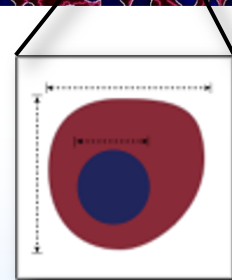


Advancing drug discovery and diagnostics through image-based AI

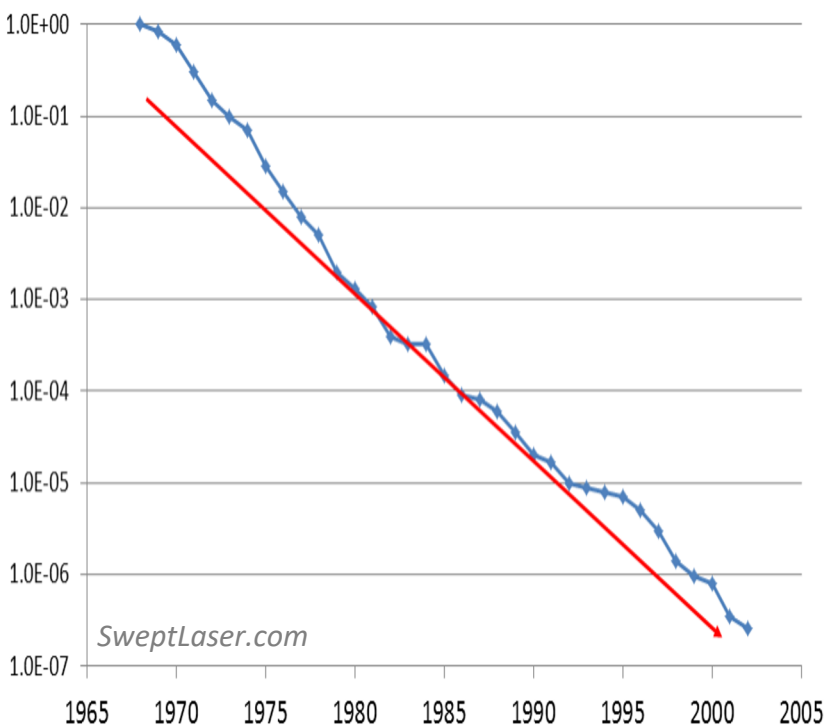
Anne E. Carpenter, PhD



Tale of two industries

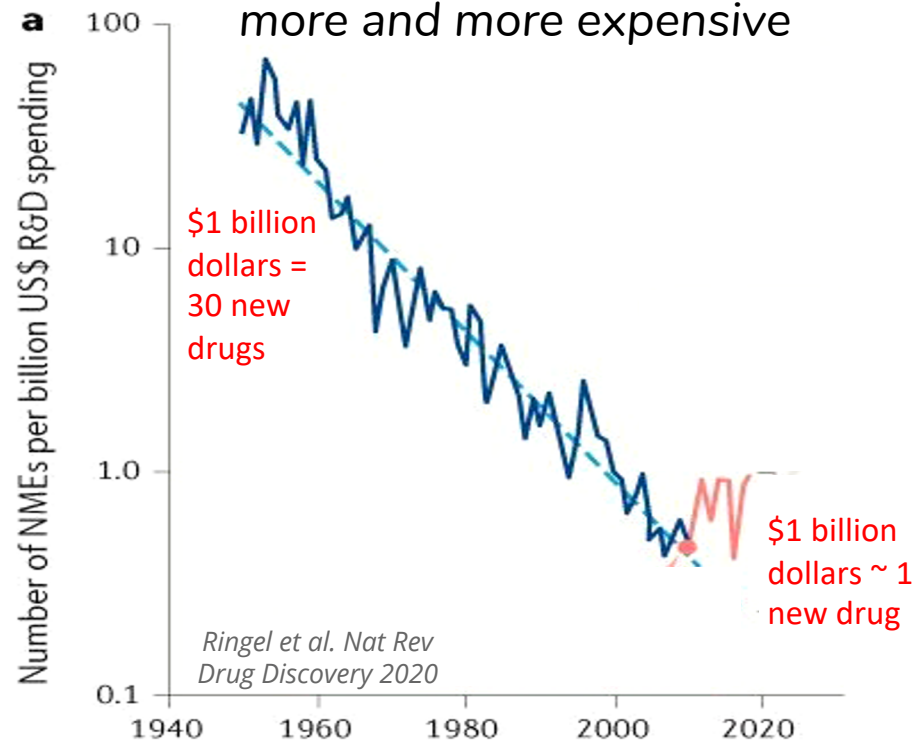
Moore's Law

Compute gets cheaper and cheaper

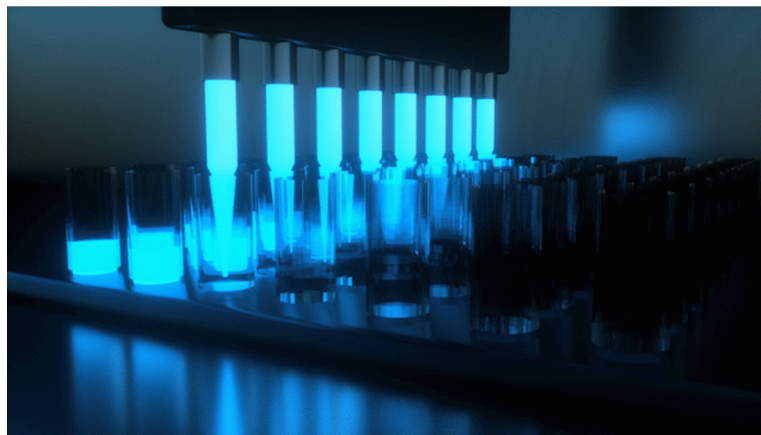
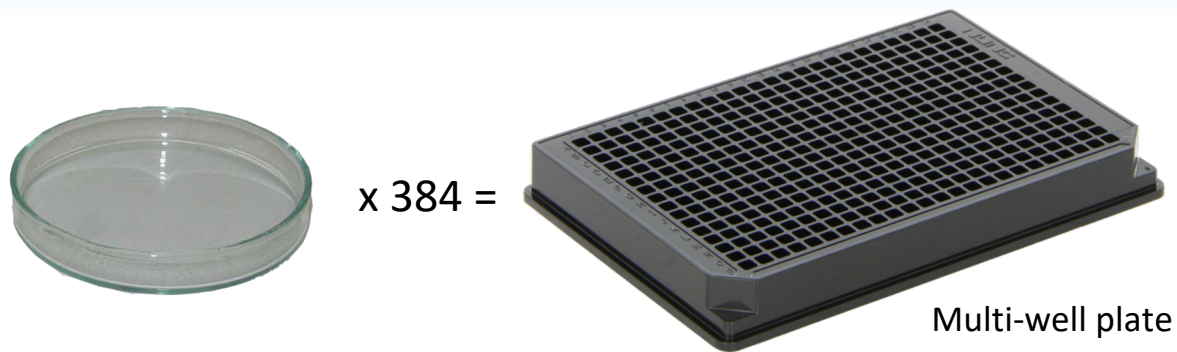


Eroom's Law

Discovering new medicines gets more and more expensive

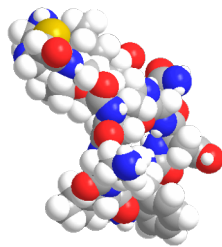


Discovering drugs in high-throughput?



Drug discovery (we wish)

