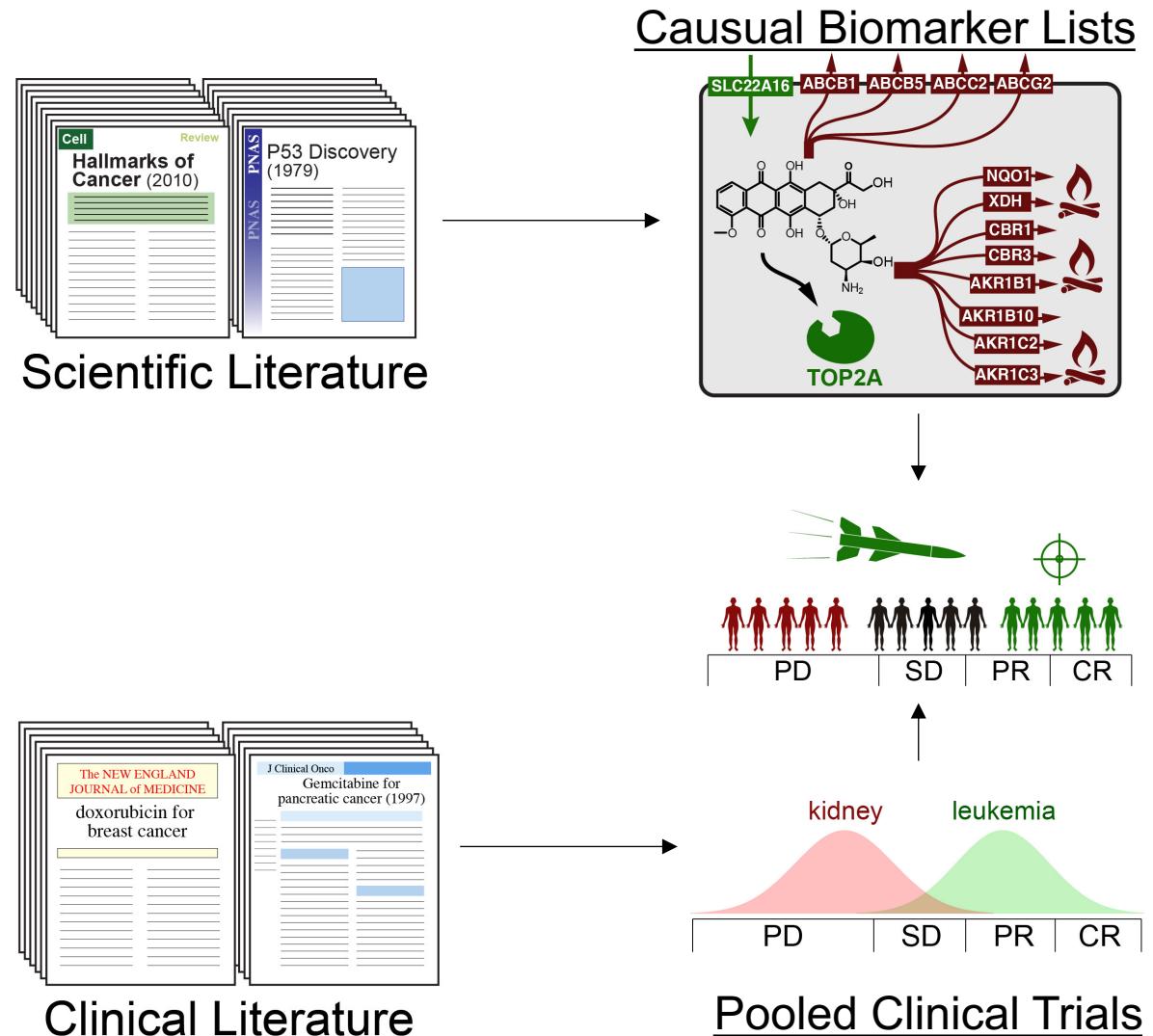
Pooled Clinical Trials

# Douglass Lab Approach: *build clinical database*

## Define Top Clinical Chemotherapies



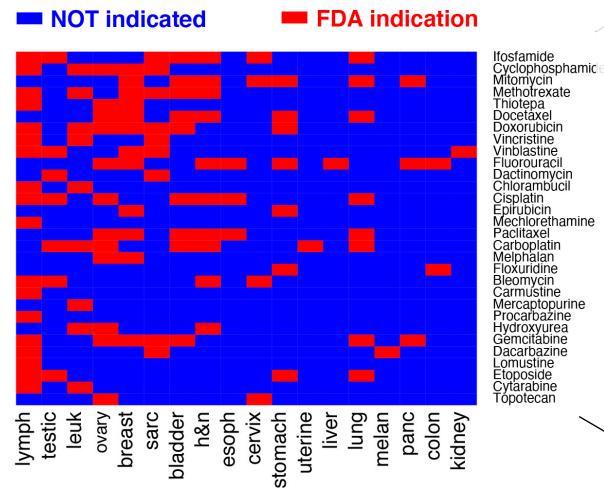
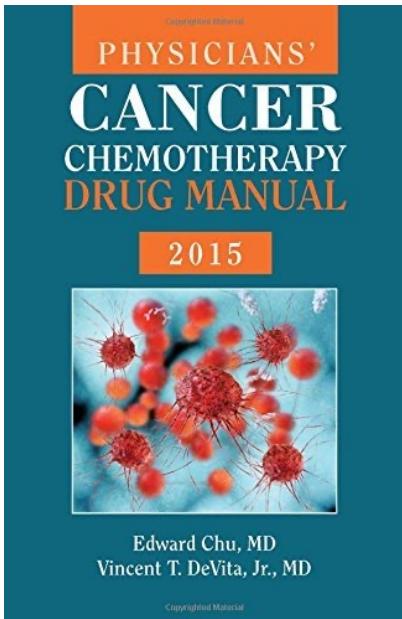
# Scientific Literature



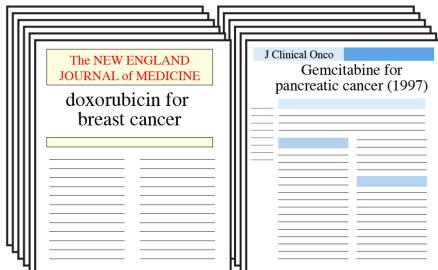
Chu, E.; et al. *Physicians' Cancer Chemotherapy Drug Manual* 2015, 2016, Jones and Bartlett

# Douglass Lab Approach: build clinical database

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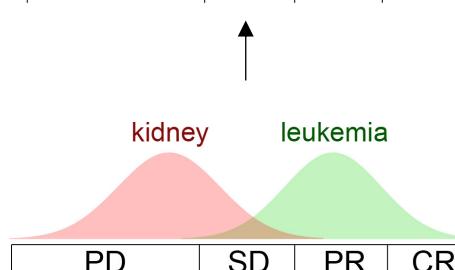
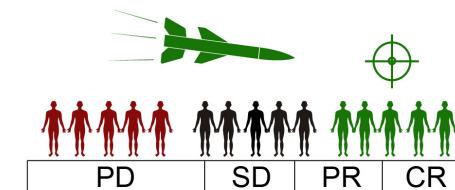
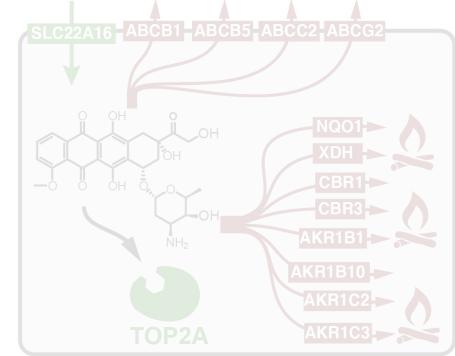


Scientific Literature



Clinical Literature

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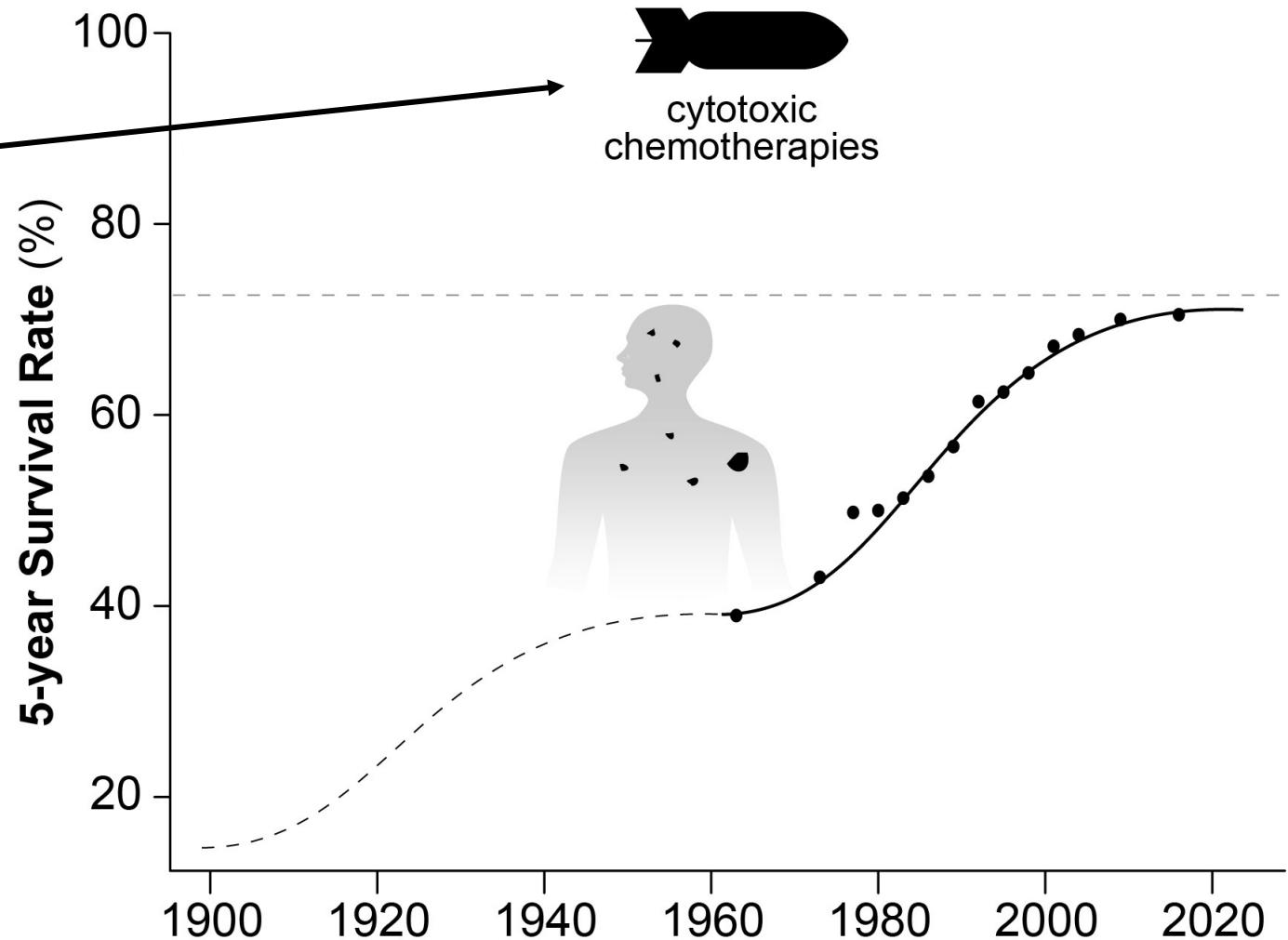


Pooled Clinical Trials

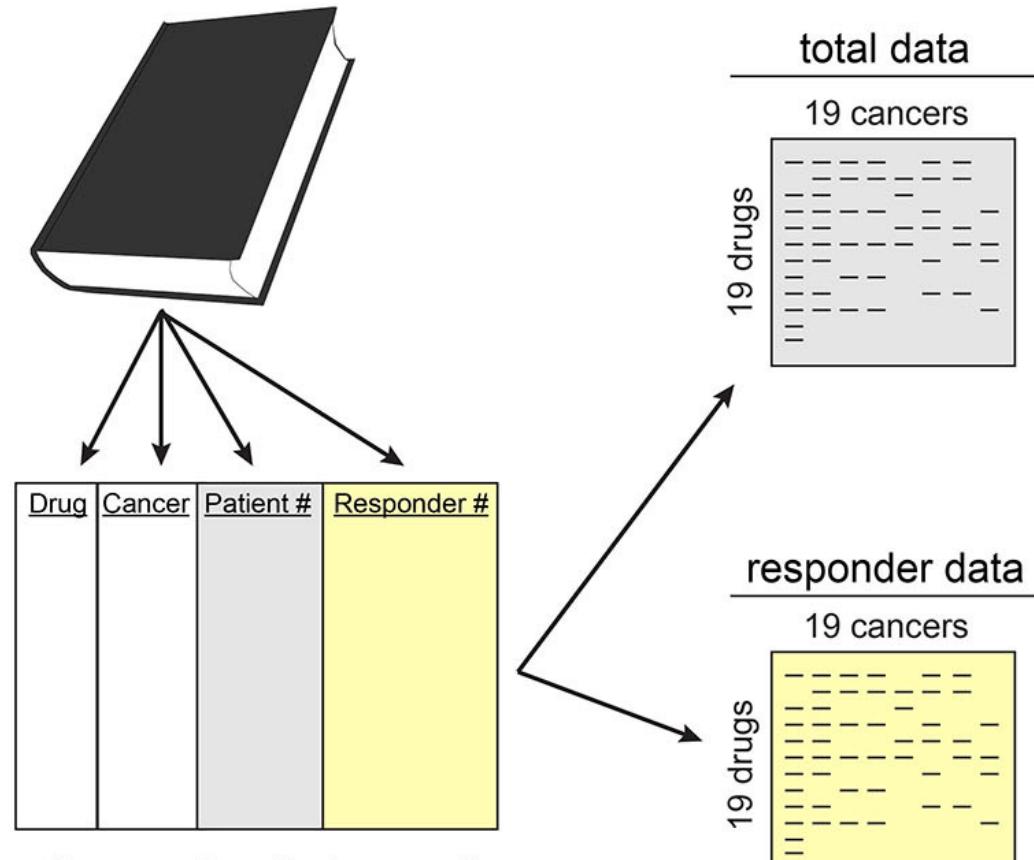
# Douglass Lab Approach: *build clinical database*



