

# Single Cell Systems Biology: Measuring Cell Signaling by Flow Cytometry

Cytobank: Manage, Analyze & Share your cytometry data from Anywhere

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Microbiology & Immunology, Stanford University

<sup>2</sup>Cytobank Inc., Mountain View CA

# What's So Great About Flow?

## Challenge

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- Study **heterogeneous** primary tissues
- Pinpoint abnormal **cell subsets**
- Identify and track **cancer stem cells**
- Look not just at 'pathways', but the broader **signaling network**
- Identify targets for **drug discovery**
- Choose, monitor, & **optimize therapies**
- Understand mechanisms of cell/cell and **disease cell/host cell interactions**
- **Detect disease earlier**

## Example Question

---

Can we begin discovery in human samples?

Can we spot pre-transformation cells?

Is there a rare, therapy-resistant subset?

Are there off-target effects of a drug?

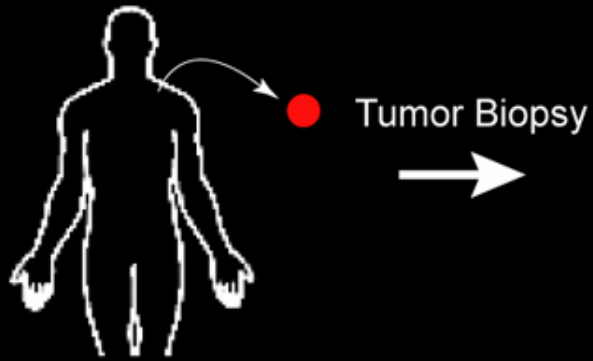
What (signaling) mechanisms enable cells to resist a particular therapy?

Do patients that share responses share profiles?

How do cancer cells interact with and alter the host microenvironment or immune system?

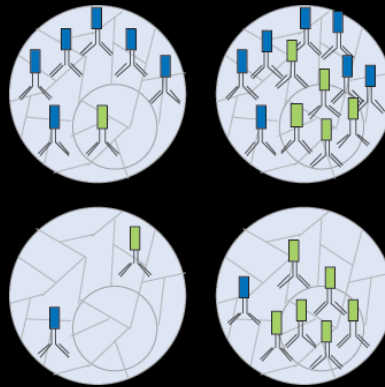
Can we detect circulating cancer cells or immune cells that encountered tumor?

# Flow cytometry allows tens of measurements *per cell*

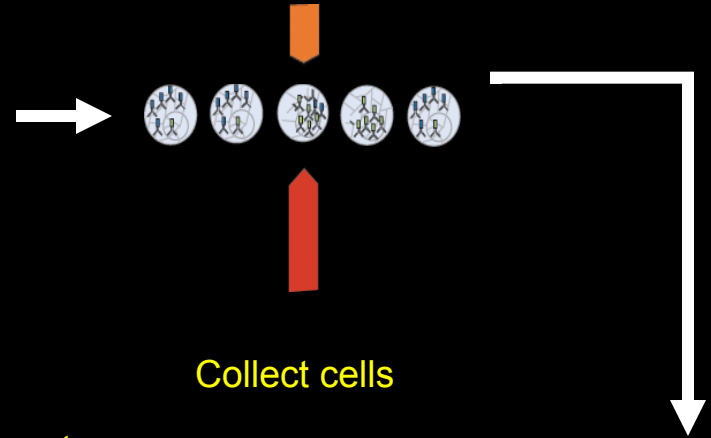


Cancer Patient

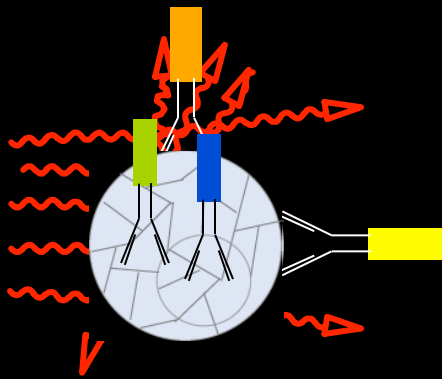
## Extract Cells



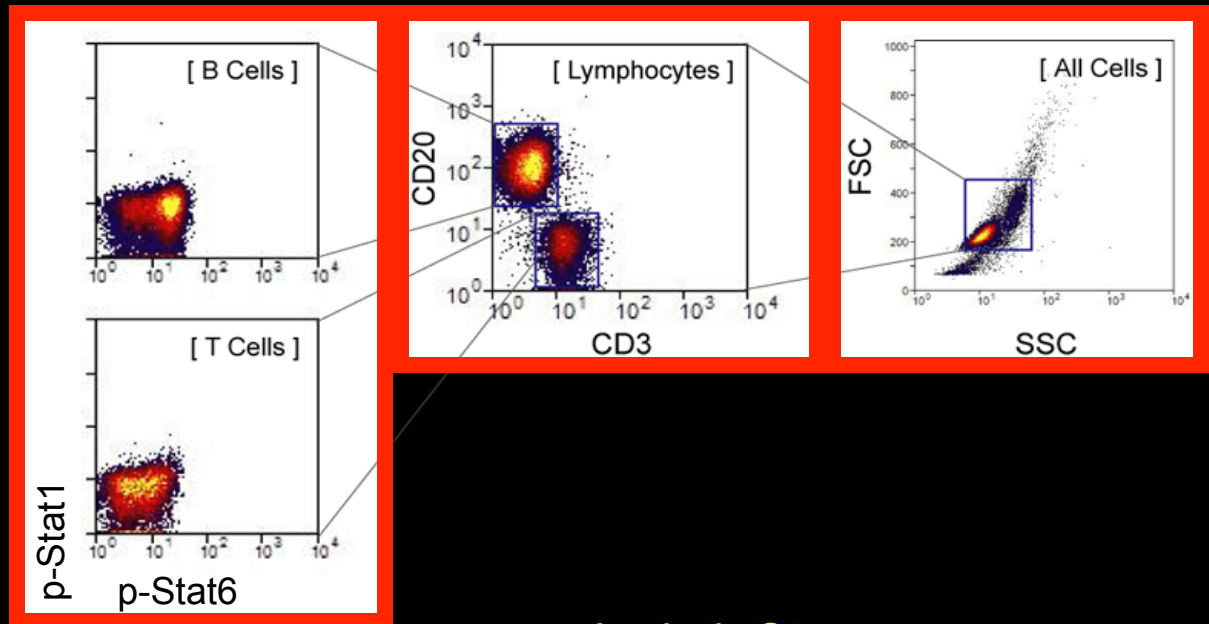
Expose to condition  
Stain with antibodies of interest