chemical



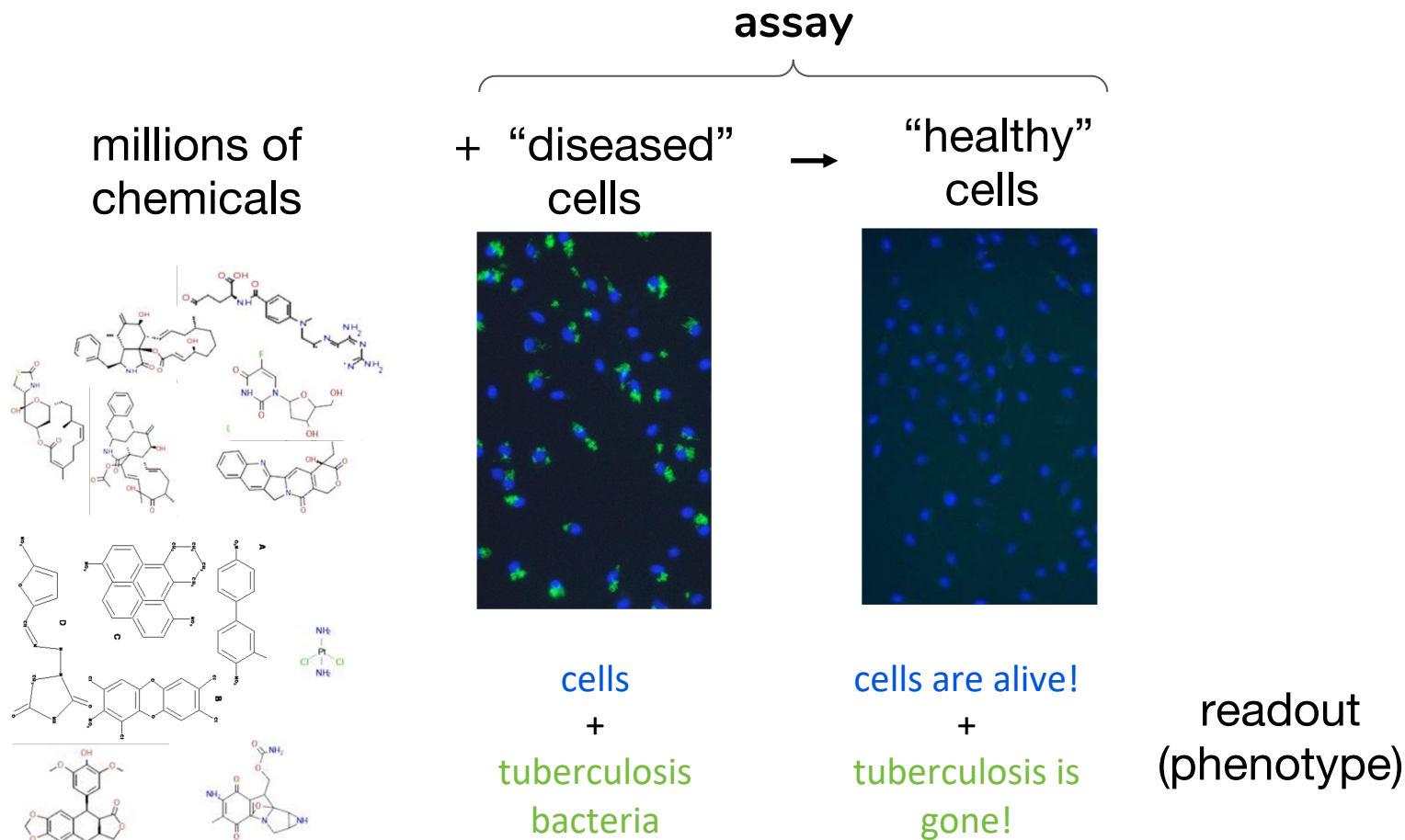
+ diseased
human



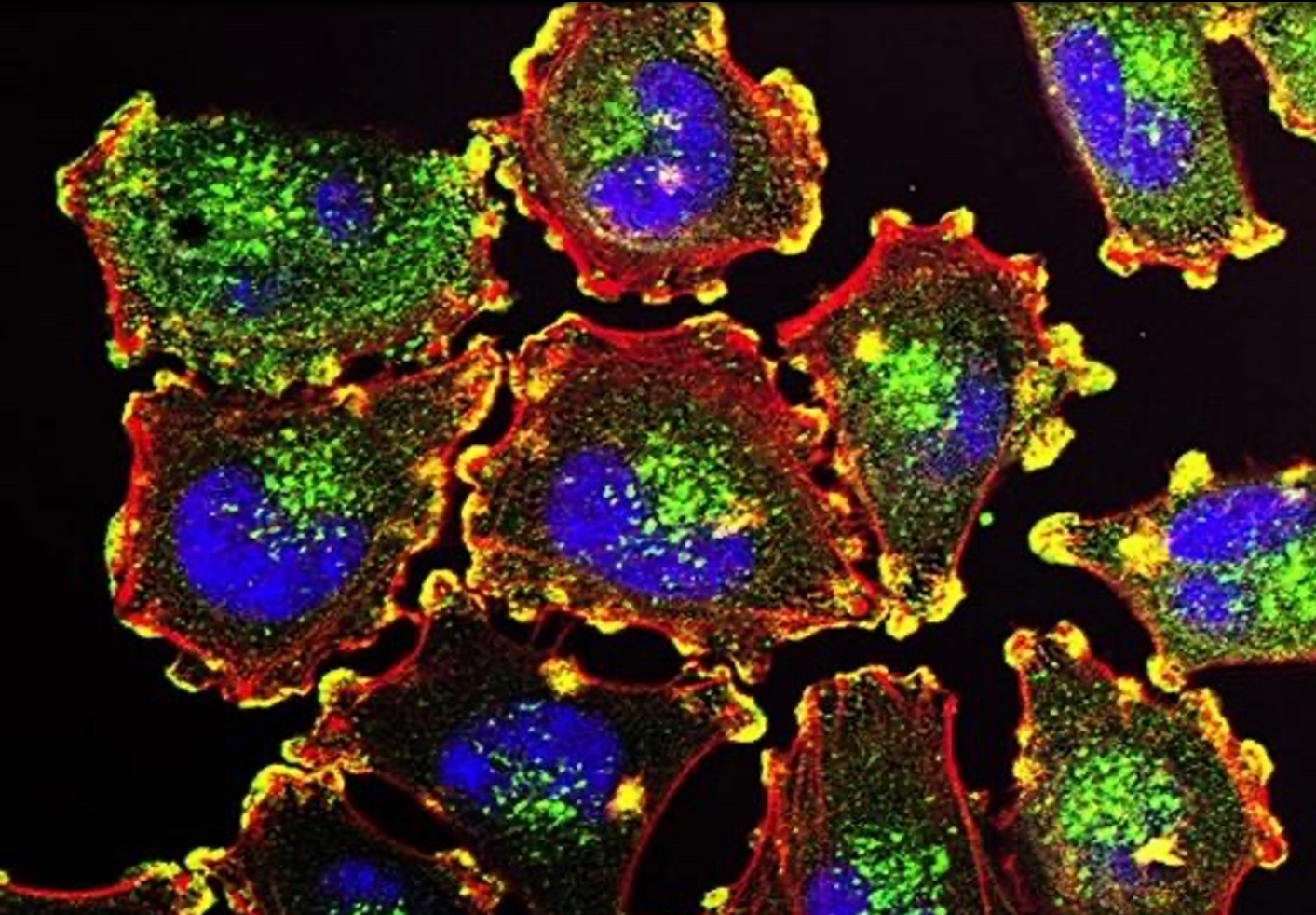
= healthy
human



Drug discovery using cells as proxies



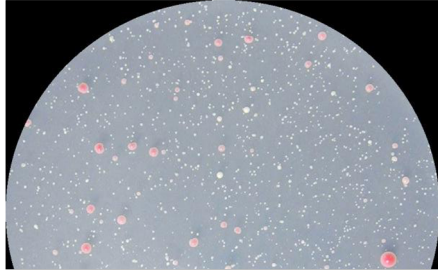
Images contain a wealth of information



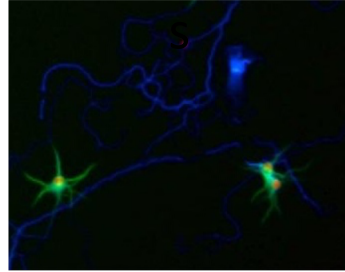
1. single-cell resolution
2. complex model systems
3. quantitative
4. multiplexed

2. Complex models can be quantified by imaging

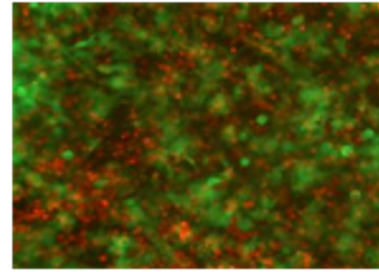
Yeast



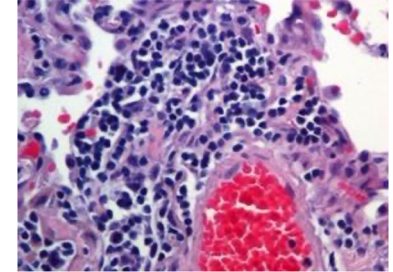
Neuron



Co-cultures



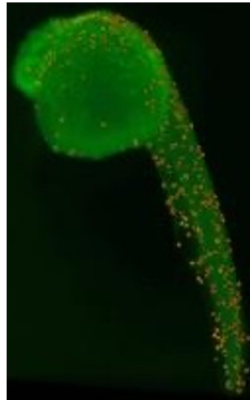
Tissue



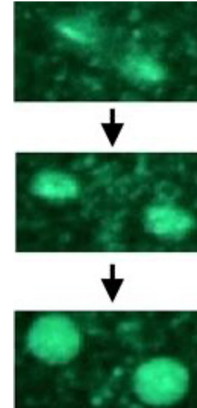
C. elegans



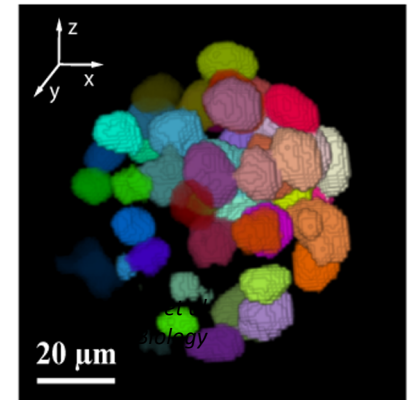
Zebrafish



Time-lapse



3D



3. Image analysis is quantitative

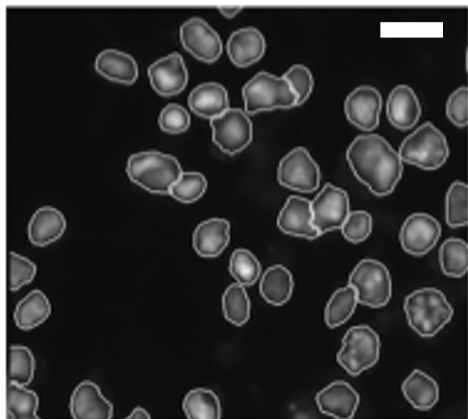


Mark Bray

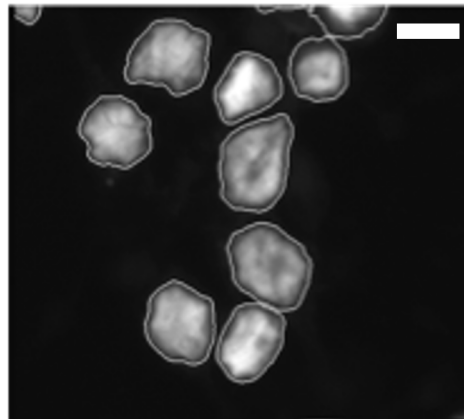


Martha Vokes

DMSO
(negative control)

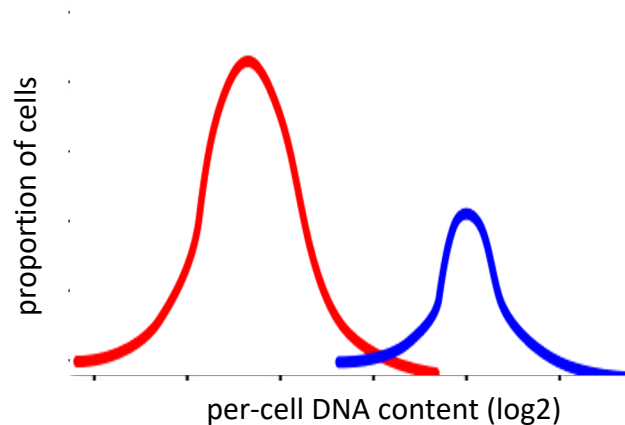


SU6656
(positive control)