**NCI Clinical Trial Database (1970)**  
786 clinical trials  
19,958 evaluable patients



# Douglass Lab Approach: *build clinical database*



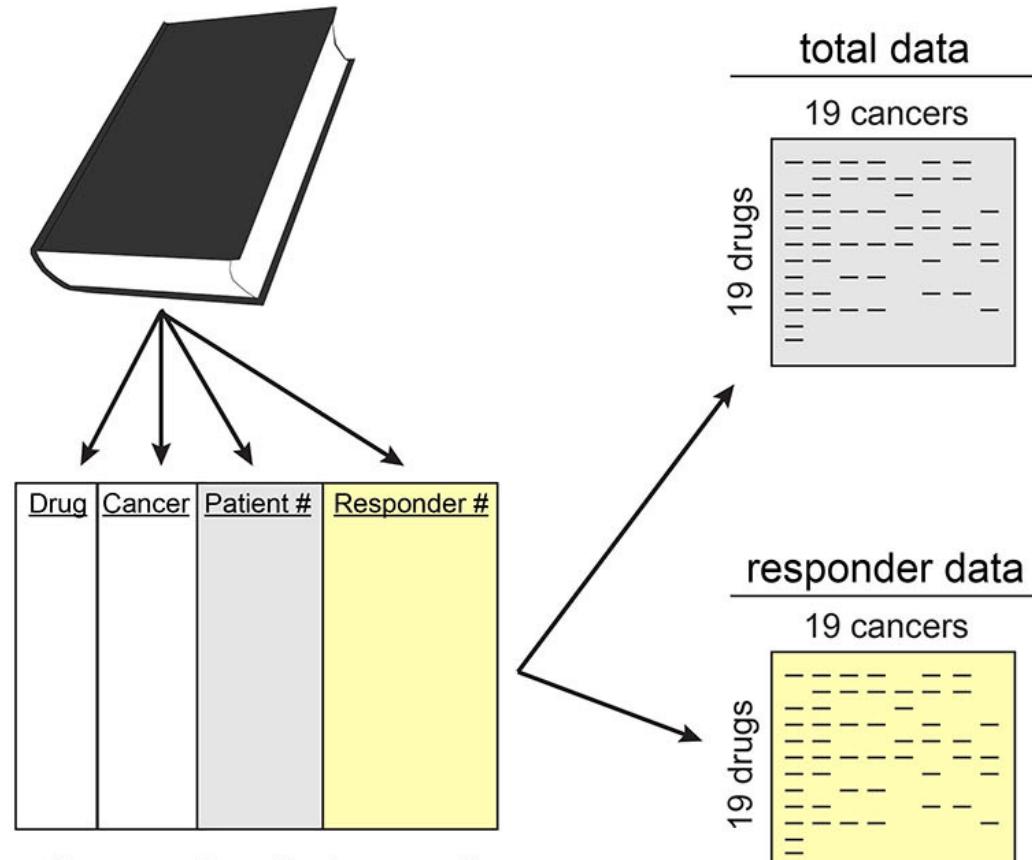
**Drug Names** = Drug Bank conventions

**Cancer Names** = TCGA abbreviations

**Patient #** = total evaluable patients

**Responder #** = WHO definitions

# Douglass Lab Approach: build clinical database

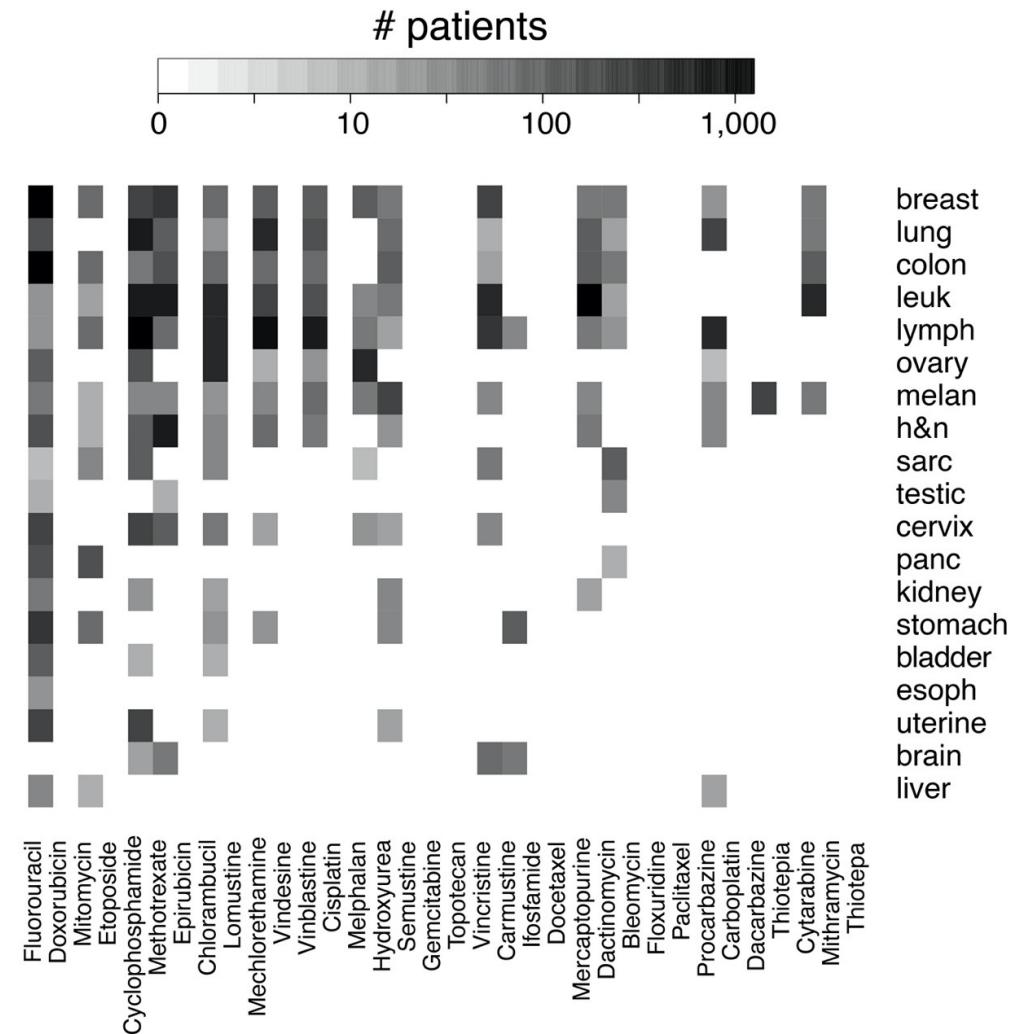


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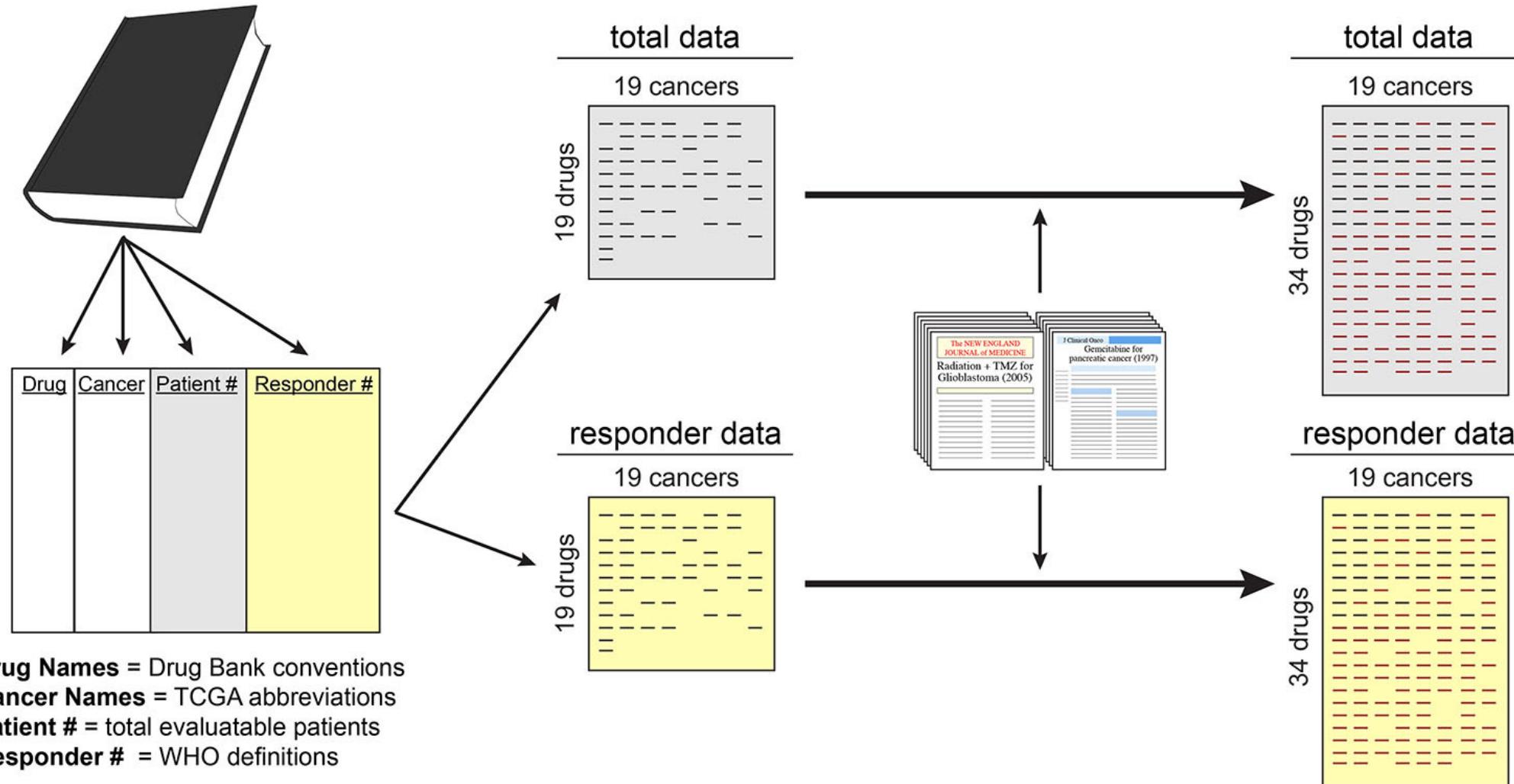
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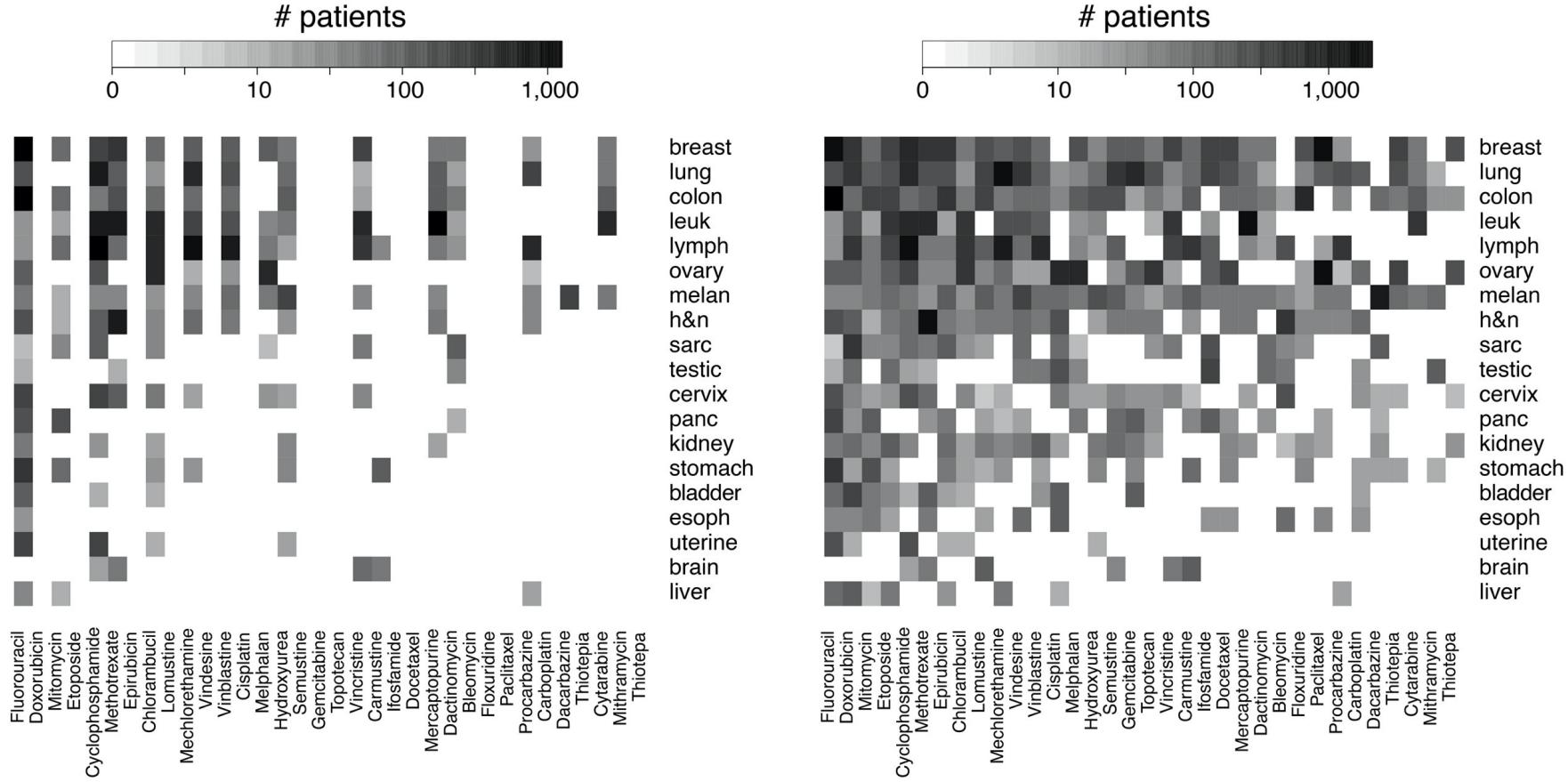
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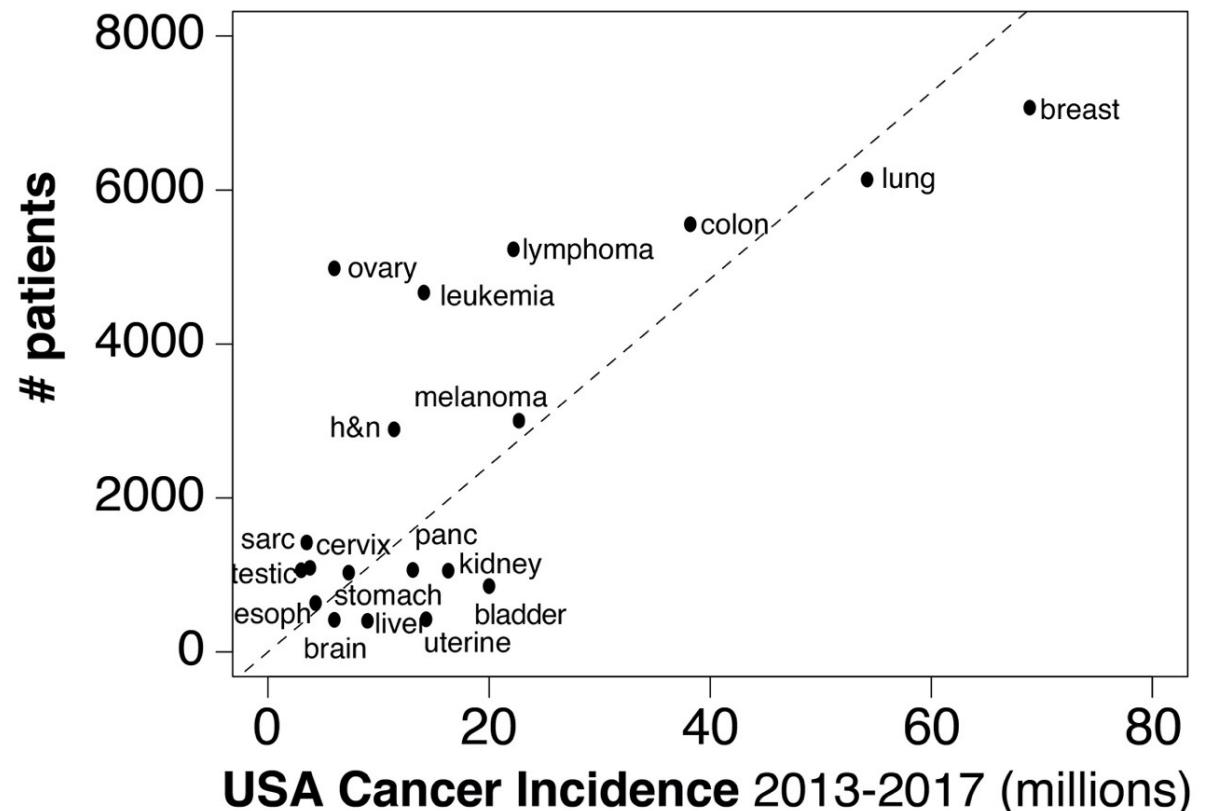
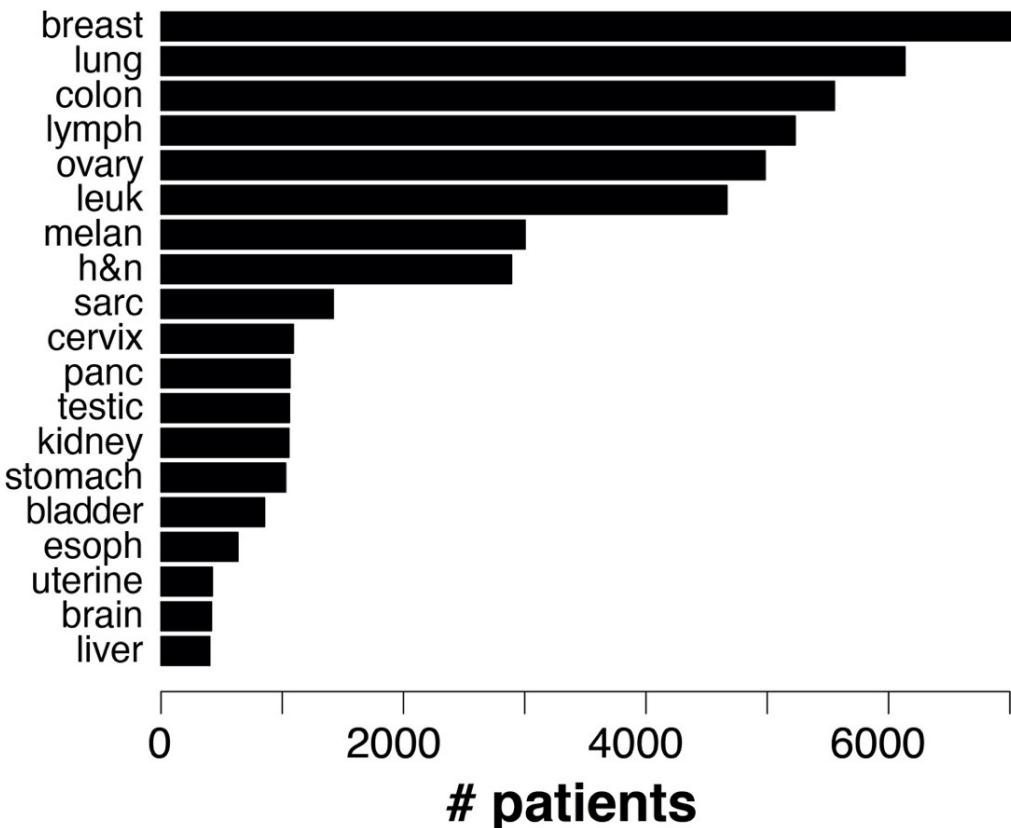
# Douglass Lab Approach: *build clinical database*



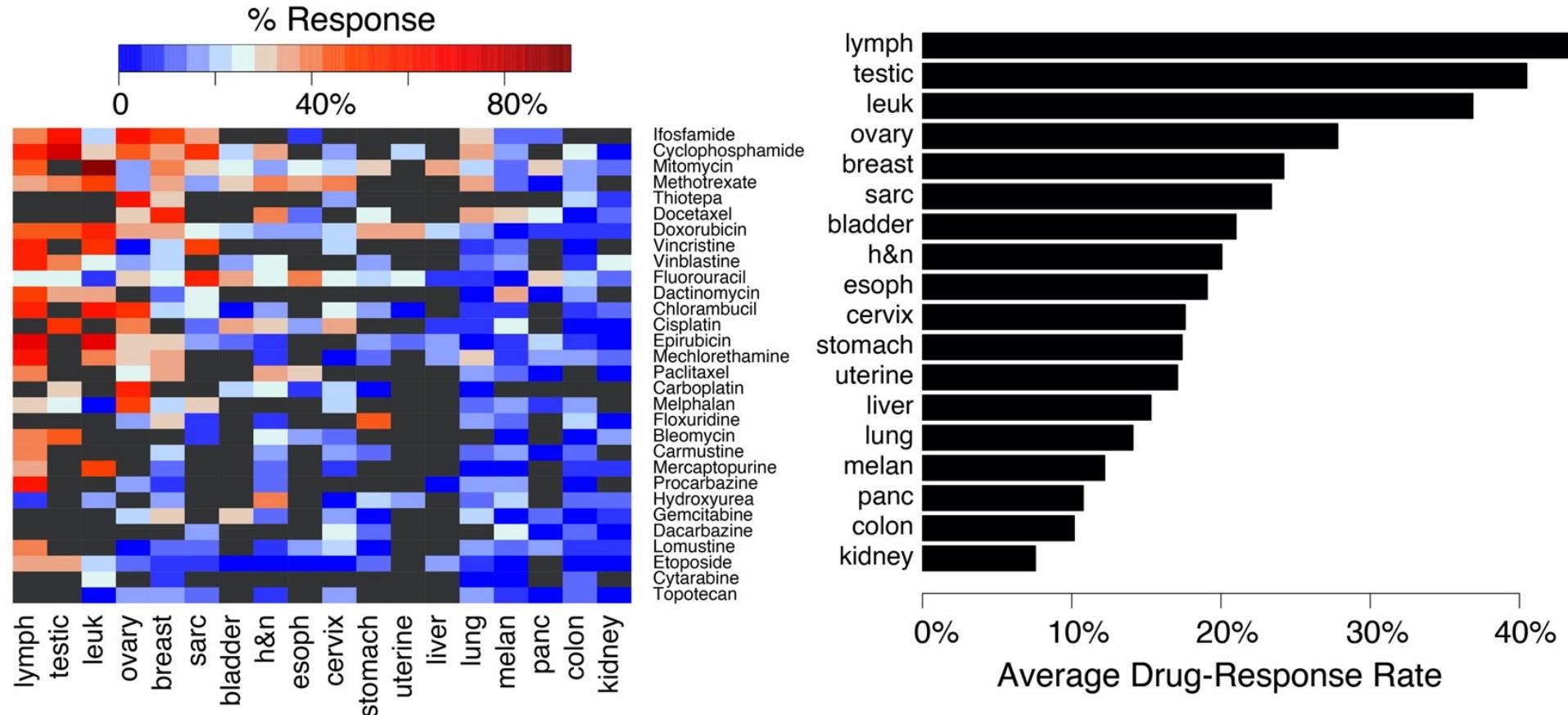
# Cancer-Type bias in clinical data:



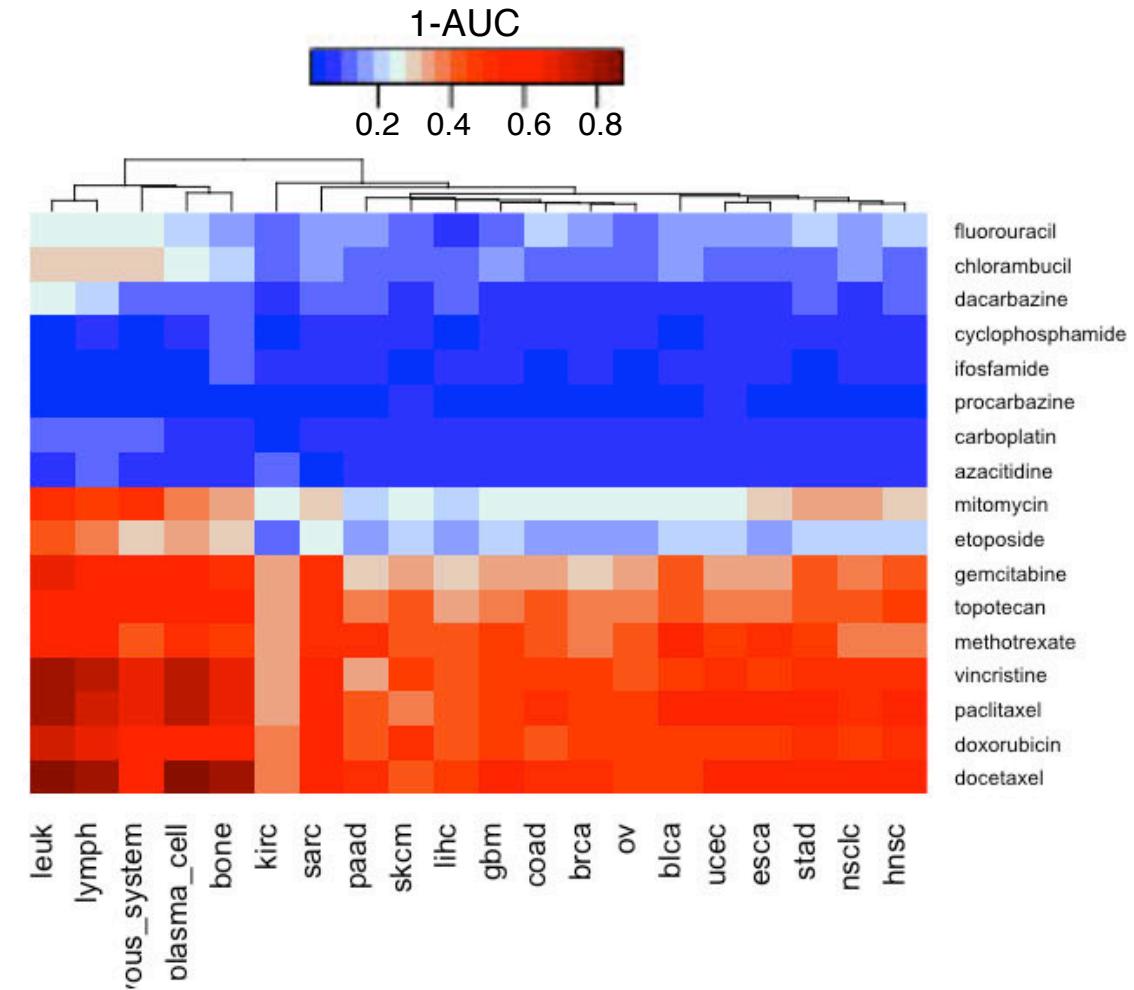
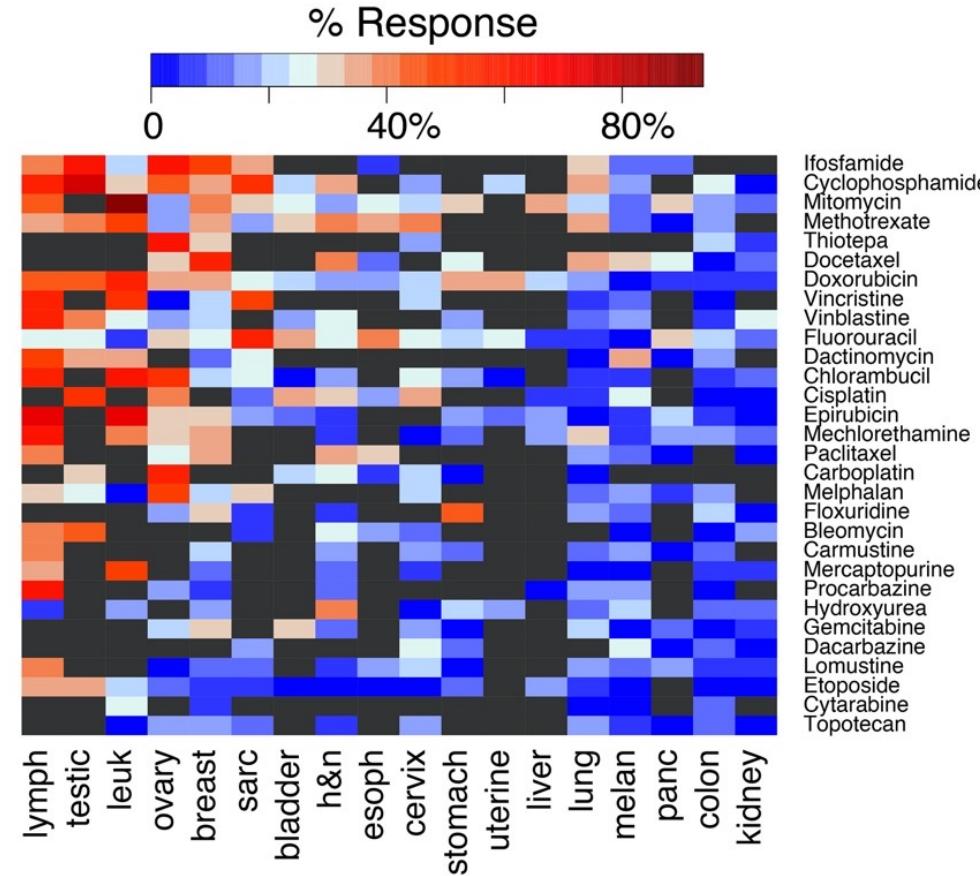
# Cancer-Type bias in clinical data: cancer incidence



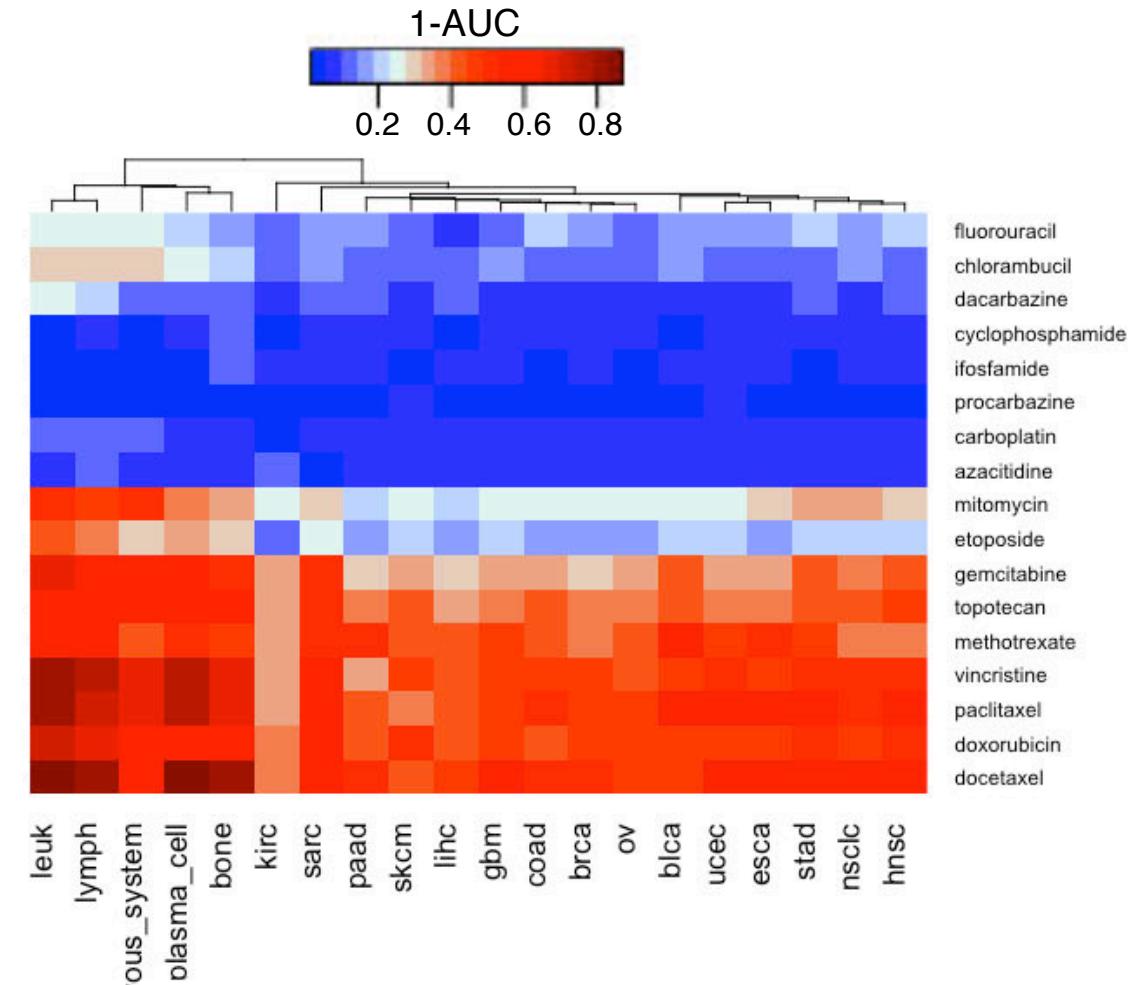
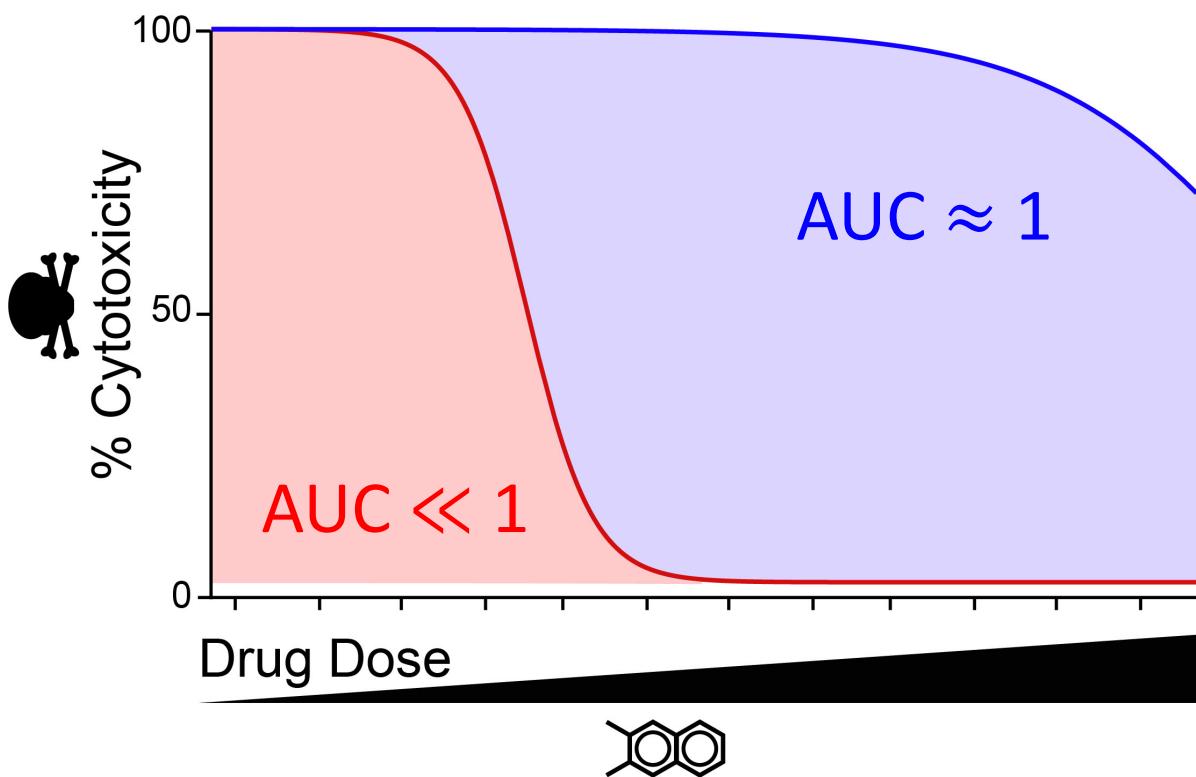
# Average chemo-sensitivity of different cancers