Collect cells

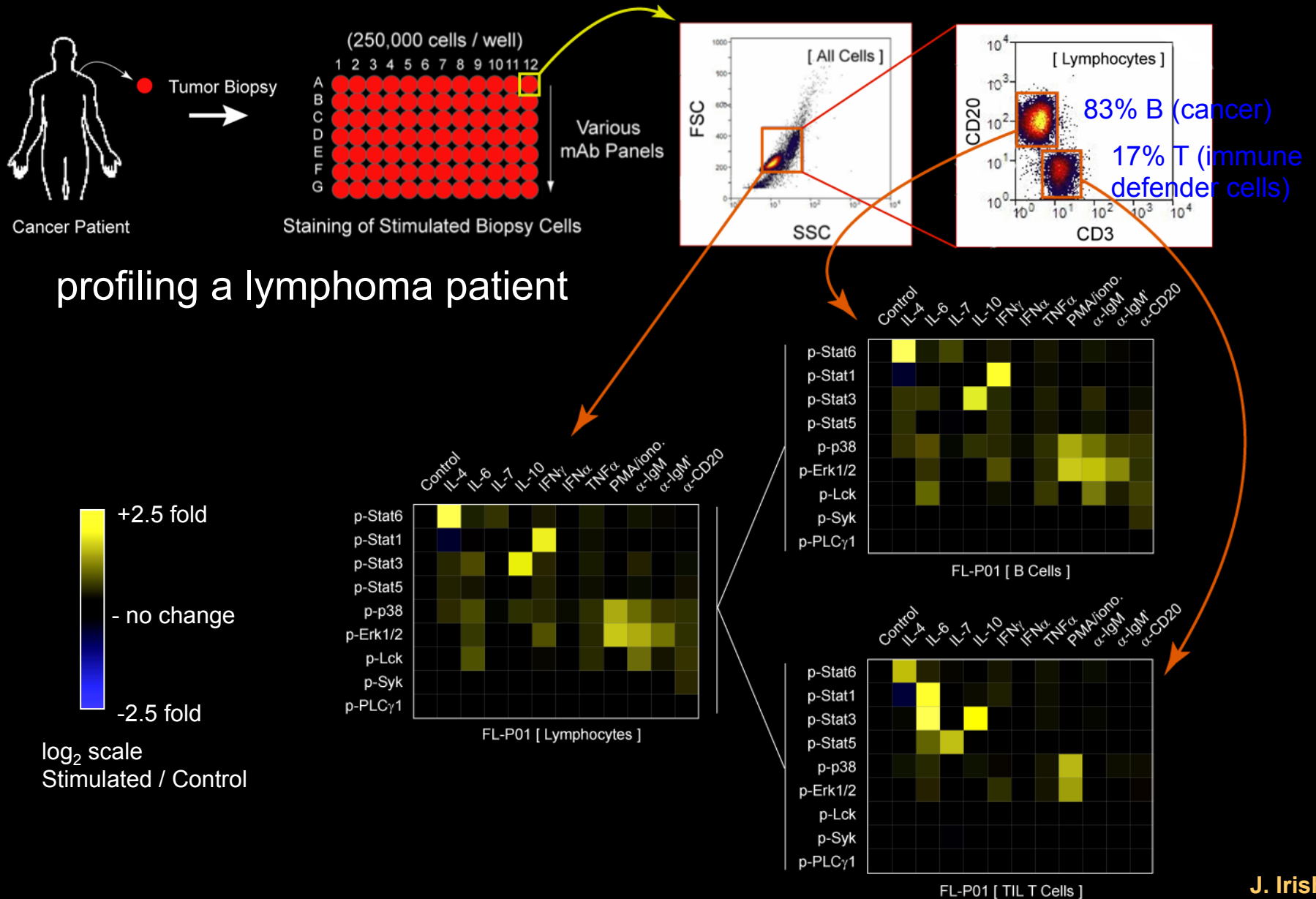


1. *Identify live cells*
2. *Identify cell type*
3. *Evaluate cell signal*



Analysis Steps

# Dramatically Different Profile in Immune Defender Cells



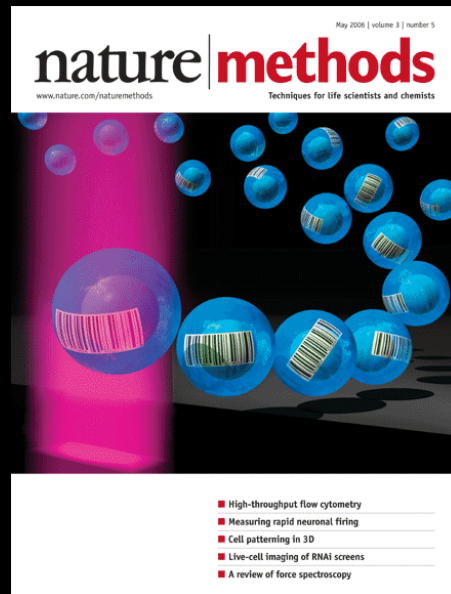


# Mechanism



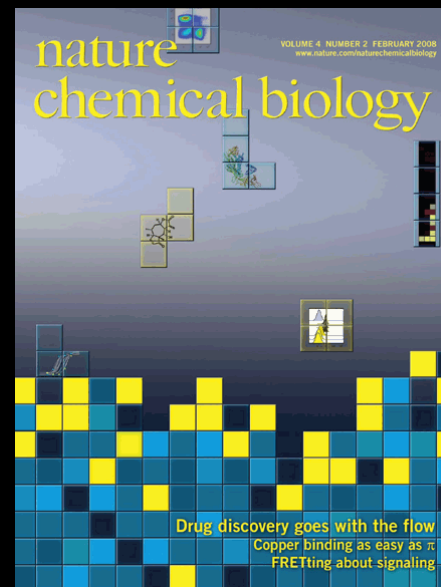
Perez, *et al.*  
Blood,

# Technique



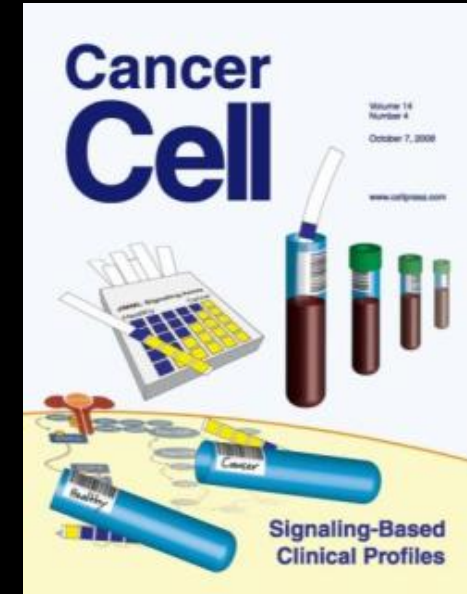
Krutzik, *et al.*  
Nat Methods

# Screening

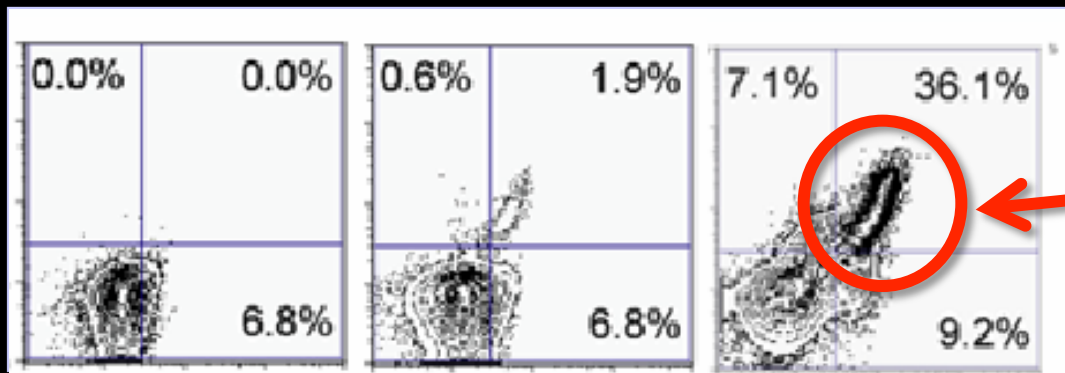


Krutzik, *et al.*  
Nat Chem Bio

# Diagnostics

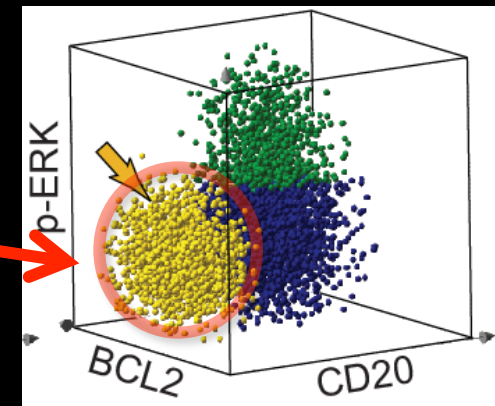


Kotecha, *et al.*  
Cancer Cell

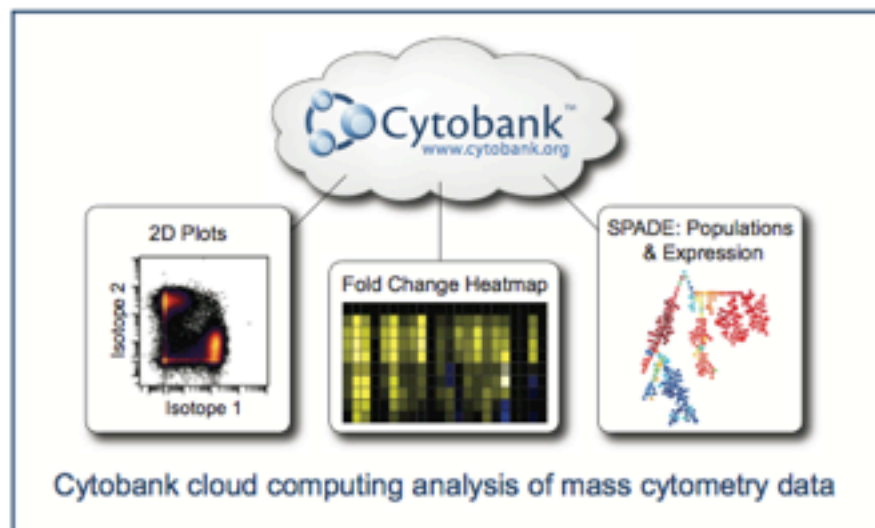


Irish, *et al.* Cell

Bad  
Guy  
Cells

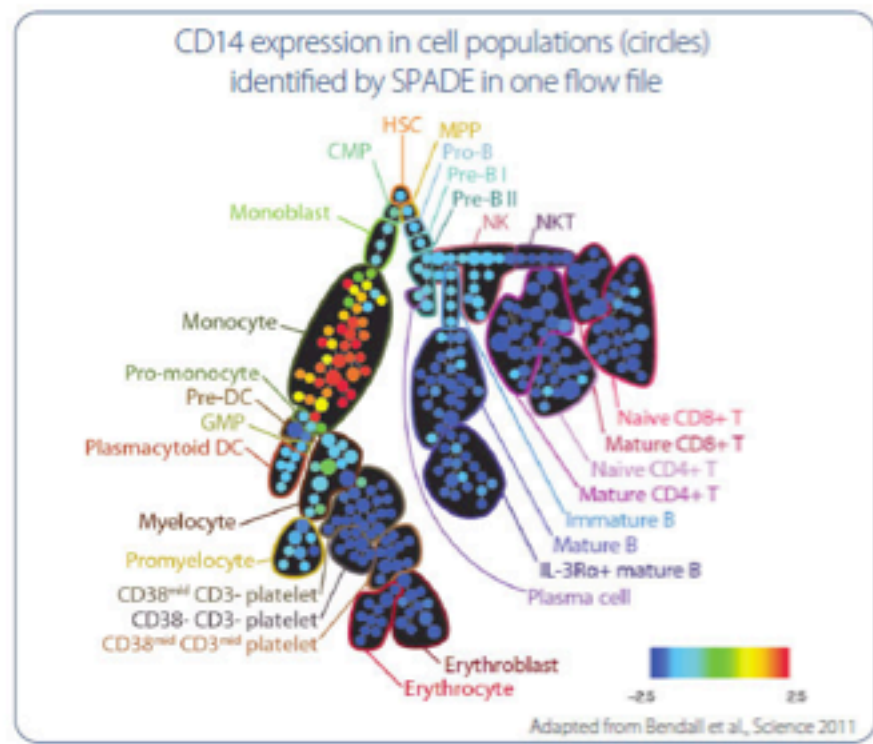


# High dimensional cytometry is here



**30+ Parameters hosted and analyzed on Cytobank**

**Bendall et al. Science 2011**



# Key Tools

## 1) Access to samples

- ideally uniform initial therapy