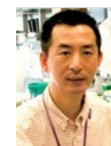
*Polyploidization of
megakaryocytes -
AMKL (leukemia)*

*Leukemic cells
(stained nuclei,
outlined by
software)*



Clinical trial success:
Alisertib for myelofibrosis
Gangat et al. Clin Cancer Research 2019

Wen, et al. Cell 2012



Jeremy Wen,
postdoc

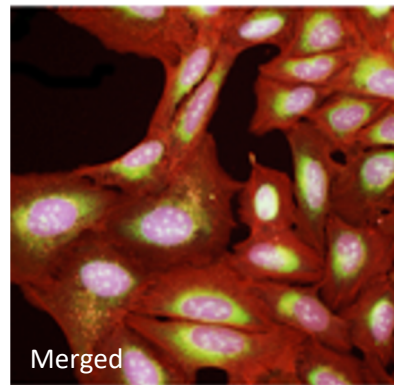
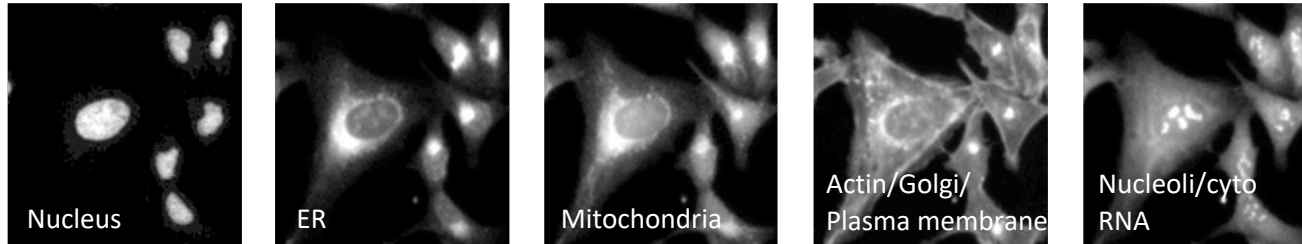


John Crispino,
Northwestern
University

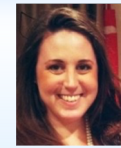
4. Imaging is highly multiplex: multiple stains

Cell Painting

6 stains, 5 channels imaged, revealing 8 constituents/organelles:



Gustafsdottir, et al. PLOS ONE 2013
Bray, et al. Nature Protocols 2016



Kate Sokolnicki



Ray Jones



Sigrun
Gustafsdottir



Paul Clemons



Aly Shamji



Stuart
Schreiber,
Broad Institute

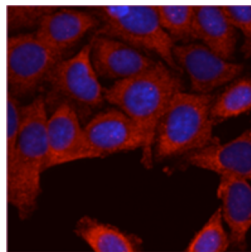
Extracting features from images

Sample prep



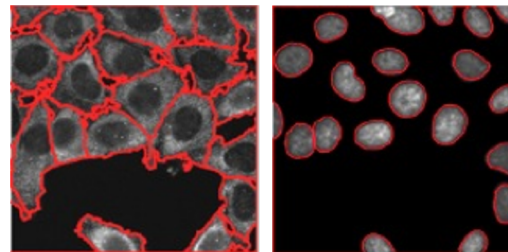
100's of 384-
well plates

Microscopy



1000's of images/ plate;
~5 channels/image

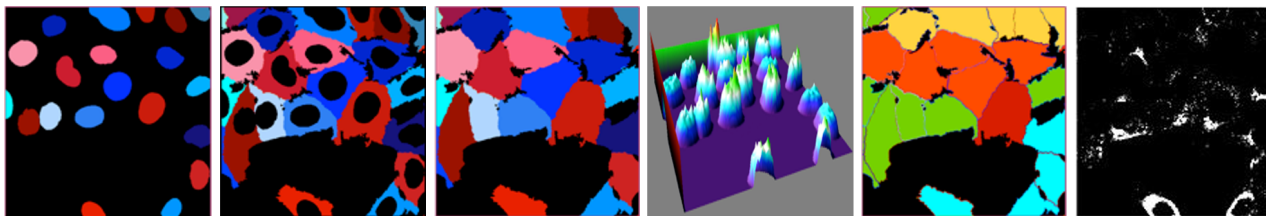
Image analysis:
Segmentation



~500 cells/ image

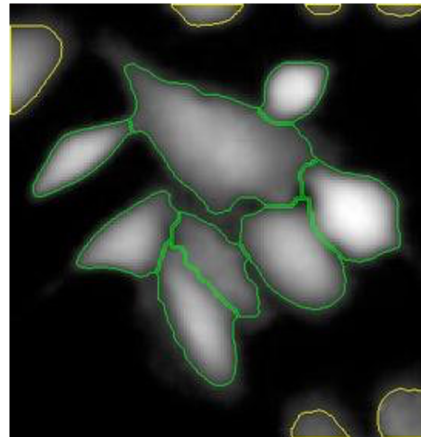
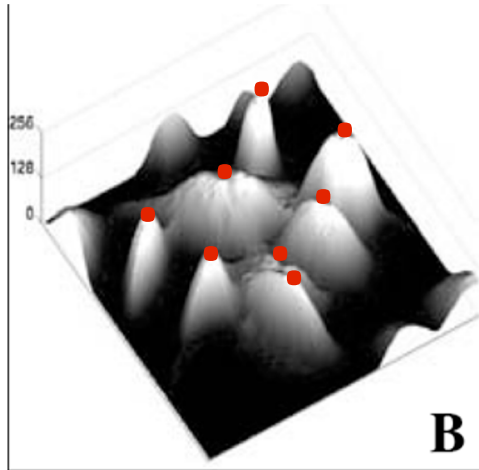
Measure everything

Counts, Shapes, Sizes, Intensities, Textures, Correlations, Relationships



~1500 features / cell

- Powerful image analysis methods
- Point-and-click
- Biologist-focused help
- Flexible, modular, extensible
- Useful for both small and large experiments
- Supports reproducible research



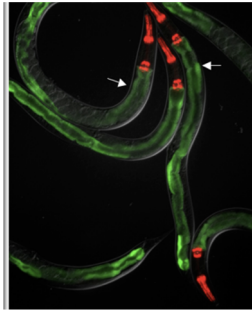
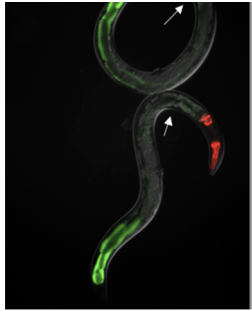
Images from Carolina Wahlby

- Free and open-source; Windows, Mac, Linux
- Cited in **2,000+** papers per year, **9,500+** total
- Ranked **most flexible** and **usable** in independent analysis (*Wiesmann et al.*)



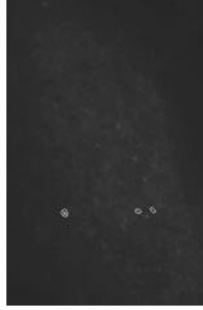
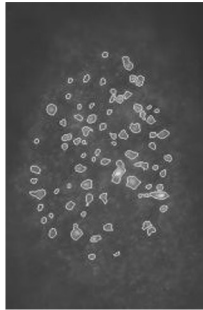
Screen all the things!

S. aureus infection



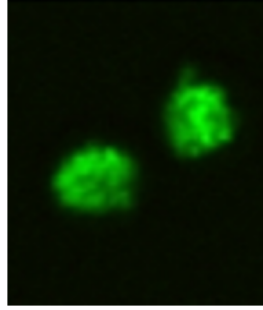
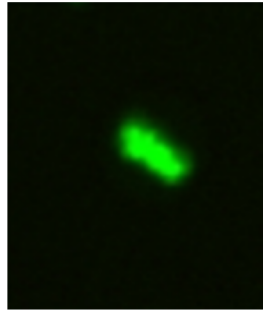
Ausubel/
Irazoqui labs

DNA damage



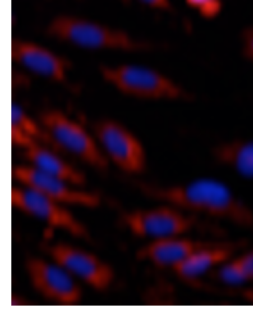
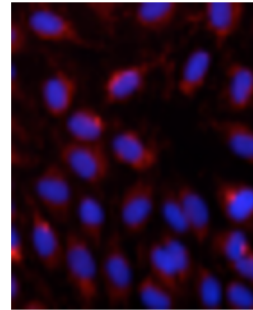
Yaffe lab

Mitosis



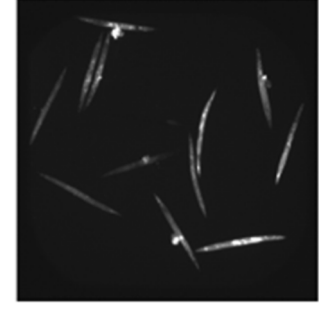
Mitchison lab

Mitochondrial
abundance



Mootha lab

E. faecalis

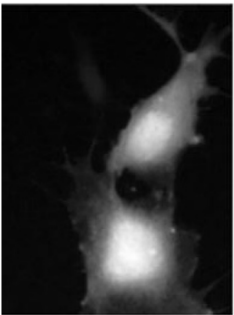
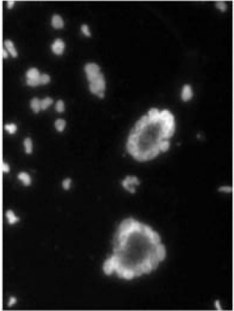


Ausubel lab



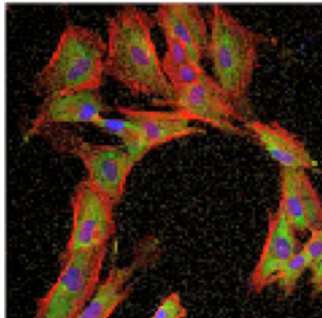
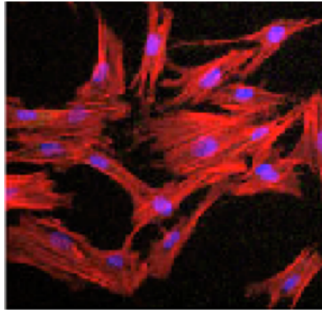
Screen all the things!