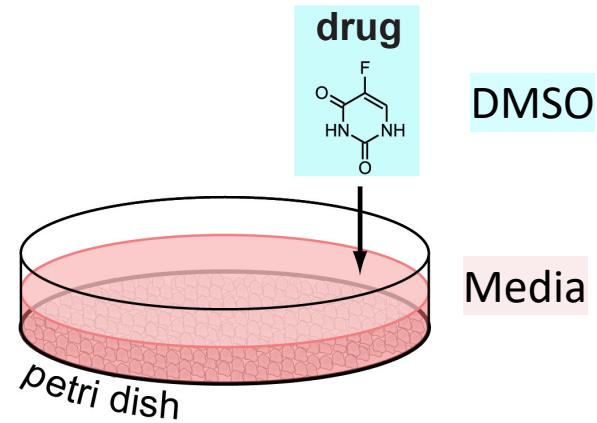
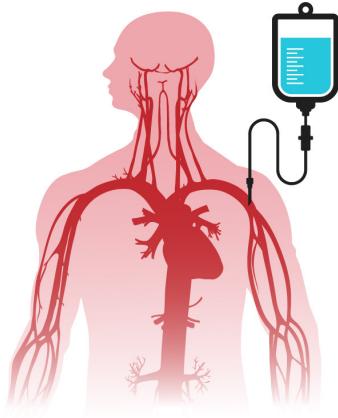
# Laboratory-Clinical Agreement: *laboratory artifacts*



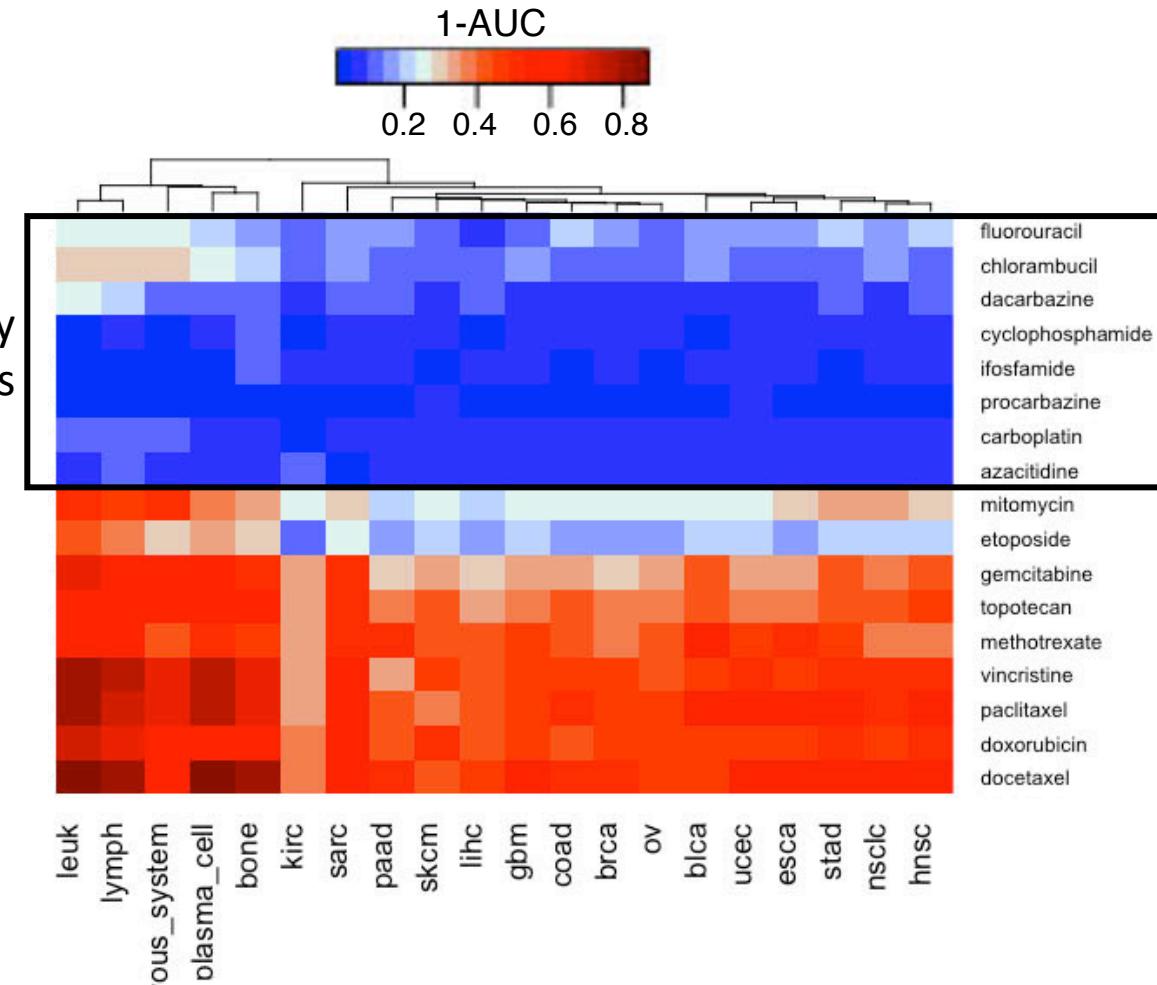
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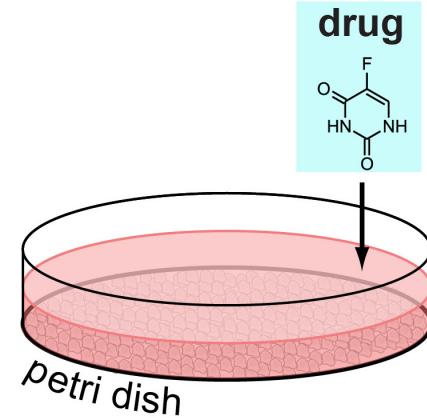
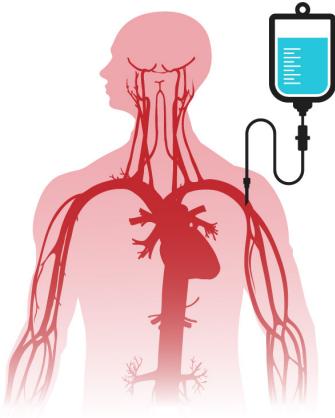
# Laboratory-Clinical Agreement: *laboratory artifacts*



Media laboratory artifacts



# Laboratory-Clinical Agreement: *laboratory artifacts*



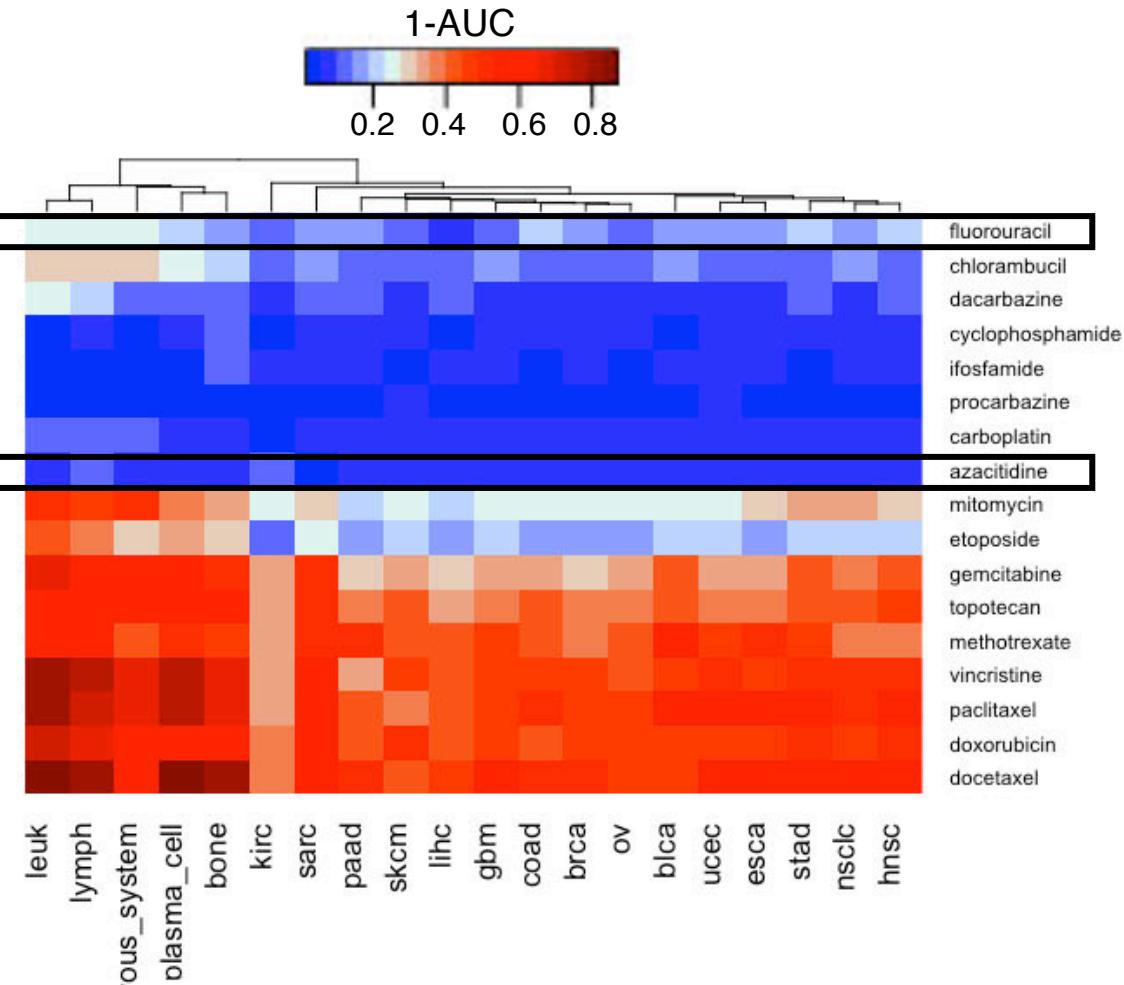
**anti-metabolite**  
chemotherapies

Article

## Physiologic Medium Rewires Cellular Metabolism and Reveals Uric Acid as an Endogenous Inhibitor of UMP Synthase

We find that uric acid directly inhibits UMP synthase and consequently reduces the sensitivity of cancer cells to the chemotherapeutic agent 5-fluorouracil.

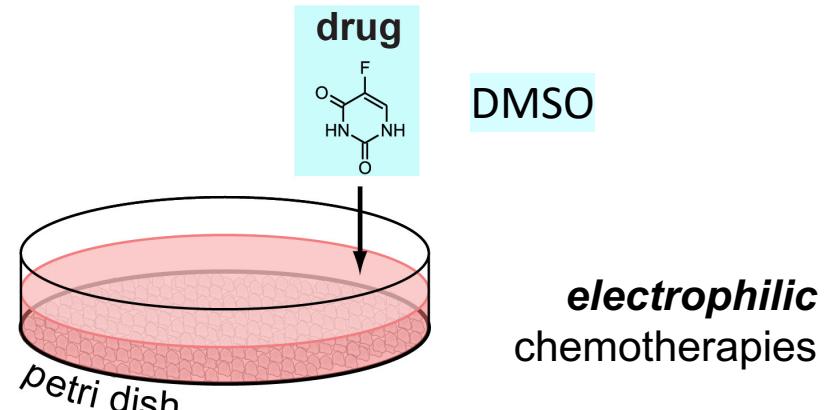
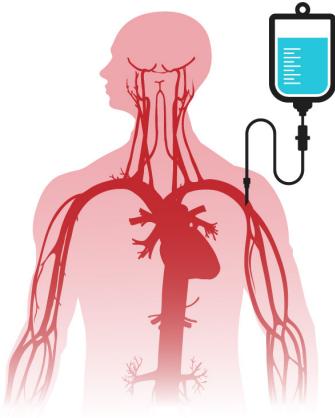
Cell, 2017, 169, 258



Cell, 2016, 166, 740

Cell, 2013, 154, 1151

# Laboratory-Clinical Agreement: *laboratory artifacts*



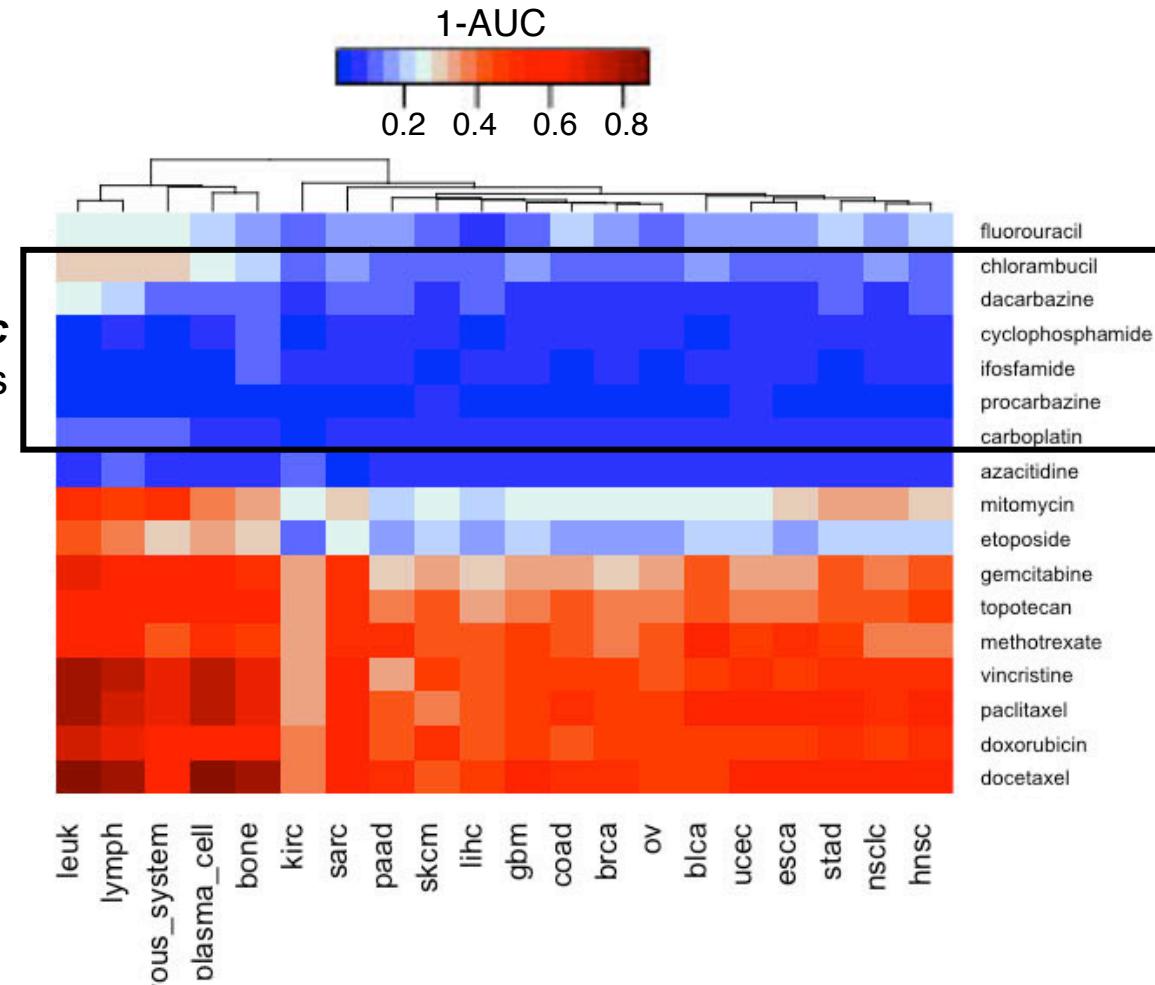
Therapeutics, Targets, and Chemical Biology

Cancer Research

**Say No to DMSO: Dimethylsulfoxide Inactivates Cisplatin, Carboplatin, and Other Platinum Complexes**

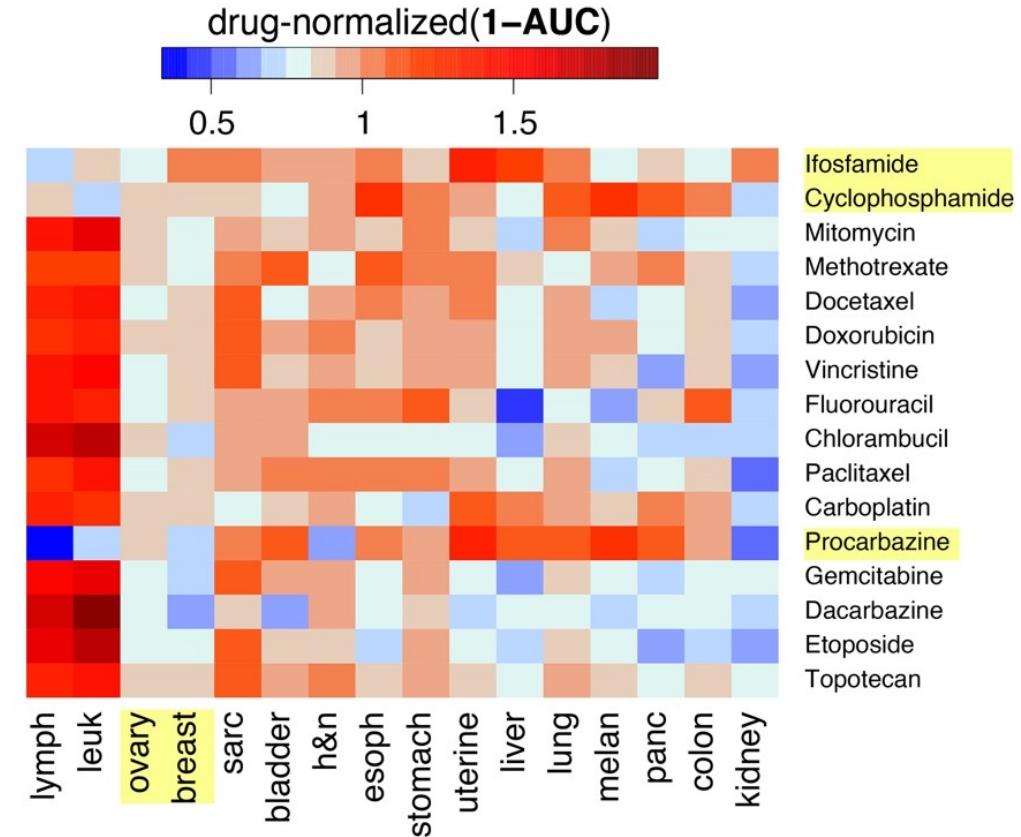
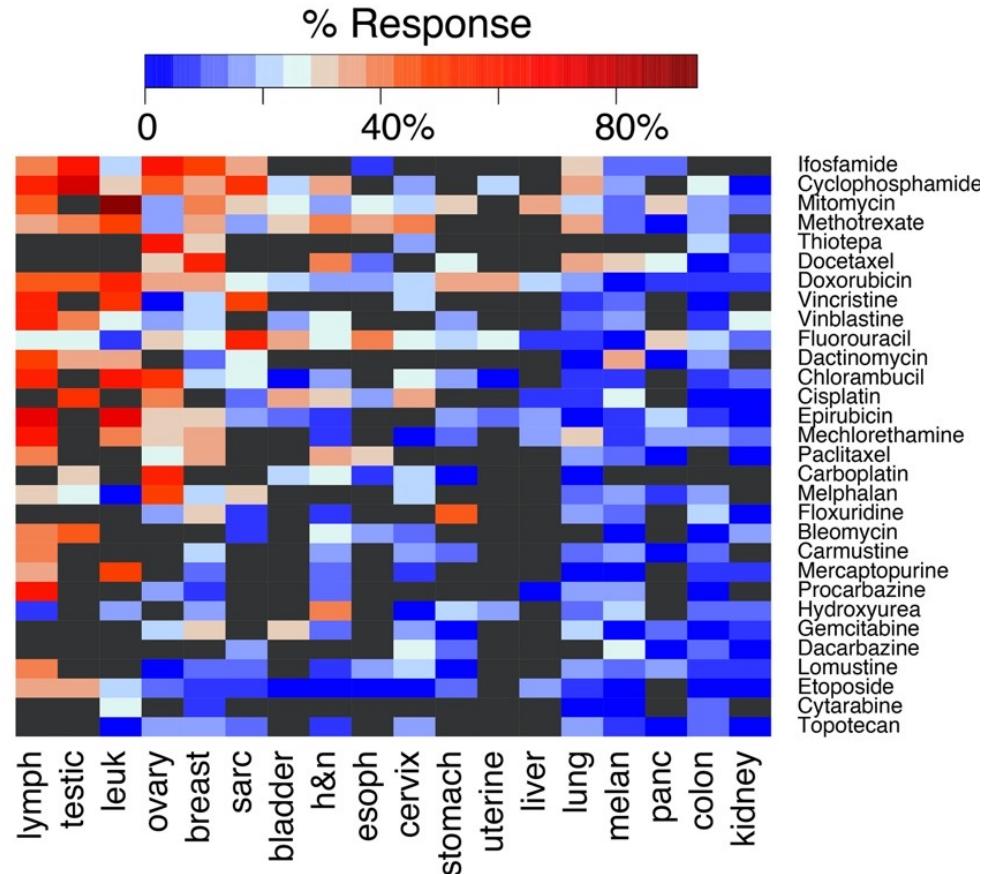
Matthew D. Hall<sup>1</sup>, Katherine A. Telma<sup>1</sup>, Ki-Eun Chang<sup>1</sup>, Tobie D. Lee<sup>1</sup>, James P. Madigan<sup>1</sup>, John R. Lloyd<sup>2</sup>, Ian S. Goldlust<sup>3</sup>, James D. Hoeschle<sup>4</sup>, and Michael M. Gottesman<sup>1</sup>

Cancer Res, 2014, 74, 3913

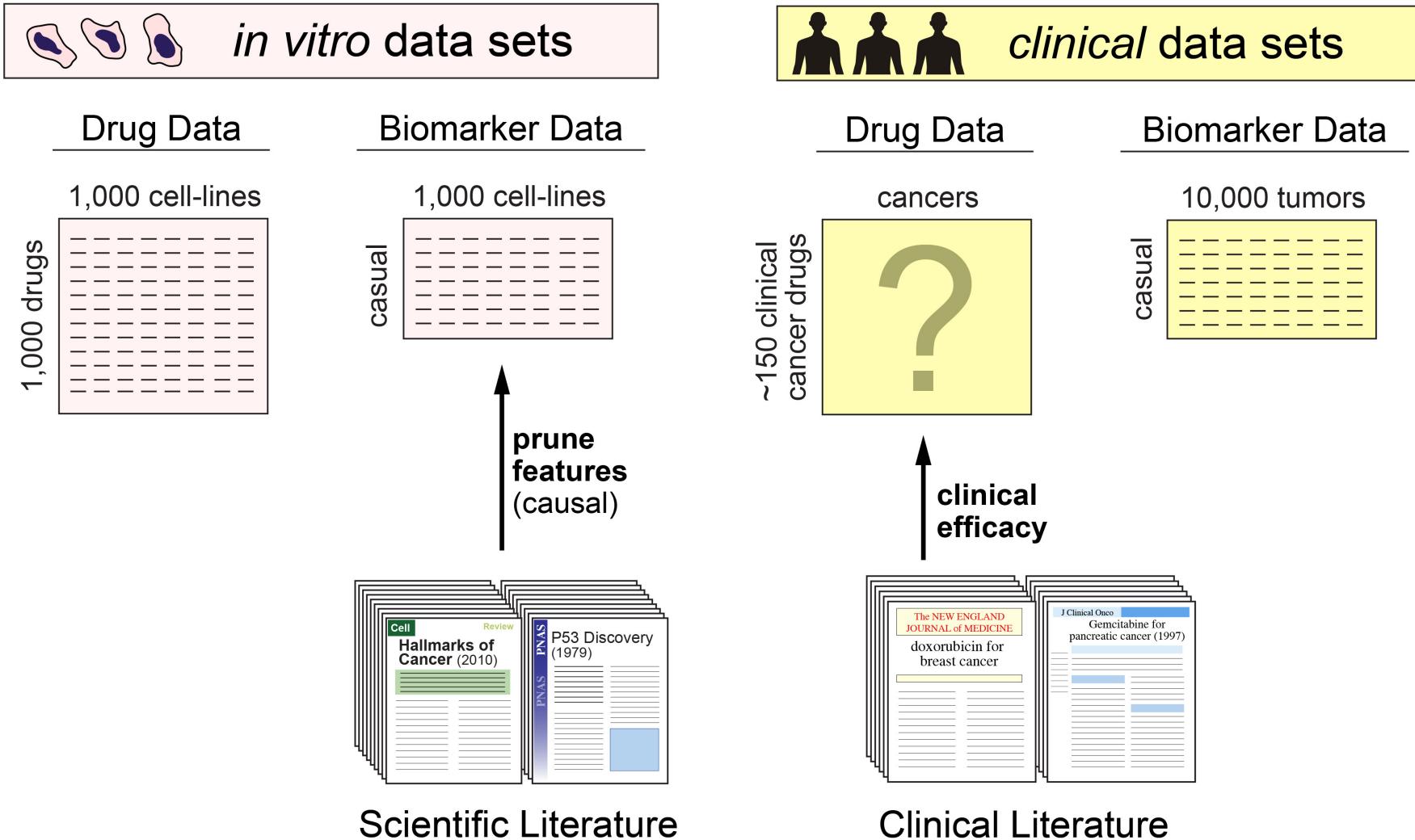


Cell, 2016, 166, 740  
Cell, 2013, 154, 1151

# Laboratory-Clinical Agreement: normalize by drug

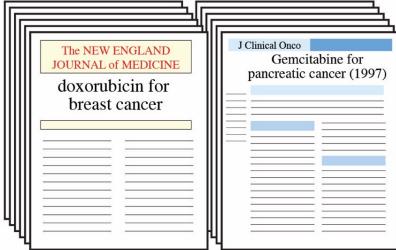


# Douglass Lab Approach: prune features & build database

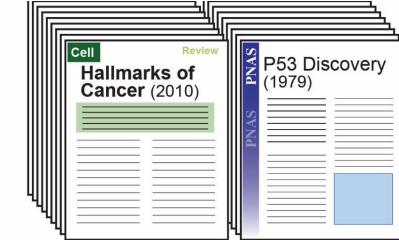
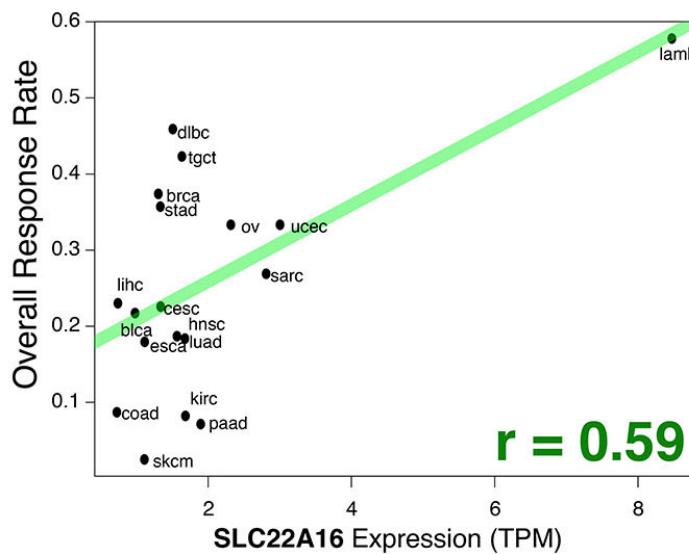