- long term clinical outcomes or paired samples
- balanced training and testing sample sets

## 2) Flow cytometry & signaling network profiles

- map signaling in every cell within a tumor specimen
- markers for tumor, non-malignant, and cell subsets
- cell sorting for follow up studies of genetics and epigenetics

## 3) Cloud computing to link all our knowledge & tools

- data storage & annotation, data sharing
- web based analysis tools for researchers
- computational analysis & modeling tools (SPADE)
- informatics (patient information, ontologies)

# Cytobank is for *managing, sharing & analyzing* flow experiments over the web



Experiment =  
Collection of FCS files



Cytobank  
Inbox Profile Help  
Welcome, [User] Logout

Create a New Experiment

Getting Started!

All Experiments (1334)

Select Labels: [None] [New Label] [Delete Experiments]

Experiment Name	PR	Project	KR	Updated		
Phospho-Flow Titration of IL-2 in Healthy PBMC	53	5	Jonathan Imph	Jan 27		
Dose experiment with kit (KR dataset 2) [Clone] - Jennifer's for user manual development	48	9	Jennifer Davis	Tutorial datasets	T-Cat	Jan 26
US37 data test	15	1	Wesley Katsche	Jan 26		
test dose mini kit	8	0	Wesley Katsche	Jan 26		
Testing stratigdm files	3	1	Wesley Katsche	Jan 26		
Testing File Uploads	15	1	Chris Coveley (personal spend)	test project	T-Cat	Jan 24
US37 experiment - testing for export	15	5	Wesley Katsche	Jan 22		

My Labels  
test (1)

My Projects  
All experiments (1334)  
My experiments (1334)  
Public experiments (1334)  
Test (1334)  
Tutorial datasets (1334)  
Experiments for training, June (1334)  
Tutorial datasets (1334)  
Tutorial datasets (1334)

Experiment Analysis



Experiment Details

Experiment [exp\_2004\_APL\_Dataset] Labels: None Primary Researcher: Wesley Katsche Public: No

Back to Inbox

My Working Illustration

Getting Started!

