

# Small Molecule Discovery in Oncology and Beyond: Challenges and Opportunities

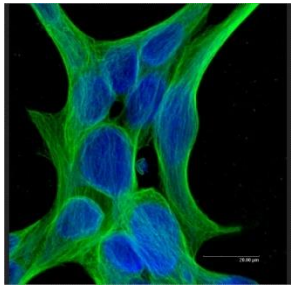
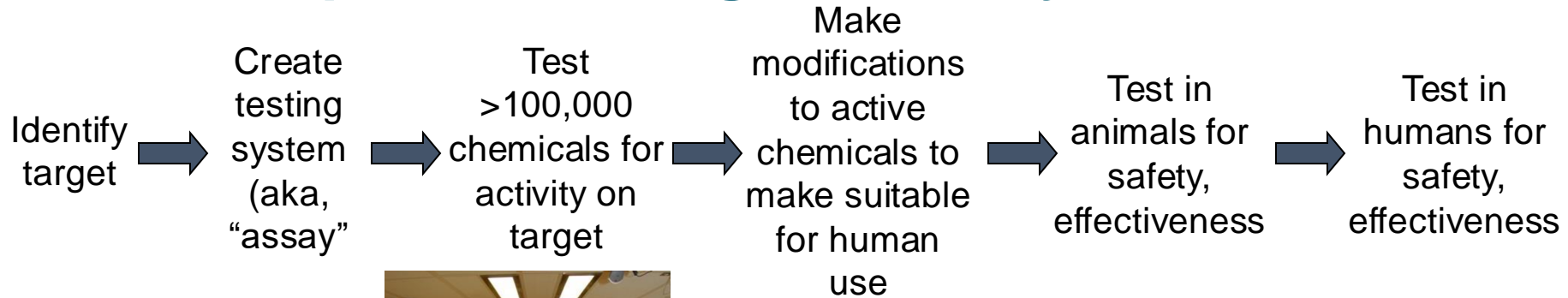
Anton Simeonov, Ph.D.

*Director, Chemical Genomics Branch, National Center for Advancing  
Translational Sciences (NCATS), National Institutes of Health (NIH)*

**TRACO Lecture  
September 30, 2024**



# Steps in the Drug Discovery Process



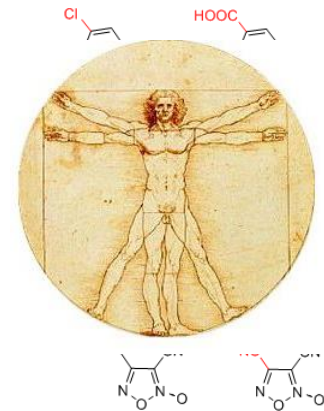
"Assay Development"



"HTS"



"Preclinical Development"



"Clinical Development"

Basic research

Target to Lead

Lead to First-in-Human

Biopharma,  
NIH CC,  
CTSA's

\$50-500K

\$500K-2M

\$5-20M

\$50-200M

(Per project cost)

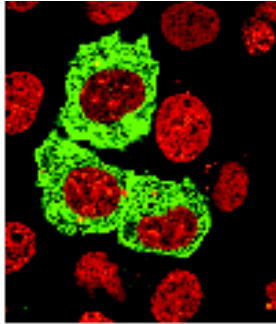


NIH National Center for Advancing Translational Sciences

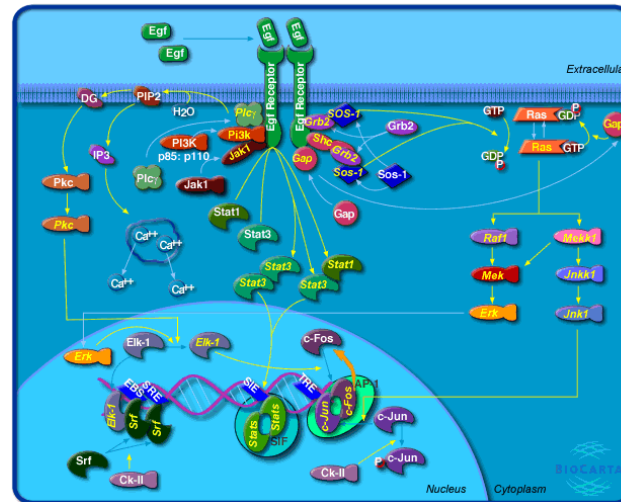
# Range of Screening Assays

*Extent of reductionism* →

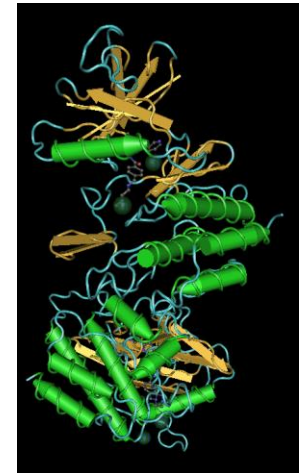
Phenotype  
(Image-based  
HCS, GFP, etc)



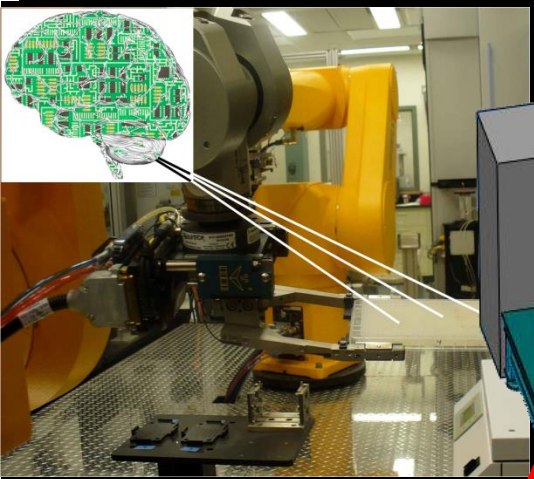
Pathway  
(Reporters, e.g., luciferase,  $\beta$ -  
lactamase)



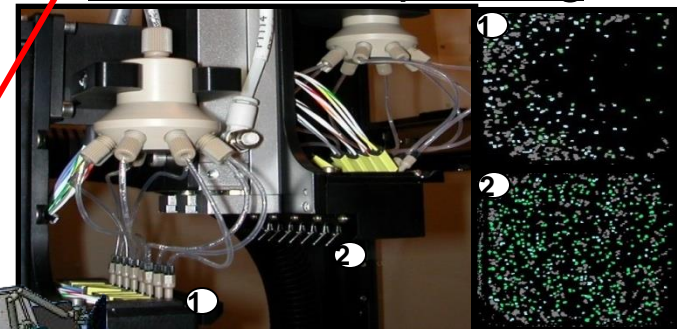
Protein  
(Enzyme readouts, interactions, etc)



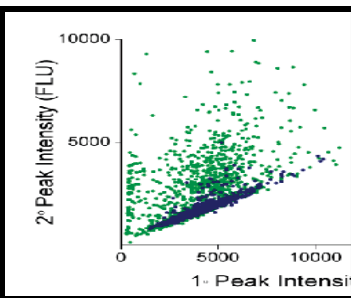
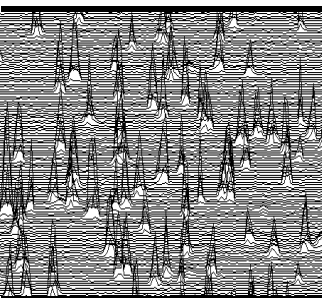
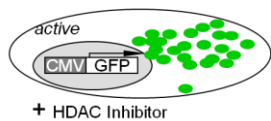
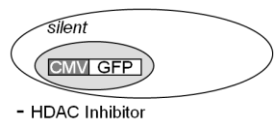
# Robotics & Informatics



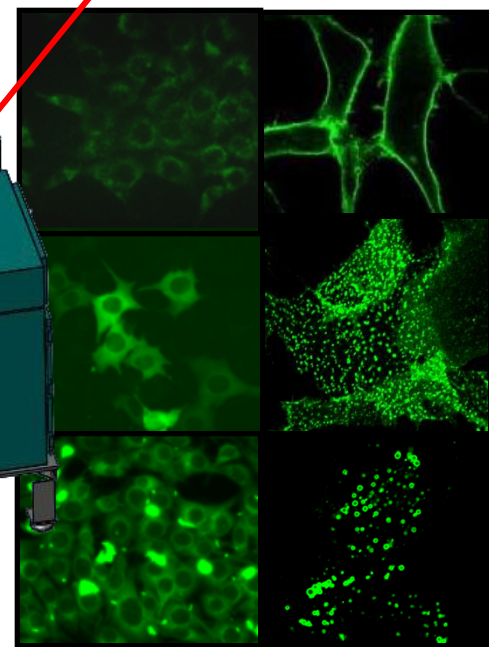
# Microliter Dispensing



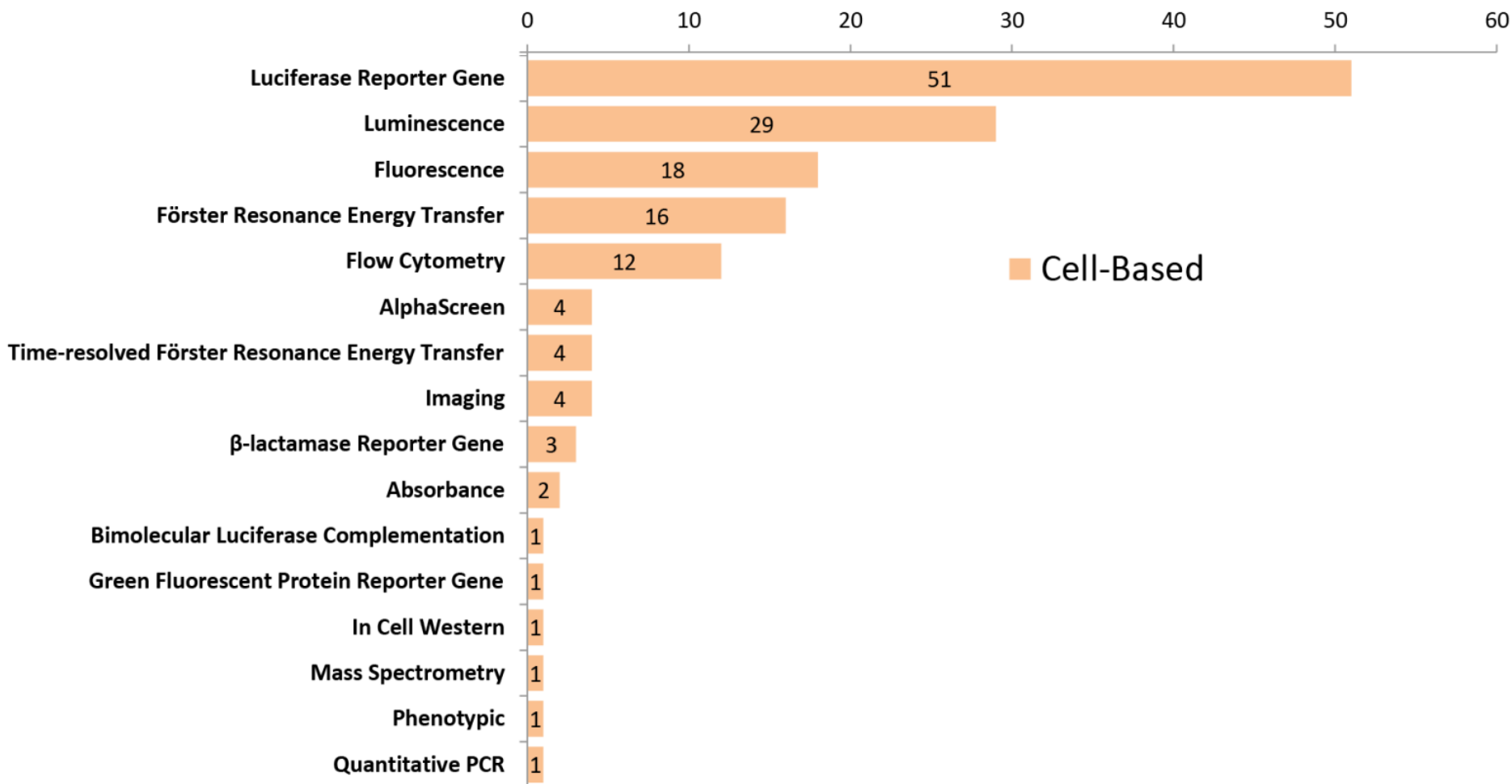
# Laser Cytometry



# Microscopy



# 149 Cancer Relevant Cell-Based HTS Assays from PubChem



Coussens, N. P., Braisted, J. C., Peryea, T., Sittampalam, S. G., Simeonov, A. and Hall, M. D. **Small Molecule Screens: A Gateway to Cancer Therapeutic Agents with Case Studies of FDA-Approved Drugs** *Pharmacological Reviews*, October 2017, 69 (4) 479-496