# Evaluate Clinical Validity of each biomarker:

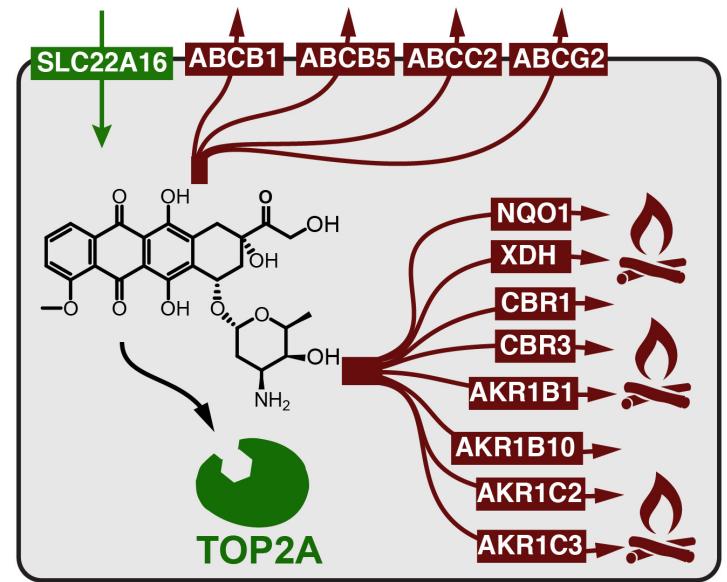
clinical data sets



Clinical Literature



Scientific Literature

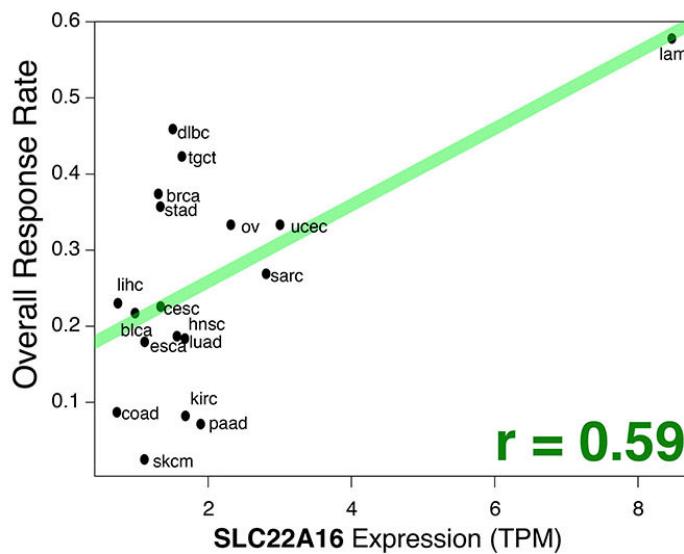


## Evaluate Clinical Validity of each biomarker:



The NEW ENGLAND JOURNAL of MEDICINE	J Clinical Oncol Gemcitabine for pancreatic cancer (1997)
doxorubicin for breast cancer	

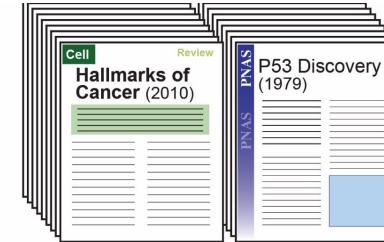
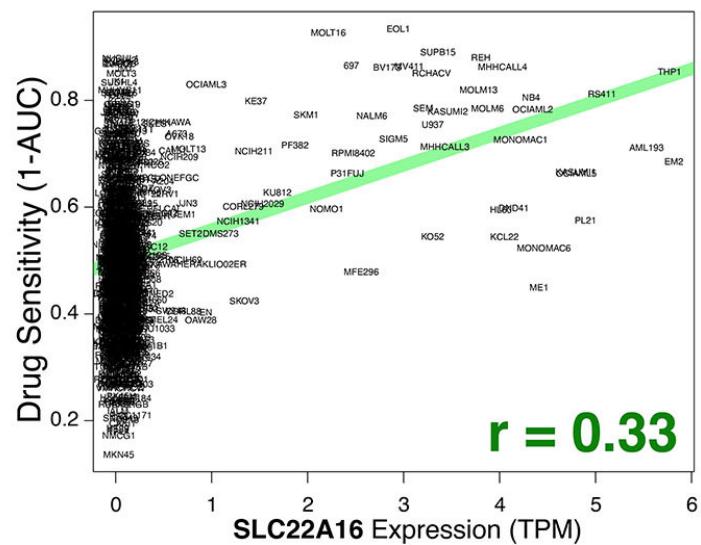
## Clinical Literature



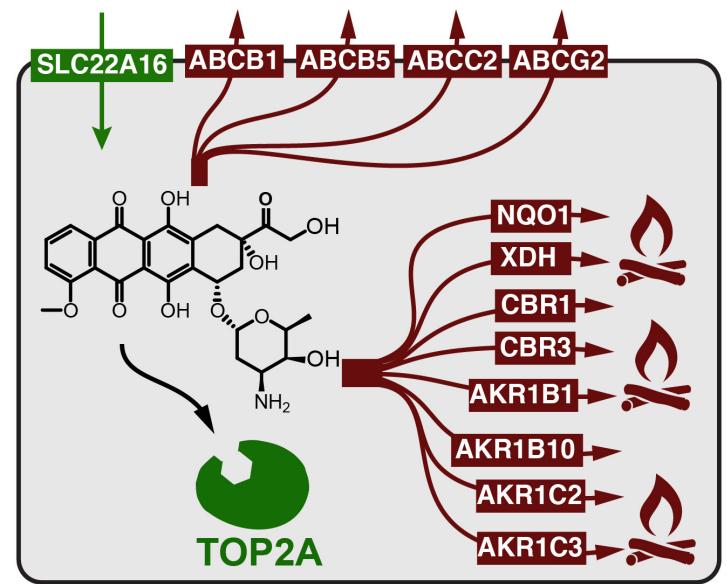
## *in vitro* data sets

A black and white illustration of a capsule-shaped object, resembling a pill or a capsule, resting on a grid of small circles. The object is oriented diagonally, with its dark, rounded end pointing towards the top right and its lighter, pointed end pointing towards the bottom left. The grid consists of numerous small, uniform circles arranged in a regular pattern across the entire frame.

# Screening Technology



## Scientific Literature



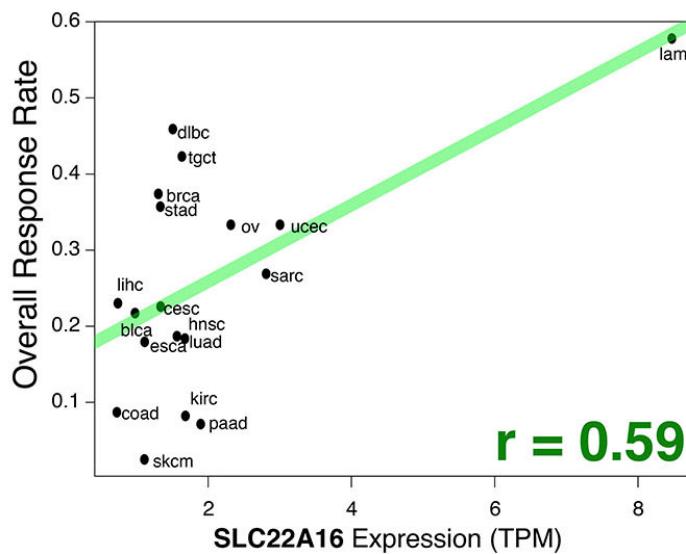
Drug Sensitivity(CTRP): *Cell*, 2013, 154, 1151  
Biomarker Expression (CCLE): *Nature*, 2019, 569, 503

# Evaluate Clinical Validity of each biomarker:



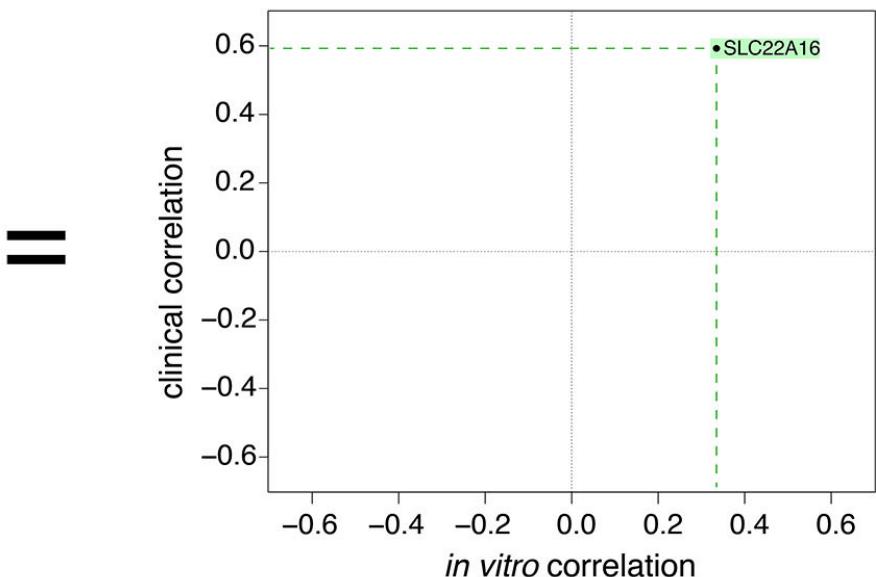
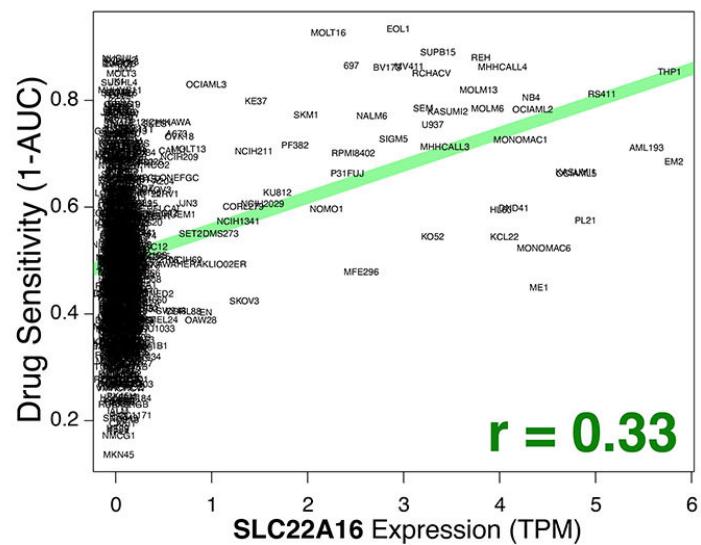
A stack of several medical journal covers from the early 1990s. The visible titles include "The NEW ENGLAND JOURNAL of MEDICINE" (1990), "J Clinical Oncol" (1991), and "Gemcitabine for pancreatic cancer (1997)". Each cover features a large, prominent title at the top, followed by several horizontal lines for article abstracts.

## Clinical Literature



A black and white graphic of a capsule-shaped object, possibly a pill or a bullet, resting on a grid of small circles.

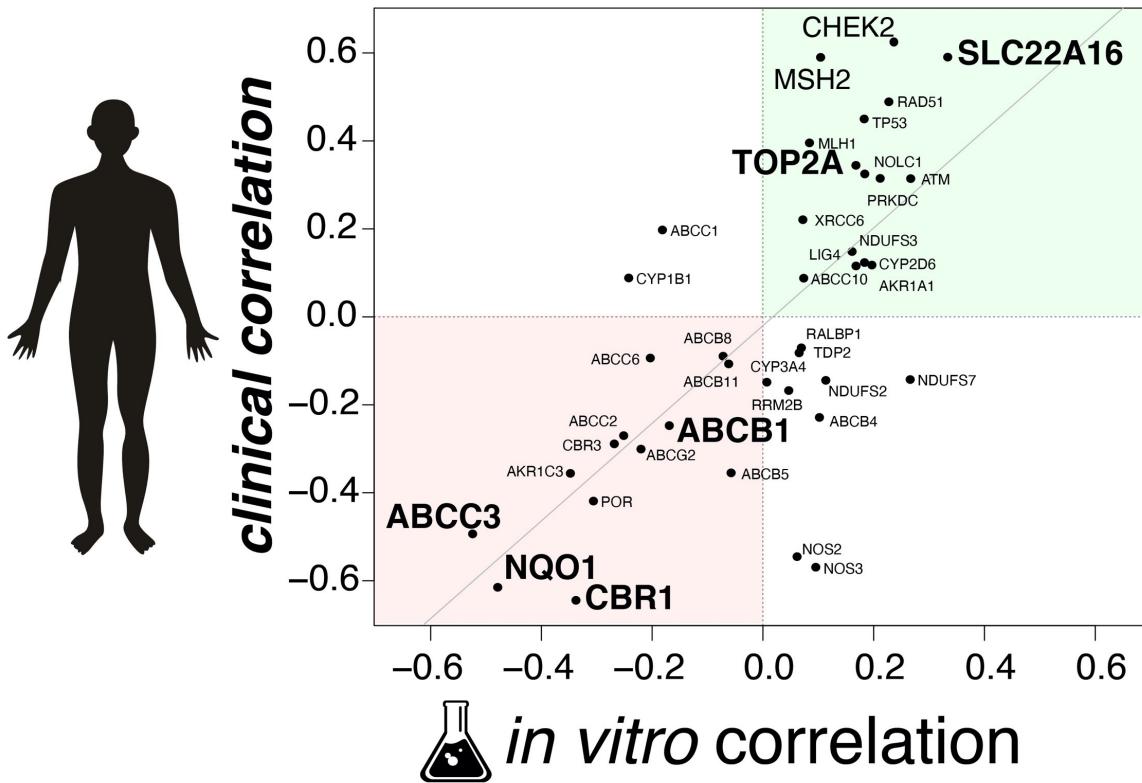
# Screening Technology



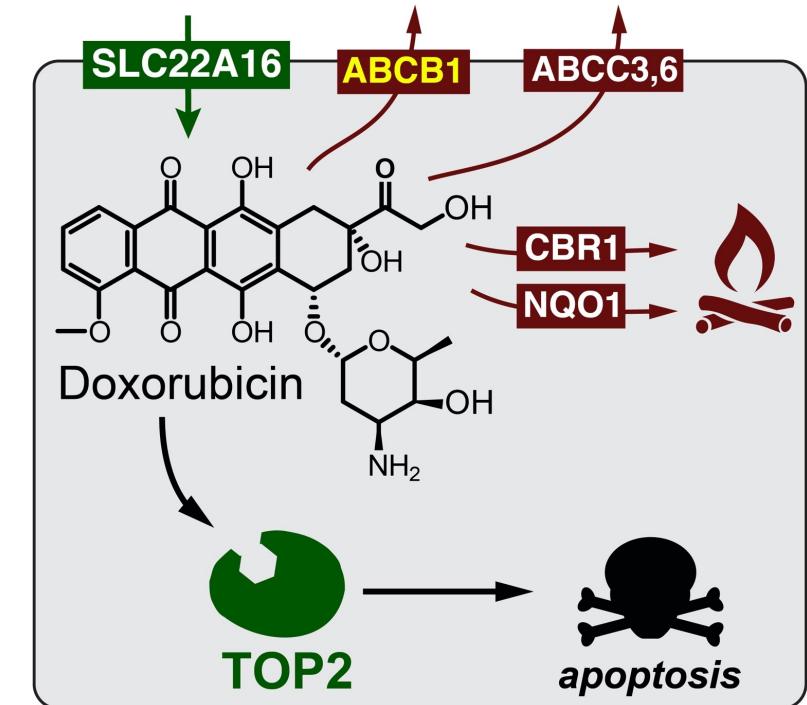
Drug Sensitivity(CTRP): *Cell*, 2013, 154, 1151  
Biomarker Expression (CCLE): *Nature*, 2019, 569, 503

# Comprehensive evaluation of anthracycline biomarkers

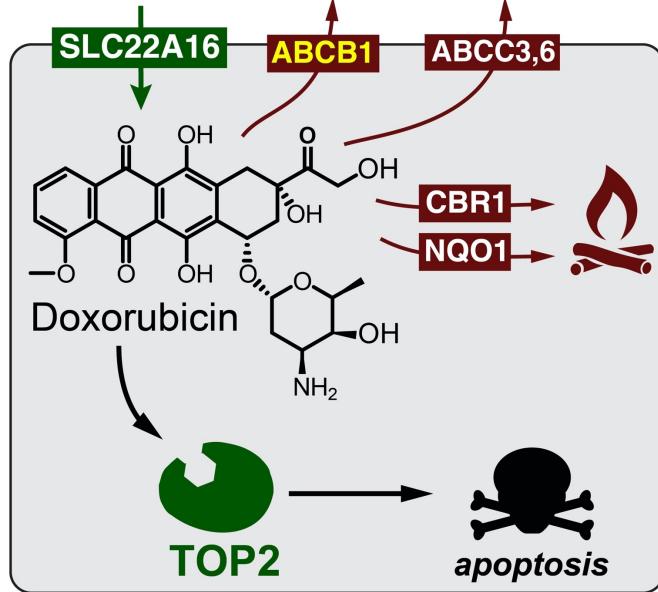
## A. Clinical Benchmark: Anthracycline Biomarkers



## B. Clinical Hypothesis: Anthracyclines



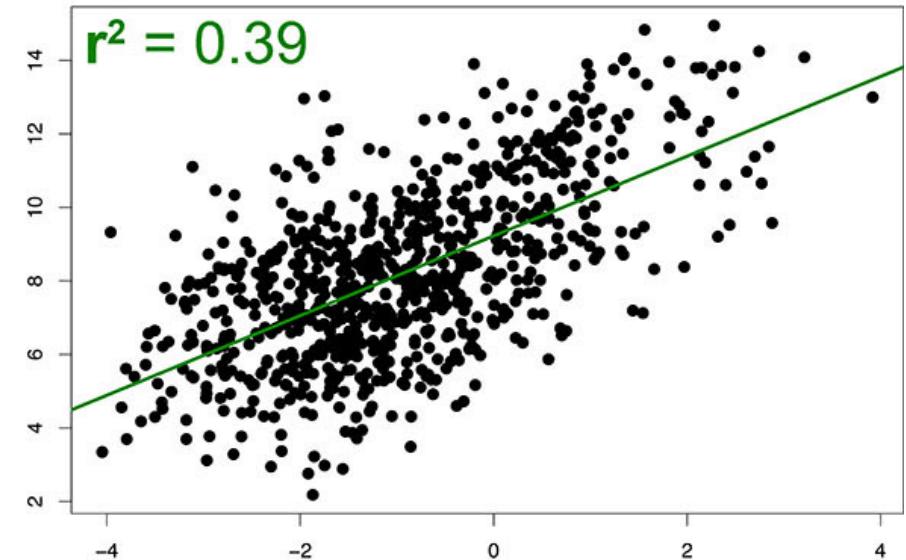
# Multi-Linear Regression: *N*-biomarker working hypothesis



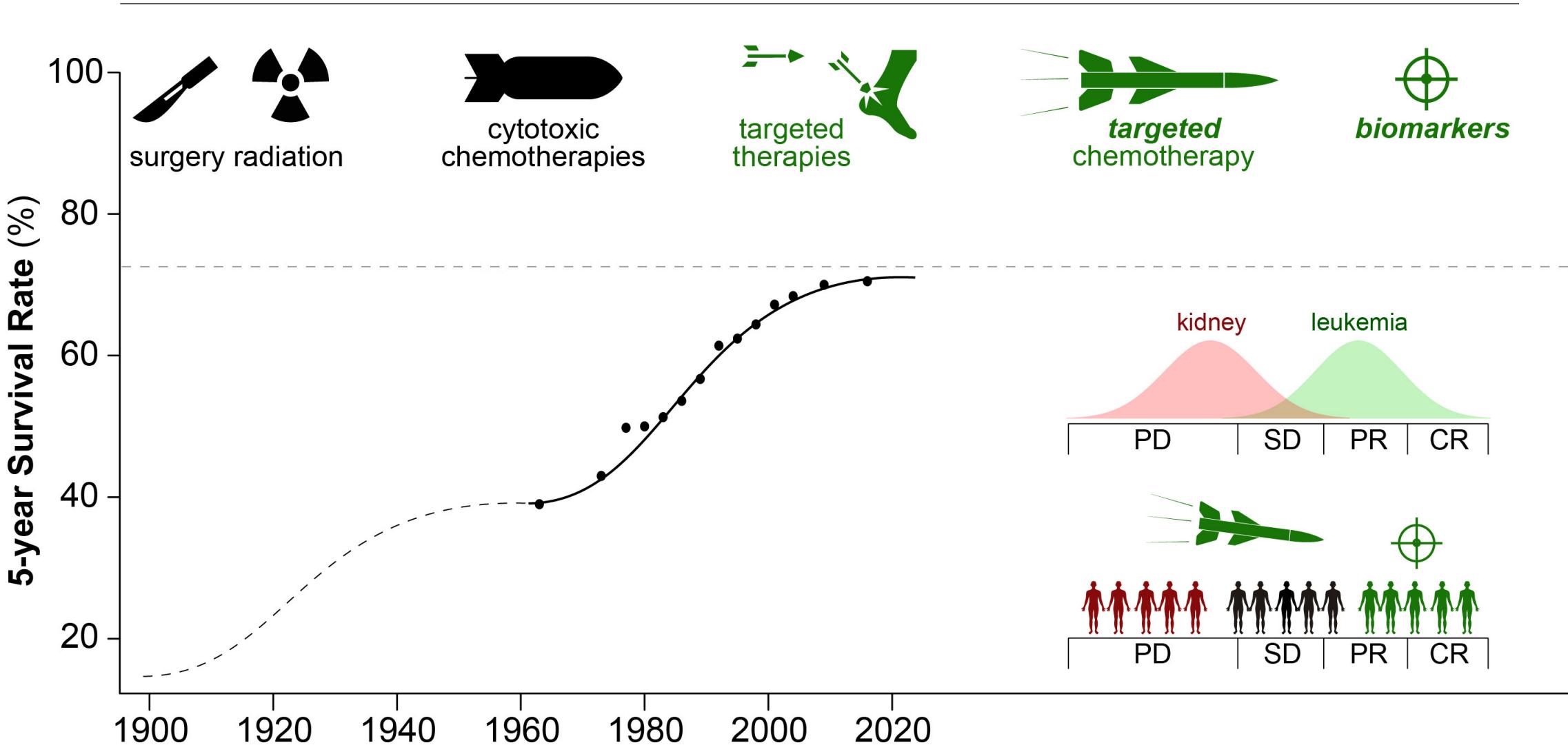
multi-linear model

$$\begin{aligned} &+0.44 \cdot SLC22A16 \\ &+0.25 \cdot NDUFS7 \\ &+0.11 \cdot TOP2A \\ &+0.11 \cdot TP53 \end{aligned}$$

$$\begin{aligned} &-0.10 \cdot CBR1 \\ &-0.11 \cdot NQO1 \\ &-0.16 \cdot ABCB1 \\ &-0.17 \cdot ABCC6 \\ &-0.22 \cdot ABCC3 \\ &-0.22 \cdot ABCC1 \end{aligned}$$