Experiment Details

Situations

Name

Histogram overlay example with four APL patient samples

Heatmap like in Figure 1, with two APL patient samples

Wesley's Working Illustration

Attachments

File Name	Date	Uploaded By	Description
Download File [exp_2004_APL_Dataset]			
File Name	Sample Name	Experiment Variants	
APL_PDL_PL_Panel1_Asc_details	1.02) ELN 204 PDL	PL_APL_PDL	
APL_PDL_PL_Panel2_Asc_details	1.08) ELN 204 PDL	PL_APL_PDL	
APL_PDL_PL_Panel3_Asc_details	1.14) ELN 204 PDL	PL_APL_PDL	
APL_PDL_G-CSF_Panel1_Asc_details	1.04) ELN 204 G-CSF	G-CSF_APL_PDL	

Full Access

Add Full Access User

Share with Everyone

Did you know?

You can print/save your Illustrations to:

# Cytobank is available in multiple forms

- Cytobank Community Version– [www.cytobank.org](http://www.cytobank.org)
  - *Hub for the Cytobank community & associated resources*
  - *Users can login to manage and analyze their data*
  - *Supported through vendor partnerships*
- Hosted Cytobank – e.g. [labx.cytobank.org](http://labx.cytobank.org)
  - *Cytobank version hosted, backed up, and maintained for lab or company X*
  - *Designated administrators regulate access and logins*
  - **Dedicated Compute Resources**
  - **Access to premium modules and functionality (e.g. SPADE)**

*Maintenance, updates and support provided by Cytobank Inc.  
([www.cytobankinc.com](http://www.cytobankinc.com))*



# Upload. Analyze. Share. Anywhere.



***A web-enabled device lets you get to your data from anywhere in the world ...***

User registered on [www.cytobank.org](http://www.cytobank.org)

See <http://blog.cytobank.org/category/user-stories/>

# What Researchers are Doing with Cytobank

- Search on “Cytobank” in Google Scholar :

- *Specific cellular signal-transduction responses to in vivo combination therapy with ATRA, valproic acid and theophylline in acute myeloid leukemia*

- Skavland et al (Norway) – Feb. 2011 Nature

**Phosflow**

- *CD137 stimulation enhances the antilymphoma activity of anti-CD20 antibodies*

- Kohrt et al (Stanford) – Oct 2011 Nature

- *Cell-to-cell communication regulates PI3K-AKT Pathway Activity in Cell Populations*

- Yuan et al (Harvard) – Jan 2011 Current Biology

**Biochemistry**

- *Oxidative Stress Induces Reactivation of Kaposi's Sarcoma-Associated Herpesvirus in T-cell Primary Effusion Lymphoma Cells*

- Li et al (University of California) – Journal of Virology

**Cancer**

- *Poor cytokine-induced phosphorylation in chronic myeloid leukemia patients at diagnosis is effectively reversed by tyrosine kinase inhibitors*

- Jalkanen et al (Finland) – Sep 2010 Experimental Hematology

**Apoptosis**

- *Computational methods for large-scale data management and analyses*

- Schumacher et al (University of California) – Sep 2010 Nature Reviews Genetics

**Informatics**

Flow Cytometry  
Examples  
(Lymphoma, Drug  
Discovery, Mass  
Cytometry)

Cytobank: A cloud-  
computing platform for  
Cytometry

Annotations using NCBO  
BioPortal

Cytobank Reports  
A new way to publish data



Flow Cytometry  
Examples  
(Lymphoma, Drug  
Discovery, Mass  
Cytometry)