# Important Considerations for Choosing an Assay

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- **Assay expense**
  - Cost per well
  - Disposal cost(s)



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  - Select the best possible assays based on the available instrumentation



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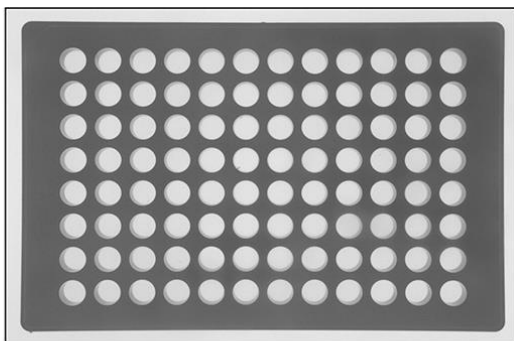
- **Assay expense**
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- **Assay throughput**
  - Miniaturization reduces the cost per well



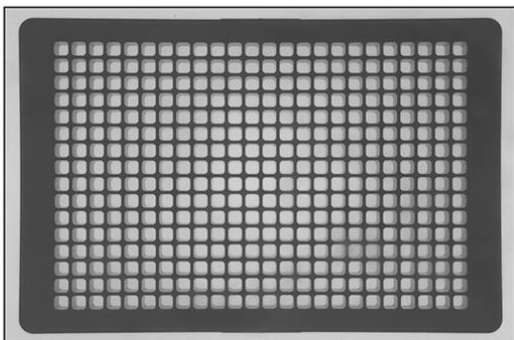


# Assay Miniaturization Saves Time and Reagents

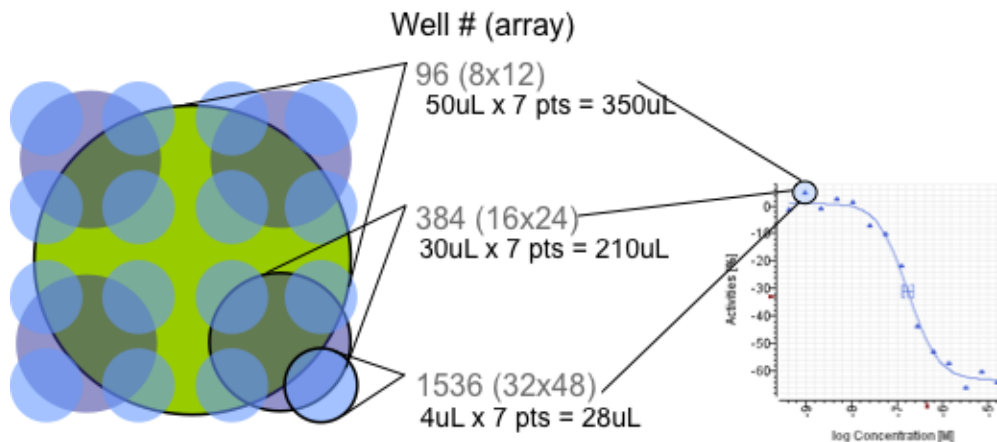
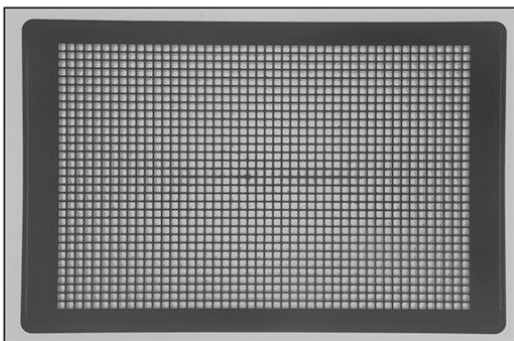
**96-well**  
~50,000 cells/well



**384-well**  
~5,000 cells/well



**1536-well**  
~500 cells/well



	96	384	1536
Plates per 100,000 compounds:	1,042	261	66
Assay volume (uL):	50-200	30-50	2-8
Adherent cell seeding density:	~10,000	~2,000	~500

Horman, Shane R. "Complex High-Content Phenotypic Screening." Special Topics in Drug Discovery. InTech, 2016.



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- **Ability to multiplex**
  - Can the response be measured by a single parameter; is multiparametric output possible?
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  - Can guide hit selection by differentiating selectivity among related targets
  - Can distinguish pathway inhibition from cytotoxicity in a cell-based assay



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- **Reagents**
  - Stability for hours is important
  - Consistency is critical (ideally obtain a large quantity from a single lot)
  - All reagents need to be validated (cell lines, antibodies, enzymatic purity, etc.)



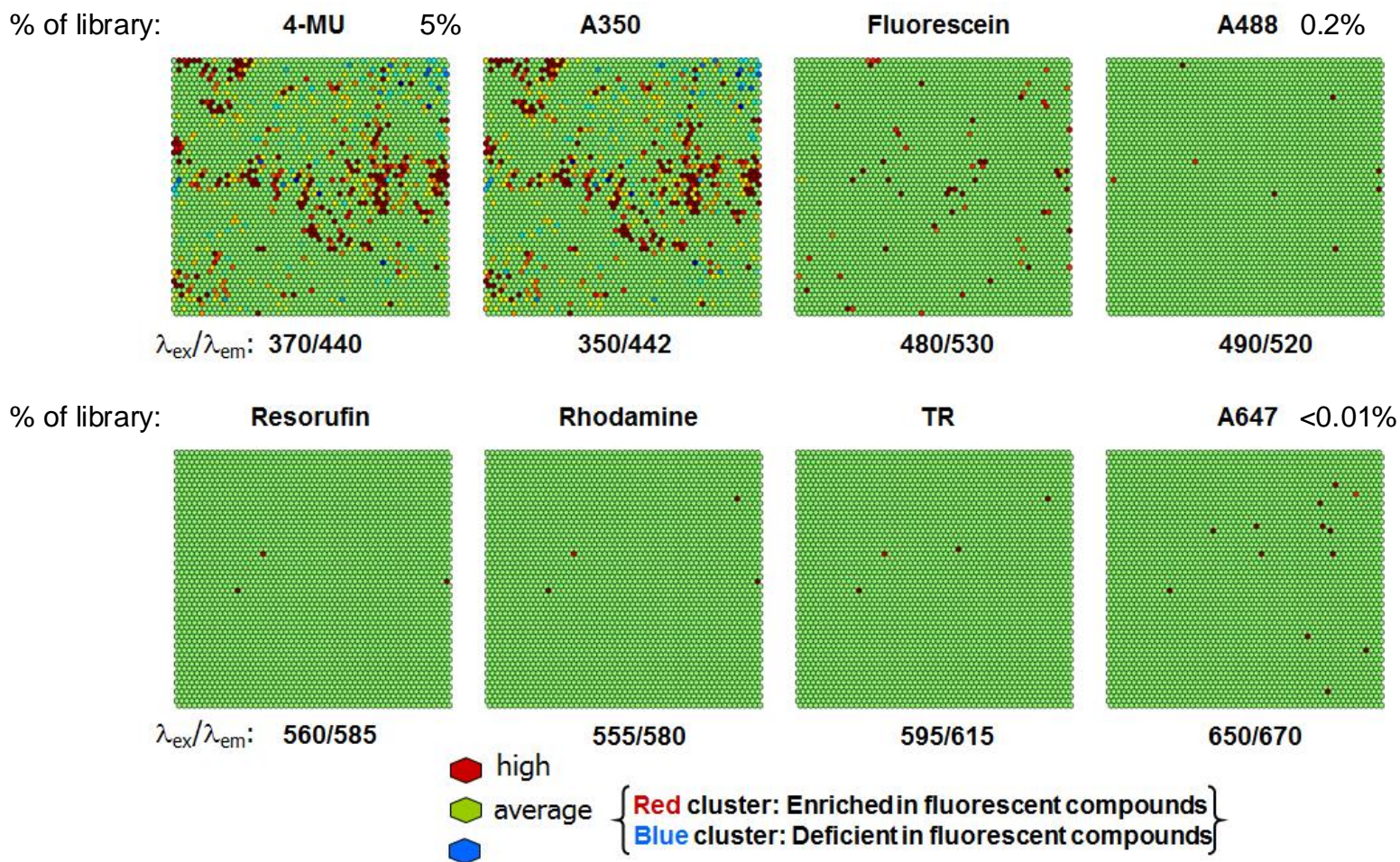
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- **Potential for assay interference**
  - Fluorescent compounds can interfere with fluorescent readouts
  - Colored compounds might interfere with luminescence