HIV neutralization



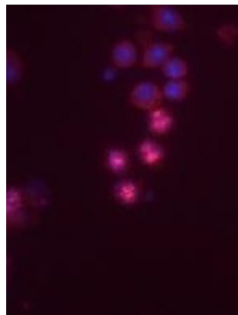
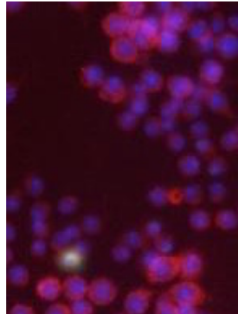
Fenyo lab

mTOR pathway
activation



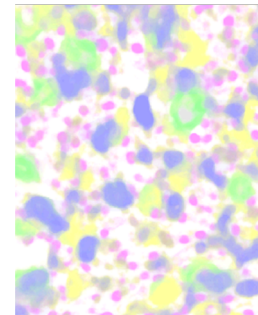
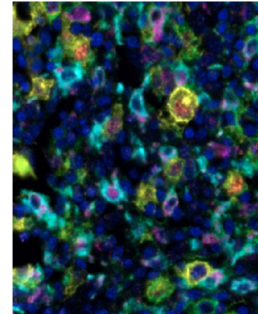
Sabatini lab

Meiosis



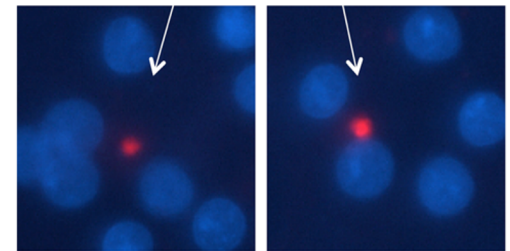
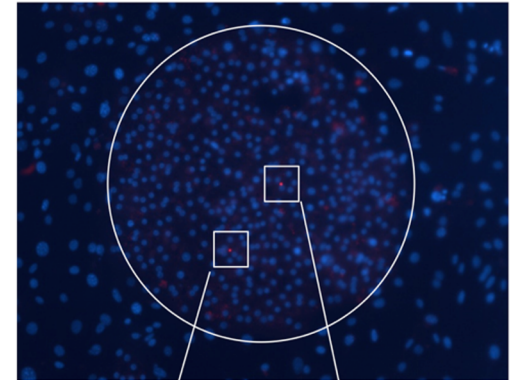
Orr-Weaver lab

Immune
environment



Shipp/Rodig
labs

Malaria



Bhatia lab



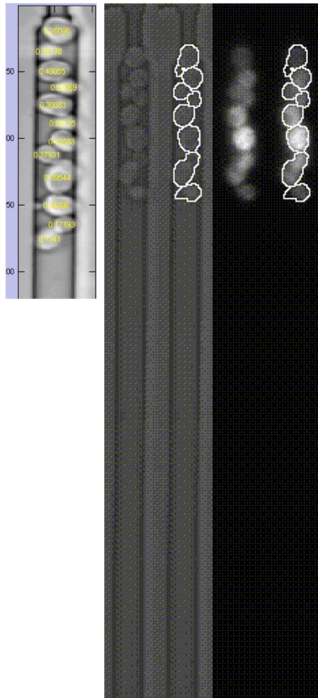
Screen all the things!

Lysosome transport,
Gelfand lab



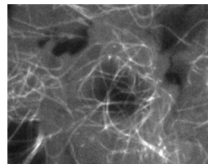
Jolly, et al. *Cell Reports*
2016

Yeast lineages,
Weitz lab

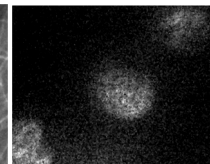


Kim, et al. *Proc. SPIE* 2010

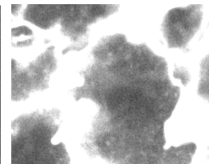
Protein localization, Allen Institute for Cell Science



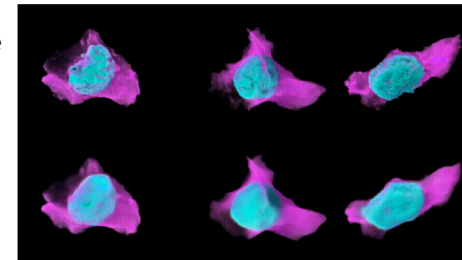
Cytoskeleton
(tubulin)



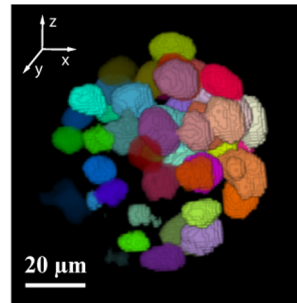
DNA
(Hoechst)



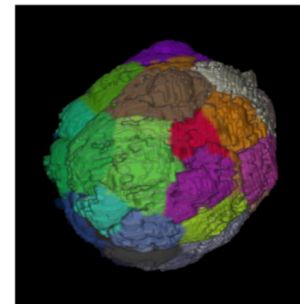
Membrane
(CellMask- membrane)



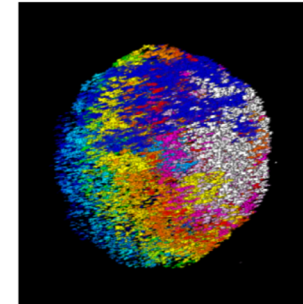
C. Segmented Nuclei



D. Segmented Cells



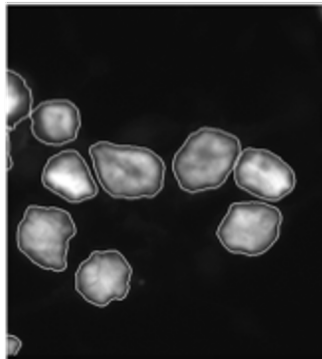
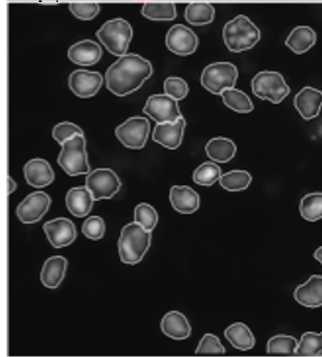
E. Segmented GAPDH



McQuin et al *PLOS Biology* 2018; Allen Institute for Cell Science

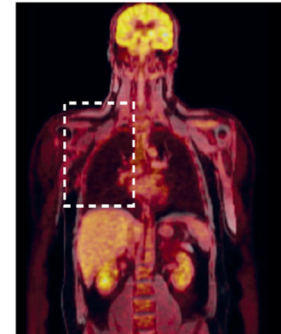
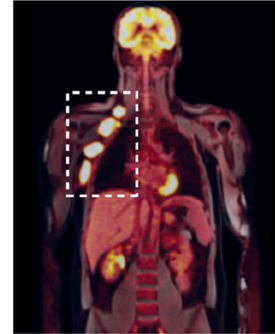
Impact on patients: successful clinical trials

Myelofibrosis



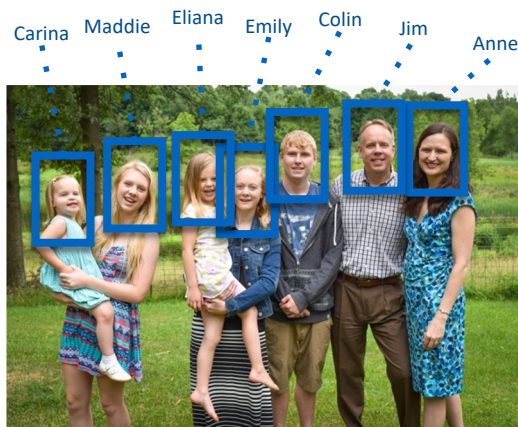
Crispino lab
(Gangat et al. *Clin Cancer Research* 2019)

Leukemias &
Lymphomas

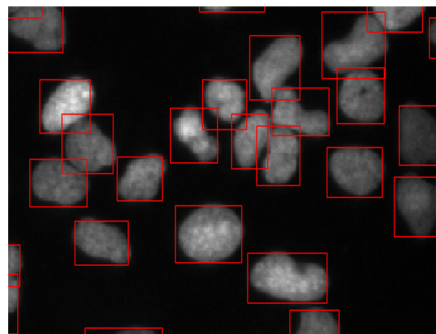


Vienna hospitals & medical
institutes, ETH Zurich
(Snijder B et al. *Lancet Haematology* 2017)

Bringing deep learning to biologists

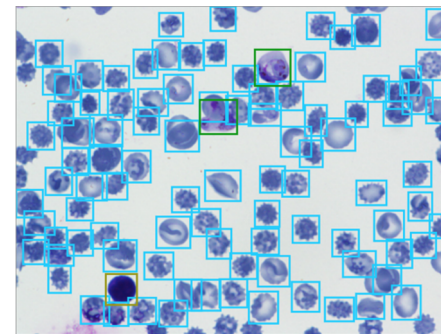


Finding objects
(e.g. nuclei)



Caicedo et al., Cytometry 2019
Caicedo et al., Nature Methods 2019
Data Science Bowl 2018

Classifying phenotypes
(e.g. stages of malaria life cycle)