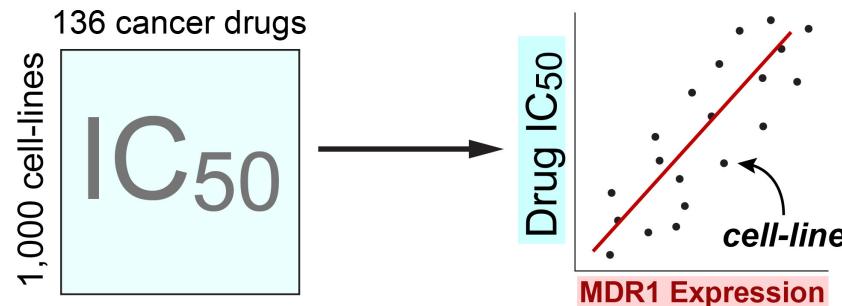


# History of Cancer Treatment: *targeted chemotherapies?*



# Douglass Laboratory: computational program

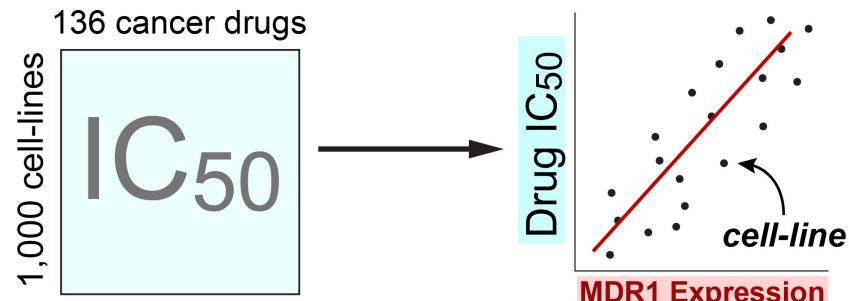
Computational Cancer Resistance Program: deconvolute N-biomarker phenotype



$$\Delta IC_{50} = \frac{k_{cat}/K_m}{k_{diff}} [ABCB1] + \frac{k_{cat}/K_m}{k_{diff}} [ABCG2] \dots$$

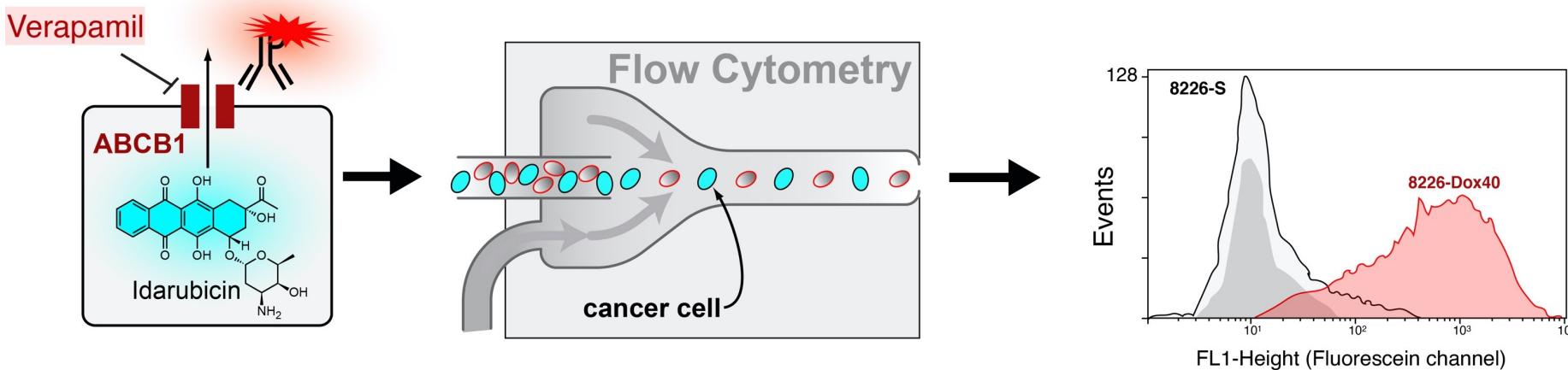
# Douglass Laboratory: experimental program

## Computational Cancer Resistance Program: deconvolute N-biomarker phenotype



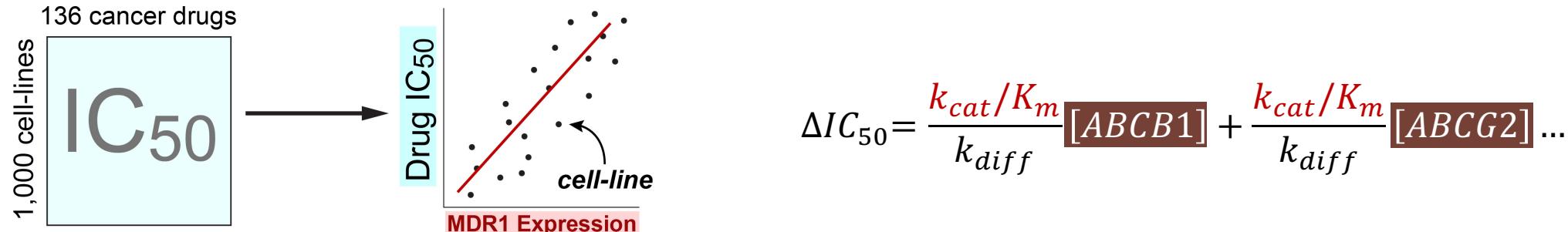
$$\Delta IC_{50} = \frac{k_{cat}/K_m}{k_{diff}} [ABCB1] + \frac{k_{cat}/K_m}{k_{diff}} [ABCG2] \dots$$

## Experimental Cancer Resistance Program: deconvolute physical constants of resistance

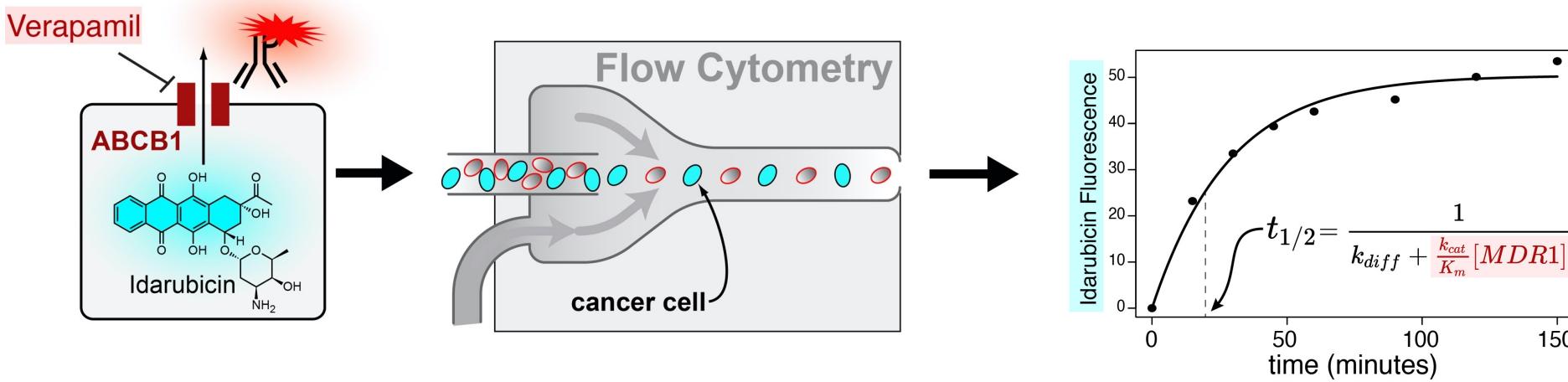


# Douglass Laboratory: experimental program

## Computational Cancer Resistance Program: deconvolute N-biomarker phenotype

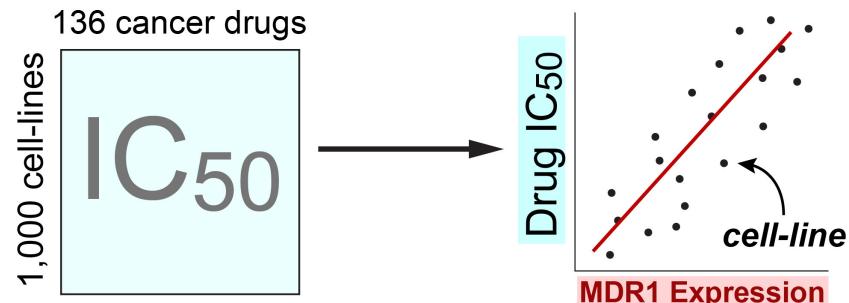


## Experimental Cancer Resistance Program: deconvolute physical constants of resistance



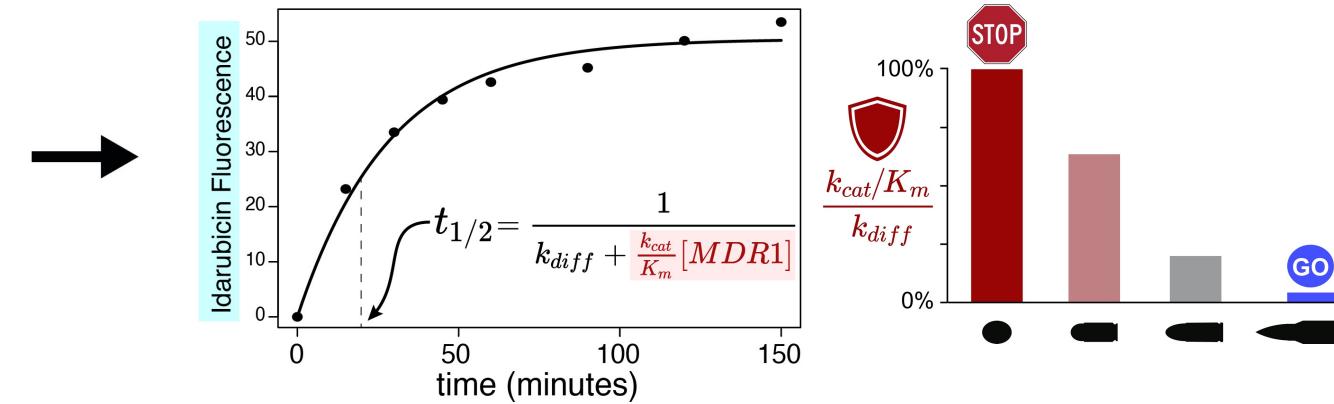
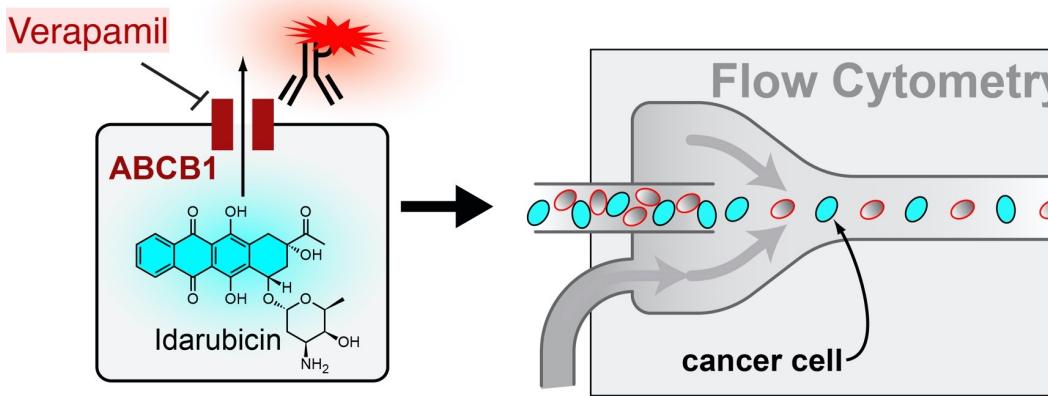
# Douglass Laboratory: experimental program

## **Computational Cancer Resistance Program: deconvolute N-biomarker phenotype**



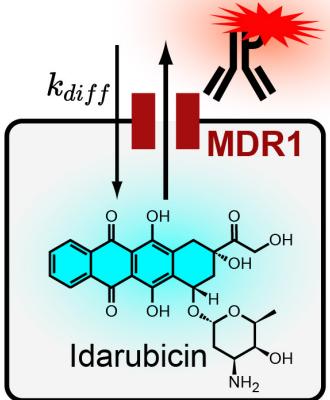
$$\Delta IC_{50} = \frac{k_{cat}/K_m}{k_{diff}} [ABCB1] + \frac{k_{cat}/K_m}{k_{diff}} [ABCG2] \dots$$

## **Experimental Cancer Resistance Program: deconvolute physical constants of resistance**



# Acknowledgements:

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Ashley Ray



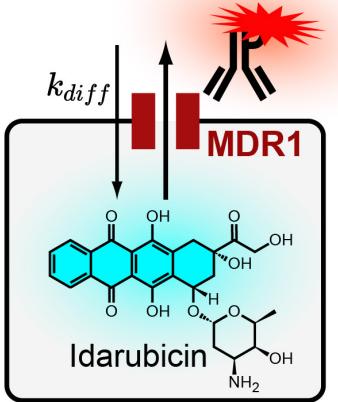
Elizabeth Hughes



George Johnson

# Acknowledgements:

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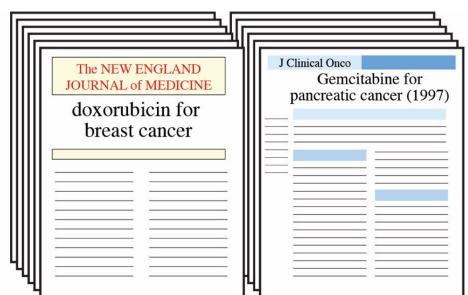
Ashley Ray



Elizabeth Hughes



George Johnson



Clinical Literature



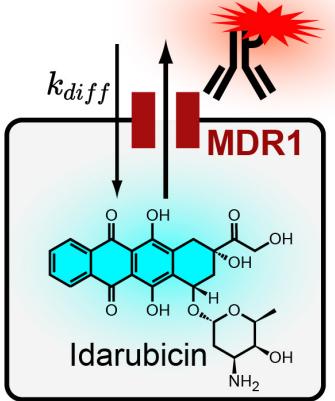
Tobin Paez



Emily Hannan

# Acknowledgements:

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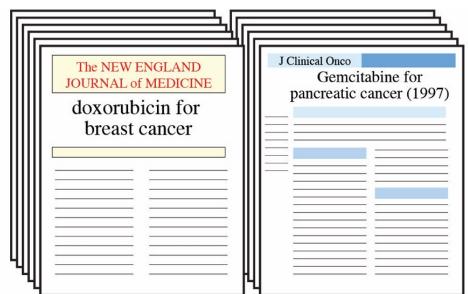
Ashley Ray



Elizabeth Hughes



George Johnson



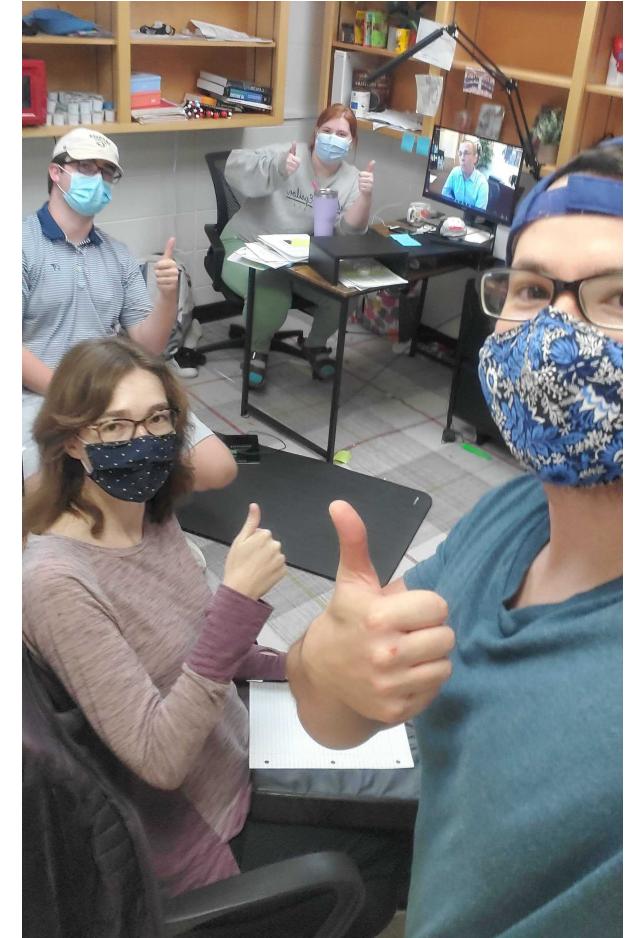
Clinical Literature



Tobin Paez



Emily Hannan

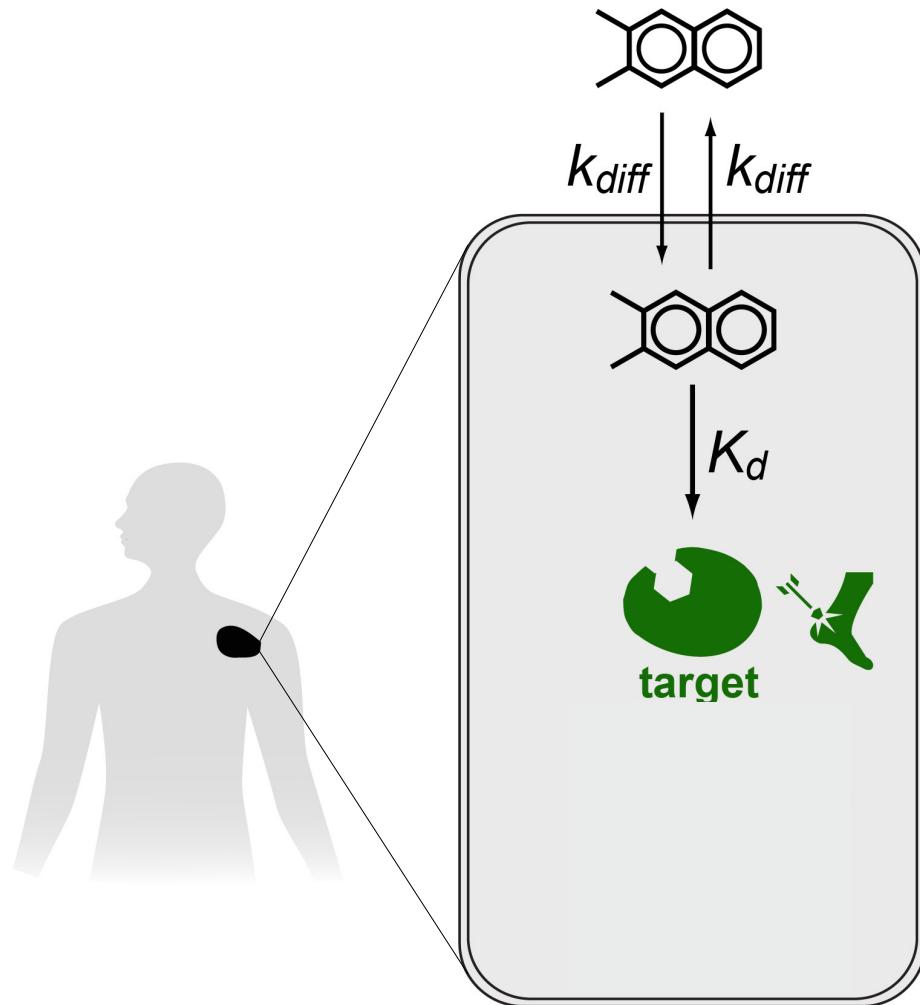


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# ***CLINICAL MEASUREMENT INTRO***

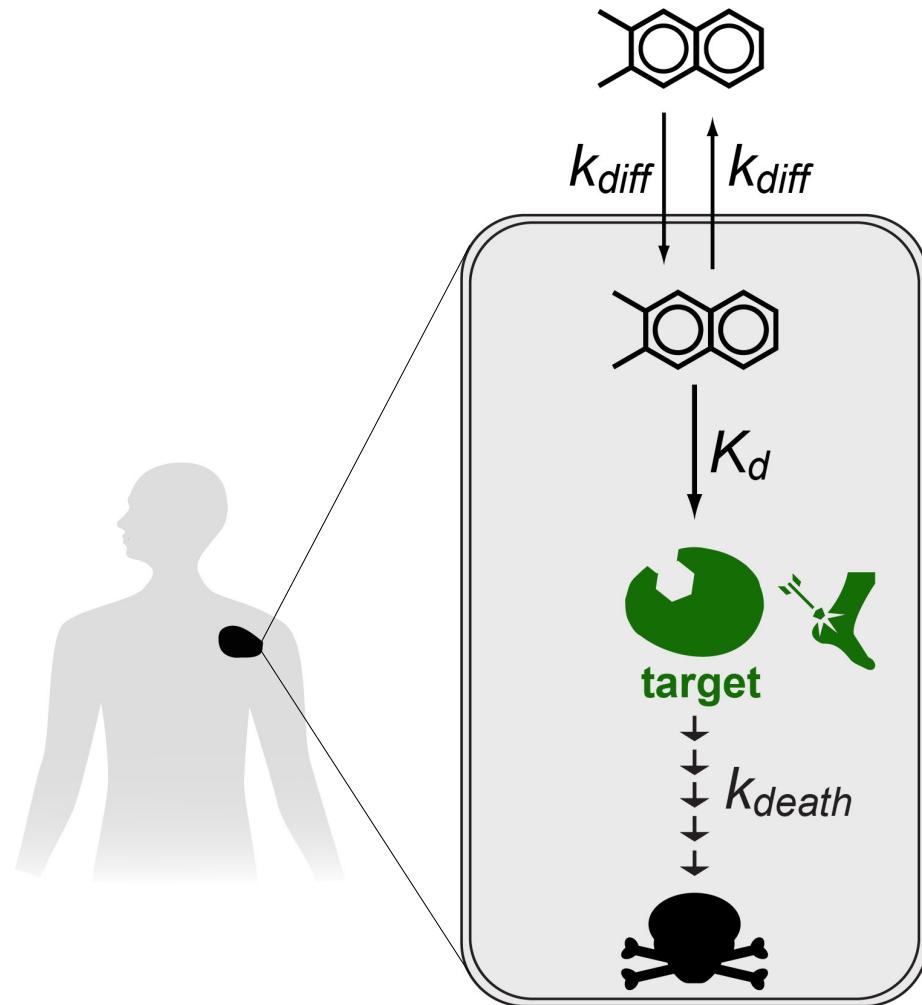
# 1-drug / 1 target paradigm: *historical artifact*