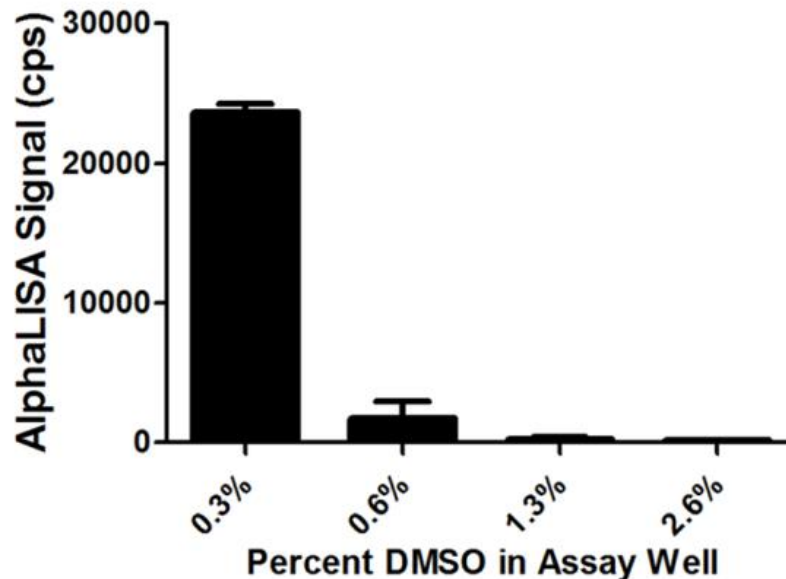
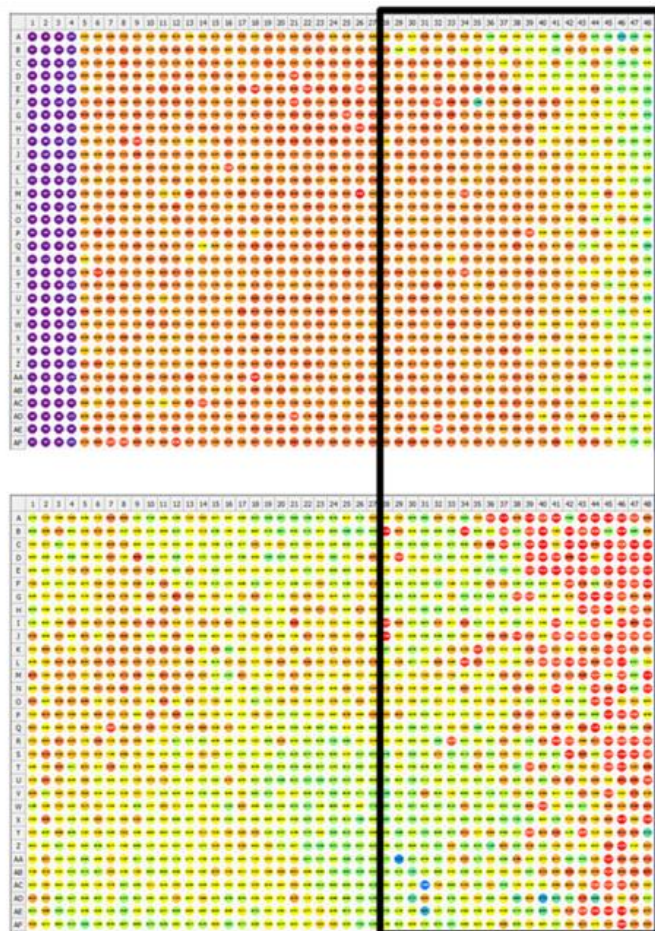
# Fluorescence Spectroscopic Profiling of Compound Libraries



Simeonov, A., Jadhav, A., Thomas, C.J., Wang, Y., Huang, R., Southall, N.T., Shinn, P., Smith, J., Austin, C.P., Auld, D.S. and Inglese, J., 2008. **Fluorescence spectroscopic profiling of compound libraries.** *Journal of Medicinal Chemistry*, 51(8), 2363-2371.



# Determination of Assay Tolerance to DMSO/Vehicle is Important



Yasgar A., Jadhav A., Simeonov A., Coussens N.P., **AlphaScreen-Based Assays: Ultra-High-Throughput Screening for Small-Molecule Inhibitors of Challenging Enzymes and Protein-Protein Interactions.** *Methods Mol Biol.* 2016;1439:77-98.

# Important Considerations for Choosing an Assay

---

- **Homogenous assay format is preferred for screening**
  - Add reagents, mix and measure (no solution removal or wash steps)
  - Automation friendly
  - Reduces variability
  - Decreases hands-on time
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- **Time required for assay**
  - Off-line reagent preparation
  - Is temperature equilibration required
  - Actual assay time
  - Kinetic versus end point read
  - Time required for data analysis and record keeping





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  - Does the response occur rapidly or within a few minutes or hours?
  - Longer signal stability allows for flexibility in automated systems
  - Longer signal stability minimizes differences among plates within a stack



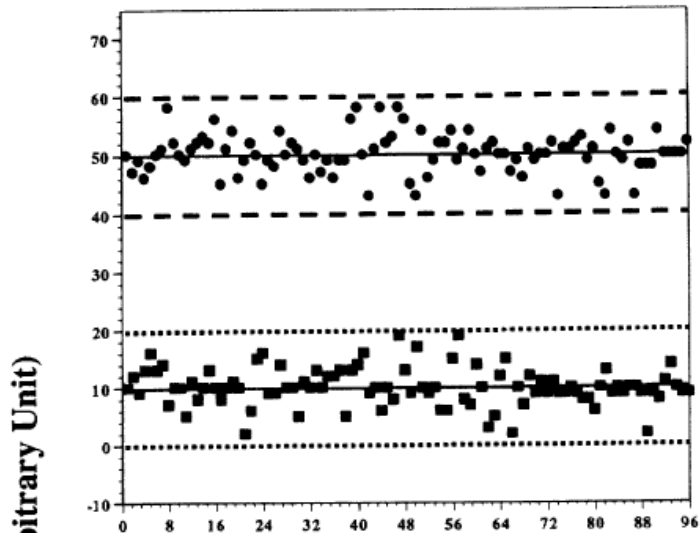
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- **Assay Sensitivity**
  - Choice of readouts is important
    - Colorimetric<fluorescent<luminescent



# Evaluating Assay Suitability for Screening



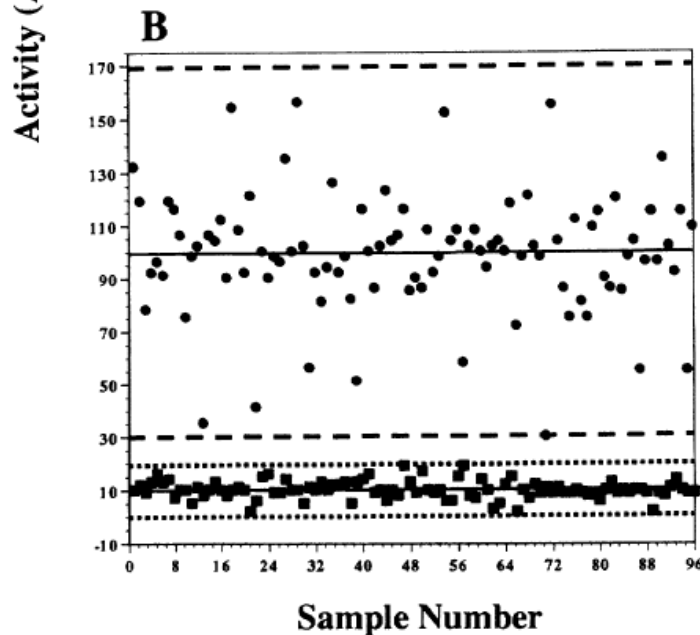
$S/N = 12$

$S/B = 5$

$z = 0.5$

$$S/N = \frac{\text{mean signal} - \text{mean background}}{\text{standard deviation of background}}$$

$$S/B = \frac{\text{mean signal}}{\text{mean background}}$$



$S/N = 27$

$S/B = 10$

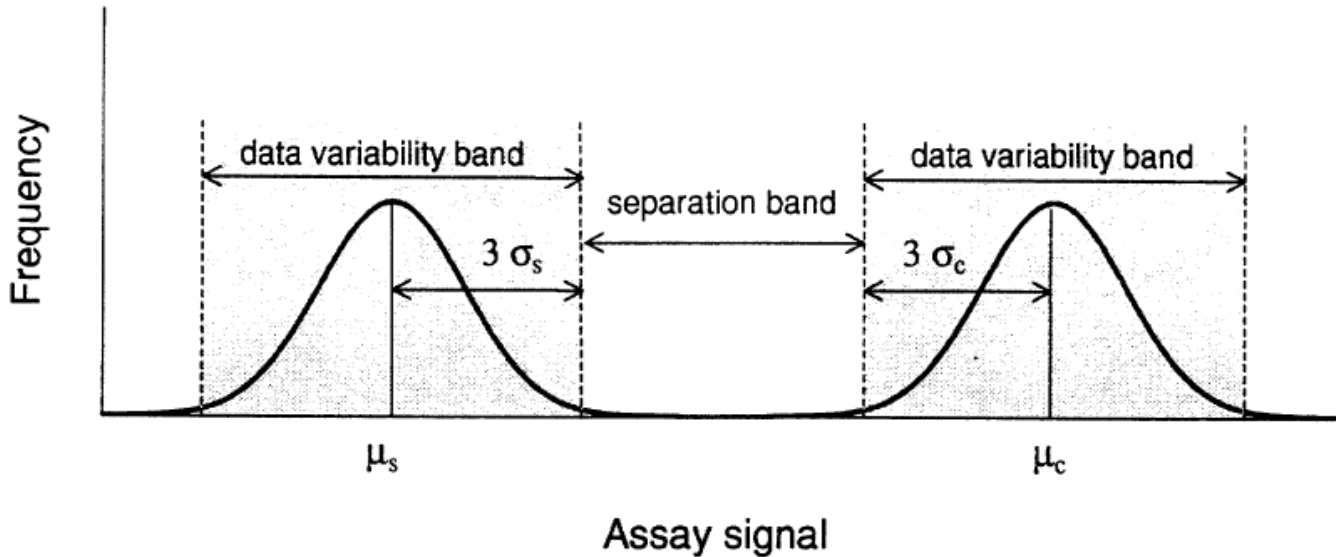
$z = 0.1$

$$Z = 1 - \frac{3SD \text{ of sample} + 3SD \text{ of control}}{|\text{mean of sample} - \text{mean of control}|}$$

A Simple Statistical Parameter for Use in Evaluation and Validation of High Throughput Screening Assays. Zhang JH, Chung TD, Oldenburg KR. J Biomol Screen. 1999;4(2):67-73.



# Evaluating Assay Suitability for Screening



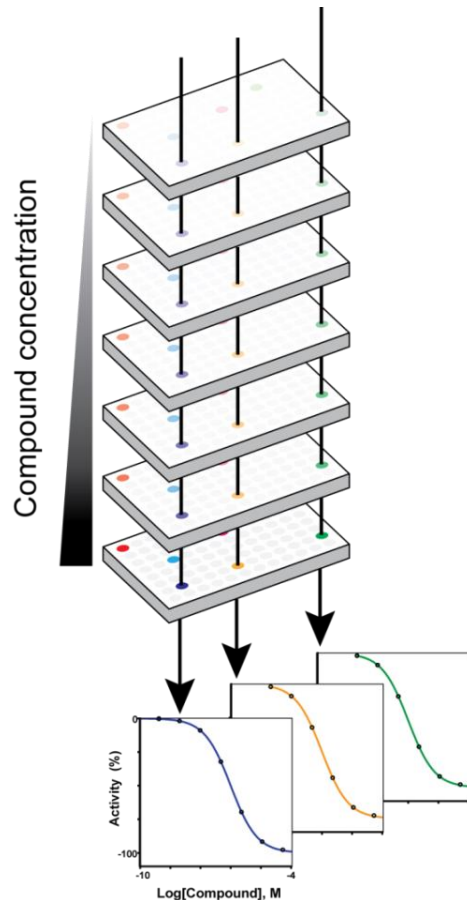
$$Z = 1 - \frac{3SD \text{ of sample} + 3SD \text{ of control}}{|\text{mean of sample} - \text{mean of control}|} *$$

<i>Z-factor value</i>	<i>Structure of assay</i>	<i>Related to screening</i>
1	SD = 0 (no variation), or the dynamic range $\rightarrow \infty$	An ideal assay
$1 > Z \geq 0.5$	Separation band is large	An excellent assay
$0.5 > Z > 0$	Separation band is small	A double assay
0	No separation band, the sample signal variation and control signal variation bands touch	A "yes/no" type assay
<0	No separation band, the sample signal variation and control signal variation bands overlap	Screening essentially impossible

A Simple Statistical Parameter for Use in Evaluation and Validation of High Throughput Screening Assays. Zhang JH, Chung TD, Oldenburg KR. J Biomol Screen. 1999;4(2):67-73.