Cytobank: A cloud-  
computing platform for  
Cytometry

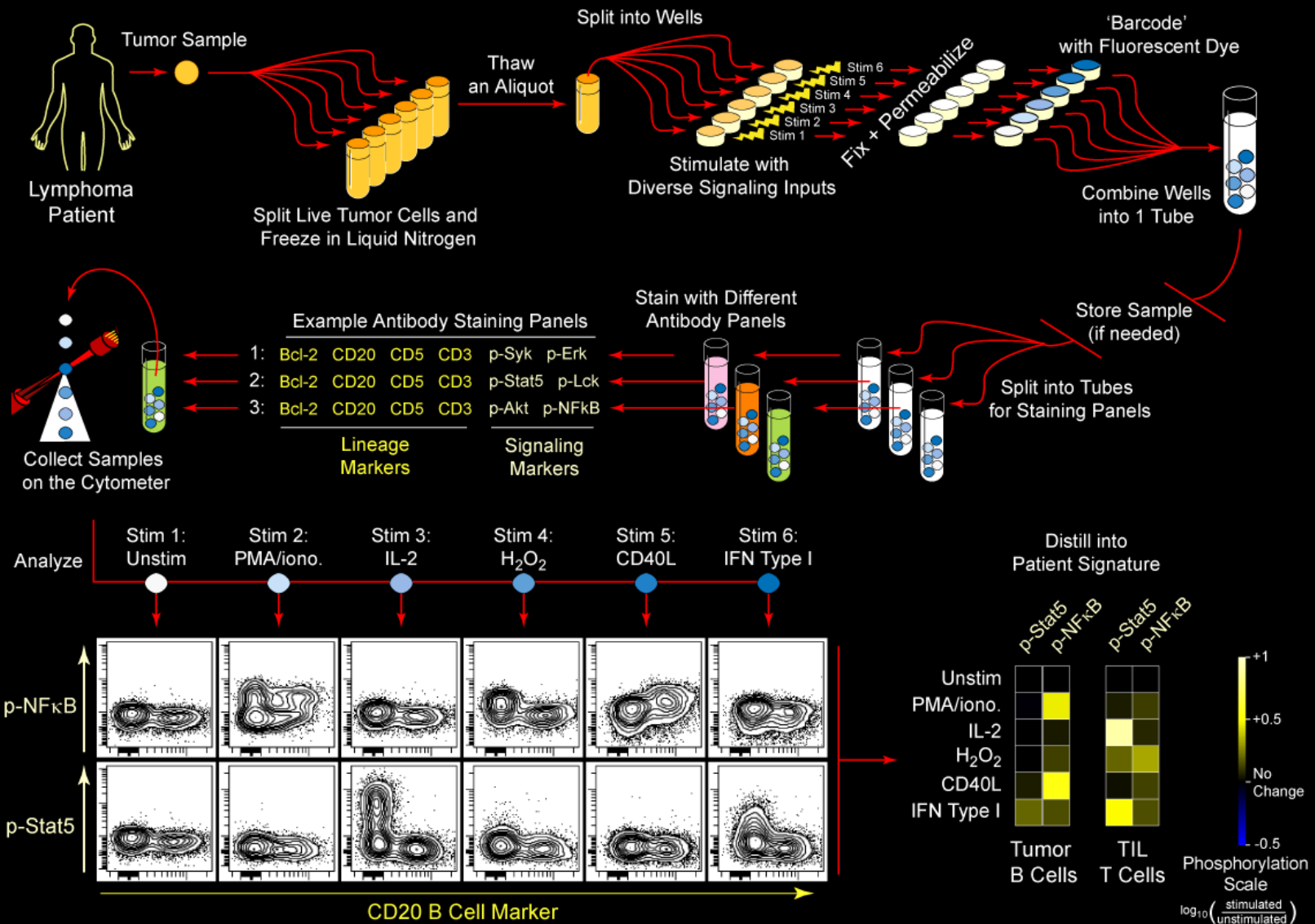
Annotations using NCBO  
BioPortal

Cytobank Reports  
A new way to publish data

## References of Examples

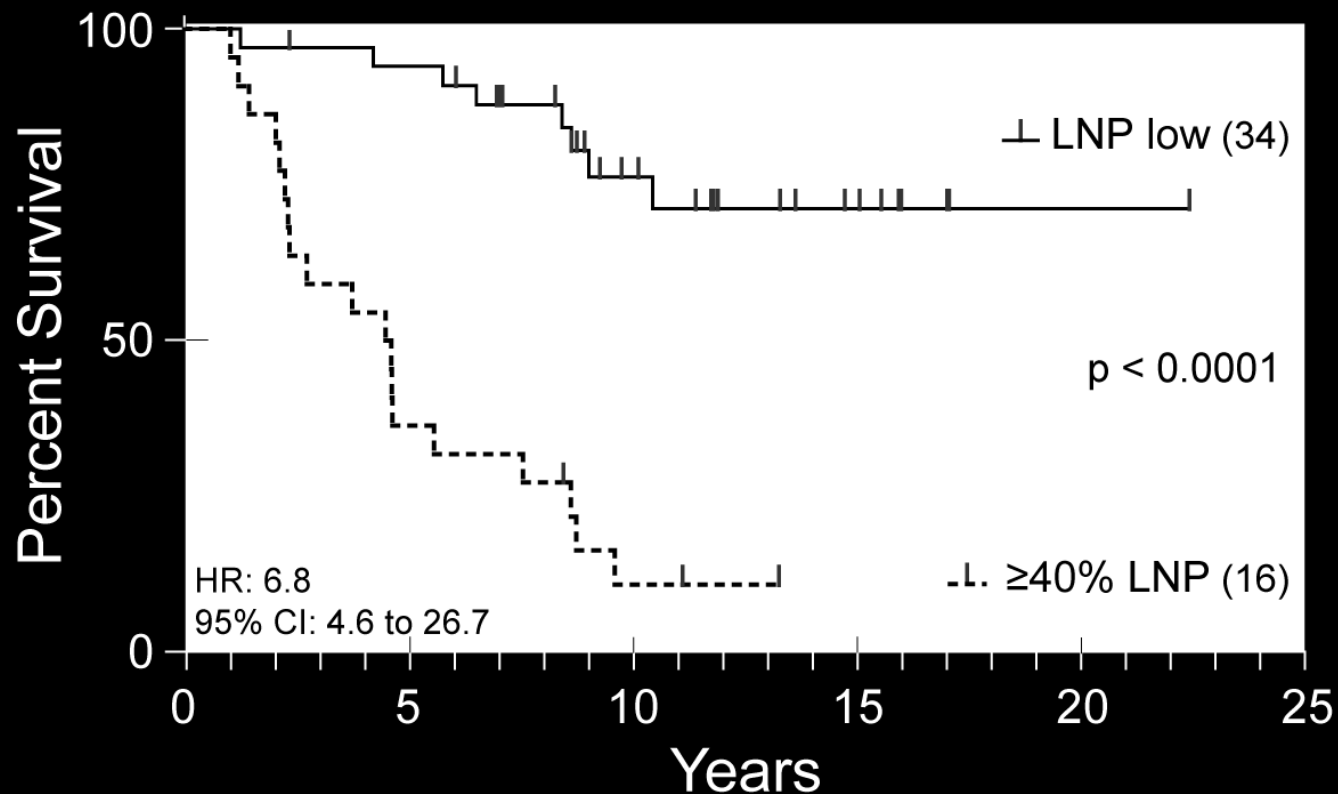
- Lymphoma
  - Irish et. al PNAS 2010
- Drug Discovery
  - Krutzik et. al Nature Chemical Biology 2008
- Mass Cytometry
  - Bendall et. al Science 2011