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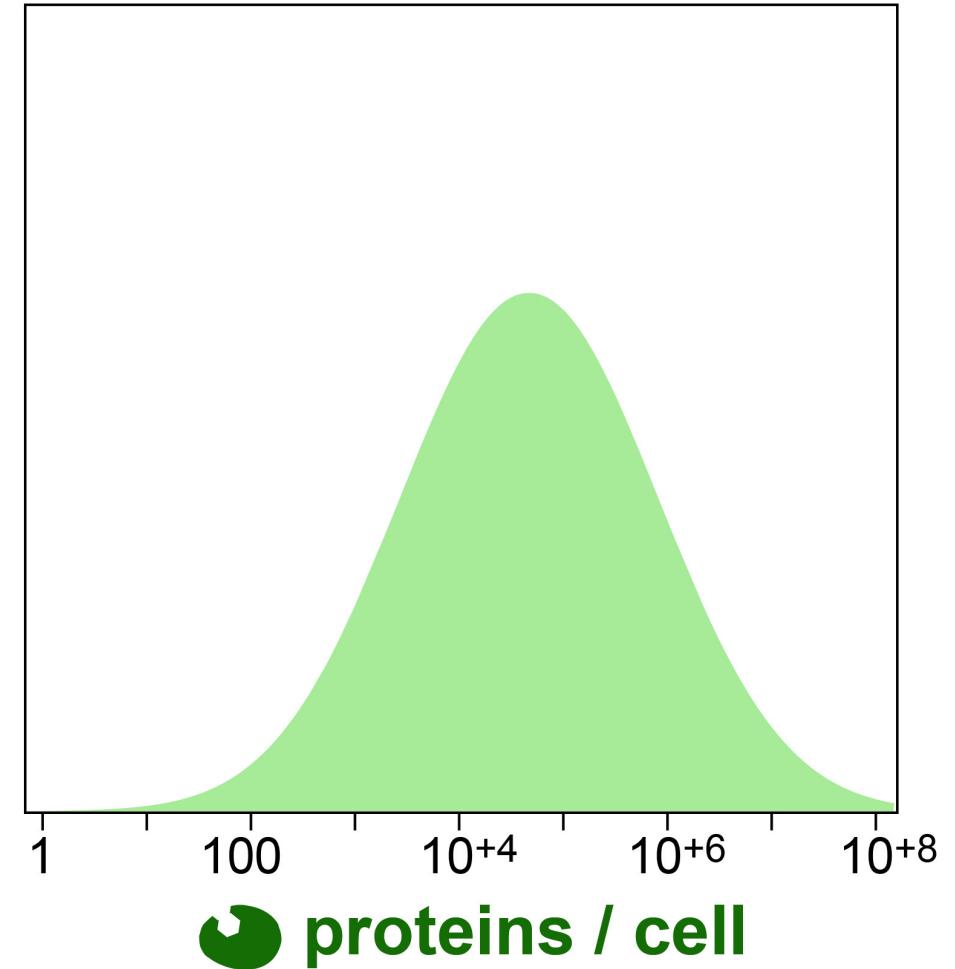
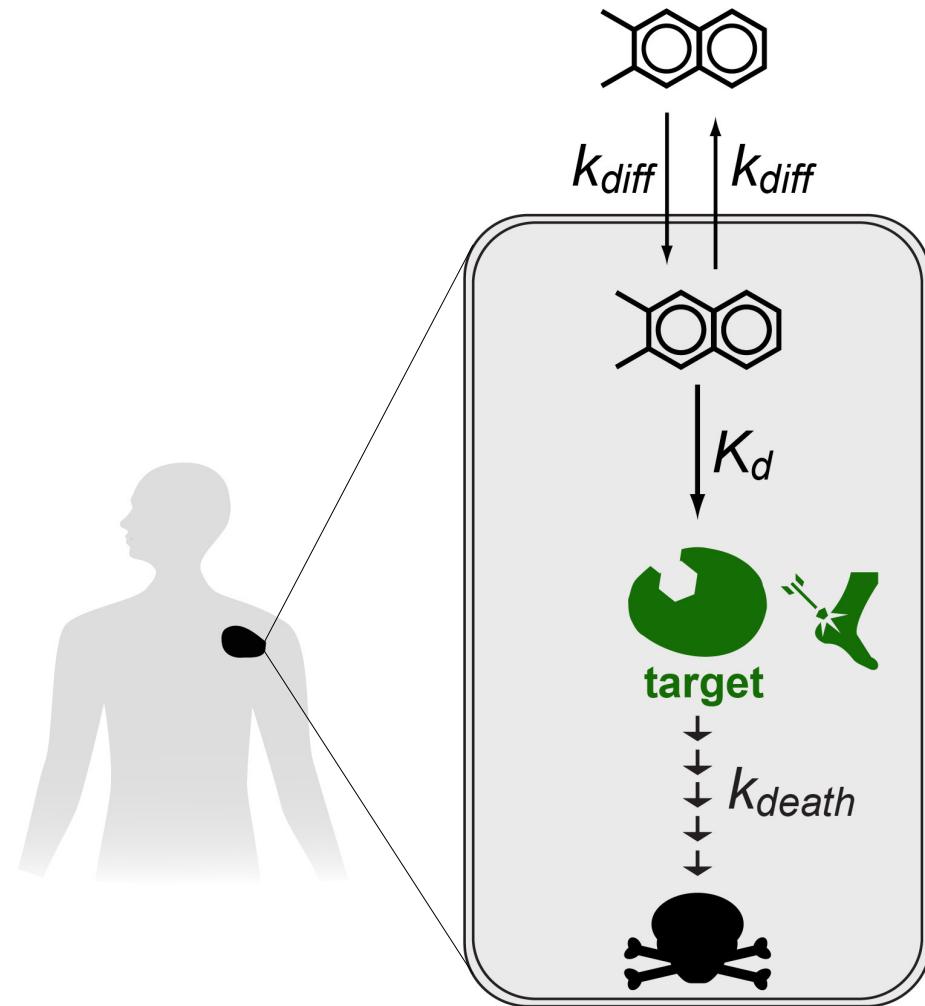


# 1-drug / 1 target paradigm: *historical artifact*

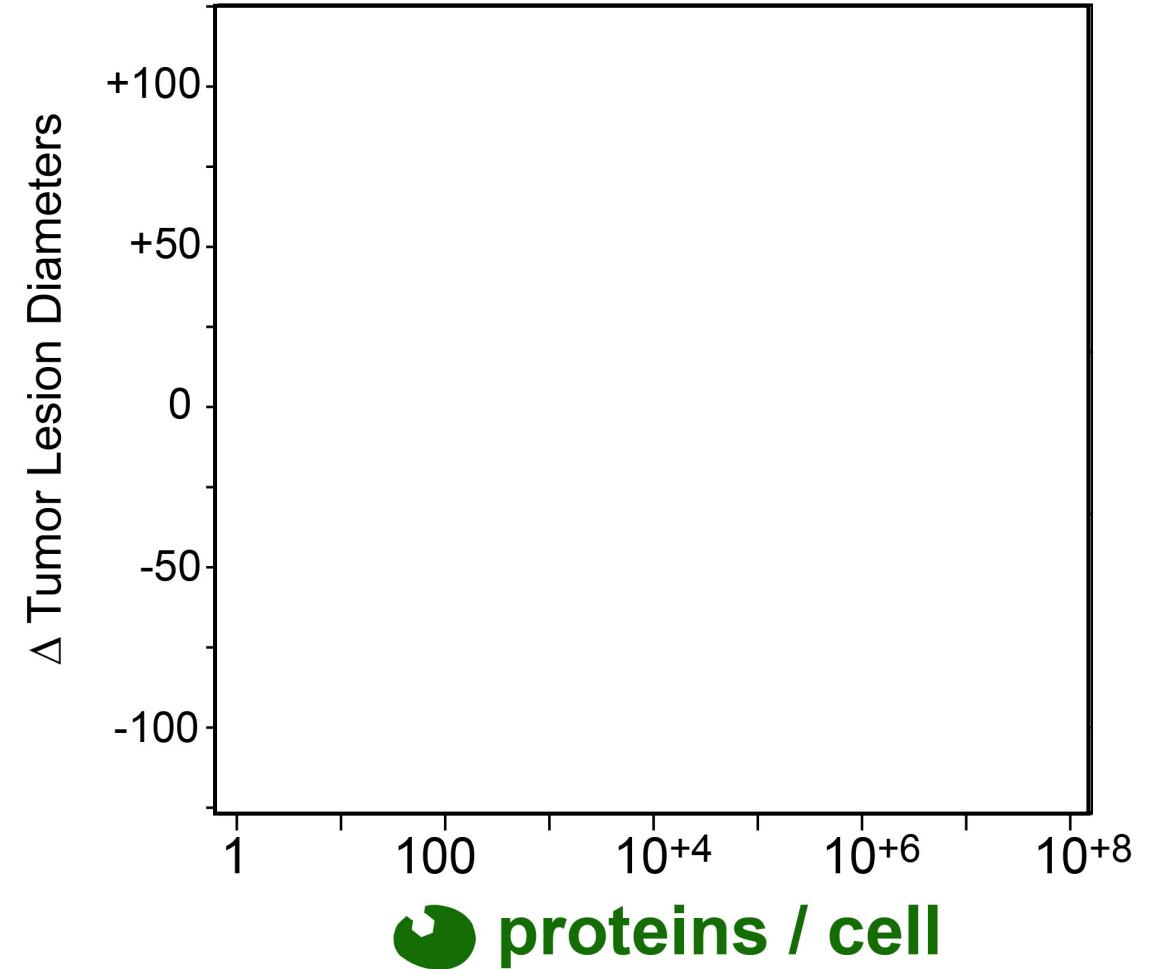
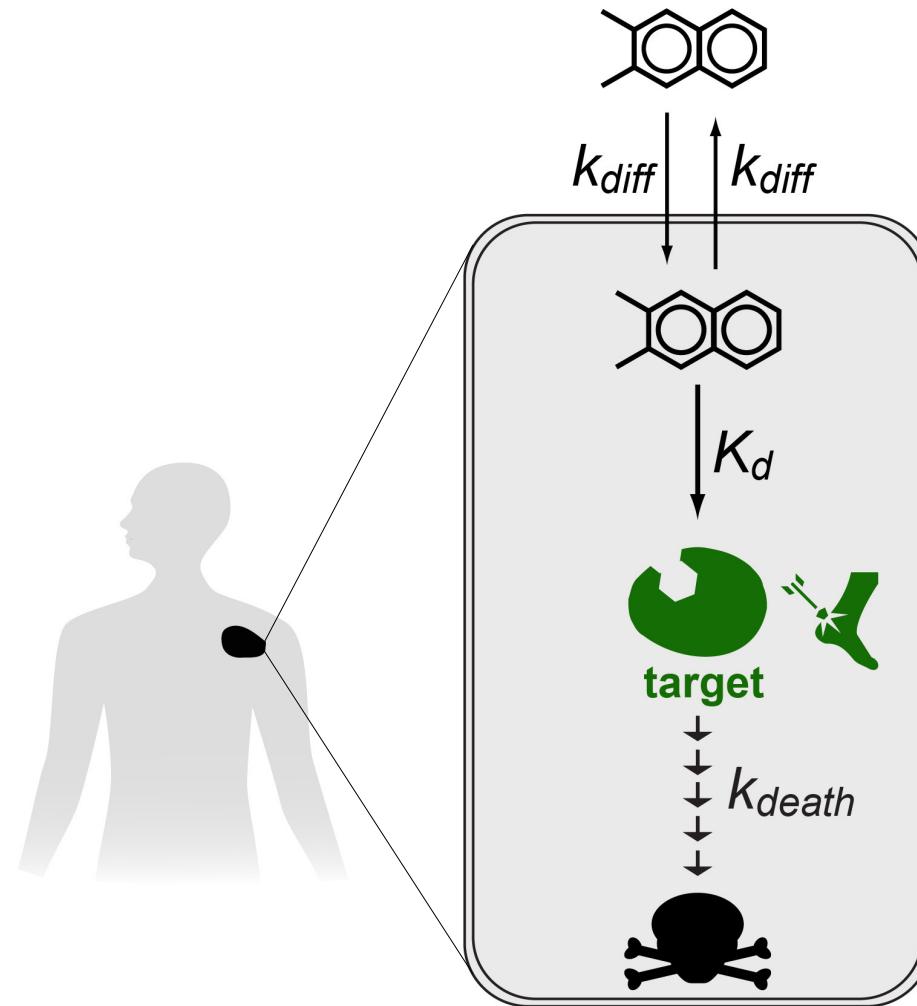
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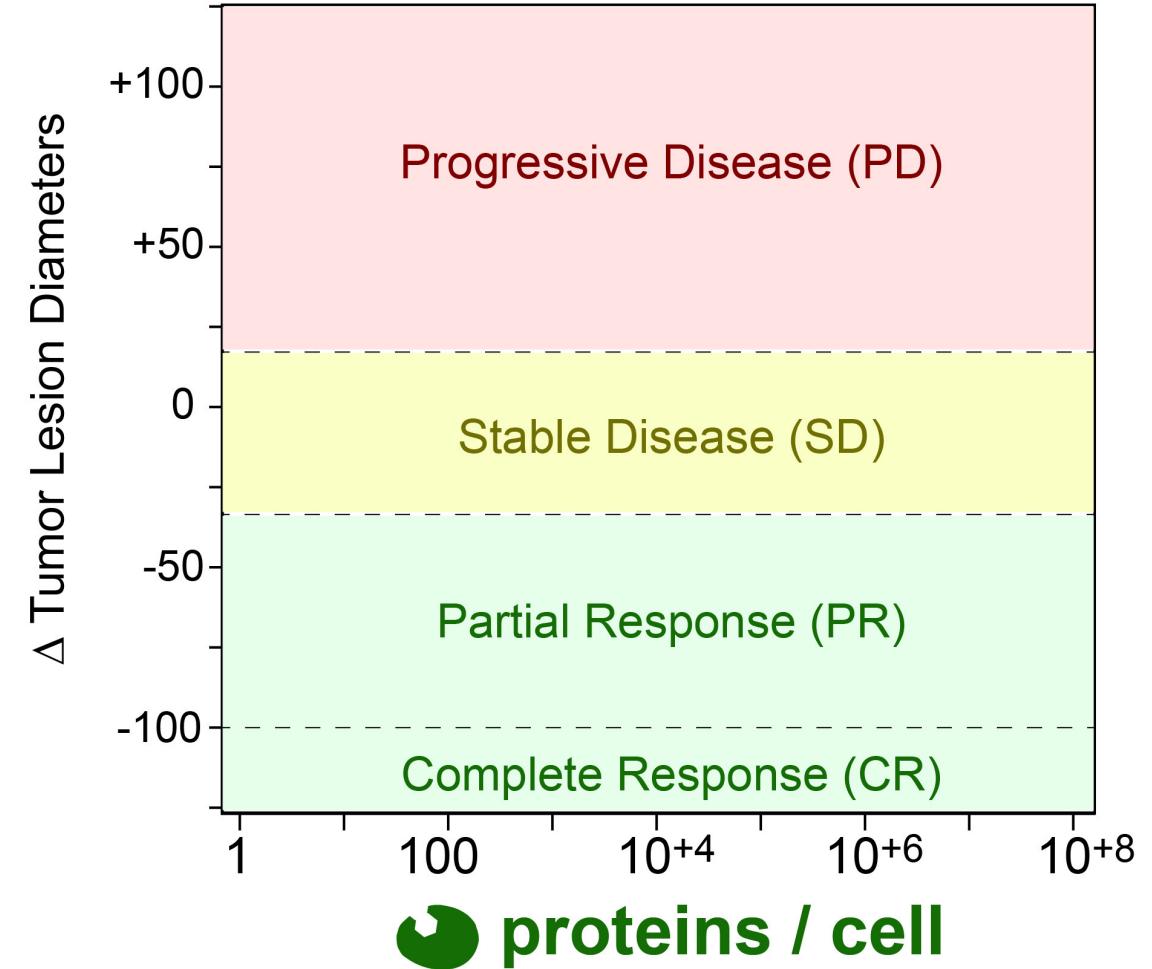
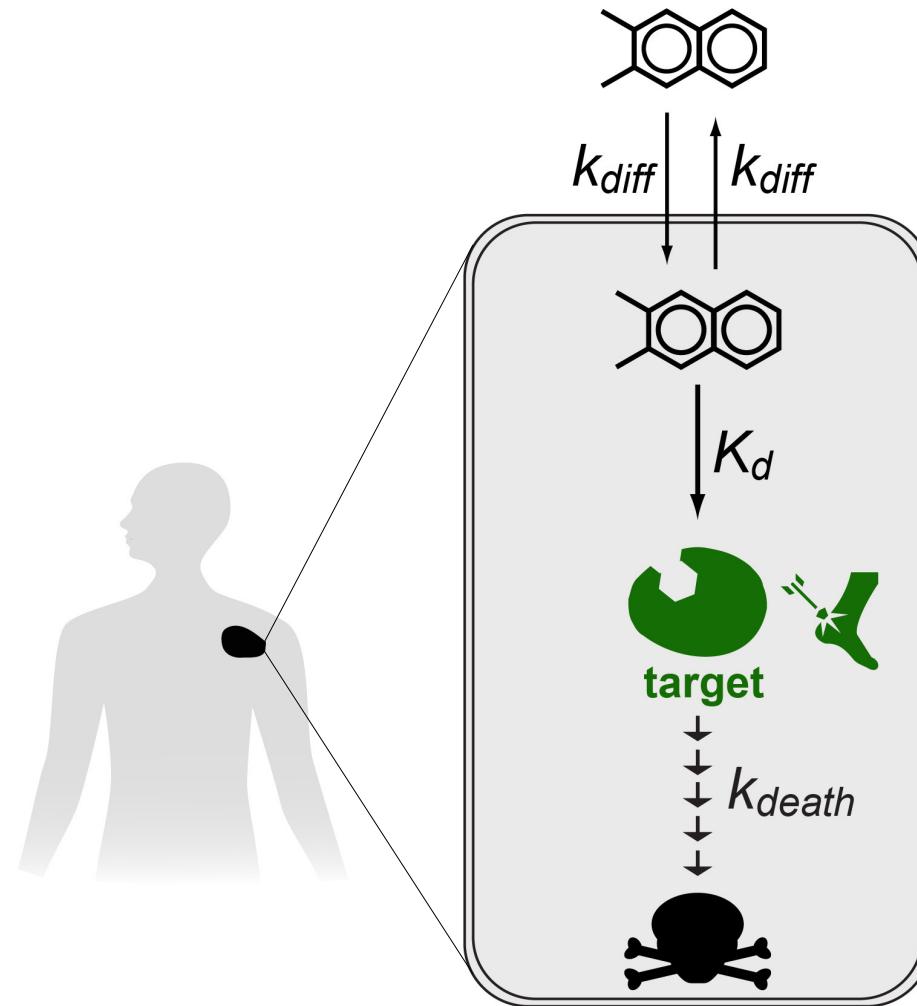
# Biomarker Expression: *log-normal distribution*



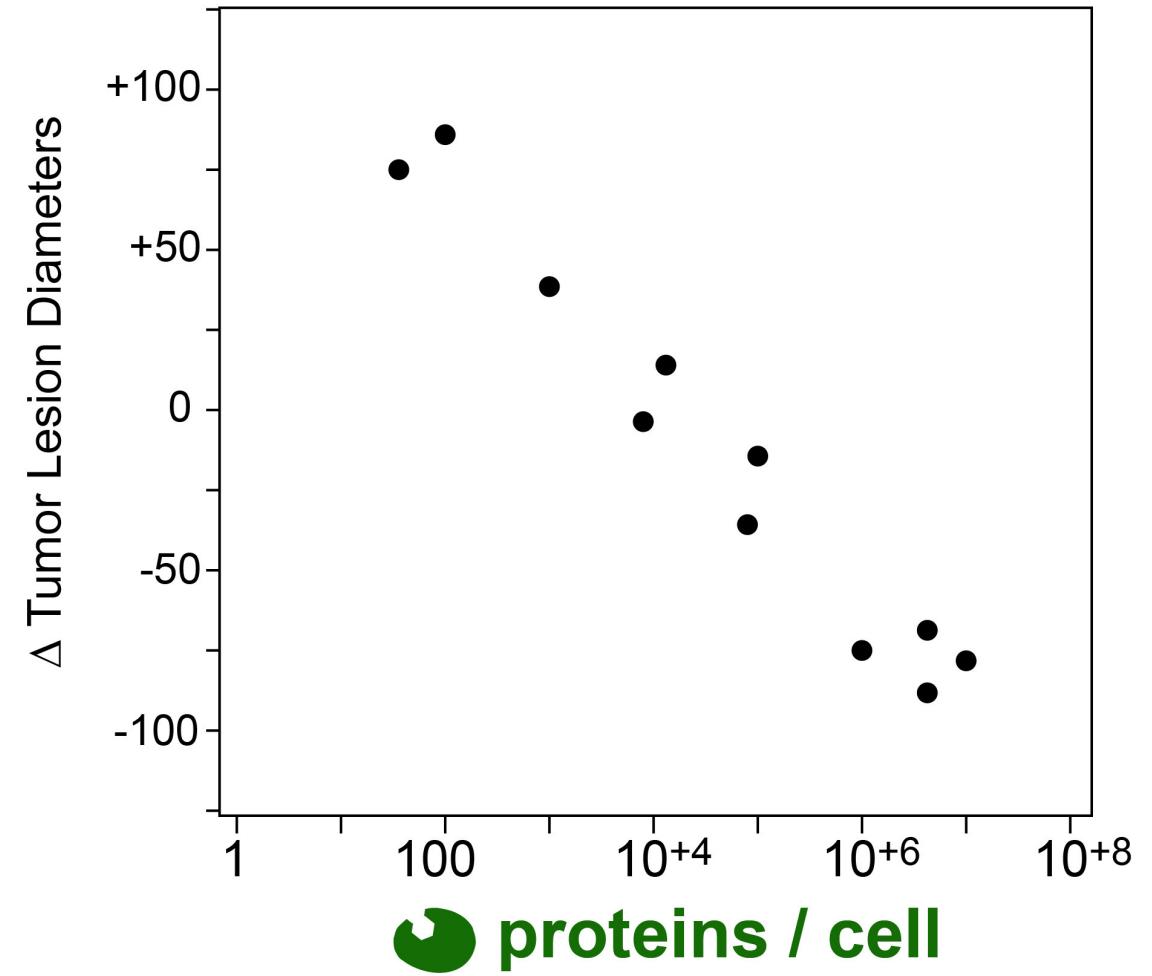
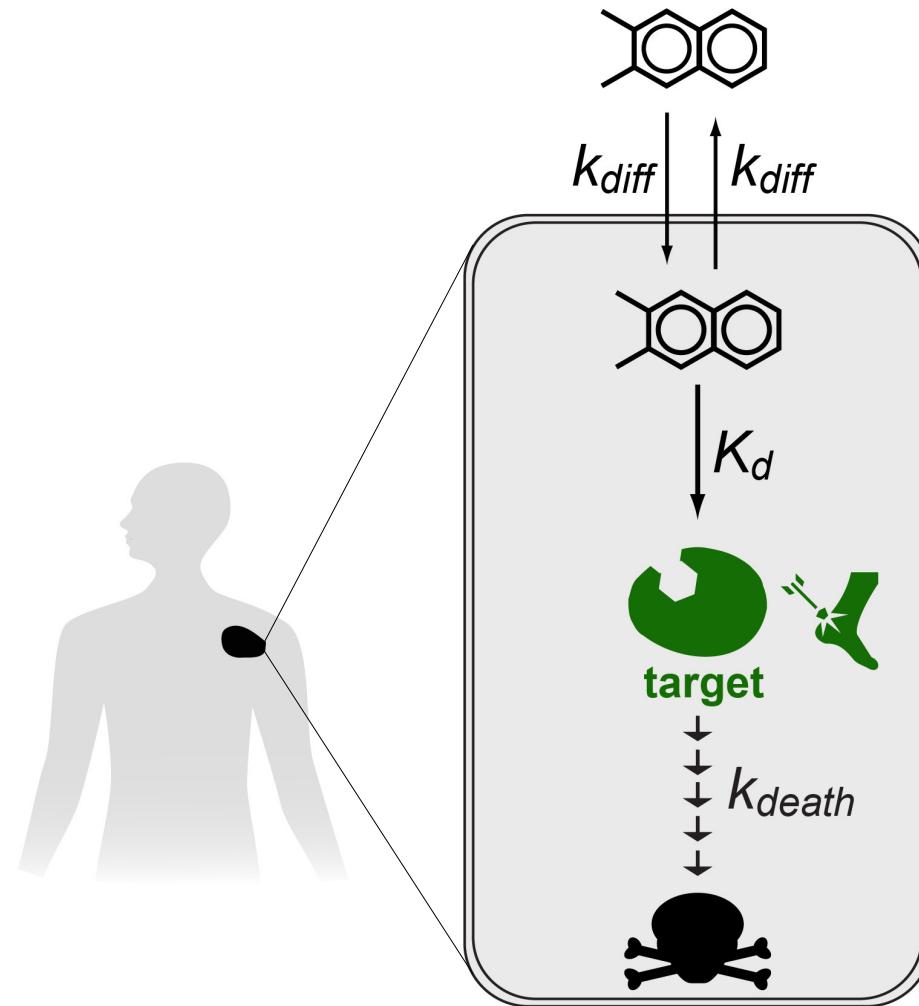
# Clinical Response Definitions: lesion diameter