# Improving the Process of Early Discovery: Quantitative High-Throughput Screening (qHTS)



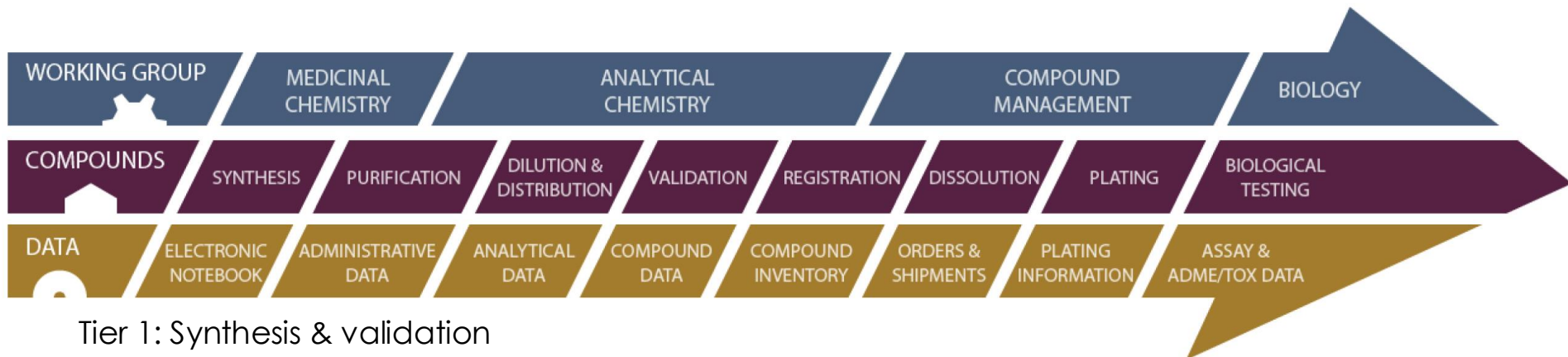
- Conventional screening done at one concentration
  - Not appropriate for potency testing – “dose makes the poison”
- qHTS tests compounds assayed at **multiple** concentrations (range: 4 logs)
- Enabled by miniaturized assay volumes (2-8  $\mu\text{L}$  per test) and informatics pipeline
- Generates *pharmacological actives* instead of statistical “hits”
  - Dramatically increases reliability
  - Dramatically reduces false positives and false negatives
- *To date, several hundred million datapoints from several hundred screens have been generated and deposited in the public domain.*

PNAS 103:11473



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Translational Sciences

# Medicinal Chemistry, an Integrated Process



## Tier 1: Synthesis & validation

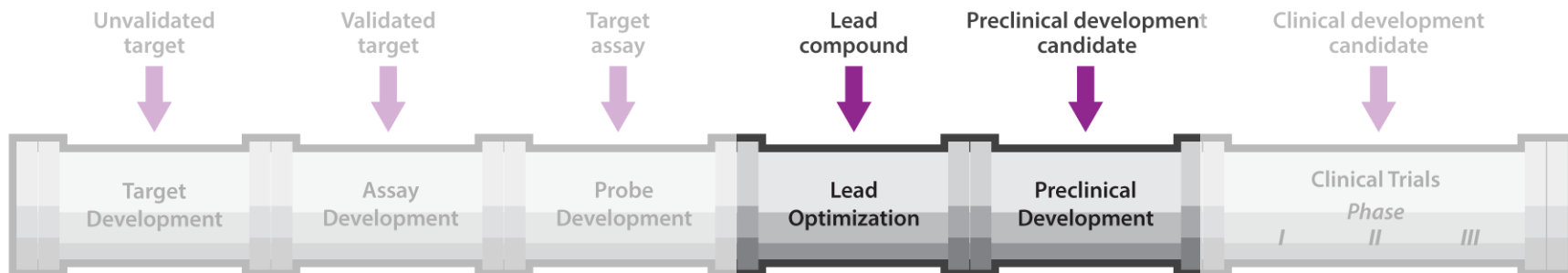
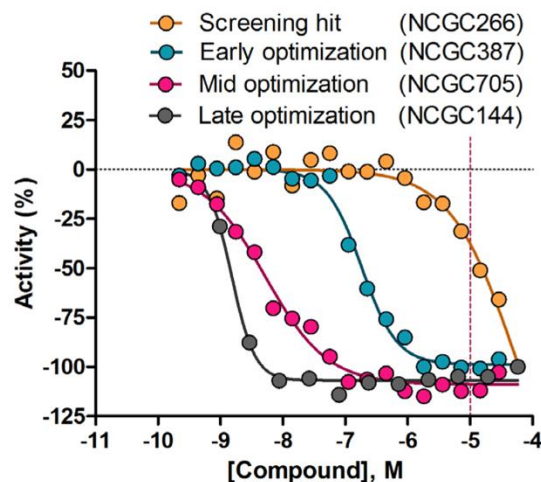
- Medicinal chemistry
- Purification
- In vitro ADME

## Tier 2: Compound profile expansion

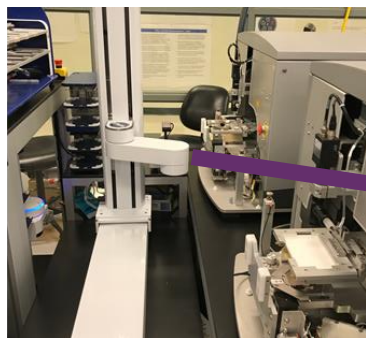
- Met ID/CYP studies
- In vitro toxicology

## Tier 3: Advanced Preclinical studies

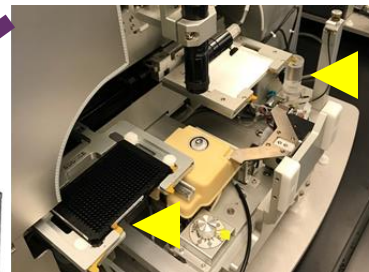
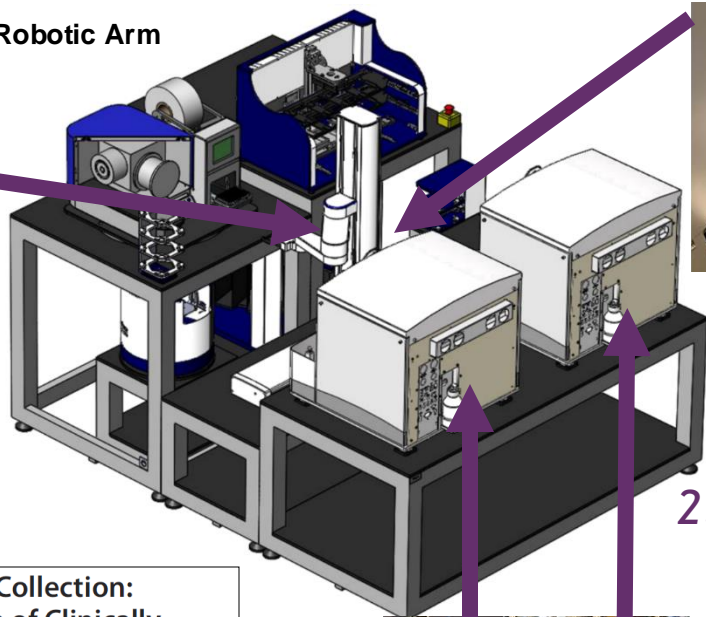
- Formulation
- Scale-up
- In vivo PK/PD
- Preclinical toxicology



# Translation Challenge: Rapid Discovery of Drug Combinations

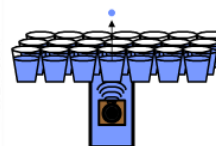


Robotic Arm



Destination Plate  
(1536, screening)

Source Plate  
(384, compounds)



1. Appropriate libraries

2. Automation/  
screening technologies

3. Informatics platform

The NCGC Pharmaceutical Collection:  
A Comprehensive Resource of Clinically  
Approved Drugs Enabling Repurposing and  
Chemical Genomics

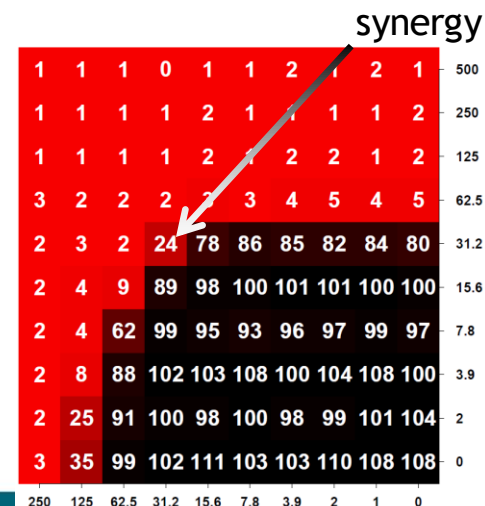
Ruilin Huang,\* Noel Southall,\* Yuhong Wang, Adam Yasgar, Paul Shinn,  
Ajit Jadhav, Dac-Trung Nguyen, Christopher P. Austin\*

Small-molecule compounds approved for use as drugs may be "repurposed" for new indications and studied to determine the mechanisms of their beneficial and adverse effects. A comprehensive collection of all small-molecule drugs approved for human use would be invaluable for systematic repurposing across human diseases, particularly for rare and neglected diseases, for which the cost and time required for development of a new chemical entity are often prohibitive. Previous efforts to build such a comprehensive collection have been limited by the complexities, redundancies, and semantic inconsistencies of drug naming within and among regulatory agencies worldwide; a lack of clear conceptualization of what constitutes a drug; and a lack of access to physical samples. We report here the creation of a definitive, complete, and nonredundant list of all approved molecular entities as a freely available electronic resource and a physical collection of small molecules amenable to high-throughput screening.