# Mapping Signaling in Every Cell using Flow Cytometry



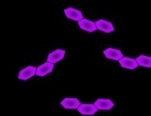
# The Subset Was Termed “Lymphoma Negative Prognostic” (LNP) Cells Because They Are Found in Patients with Poor Clinical Outcomes

Every 1% LNP cells in the tumor at diagnosis increased the patient's risk of death in the next year by 2.5%

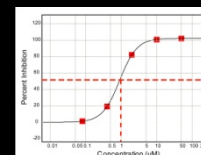


# Primary Cell Screening using Phosphoflow

4 natural products + 4 commercial inhibitors



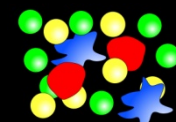
Titrate 6 concentrations of each compound



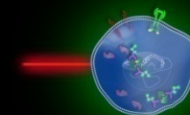
Stimulate with IFN $\gamma$ , IL-4, IL-6, IL-7, IL-10, IL-15



Analyze B cells, CD4 $^{+}$  and CD4 $^{-}$  T cells,  
CD11b $^{hi}$  neutrophils, CD11b $^{int}$  macrophages



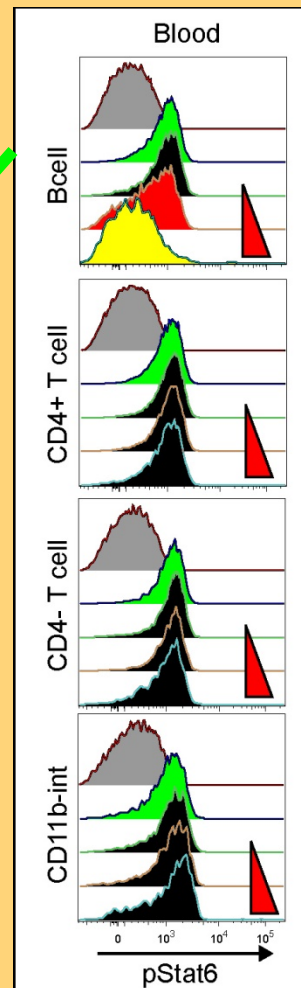
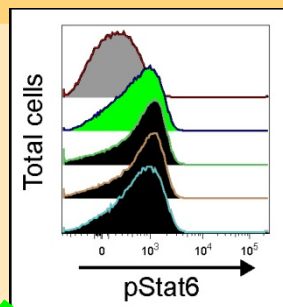
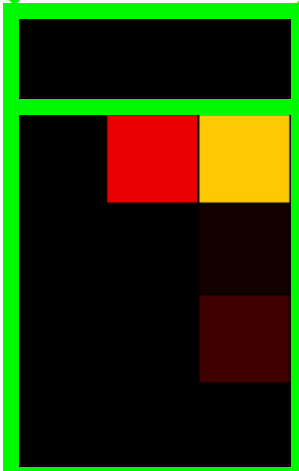
Measure Stat1, Stat3, Stat5, Stat6 phosphorylation



Phospho  
Flow

pStat6

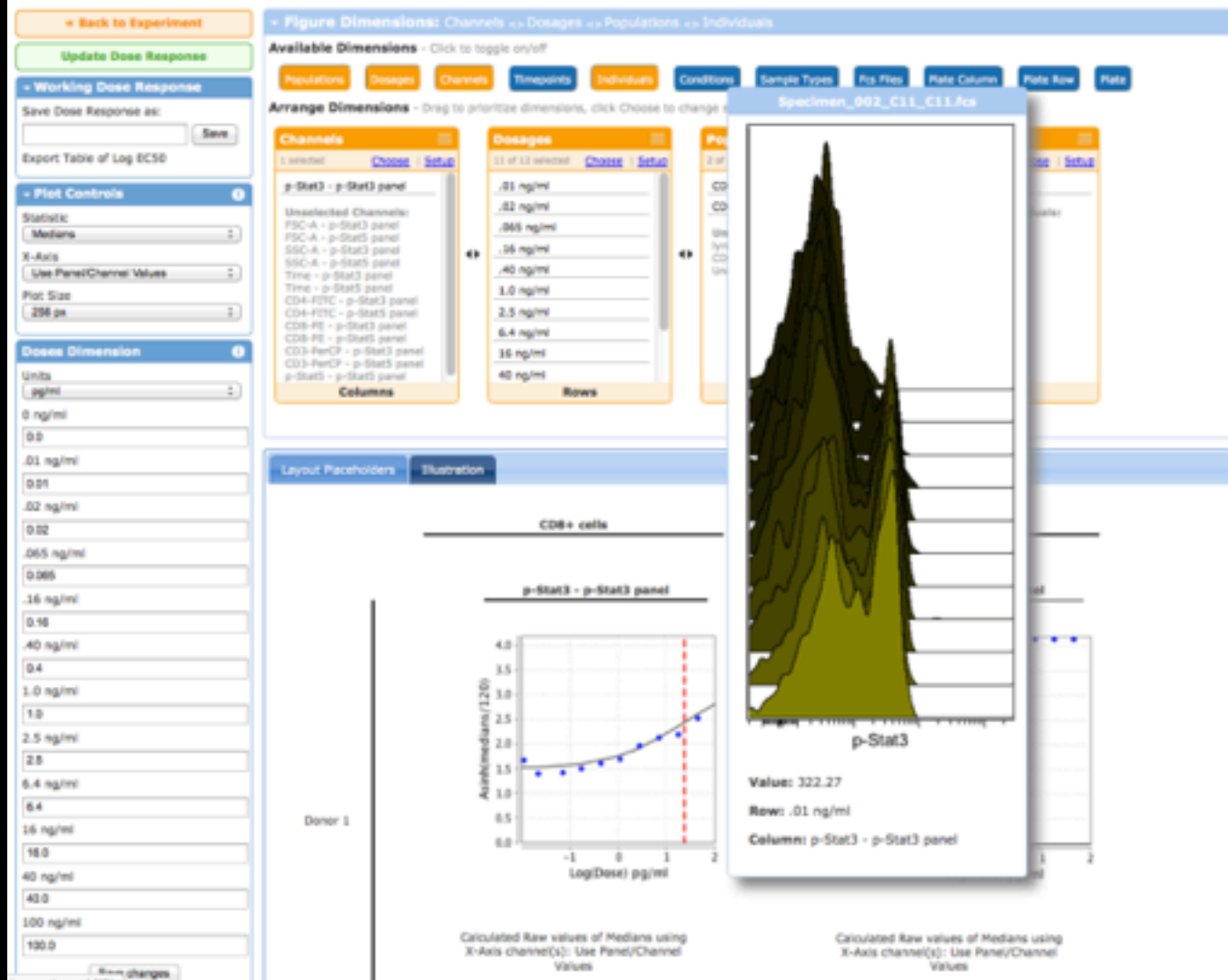
Total cells	
B cell	
CD4+ T	
CD4- T	
CD11b-int	