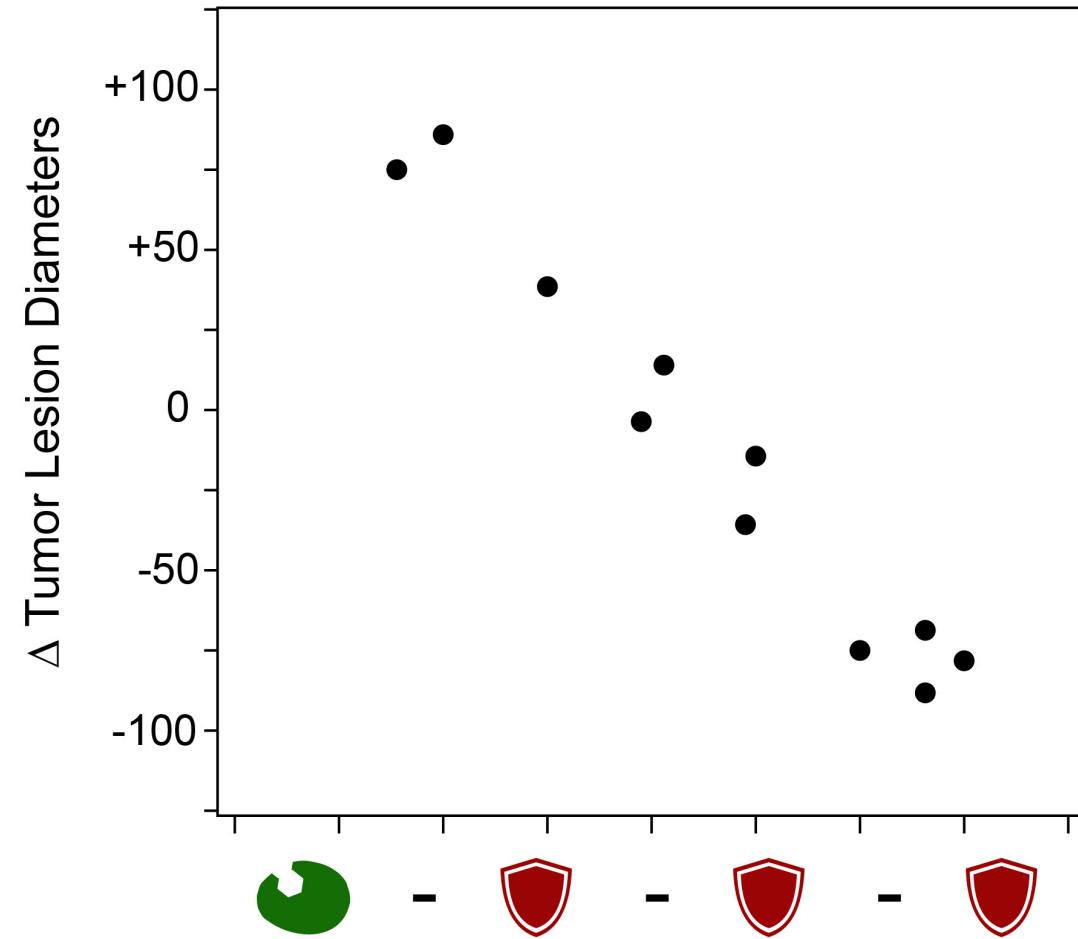
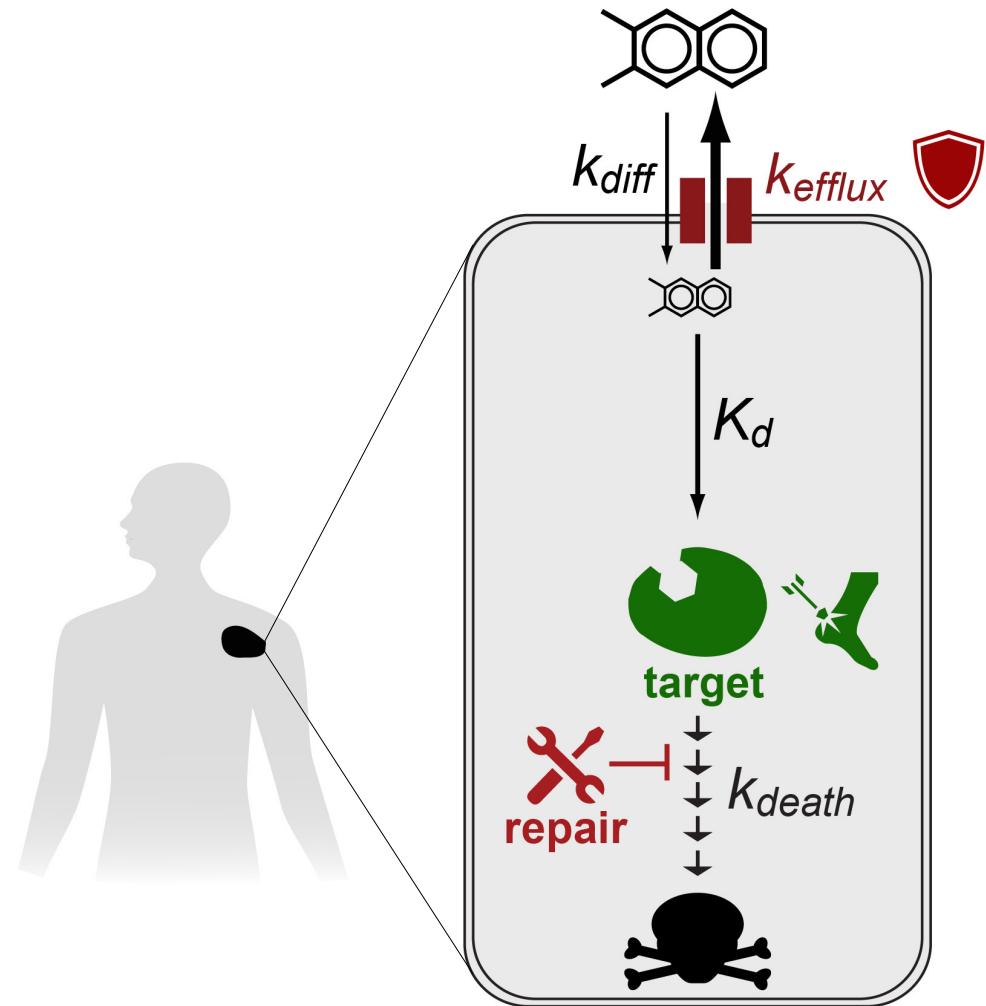
# Clinical Response Definitions: lesion diameter



# Clinical Biomarker Research: focus on drug-target

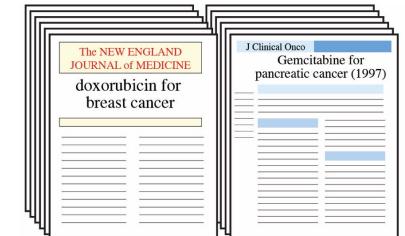
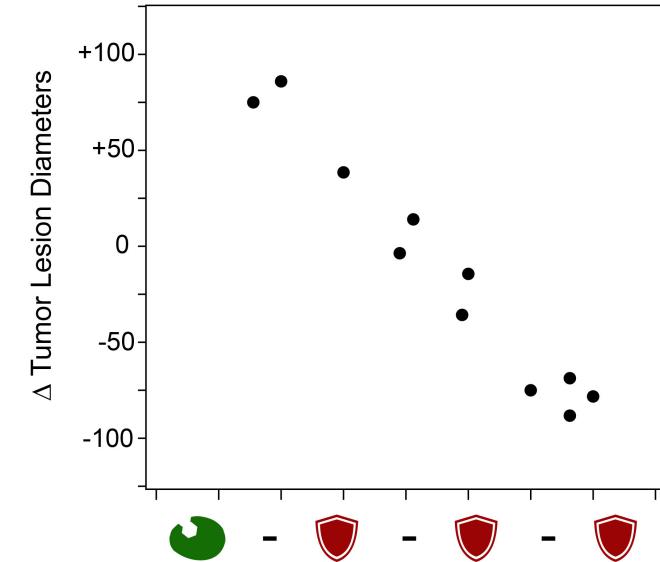
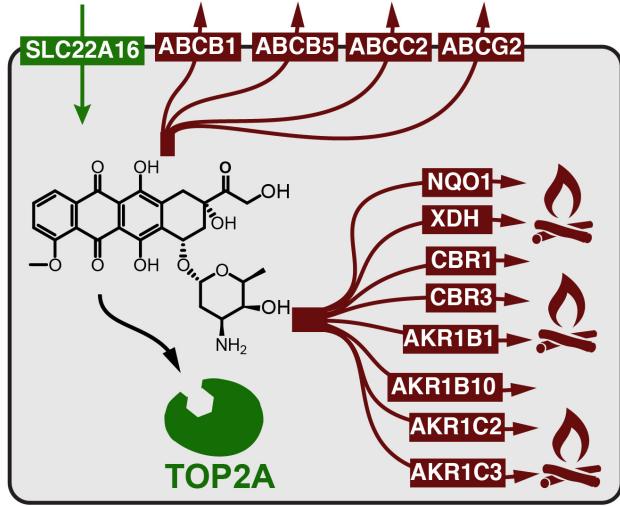
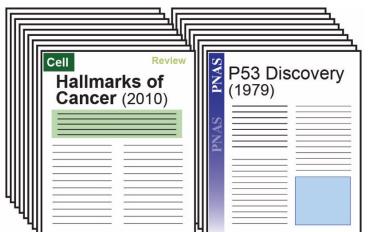


# Cytotoxic chemotherapies: 1-drug / N-biomarkers



# **EXPERIMENTAL WORK**

# How do you combine N-biomarkers?

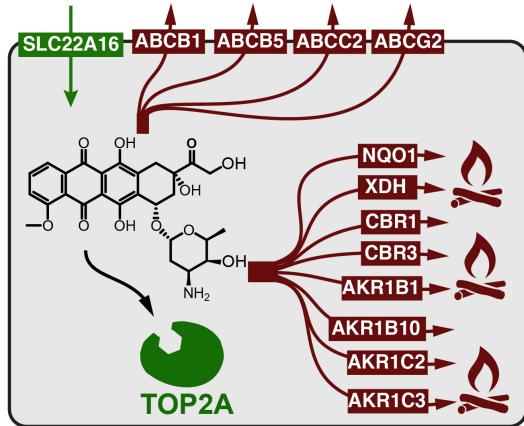


$$\Delta IC_{50} = \frac{k_{cat}/K_m}{k_{diff}} [ABCB1] + \frac{k_{cat}/K_m}{k_{diff}} [ABCG2] \dots$$

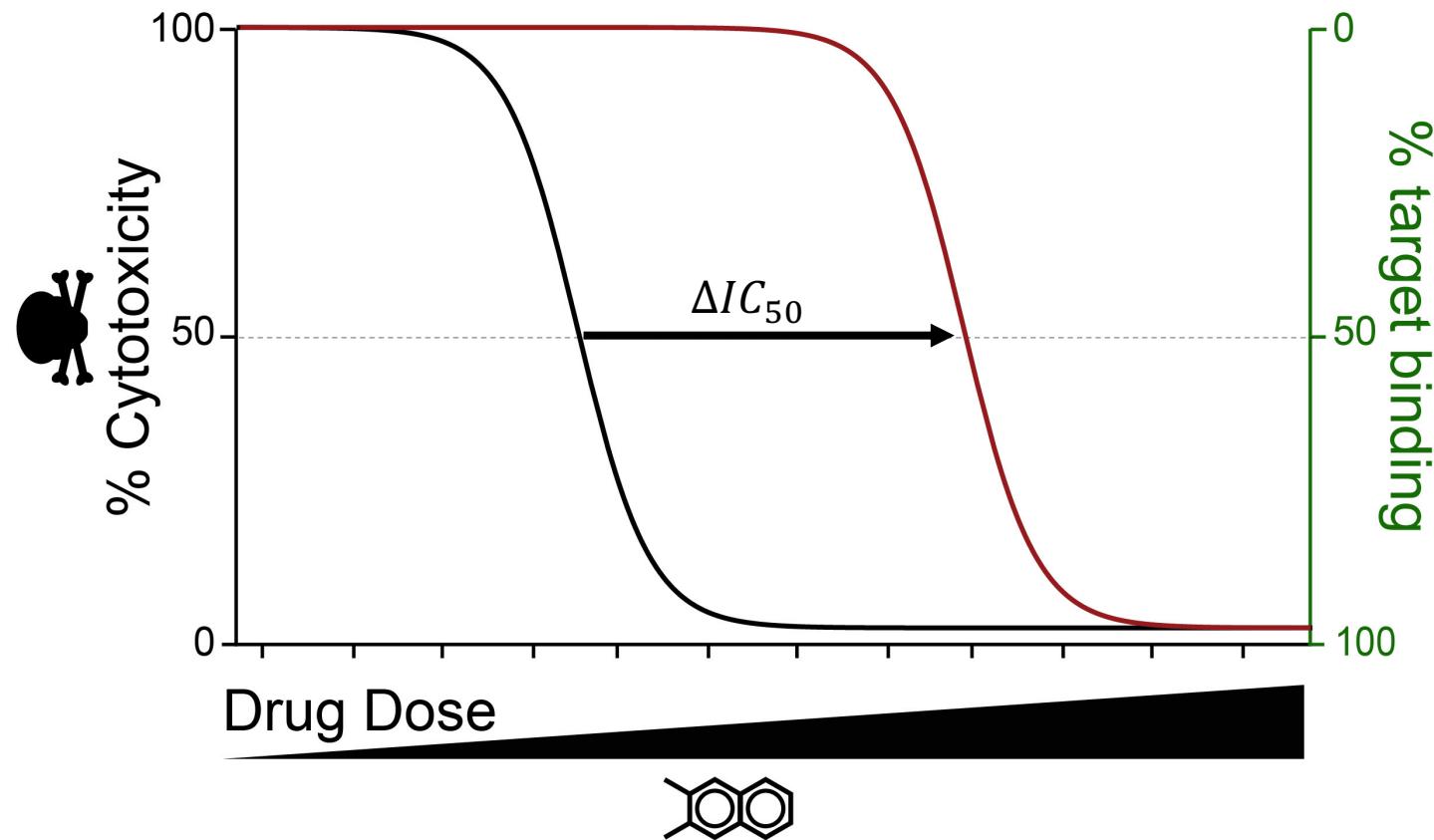
$$\Delta IC_{50} = \frac{k_{resist}}{k_{diff}} = \sum_{i=1}^N \frac{k_{cat}/K_m}{k_{diff}} [E]_t$$

$$\Delta\Delta IC_{50} = \frac{k_{cat}/K_m}{k_{diff}} \Delta [ABCB1]$$

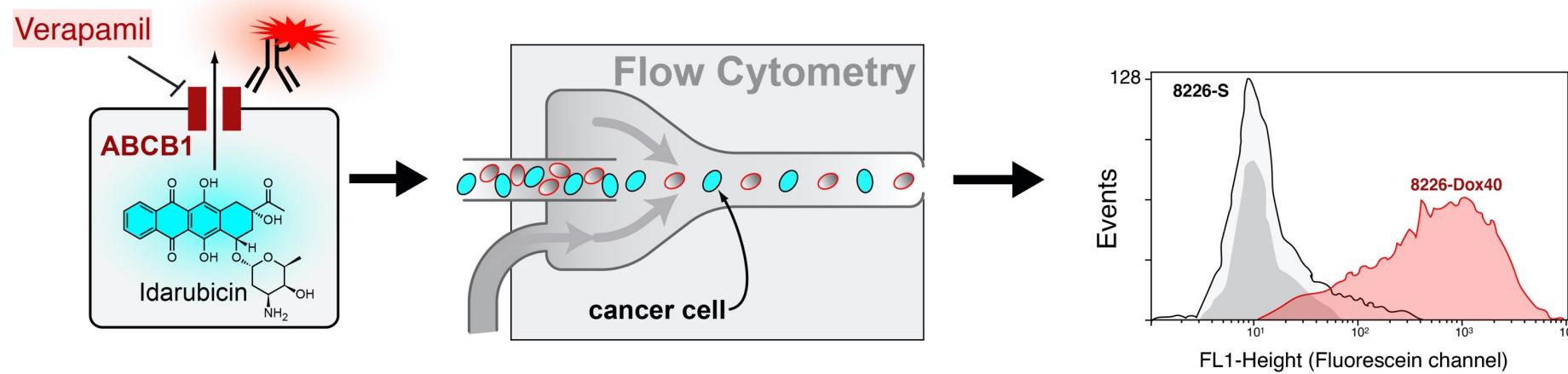
# Tool to study contribution of individual biomarkers



$$\frac{\Delta IC_{50}}{\Delta [ABCB1]} = \frac{k_{cat}/K_m}{k_{diff}}$$

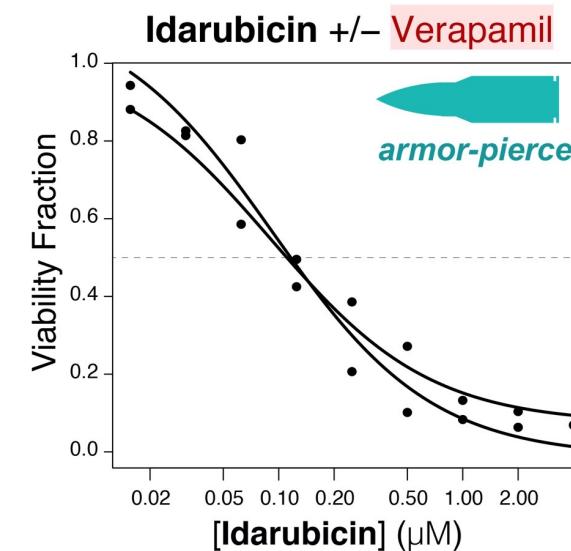
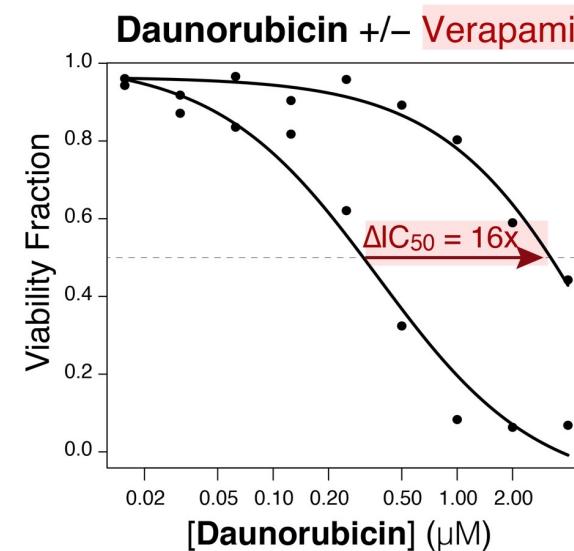
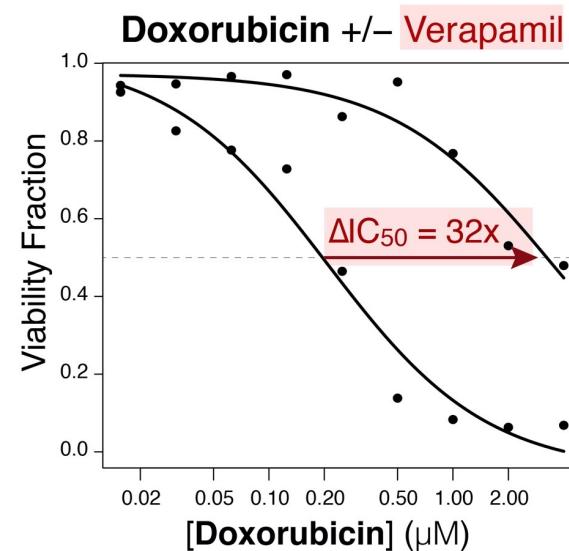
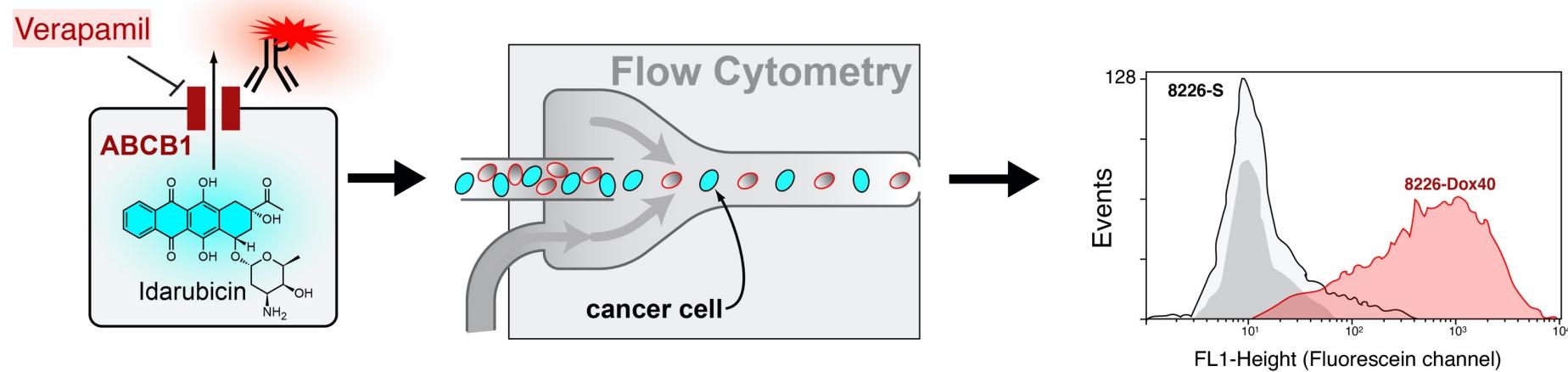


# Measure Biomarker Expression: *Flow cytometry*

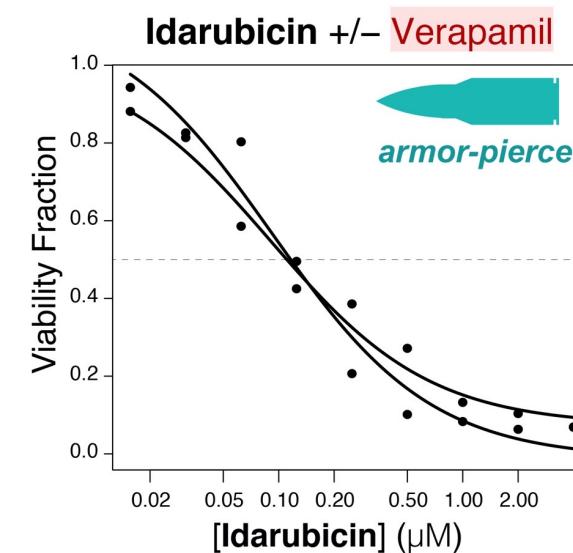
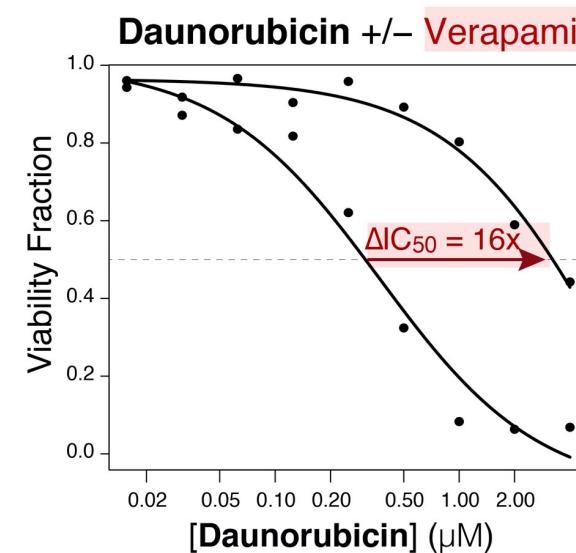
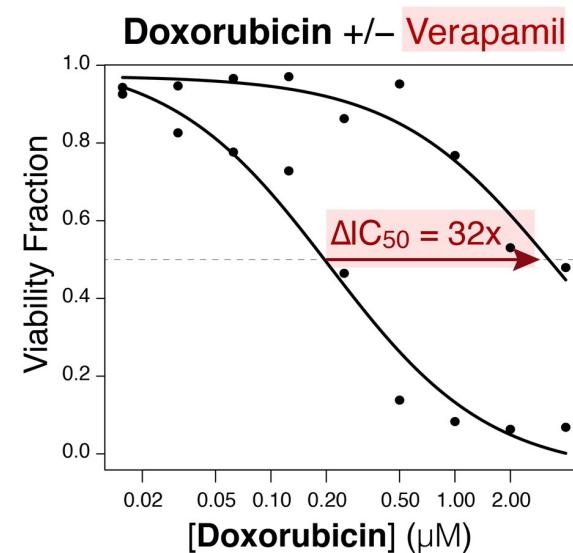
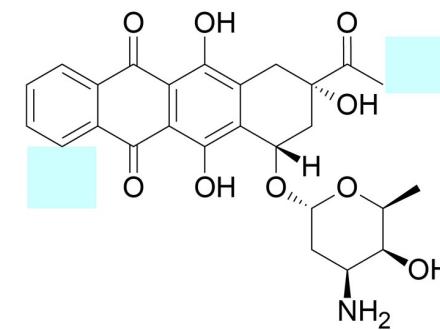
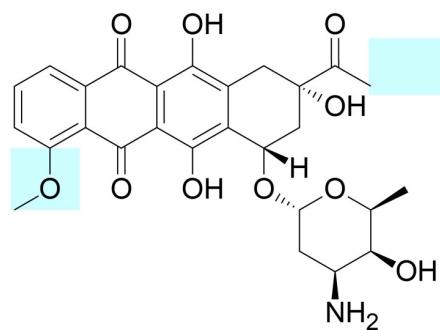
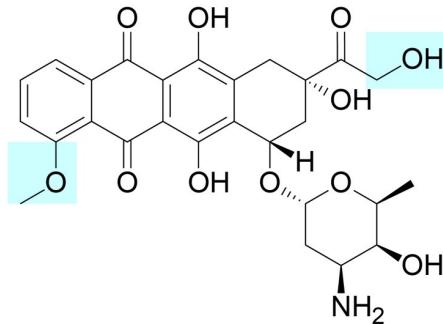


$$\frac{\Delta IC_{50}}{\Delta [ABCB1]} = \frac{k_{cat}/K_m}{k_{diff}}$$

# Measure Biomarker Expression: *Flow cytometry*

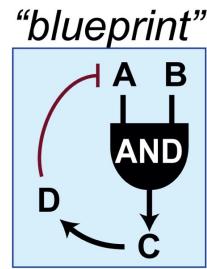


# Design “Armor Piercing Chemo”: Anthracyclines

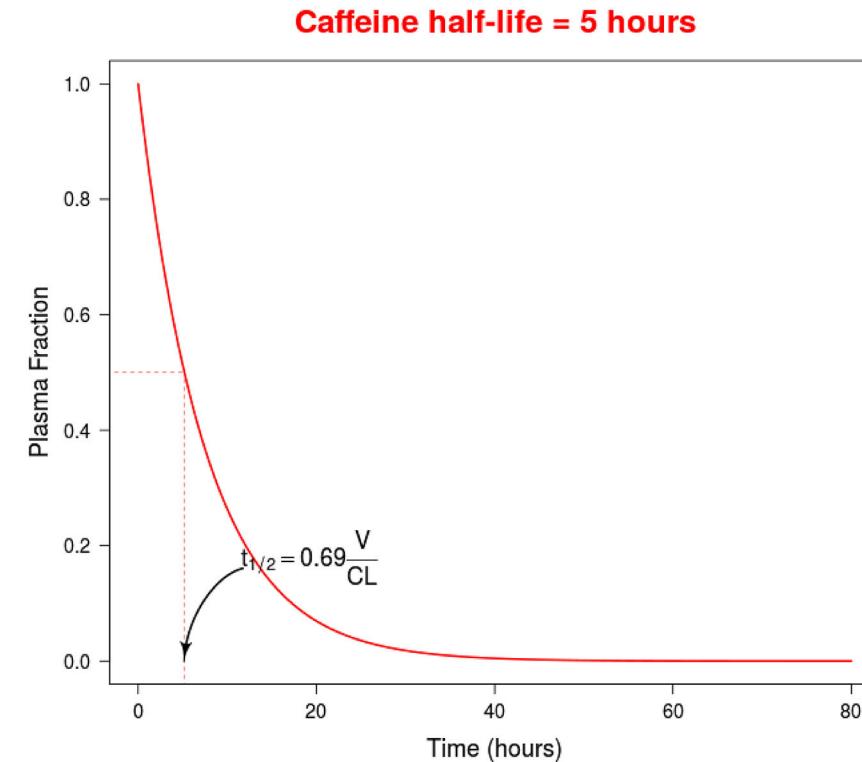
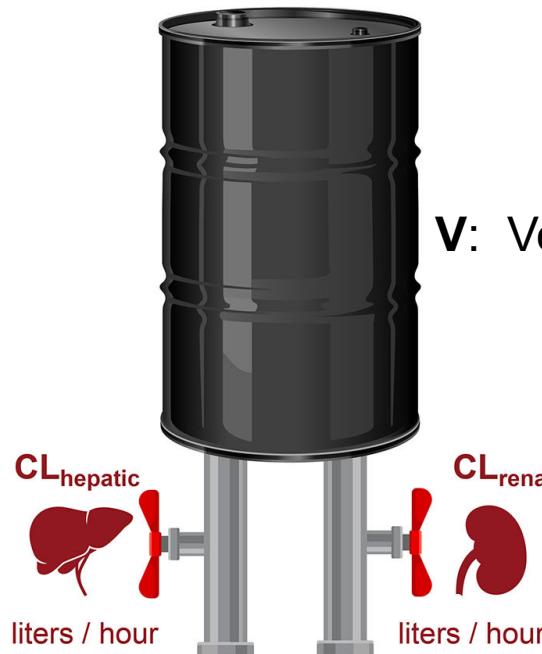