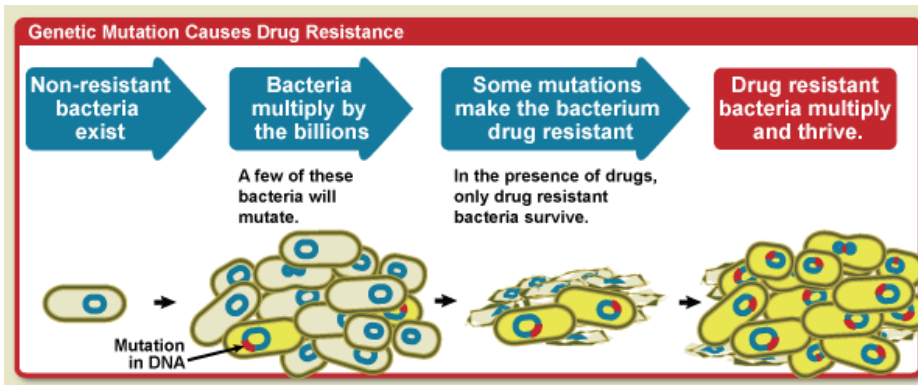
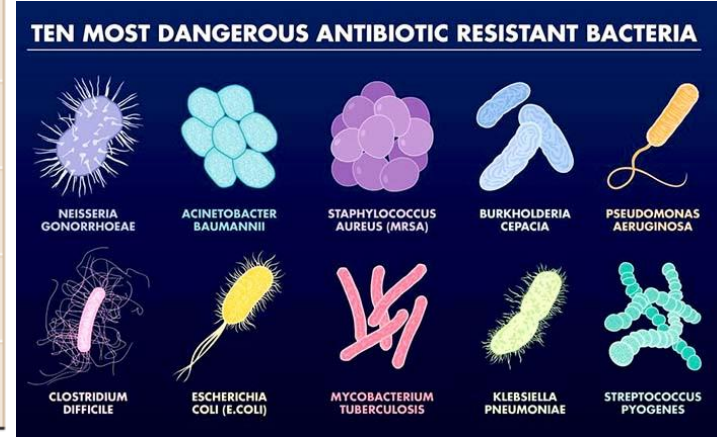
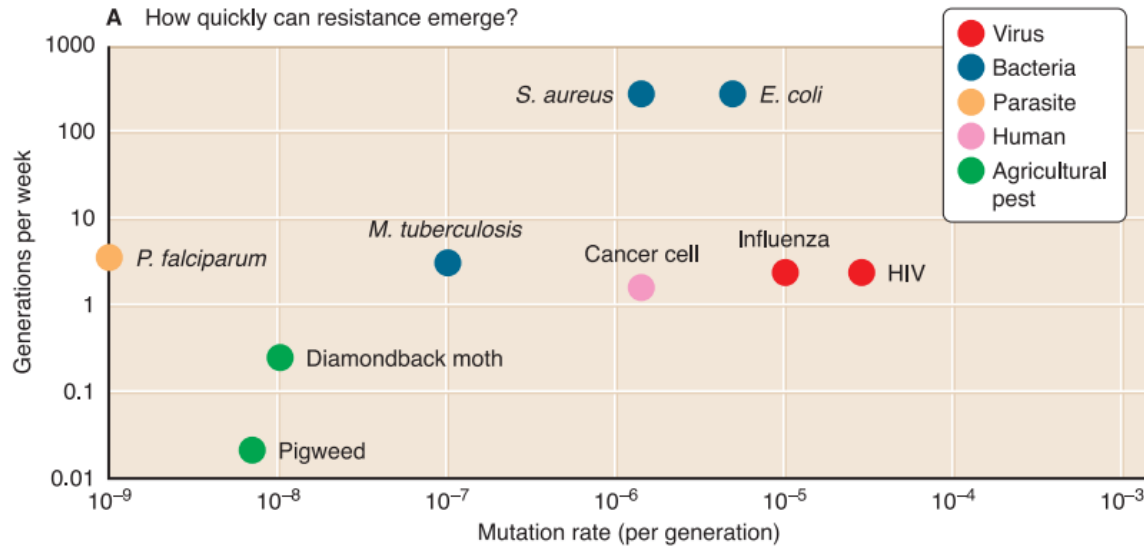
[www.ScienceTranslationalMedicine.org](http://www.ScienceTranslationalMedicine.org) 27 April 2011 Vol 3 Issue 80 80ps16



Acoustic  
Dispensers



# Application of Drug Combinations to Address Resistance





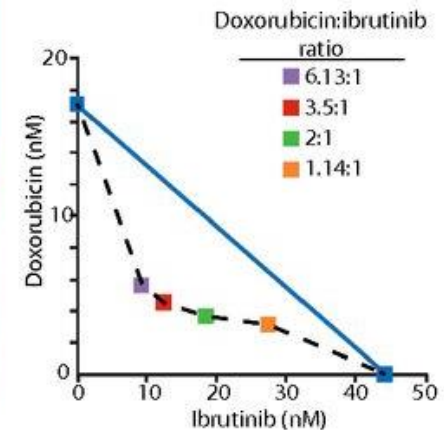
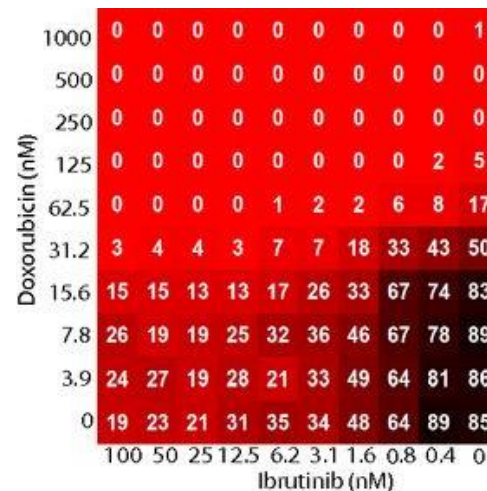
# Dissemination of technology: combination screening to overcome drug resistance in cancer cells

- Applied to ABC subtype of Diffuse Large B-Cell Lymphoma (ABC-DLBCL)
- Ibrutinib is a BTK inhibitor that has activity against ABC DLBCL
- Lead investigators: Craig Thomas (NCATS) and Louis Staudt (NCI)
- Study evaluated 459 drugs *in combination* with Ibrutinib
  - » 6 x 6 concentration-response “matrix blocks”, validation in 10 x 10 concentration-response matrix blocks
- DNA-damaging agents identified as synergizing with Ibrutinib in killing ABC DLBCL cell lines
- **Dissemination:**
  - » Protocols
  - » Source code for dispense

High-throughput combinatorial screening identifies drugs that cooperate with ibrutinib to kill activated B-cell-like diffuse large B-cell lymphoma cells

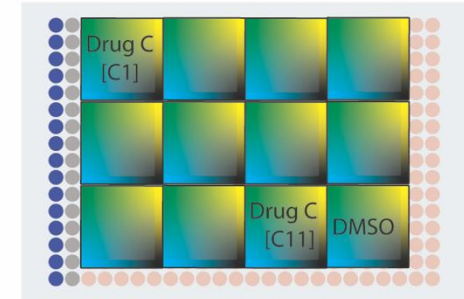
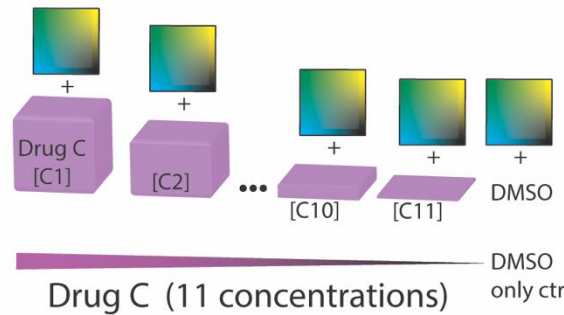
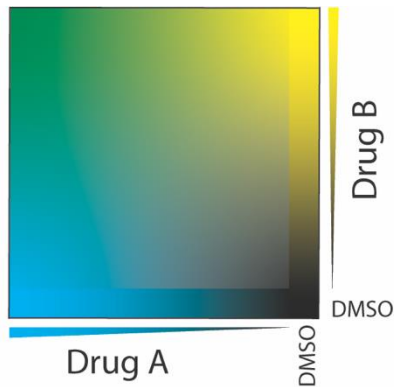
Lesley A. Mathews Griner<sup>a,1</sup>, Rajarshi Guha<sup>a,1</sup>, Paul Shinn<sup>a,1</sup>, Ryan M. Young<sup>b,1</sup>, Jonathan M. Keller<sup>a</sup>, Dongbo Liu<sup>a</sup>, Ian S. Goldlust<sup>a</sup>, Adam Yasgar<sup>a</sup>, Crystal McKnight<sup>a</sup>, Matthew B. Boxer<sup>a</sup>, Damien Y. Duveau<sup>a</sup>, Jian-Kang Jiang<sup>a</sup>, Sam Michael<sup>a</sup>, Tim Mierzwa<sup>a</sup>, Wenwei Huang<sup>a</sup>, Martin J. Walsh<sup>a</sup>, Bryan T. Mott<sup>a</sup>, Paresma Patel<sup>a,c</sup>, William Leister<sup>a</sup>, David J. Maloney<sup>a</sup>, Christopher A. Leclair<sup>a</sup>, Ganesha Rai<sup>a</sup>, Ajit Jadhav<sup>a</sup>, Brian D. Peysers<sup>d</sup>, Christopher P. Austin<sup>a</sup>, Scott E. Martin<sup>a</sup>, Anton Simeonov<sup>a</sup>, Marc Ferrer<sup>a</sup>, Louis M. Staudt<sup>b,2</sup>, and Craig J. Thomas<sup>a,2</sup>

<sup>a</sup>Division of Preclinical Innovation, National Institutes of Health Chemical Genomics Center, National Center for Advancing Translational Sciences, <sup>b</sup>Metabolism Branch, Center for Cancer Research, and <sup>c</sup>Developmental Therapeutics Program, Division of Cancer Treatment and Diagnosis, National Cancer Institute, National Institutes of Health, Bethesda, MD 20892; and <sup>d</sup>Basic Science Program, SAIC-Frederick, Inc., Chemical Biology Laboratory, Frederick National Laboratory for Cancer Research, Frederick, MD 21702

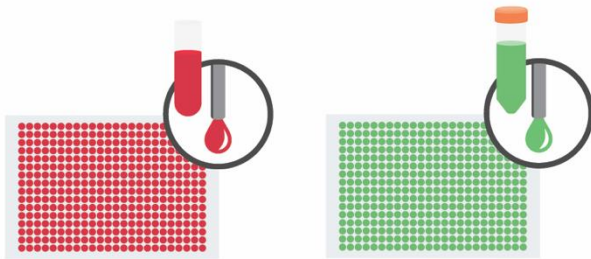


# Example: triple drug combination screening to tackle resistance against artemisinin-based combination therapies in malaria

ACS Pharmacol. Transl. Sci. 2020, <https://dx.doi.org/10.1021/acspsci.0c00110?ref=pdf>

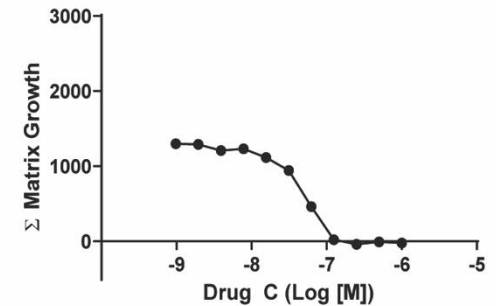
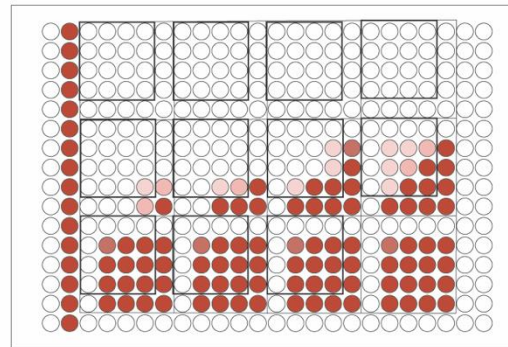


- ① Drugs A and B are acoustically dispensed in a 10x10-well matrix, 12 replicate blocks per plate. Single drug responses, bottom row (Drug A) and right column (Drug B).



- ② To each replicate block, serial dilutions of Drug C is acoustically dispensed, with the final block serving as a DMSO control

- ③ Plate view of triple combination screening plate with positive control (artesunate, blue) and neutral controls (DMSO, grey) also shown.



- ④ Dispense *P. falciparum* and erythrocytes, incubate 72 hr

- ⑤ Dispense 2  $\mu\text{L}$  of SYBRGreen1 and lysis solution, incubate overnight. Fluorescence quantified

- ⑥ Parasite proliferation response is normalized to artesunate and DMSO controls. For each concentration Drug C block, response of Drug A + Drug B wells is summed.

- ⑦ Triple drug response is analyzed as a function of Drug C concentration.

**Venetoclax, Ibrutinib, Prednisone, Obinutuzumab, and Revlimid (ViPOR) in Relapsed/Refractory B-cell Lymphoma**

ClinicalTrials.gov Identifier: NCT03223610

Recruitment Status ⓘ : Recruiting  
 First Posted ⓘ : July 21, 2017  
 Last Update Posted ⓘ : July 7, 2022  
 See [Contacts and Locations](#)



The safety and scientific validity of this study is the responsibility of the study sponsor and investigators. Listing a study does not mean it has been evaluated by the U.S. Federal Government. [Know the risks and potential benefits](#) of clinical studies and talk to your health care provider before participating. Read our [disclaimer](#) for details.