Hung et al., CVPR workshop 2017



Anne
Carpenter



Allen
Goodman



Kyle Karhohs



Jeanelle
Ackerman



Beth
Cimini



Minh
Doan



Tim
Becker



Shantanu
Singh



Juan
Caicedo



Jonathan
Roth

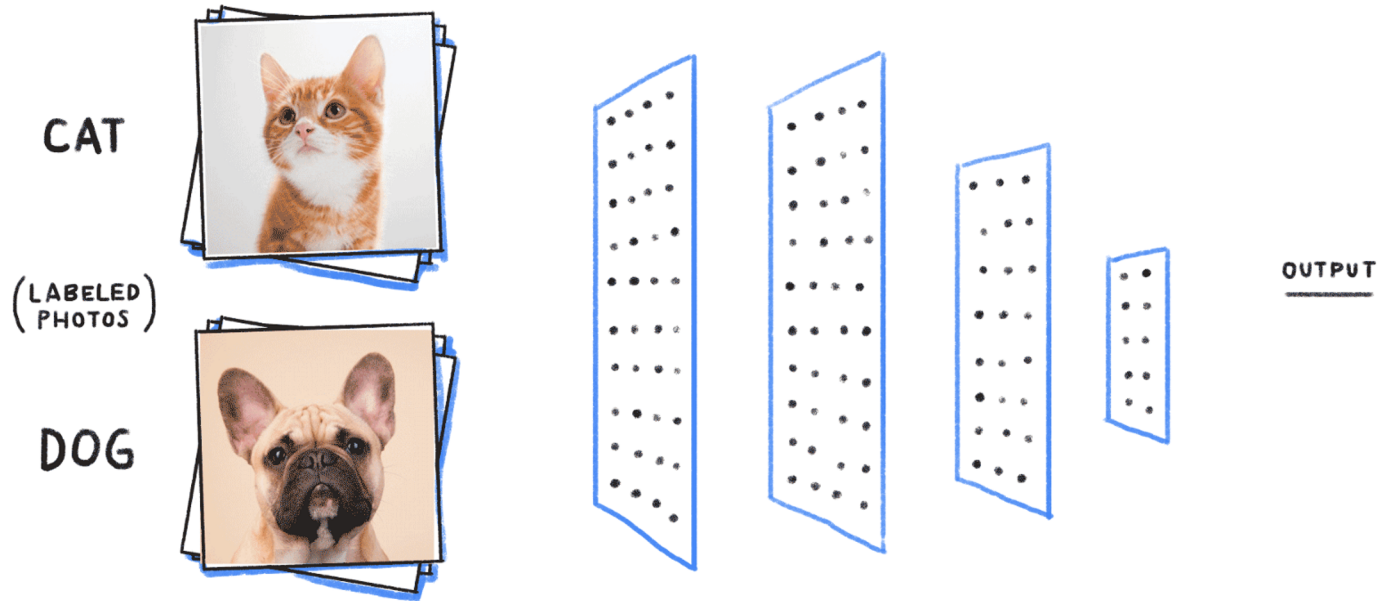


Jane
Hung

What is deep learning?

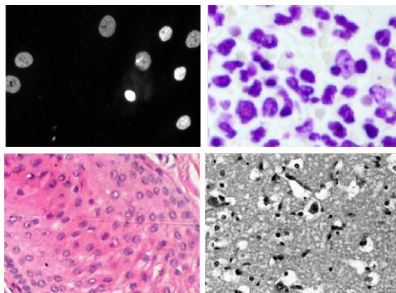
Machine learning: builds an algorithm/model/set of rules that can predict outcomes (pixels / structures / phenotypes) given training examples to learn from

Deep learning: a type of machine learning using many layers of processing



Deep learning excels at segmentation (finding things)

2018 Data Science Bowl: Towards a Universal Nucleus Finder



From Jan to Apr 2018
700 images
~37,333 nuclei
3,919 teams
65,333 submissions



Anne
Carpenter



Allen
Goodman



Kyle Karhohs



Jeanelle
Ackerman



Beth
Cimini



Minh
Doan



Tim
Becker



Shantanu
Singh



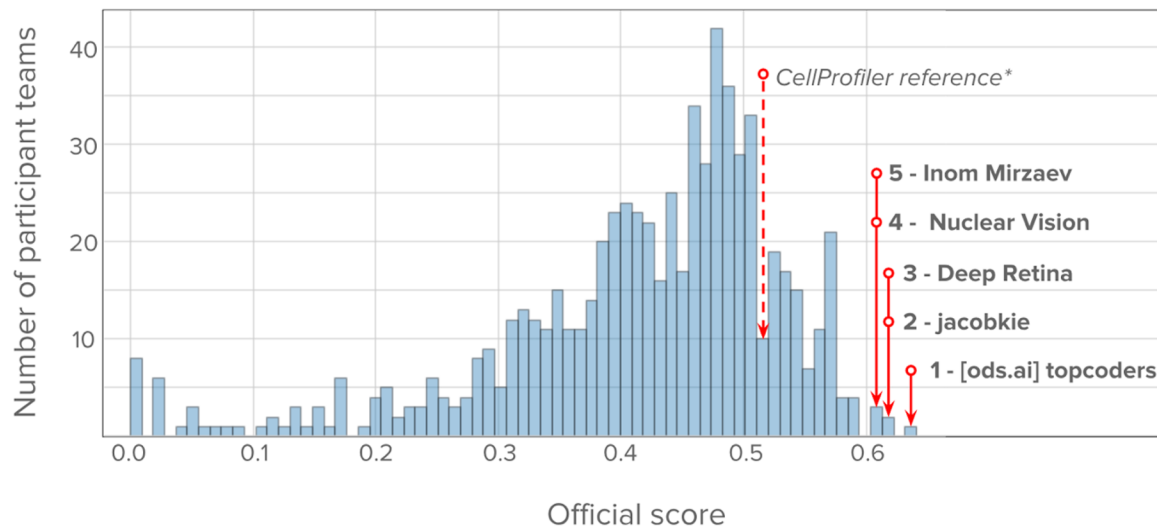
Juan
Caicedo

Caicedo et al., *Nature Methods* (2019)

Deep learning excels at segmentation (finding things)

2018 Data Science Bowl: Towards a Universal Nucleus Finder

Distribution of scores in second-stage evaluation



Anne
Carpenter



Allen
Goodman



Kyle Karhohs



Jeanelle
Ackerman



Beth
Cimini



Minh
Doan



Tim
Becker



Shantanu
Singh

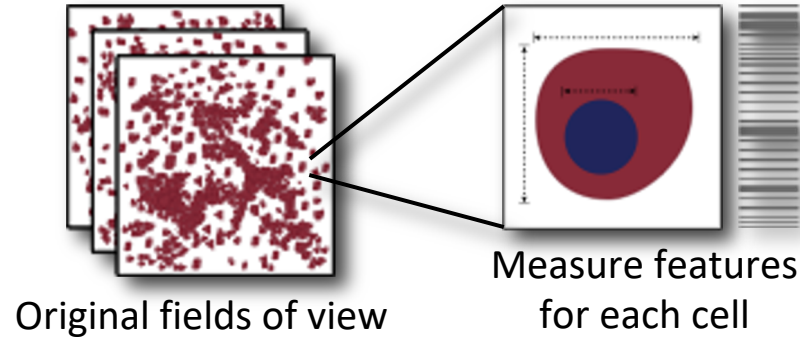


Juan
Caicedo

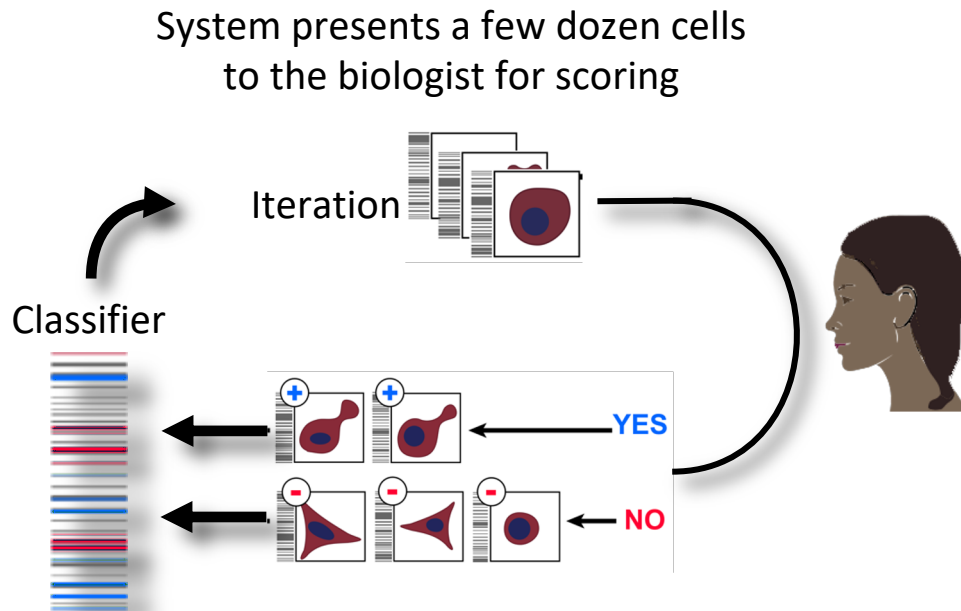
Horvath lab: www.NucleAlzer.org

(supervised classification)