***Doctoral Student Educational Approach:***  
***experimental scientists***

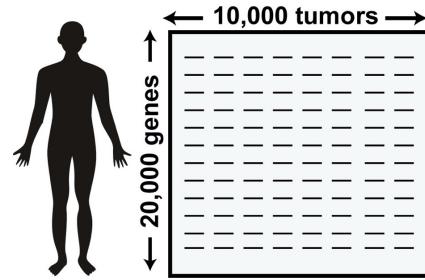
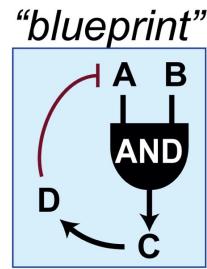
# *Pharmacist and Scientists: improving data-numeracy*



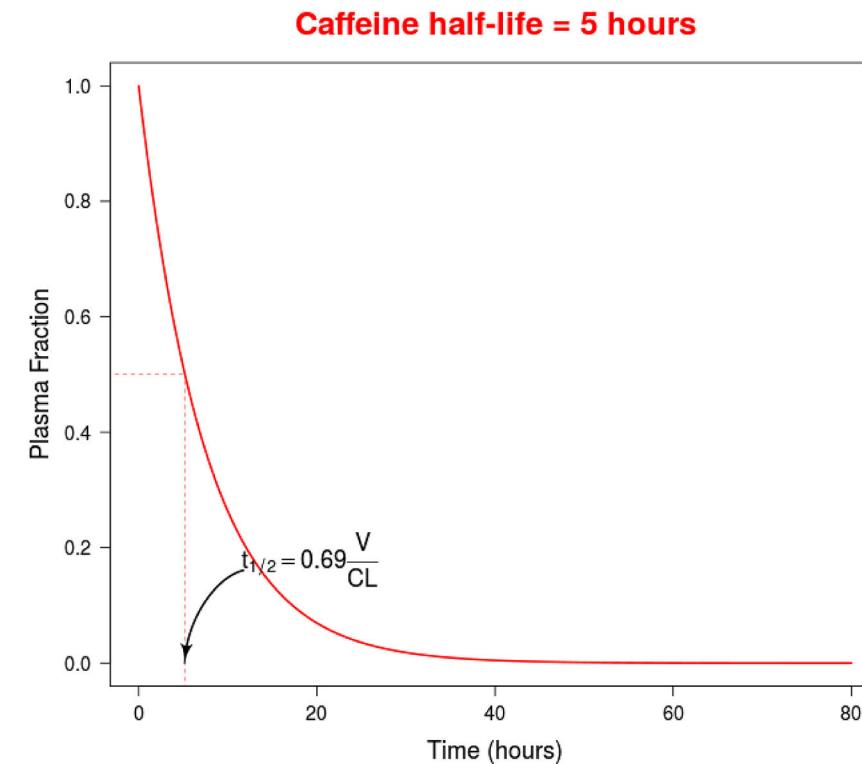
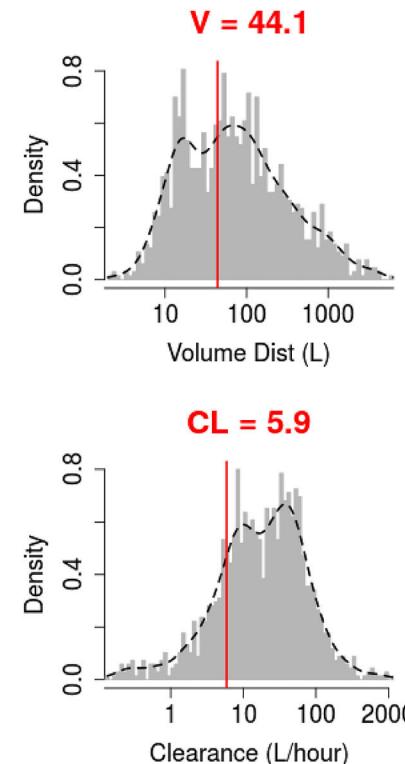
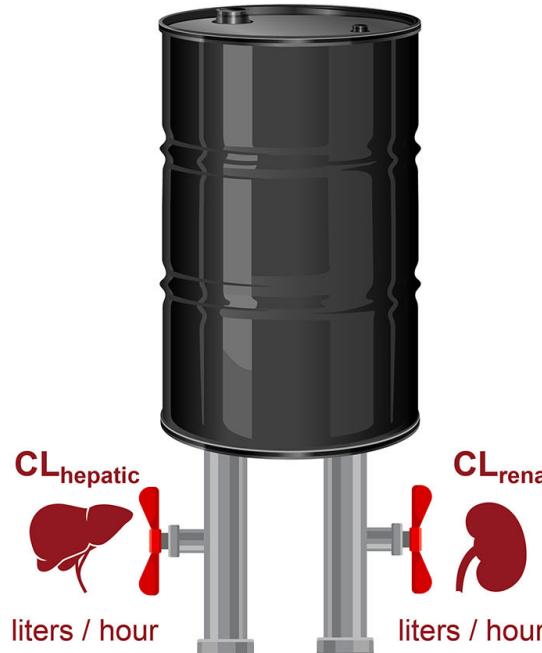
$$\text{Plasma Fraction} = e^{time \cdot CL/V}$$



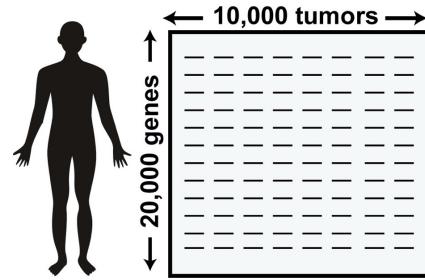
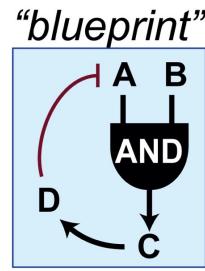
# Pharmacist and Scientists: improving data-numeracy



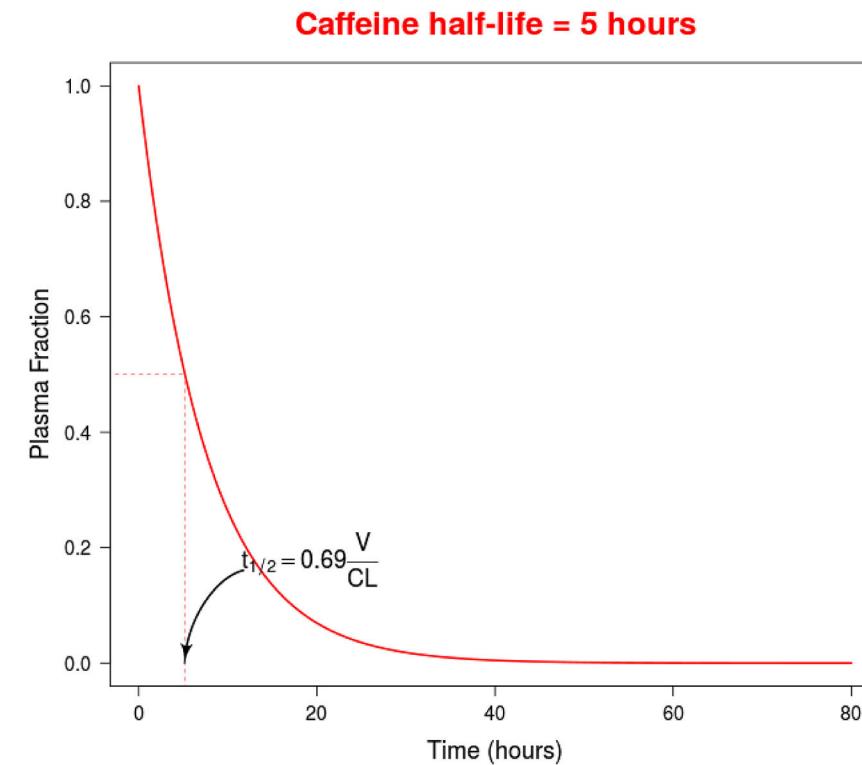
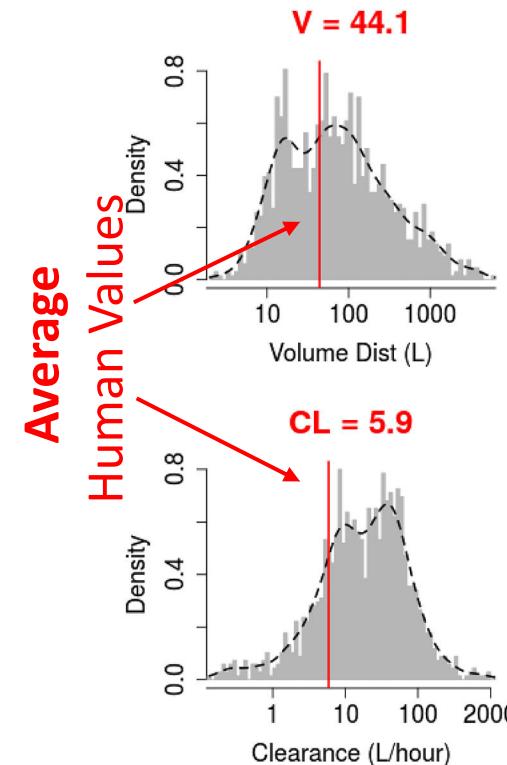
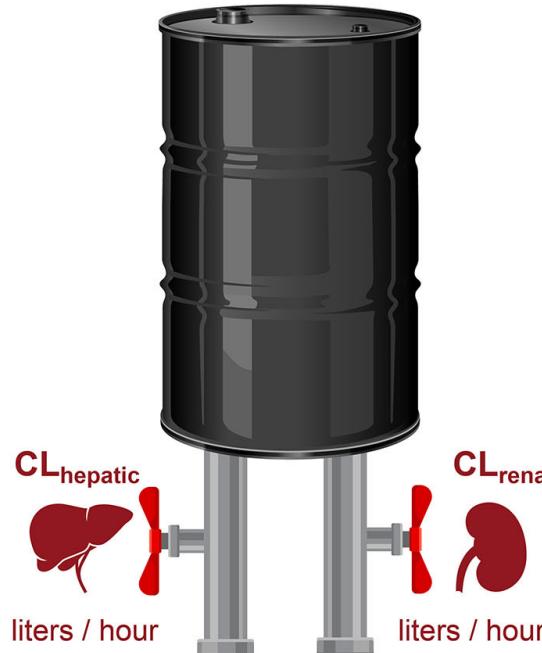
$$\text{Plasma Fraction} = e^{\text{time} \cdot CL/V}$$



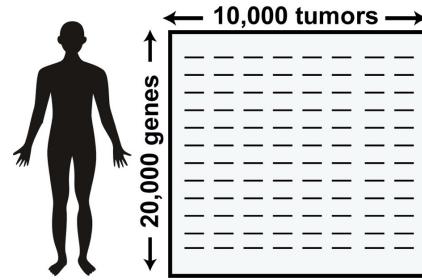
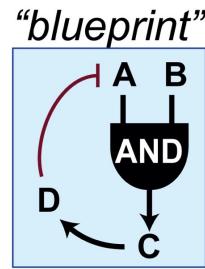
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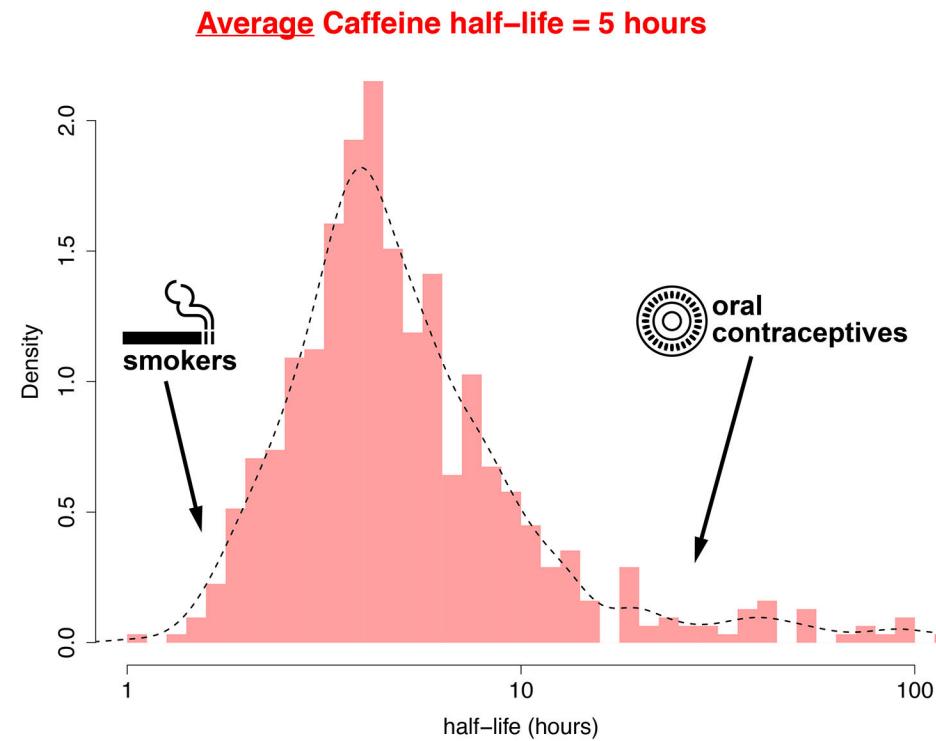
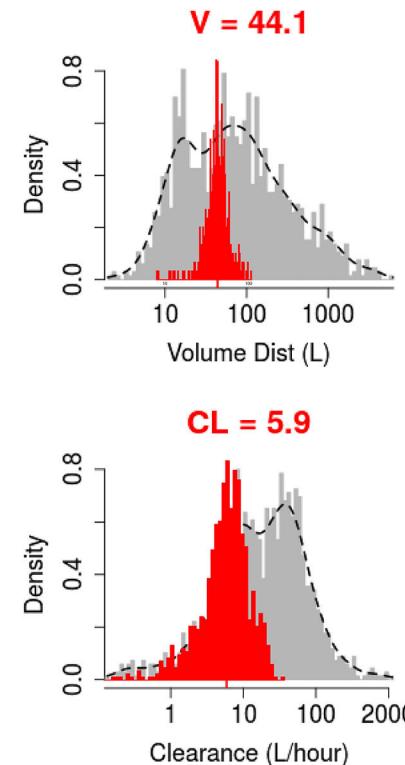
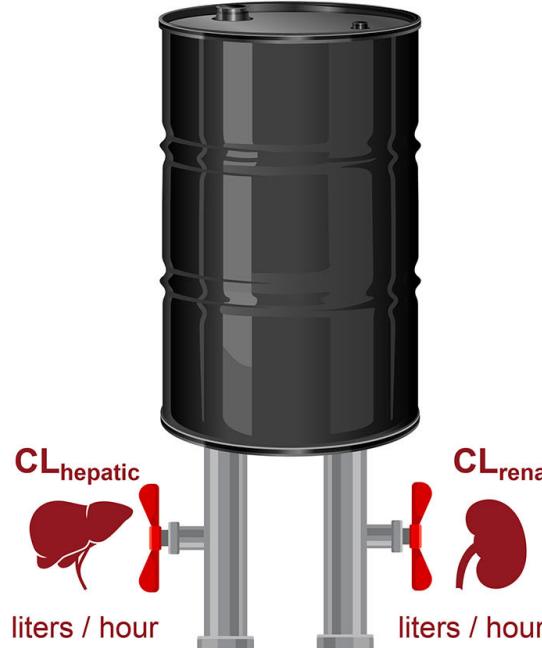
$$\text{Plasma Fraction} = e^{\text{time} \cdot CL/V}$$



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$$\text{Plasma Fraction} = e^{time \cdot CL/V}$$



# Webtool: <https://douglasslab.com/pk-db-iv/>

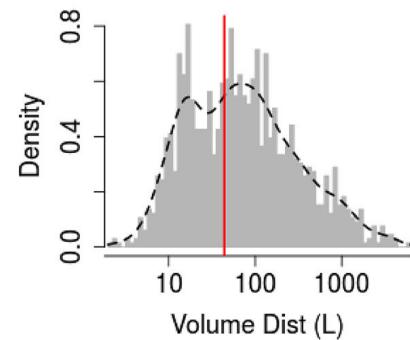
## PK DATABASE:

The plot below illustrates the IV-kinetics of ~1000 drugs whose pharmacokinetic parameters from a were taken from a database published in 2018 (Drug Metab Dispos 2018, 46, 1466)

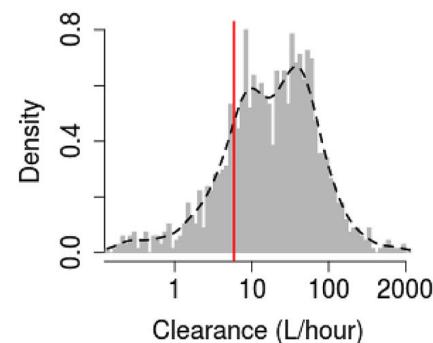
Type Drug Name below

Caffeine

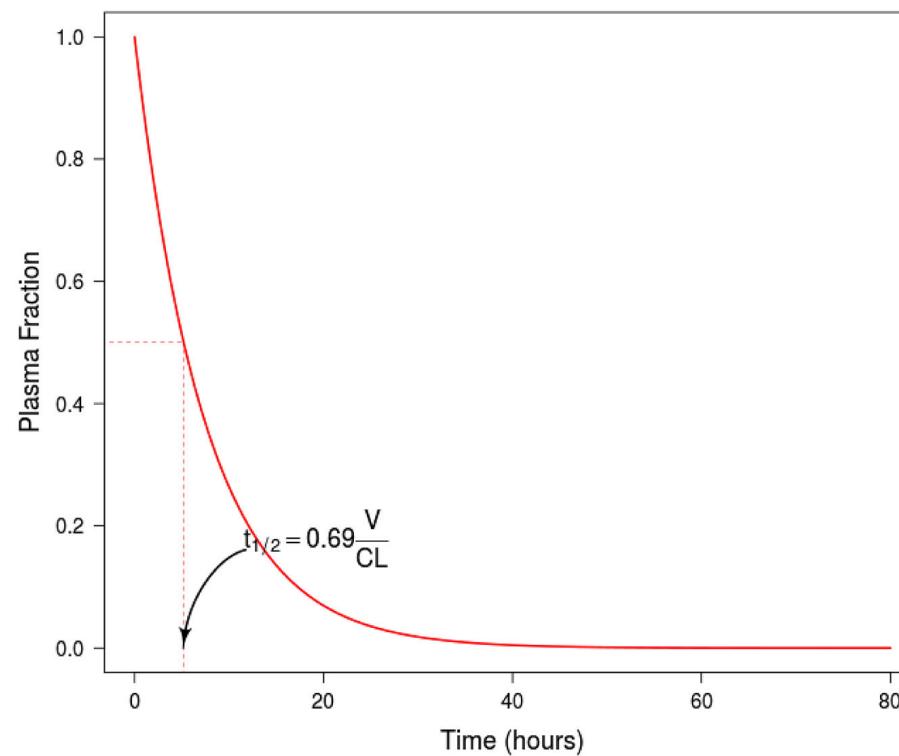
**V = 44.1**



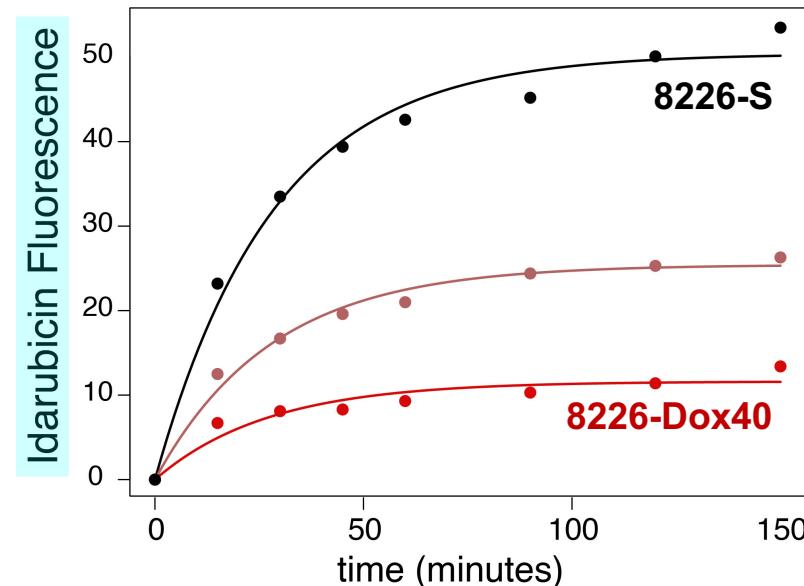
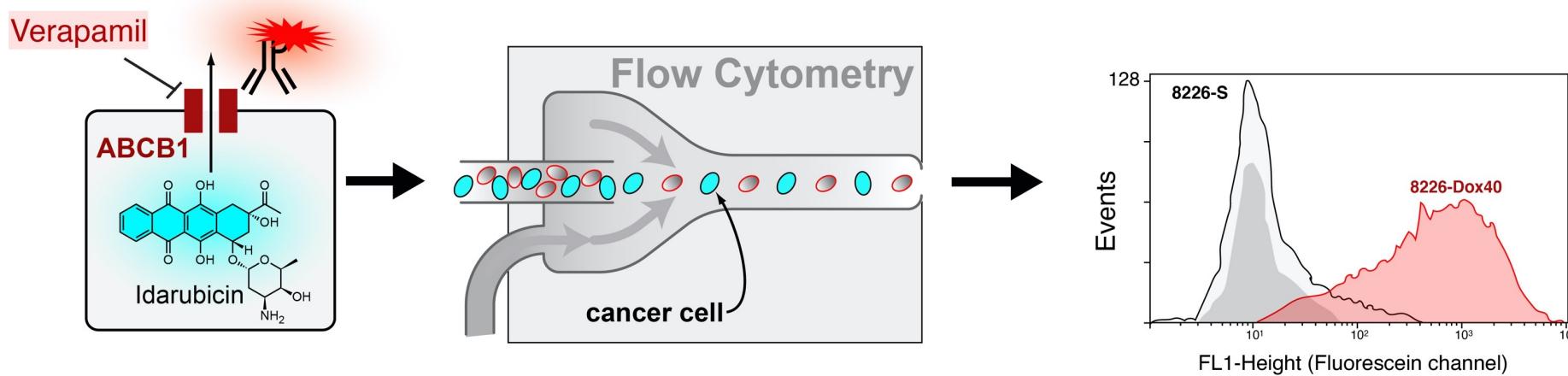
**CL = 5.9**



**Caffeine half-life = 5 hours**



# Deconvolute diffusion & enzyme flux



$$[IDA]_{intra} = [IDA]_{extra} \frac{k_{diff} \cdot K_m}{k_{cat} \cdot [ABCB1]}$$

$$t_{1/2} = \frac{1}{k_{diff} + [ABCB1] \cdot k_{cat}/K_m}$$