**Sponsor:**  
 National Cancer Institute (NCI)

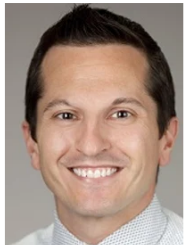
**Information provided by (Responsible Party):**  
 National Institutes of Health Clinical Center (CC) ( National Cancer Institute (NCI) )

**ViPOR Regimen Is Safe, Shows Impressive Activity in Relapsed/Refractory DLBCL**

December 21, 2020  
 Gina Mauro



December 21, 2020 — The 5-drug regimen of venetoclax, ibrutinib, prednisone, obinutuzumab, and lenalidomide showed a tolerable safety profile and encouraging antitumor activity with complete responses in patients with relapsed/refractory diffuse large B-cell lymphoma.



Christopher J. Melani, MD

The 5-drug regimen of venetoclax (Venclexta), ibrutinib (Imbruvica), prednisone, obinutuzumab (Gazyva), and lenalidomide (Revlimid; ViPOR) showed a tolerable safety profile and encouraging antitumor activity with complete responses (CRs) in patients with relapsed/refractory diffuse large B-cell lymphoma (DLBCL), according to phase 1b/2 findings that were presented during the 2020 ASH Annual Meeting and Exposition.

Results showed that of 52 evaluable patients, the best overall response was 71%, with a complete response (CR) rate of 52% and a 19% partial response (PR) rate. Specifically, in relapsed patients (n = 30), the objective response rate (ORR) was 83% with a 70% and 13% CR and PR rate, respectively. The ORR was 55% in refractory patients (n = 22), with a 27% CR rate and a 27% PR rate.

<https://www.onclive.com/view/vipor-regimen-is-safe-shows-impressive-activity-in-relapsed-refractory-dlbcl>

**ViPOR Regimen Signals Benefit in Patients With Mantle Cell Lymphoma**

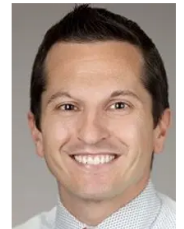
February 7, 2022  
 Nichole Tucker



In an interview with Targeted Oncology, Christopher Melani, MD, discussed the ongoing ViPOR study exploring a Bruton's tyrosine kinase inhibitor and BCL2 inhibitor, and NF-κB survival pathway activating combination.

Treatment with the ViPOR regimen consisting of venetoclax (Venclexta), ibrutinib (Imbruvica), prednisone, and lenalidomide (Revlimid), has thus far appeared safe for use in patients with mantle cell lymphoma (MCL) and has demonstrated preliminary activity.

Results from the phase 1 portion of the ViPOR study (NCT03223610) were presented during the 63rd American Society of Hematology (ASH) Annual Meeting & Exposition. Of the 11 patients who were treated, the ORR was 100% and the complete remission (CR) rate was 80%. Nine patients were evaluable for safety and no dose-limiting toxicities were observed. There were few grade 3 and 4 adverse events (AEs), but the hematologic grade 3/4 AEs included neutropenia (13%), anemia (11%), and thrombocytopenia (9%). The non-hematologic grade 3/4 AEs included hypokalemia (33%) along with fatigue, hypomagnesemia, elevated bilirubin, atrial fibrillation, lung infection, and syncope occurring in 11% of patients each.



Christopher Melani, MD

<https://www.targetedonc.com/view/vipor-regimen-signals-benefit-in-patients-with-mantle-cell-lymphoma>



# First matrix screen enables a translation journey in lymphoma

1st Matrix Paper  
Feb, 2014

**High-throughput combinatorial screening identifies drugs that cooperate with ibrutinib to kill activated B-cell-like diffuse large B-cell lymphoma cells**

Lesley A. Mathews Grimes<sup>1\*</sup>, Rajarshi Guha<sup>1\*</sup>, Paul Shih<sup>1\*</sup>, Ryan M. Young<sup>1</sup>, Jonathan M. Kelle<sup>1</sup>, George Liu<sup>1</sup>, Ian S. Goldfarb<sup>1</sup>, Adam Yagoe<sup>1</sup>, Crystal McKeighan<sup>1</sup>, Matthew B. Roemer<sup>1</sup>, Damien V. Doreau<sup>1</sup>, Jan-Kang Bang<sup>1</sup>, Sam Michael<sup>1</sup>, Tom Mirzavand<sup>1</sup>, Myoung Hoang<sup>1</sup>, Marissa J. Walsh<sup>1</sup>, Bryan T. Meier<sup>1</sup>, Poojara Kashi<sup>1</sup>, William Lester<sup>1</sup>, David J. Maloney<sup>1</sup>, Christopher A. Leika<sup>1</sup>, Suresha Reddy<sup>1</sup>, Raj Jadhav<sup>1</sup>, Brian D. Pappas<sup>1</sup>, Christopher F. Austin<sup>1</sup>, Scott E. Marder<sup>1</sup>, Andrew Simonovich<sup>1</sup>, Isaac Fierer<sup>1</sup>, Linah M. Sica<sup>1</sup>, and George C. Thomas<sup>1</sup>

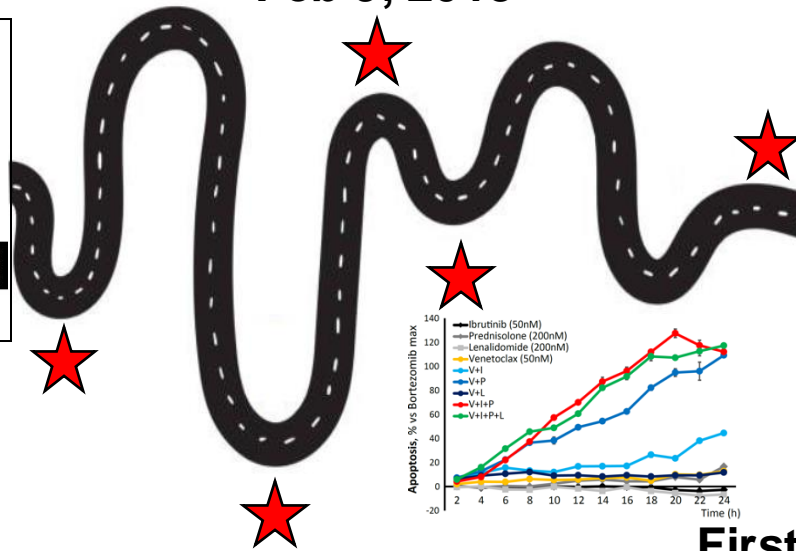
\*Division of Preclinical Innovation, National Institutes of Health Chemical Genomics Center, National Center for Advancing Translational Sciences, National Institutes of Health, Bethesda, MD 20892; <sup>1</sup>Translational Research Program, Division of Cancer Treatment and Diagnosis, National Cancer Institute, National Institutes of Health, Bethesda, MD 20892; and <sup>2</sup>Cell Science Program, NCI Frederick, 94, Chemical Biology Laboratory, National Cancer Institute for Cancer Research, Bethesda, MD 20892

Editor: Jay Parker, National Cancer Institute, National Institutes of Health, Bethesda, MD, and approved December 27, 2013 (received for review June 25, 2013)

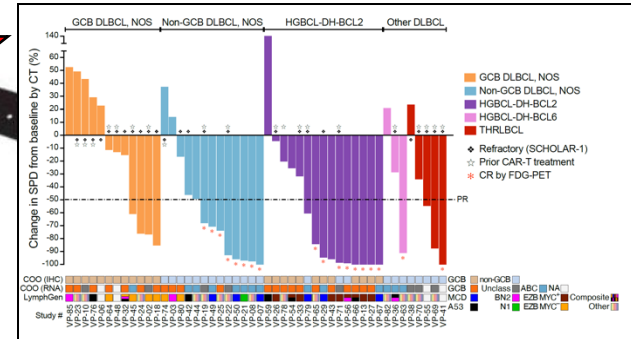
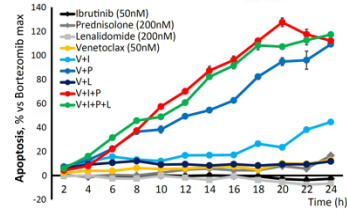
The clinical development of drug combinations is typically achieved through trial-and-error or via insight gained through a limited molecular understanding of dysregulated signaling pathways in a specific cancer type. Unbiased small-molecule combination function screening represents a high-throughput means to explore hundreds and even thousands of drug-drug pairs for potential investigation and translation. Here, we describe a high-throughput screening platform capable of testing compounds in complex systems under the influence of a specific inhibitor. We use this platform to define potential drug combinations that cooperate with the BTK inhibitor ibrutinib to kill large B-cell lymphoma (DLBCL). We identify drugs with synergistic activity, and explore with the BTK-specific kinase inhibitor venetoclax, which targets the chronic active B-cell receptor signaling that characterizes ABC-DLBCL. Venetoclax cooperates synergistically with a wide range of compounds, including inhibition of the PI3K-Akt-mTOR pathway target of mTOR signaling, other BCR receptor pathway inhibitors, BCL2 family inhibitors, and several components of transcription that is the standard of care for DLBCL.

**Significance**  
The treatment of cancer is highly reliant on drug combinations. Non-genotoxic, targeted therapies are demonstrating to

VIPOR Trial  
Initiated  
Feb 9, 2018



VIPOR Clinical Study  
Accepted for Publication  
April 8, 2024



0	0	1	1	1	2	2	3	3	3
0	0	0	1	1	1	2	2	3	4
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0	0	0	1	3	8	21	32	38	45
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1	1	6	12	26	37	52	61	67	77
2	9	16	27	40	49	61	71	72	81
9	19	31	39	49	66	65	78	82	91
16	24	35	45	53	59	69	79	91	93
24	34	42	50	59	71	84	95	100	100

Ibrutinib  
BTK  
Inhibitor

DA-TEDDI-R  
Clinical Report  
Published  
June 2017

First MCTC  
Experiment  
March, 2020

VIPOR: Venetoclax, Ibrutinib, Prednisone, Obinutuzumab, Revlimid

# The NEW ENGLAND JOURNAL of MEDICINE

ESTABLISHED IN 1812

JUNE 20, 2024

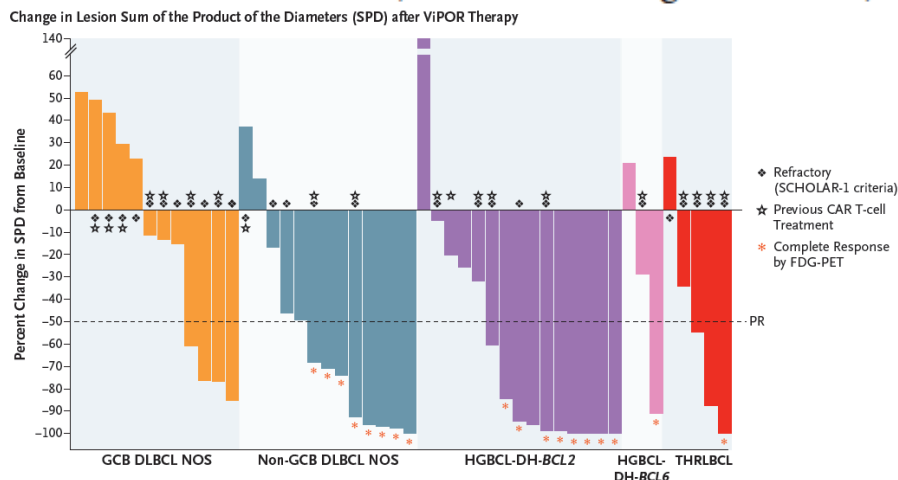
VOL. 390 NO. 23

## Combination Targeted Therapy in Relapsed Diffuse Large B-Cell Lymphoma

C. Melani, R. Lakhota, S. Pittaluga, J.D. Phelan, D.W. Huang, G. Wright, J. Simard, J. Muppidi, C.J. Thomas, M. Ceribelli, F.A. Tosto, Y. Yang, W. Xu, T. Davies-Hill, S.D. Pack, C.J. Peer, O. Arisa, E. Mena, L. Lindenberg, E. Bergvall, C.A. Portell, R.J. Farah, S.T. Lee, A. Pradhan, C. Morrison, A. Tadese, A.M. Juanitez, C. Lu, A. Jacob, H. Simmons, W.D. Figg, S.M. Steinberg, E.S. Jaffe, M. Roschewski, L.M. Staudt, and W.H. Wilson

### CONCLUSIONS

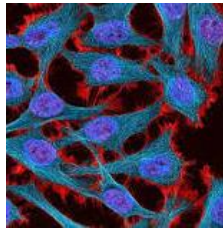
Treatment with ViPOR was associated with durable remissions in patients with specific molecular DLBCL subtypes and was associated with mainly reversible adverse events. (Funded by the Intramural Research Program of the National Cancer Institute and the National Center for Advancing Translational Sciences of the National Institutes of Health and others; ClinicalTrials.gov number, NCT03223610.)