Step 1: Identify & measure cells

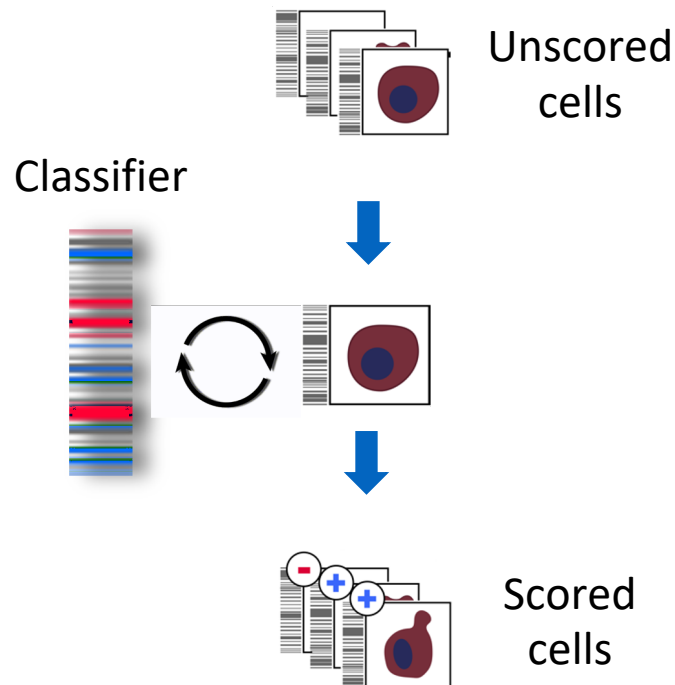


Step 2: Train the classifier



System defines classifier based on cell features

Step 3: Score cells



User-friendly machine learning for phenotype classification



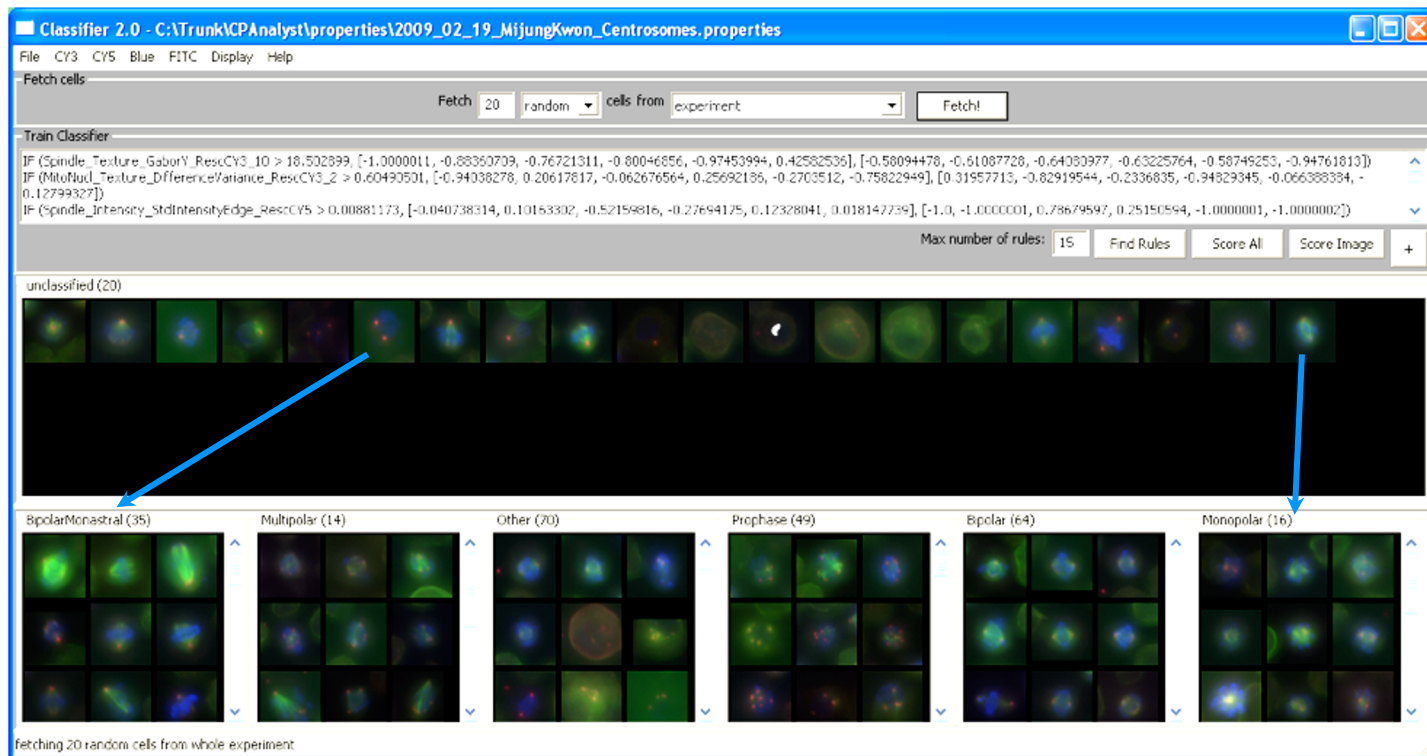
CellProfiler Analyst
data exploration software



ADVANCED
CELL CLASSIFIER

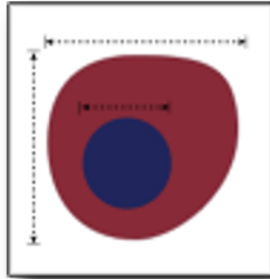
Coming soon!

Piximi: *deep learning, web-based app for classification*



A revolution in how we use images

Measure known
phenotypes



Profile to characterize
samples

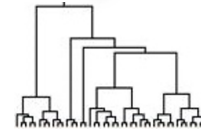
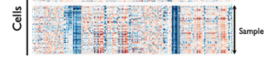
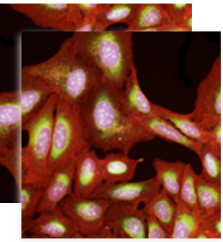


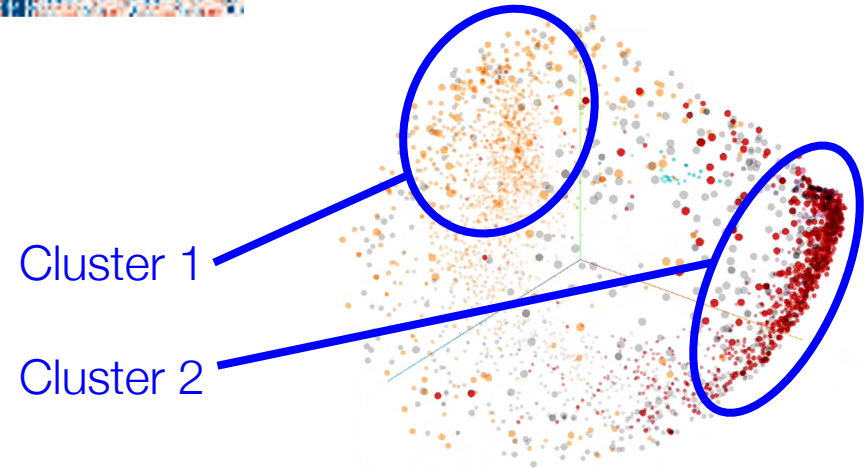
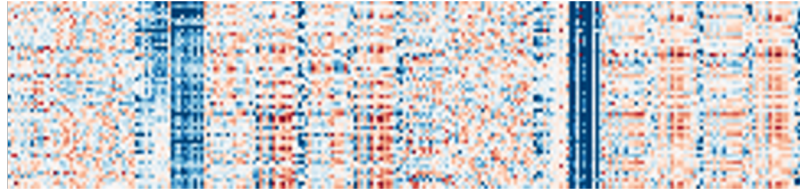
Image-based profiling

“Measure everything, ask questions later”



single cells

1,000+ morphology features



Transforming drug discovery via image-based profiling

Assay

Small molecule library

Hits → leads

Preclinical studies
Clinical trials

Effective drug

How might the rich information
in images speed up the drug
discovery process?

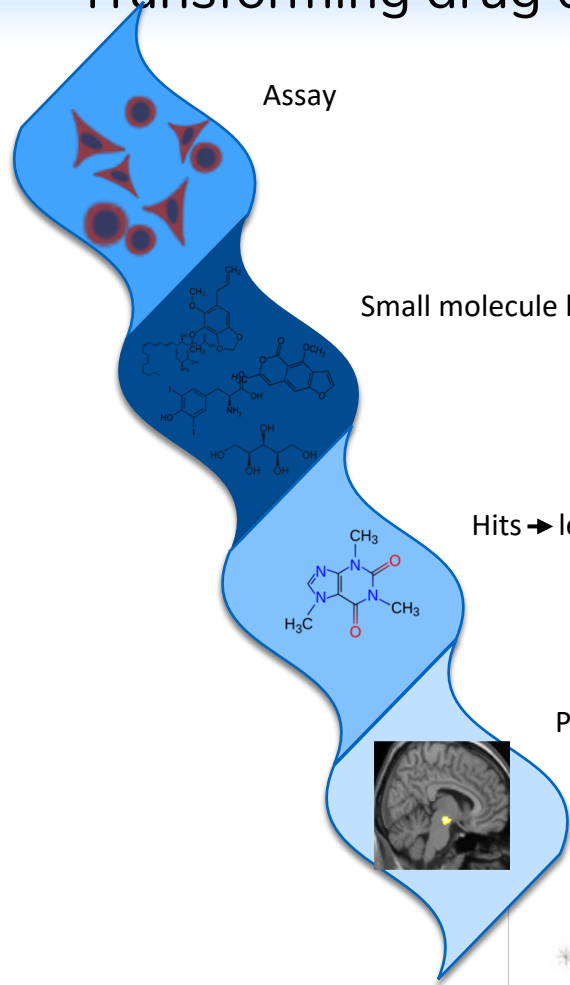


Image-based profiling: accelerating many steps of drug discovery

Annotate function of
disease-associated genes

(Rohban eLife 2016)

Disease-relevant assays
based on causative
human alleles

(Gibson Circulation 2015)

Virtual screening by
computationally predicting
assay activity

(Simm Cell Chem Bio 2018)

Small molecule libraries
enriched for phenotypically
diverse compounds

(Wawer PNAS 2014)

Determining
targets/mechanism-of-action
of lead compounds

*(Gustafsdottir PLOS One
2013, Ljosa JBS 2013)*

Structure-activity-relationship
(SAR) for hit-to-lead

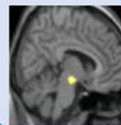
*(Gerry JACS 2016, Nelson Org
Letters 2016, Melillo JACS 2018)*

Lead-hopping

Predicting drug toxicity,
side effects

*(Nyffeler Tox Appl
Pharmacol 2019)*

Stratify patients
for precision
medicine



Effective drug

Transforming drug discovery via image-based profiling

Assay

Can we identify signatures of disease?
Then screen drugs to reverse the signature?