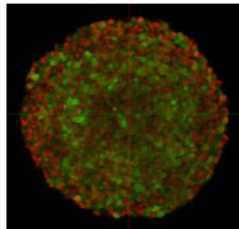
NIH National Center for Advancing Translational Sciences

# Increasing the predictivity of *in vitro* assays: a continuum of 3D models of healthy and diseased tissues

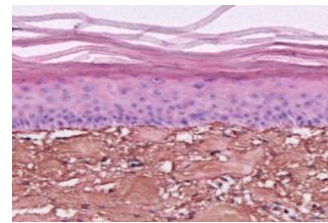
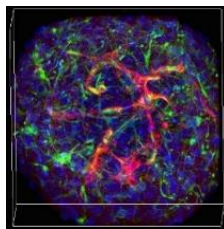
2D



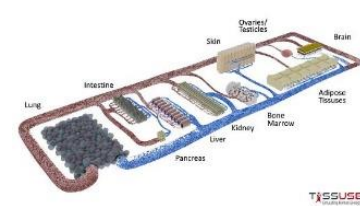
Spheroids



Organoids

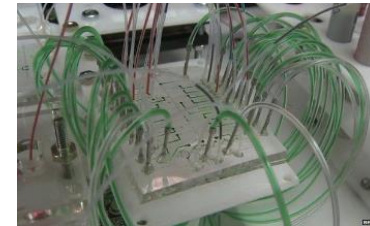
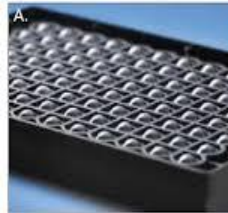
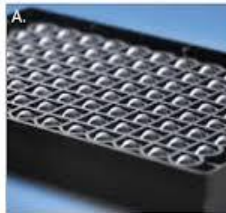


Organ-on-a-chip

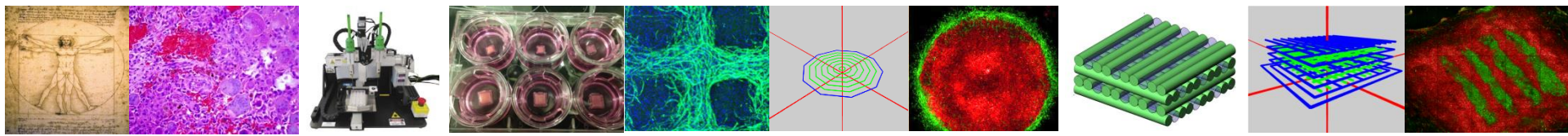


HTS compatibility

Physiological complexity

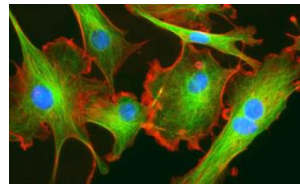


# 3D Tissue Bioprinting



Gel

+



Cells

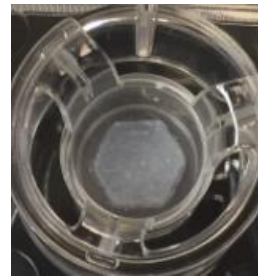


Syringe

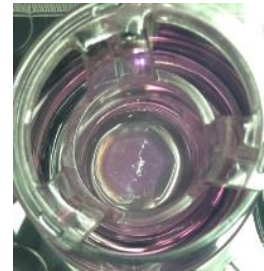


Printer

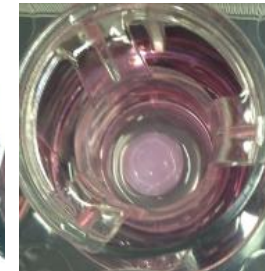
Hydrogel polymer is mixed with cells and loaded into syringe.



Printed construct



1 day



1 week



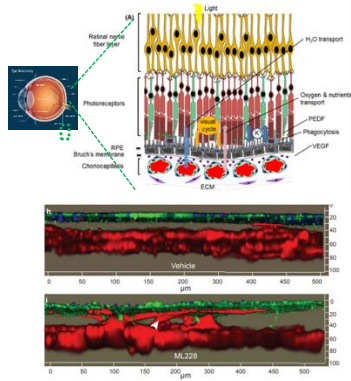
2 weeks

The printer “3D prints” the cell/gel mixture in a layer by layer approach.

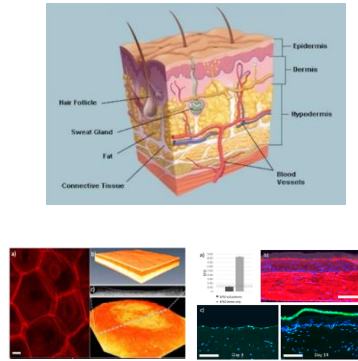
The printed construct is incubated to allow the cells to form a tissue, and to enable proper cell differentiation.

# Current portfolio of engineered 3D tissue models

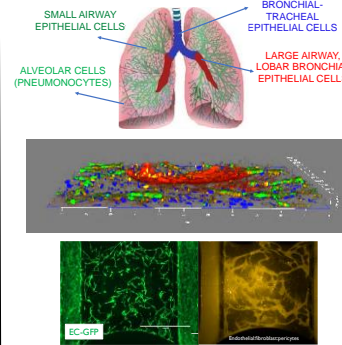
## Retina



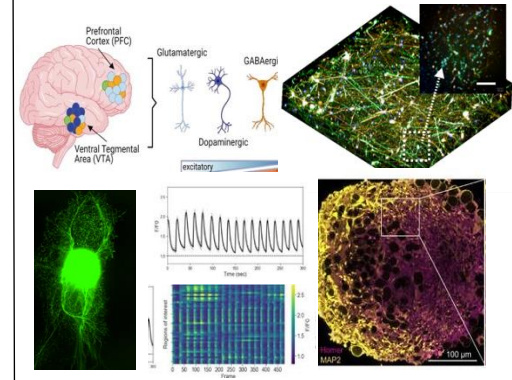
## Skin



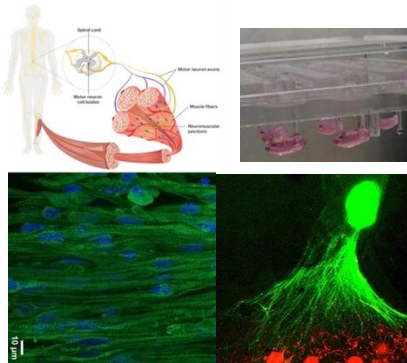
## Lung



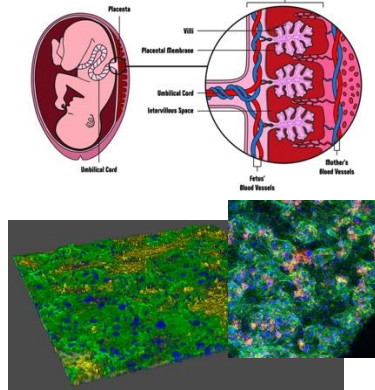
## CNS/PNS



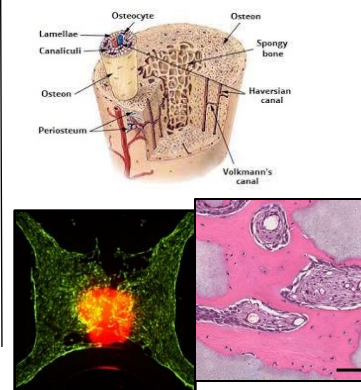
## Muscle/ Neuromuscular Junction



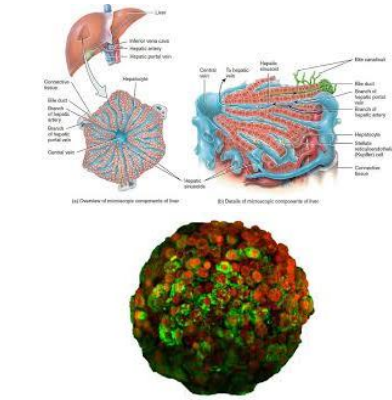
## Placenta barrier



## Bone



## Liver



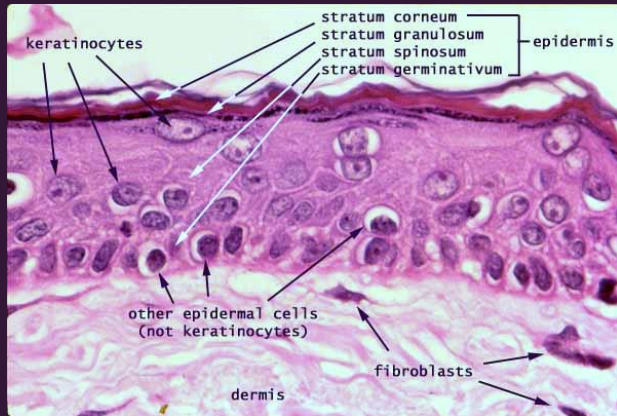
Program Director Marc Ferrer, Ph.D. <https://ncats.nih.gov/bioprinting>





# Skin biofabrication

## Native Skin



## 3D-Bioprinted Skin

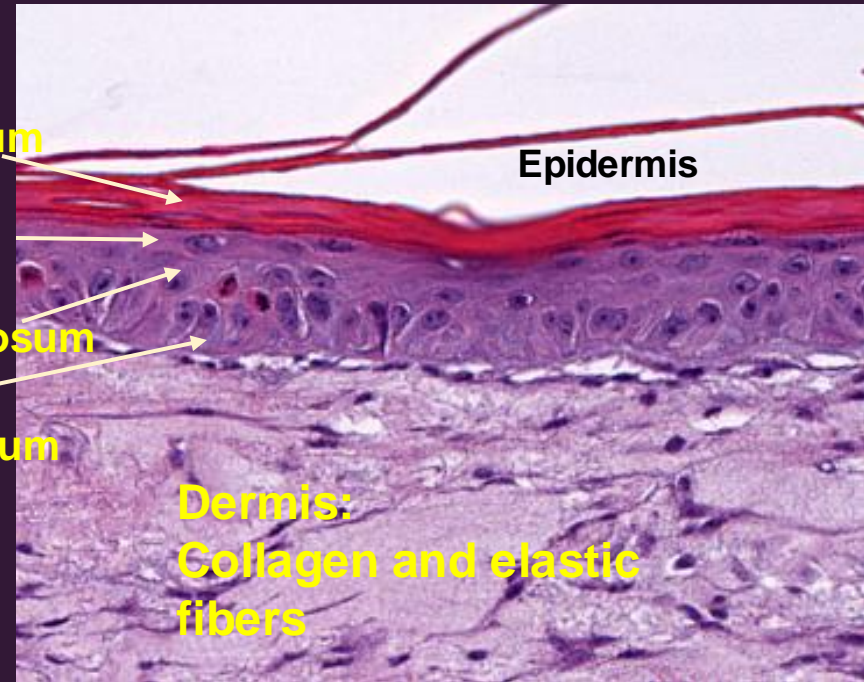
Stratum corneum

Stratum granulosum

Stratum spinosum

Stratum germinativum

Dermis:  
Collagen and elastic  
fibers



<http://www.siumed.edu/~dking2/intro/IN005b.htm>



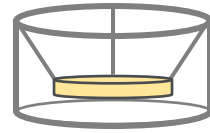
NIH National Center  
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Translational Sciences

# Generation of bioprinted skin tissues

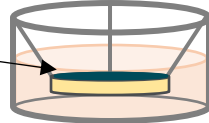
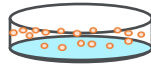
## Full thickness skin tissue (FTS)

### Reconstructed human epidermis (RhE)

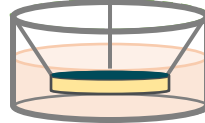
1. Coat the 96-well transwell insert membrane with collagen



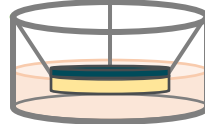
2. Add keratinocytes



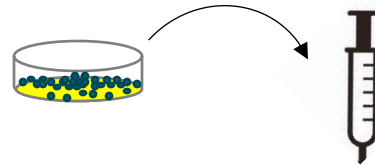
3. Submerge culture for 3 days



4. Air-liquid interface culture for 8 days

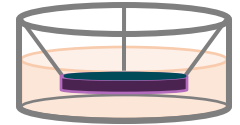
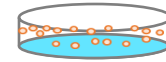


2. Bioprint fibroblast bioink to a 3-layer U shape on bottom side of 96-well transwell insert membrane

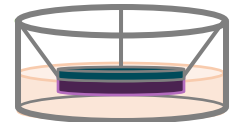


4. Submerge bioprinted tissue in medium for 7 days

5. Add keratinocytes and submerge culture for 3 days



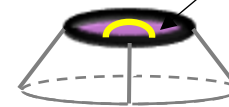
6. Air-liquid interface culture for 8 days



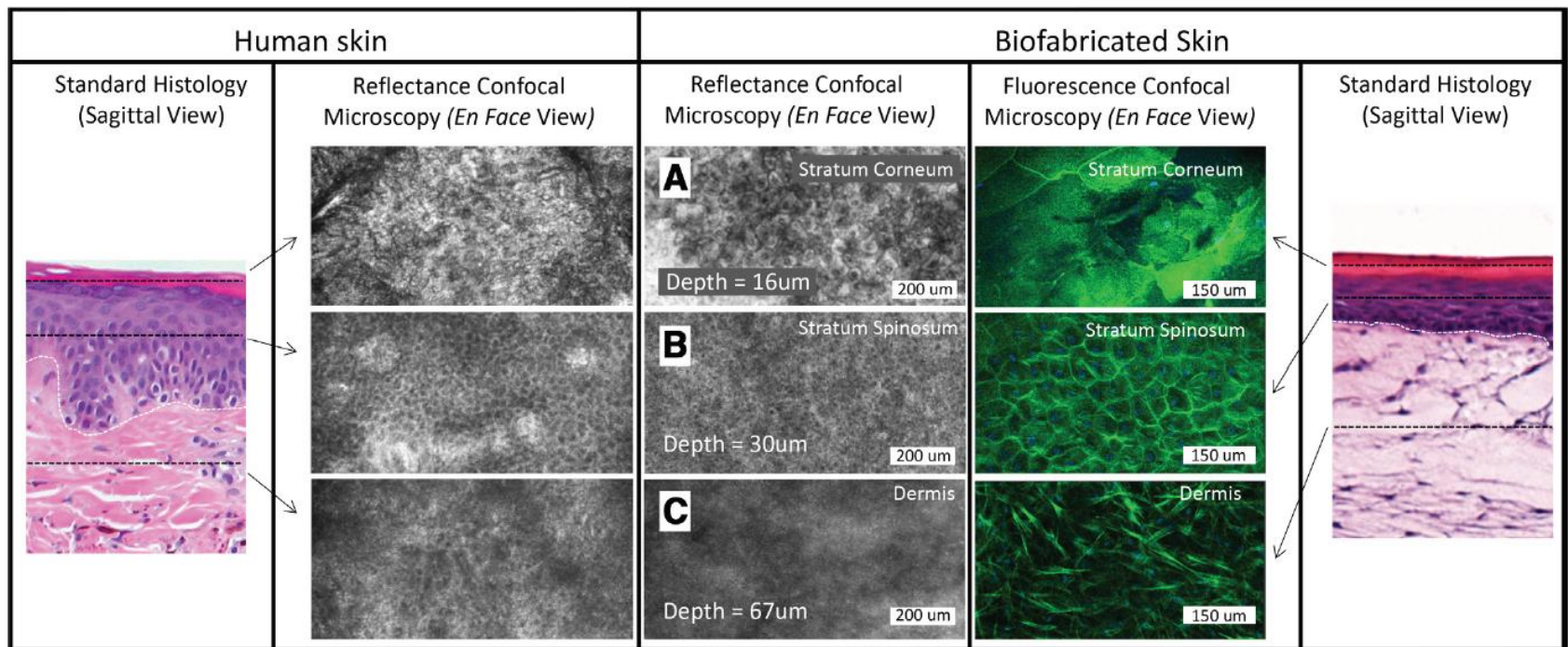
1. Suspend fibroblasts in bioprinting gel



3. Add bioprinting gel to cover the U shape



# A 3D biofabricated cutaneous squamous cell carcinoma tissue model with multi-channel confocal microscopy imaging biomarkers to quantify antitumor effects of chemotherapeutics in tissue



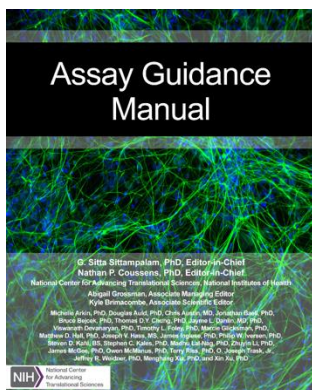
Collaboration between NCATS (Marc Ferrer) and Rockefeller University (Daniel Gareau)



**Where do I go for more  
information about assay  
development?**



# Sharing internal know-how: Assay Guidance Manual (47 chapters/ 1,338 printed pages)



<https://ncats.nih.gov/agm-video>

## August 7<sup>th</sup> Videos

1. Austin, CP: **Welcome to the Assay Guidance Manual (AGM) Workshop**
2. Coussens, NP: **Strategies for Assay Selection & Robust Biochemical Assays**
3. Riss, T: **Treating Cells as Reagents to Design Reproducible Screening Assays**
4. Trask, OJ: **Assay Development Considerations for High Content Imaging**
5. Auld, DS: **Studies in Mechanisms and Methods in Assay Interferences**
6. Dahlin, JL: **Assay Interference by Chemical Reactivity**
7. Chung, TDY: **Basic Assay Statistics, Data Analysis & Rules of Thumb**
8. Devanarayan, V: **Reproducibility & Differentiability of Potency Results**
9. Sittampalam, GS: **Avoiding Artifacts & Interferences in Assay Operations**

## March 26-27<sup>th</sup> Videos

1. Austin, CP: **Welcome to the Assay Guidance Manual (AGM) Workshop**
2. Coussens, NP: **Robust Assays Define Success in Preclinical Research**
3. Lal-Nag, M: **Target Identification & Validation in Translational Discovery**
4. Foley, TL: **Development & Validation of Cell-Based and Biochemical Assays**
5. Riss, T: **Treating Cells as Reagents to Design Reproducible Screening Assays**
6. Trask, OJ: **Assay Development for HCS & Best Practices for 3D HCS**
7. Roth, KD: **Mass Spectrometry for Drug Screening and Lead Optimization**
8. Dahlin, JL: **Bioassay Interference by Aggregation and Chemical Reactivity**
9. Patnaik, S: **Lead Selection and Optimization by Medicinal Chemistry**
10. Xia, M: ***In Vitro* Toxicological Testing Using a qHTS Platform**
11. Xu, X: ***In Vitro* Assessment of ADME Properties of Lead Compounds**
12. Kahl, SD: **Statistical Design of Experiments for Assay Development**
13. Guha, R: **Pharos Application to Target Evaluation and Drug Discovery**
14. Weidner, JR: **Assay Operations: Keeping Assays Robust and Reproducible**

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<b>Assay Artifacts and Interferences</b>	4 Chapters
<b>Assay Validation, Operations and Quality Control</b>	5 Chapters
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<b>Glossary of Quantitative Biology Terms</b>	1 Chapter

Website: <https://ncats.nih.gov/expertise/preclinical/agm>

Email us: [NCATS\\_AGM\\_Editors@mail.nih.gov](mailto:NCATS_AGM_Editors@mail.nih.gov)



Facebook: [www.facebook.com/assayguide](http://www.facebook.com/assayguide)



LinkedIn: [www.linkedin.com/groups/7437344](http://www.linkedin.com/groups/7437344)



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# Assay Guidance Manual Training Workshops

- [Online Training Modules](#)
- [Upcoming Workshops](#)
- [Past Workshops](#)

NCATS offers a variety of [Assay Guidance Manual \(AGM\)](#) training workshops throughout the year designed to share best practices and advice on robust assay design, development and implementation for researchers involved in the drug discovery process.



## Online Training Modules

NCATS offers an online AGM training workshop in addition to the in-person AGM workshops held throughout the year. The online training workshop also features experts sharing best practices and expert advice on assay design, development and implementation. [View the video modules.](#)



## Assay Guidance Workshop on 3D Tissue Models for Antiviral Drug Development

🕒 Tue Jun 7, 11:00 AM - Wed Jun 8, 5:15 PM (EDT)

📍 Zoom

📅 Add to calendar

🔗 Share

### THIS EVENT HAS ENDED

Video of the workshop is available at the links below:

[Assay Guidance Workshop on 3D Tissue Models for Antiviral Drug Development \(Day 1\)](#) (June 7, 2022)

[Assay Guidance Workshop on 3D Tissue Models for Antiviral Drug Development \(Day 2\)](#) (June 8, 2022)

### About the Workshop

The [National Center for Advancing Translational Sciences \(NCATS\)](#) [Assay Guidance Manual \(AGM\)](#) program is hosting a two-day workshop that will cover a broad range of critical concepts, including practical approaches and best practices, for developing standardized 3D cellular assays with the hope of helping the community to successfully develop therapeutics for future pandemic threats. This workshop is jointly organized by NCATS, the [National Institute of Allergy and Infectious Diseases \(NIAID\)](#) and the [Bill & Melinda Gates Foundation](#). The overall goal of this workshop is to help scientists establish robust, reproducible, scalable, consistent, advanced 3D tissue models to study pandemic threat viruses.

<https://ncats.nih.gov/expertise/preclinical/agm/training>



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Translational Sciences