Small molecule library

Hits → leads

Preclinical studies
Clinical trials

Effective drug

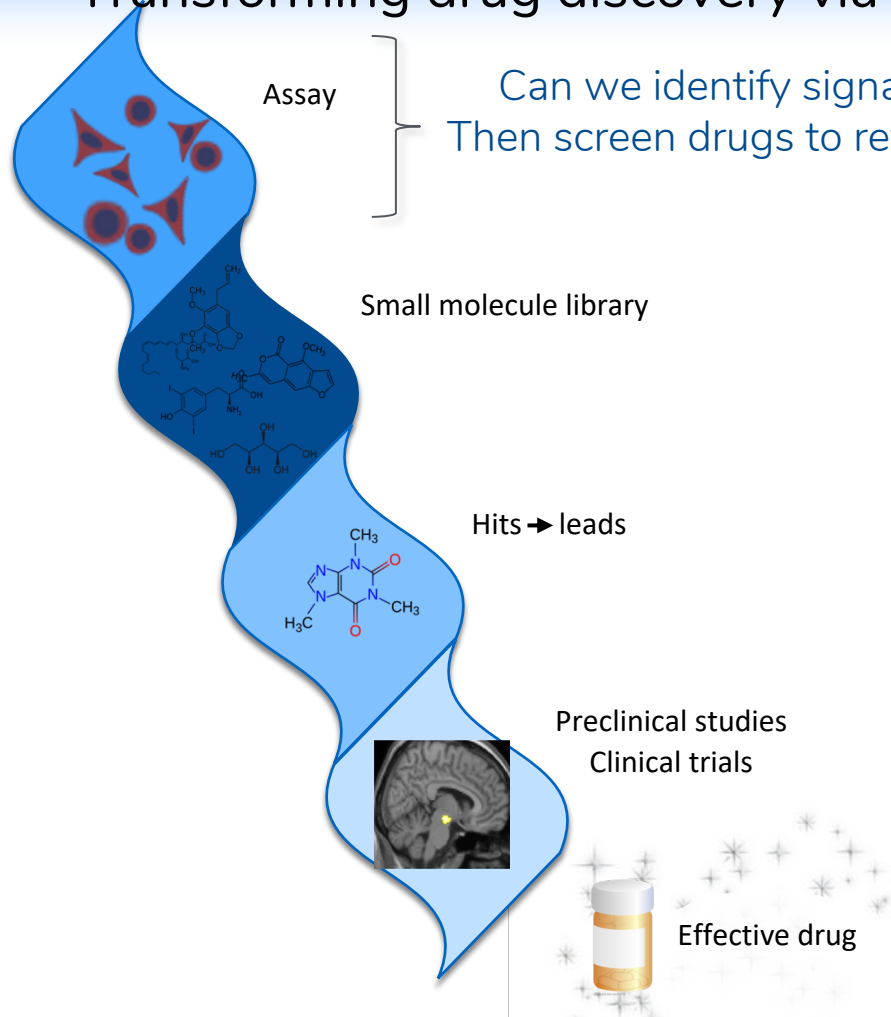
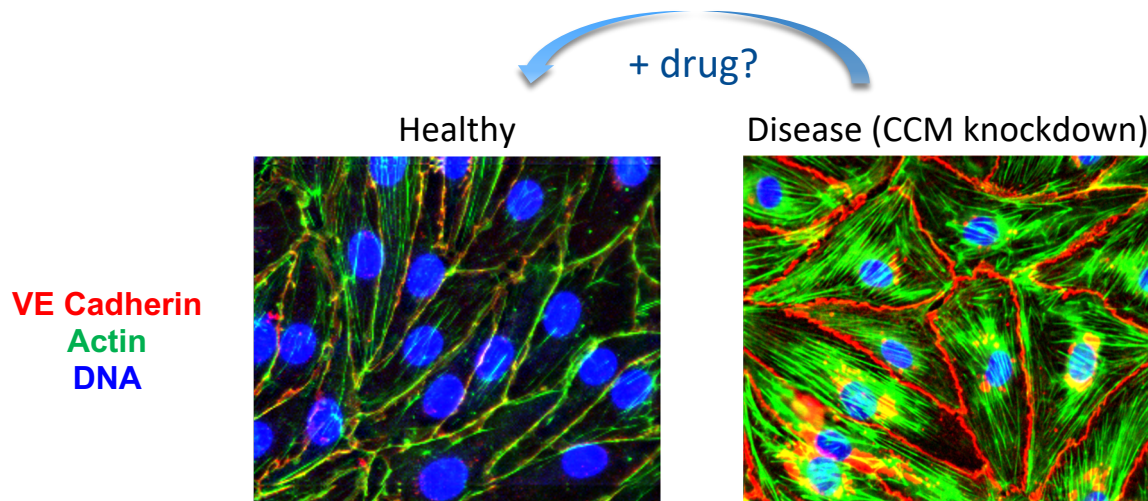


Image-based profiling can identify drugs for disease



Drug chosen as hits based on automated analysis outperformed those chosen by expert visual analysis.

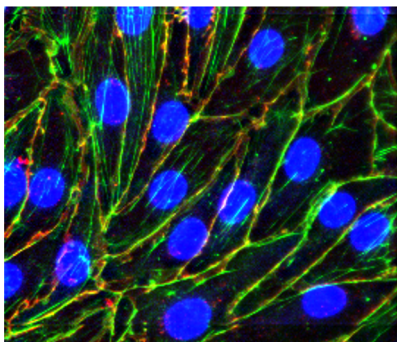
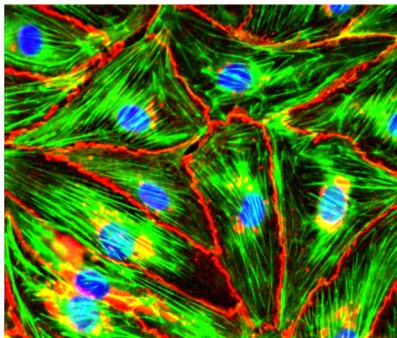
Now: 500+ disease models available for screening in parallel.



*Disclosure: I serve on
Recursion's Scientific
Advisory Board*

Impact on patients: in-progress clinical trials

(1) Cerebral cavernous malformation

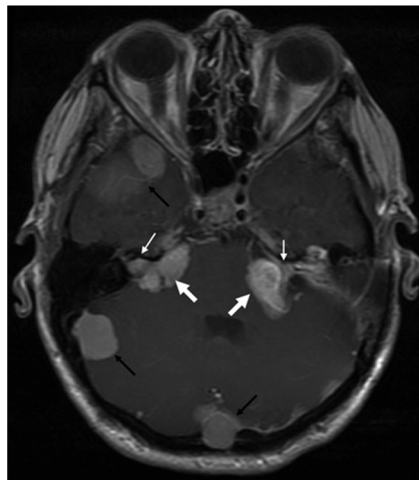


University of Utah
(Gibson et al. *Circulation*
2015)

(2) GM2 Gangliosidosis (Tay-Sachs)

(3) Solid Tumors

(4) Neurofibromatosis Type 2



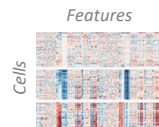
Recursion



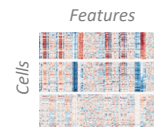
*Disclosure: I serve on Recursion's
Scientific Advisory Board*

Disease states can be morphologically distinct

Cells from patient without mental illness

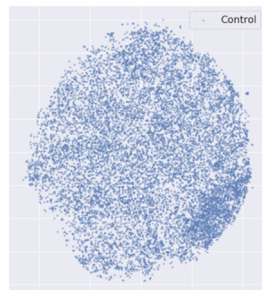


Cells from patient with a disorder

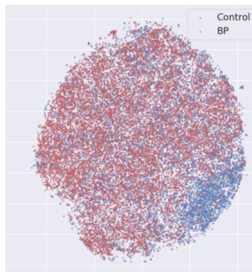
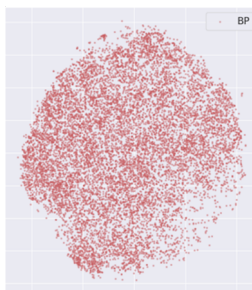


Detectable
phenotypic
difference?

Controls



Bipolar



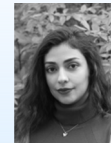
Project in progress



Kyle Karhohs



Mohammad
Rohban



Marzieh
Haghighi



Bruce Cohen,
McLean
Hospital



Donna McPhie

Transforming drug discovery via image-based profiling

Assay

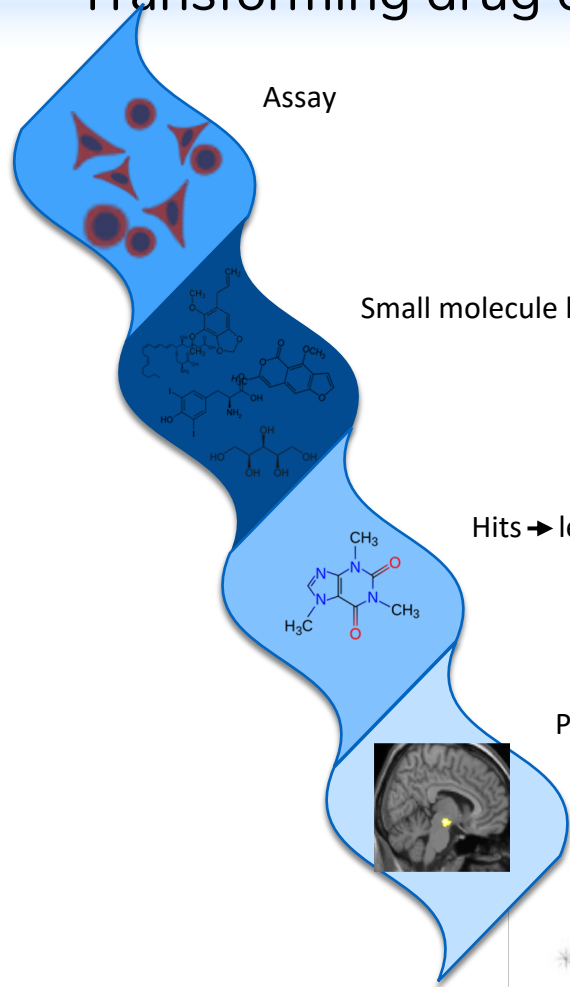
Small molecule library

Hits → leads

Preclinical studies
Clinical trials

Can we make better diagnostics?

Effective drug

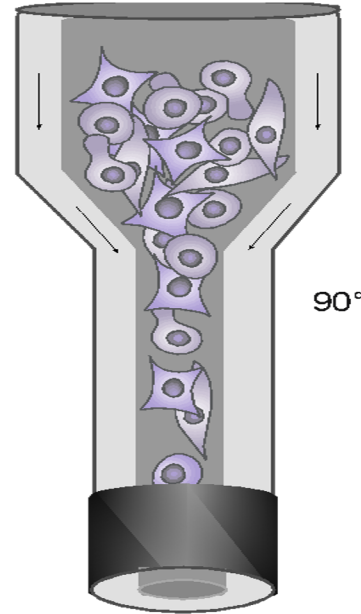


Imaging flow cytometry


Imaging flow cytometry



Light
sources



Darkfield
detect



Brightfield
detect

+ fluorescence
channels

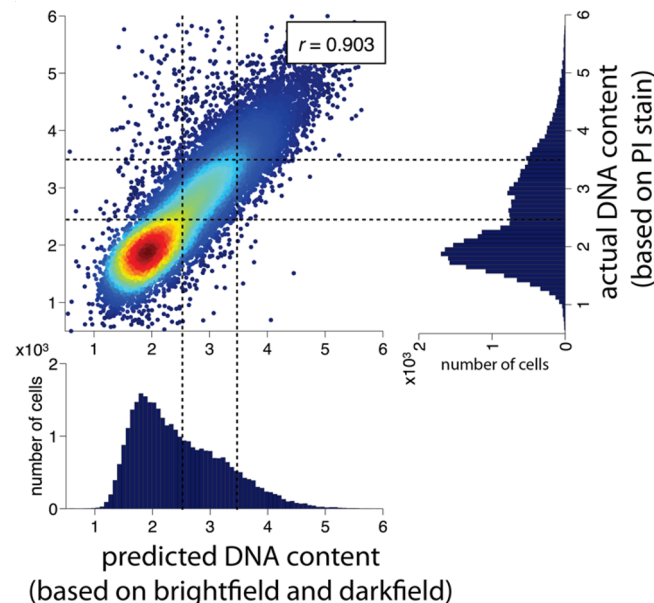
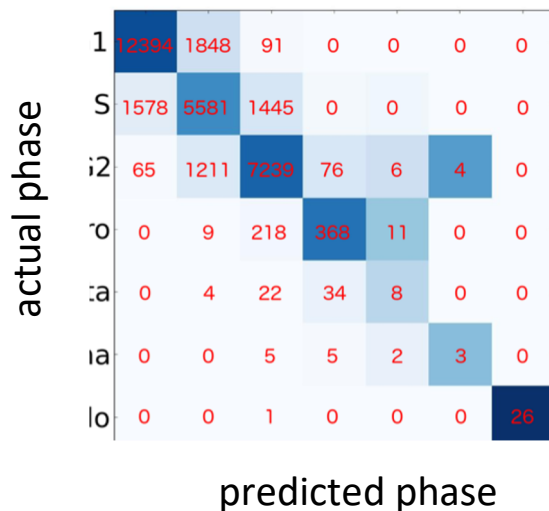
Hidden information: Classifying cell cycle stage label-free



Holger Hennig



Thomas Blasi



Paul Rees,
Swansea Univ.



Fabian Theis,
Helmholtz
Zentrum München



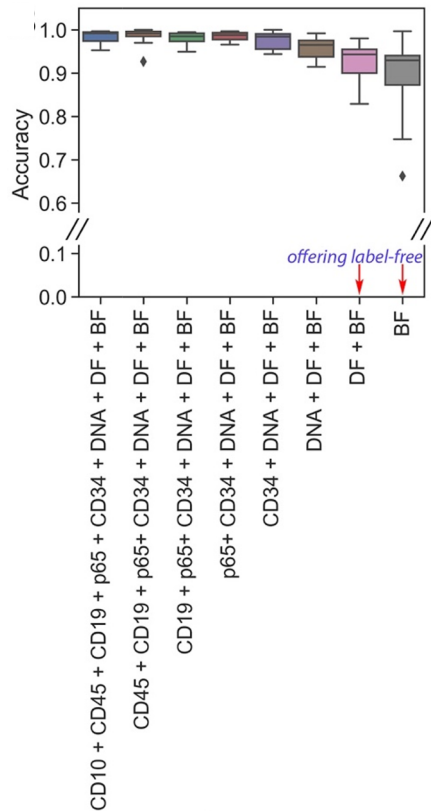
Andrew Filby,
Newcastle
Univ.



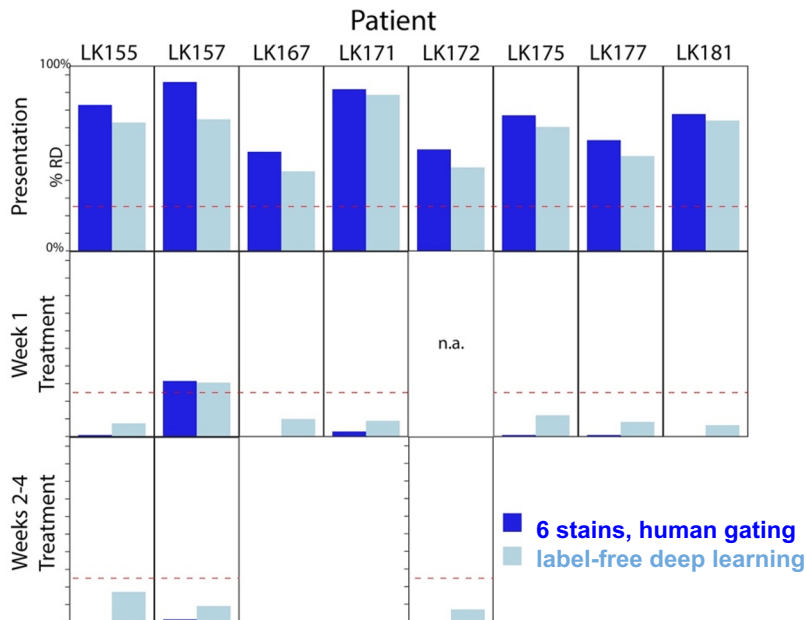
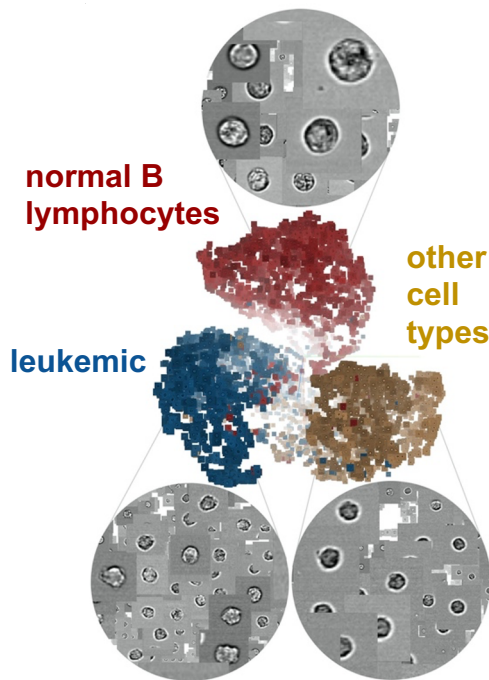
Huw Summers,
Swansea Univ.

Detecting leukemia (ALL) in blood samples, label-free

Minh Doan Marian Case Dino
Masic Holger Hennig Claire
McQuin Juan Caicedo Shantanu
Singh Allen Goodman Olaf
Volkenhauer Huw D. Summers
David Jamieson Frederik W van
Delft Andrew Filby Paul Rees Julie
Irving



Can we replace a 4 to 10-antibody flow cytometry assay with images captured label-free?

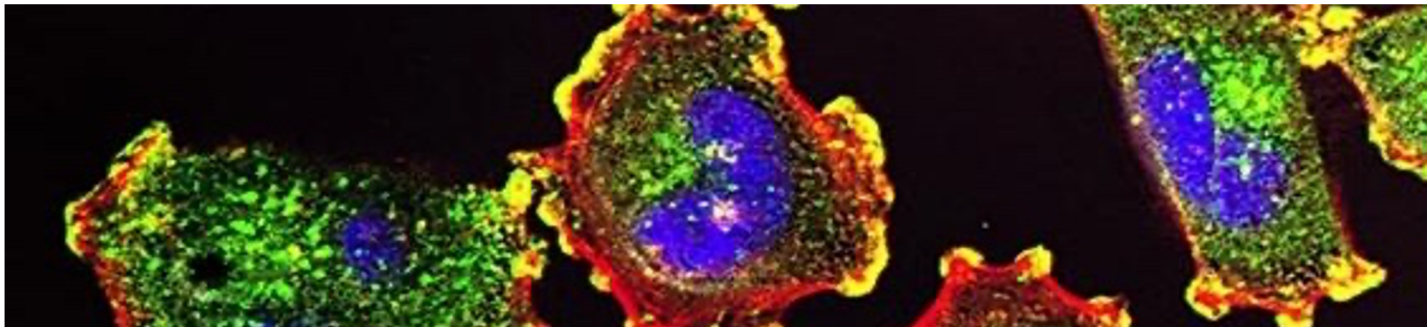


What's needed to use AI well?

- **Identifying** situations appropriate for deep learning
- **Lots of examples** of right & wrong answers are needed for training
 - (for example, collecting manually-set parameters from users)
- Examples must come from diverse sources and conditions that **reflect reality** (what will be seen in the wild)
- Focus on **user needs** and workflow
- **Imagination**: willingness to trust AI can see more than we can (with controls!)

What will AI increase demand for?

- Instruments & reagents that produce **consistent** data are highly valued
- Large-scale, high-throughput experiments fuel advanced AI applications





Lab members

Hamdah Abbasi
Adeniyi Adeboye
Niranj Chandrasekaran
Yuen Ler Chow
Beth Cimini
Allen Goodman
Marzieh Haghighi
Nasim Jamali

Becki Ledford
Alice Lucas
Saketh Mynampati
Pearl Ryder
Shantanu Singh
David Stirling
Hillary Tsang
Gregory Way
Erin Weisbart

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- NSF/BBSRC Bio award (Carpenter/Rees)

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