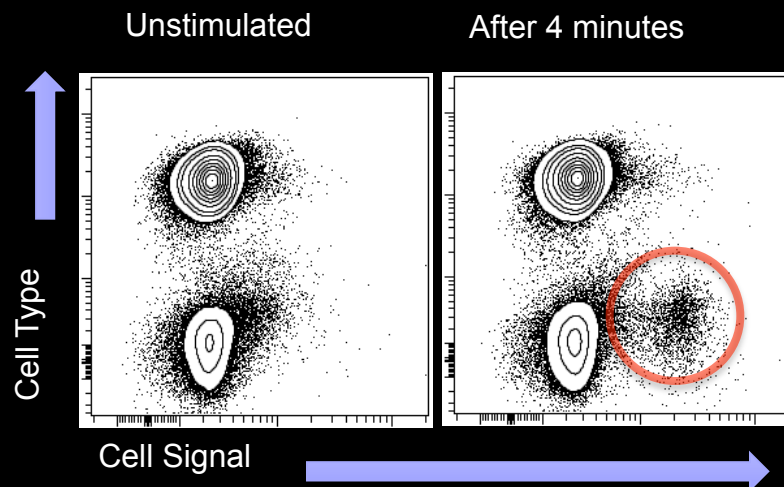
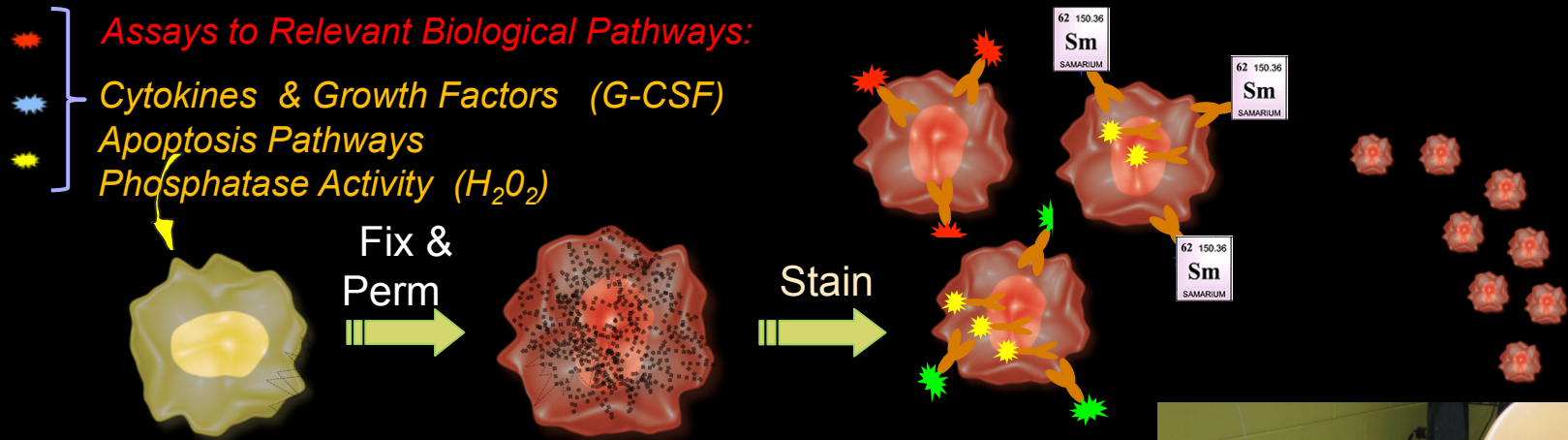
## Blood

- Total cell measure from the blood indicates no drug effect.
- Looking at individual cell types shows B cells are potentially inhibited

# Drug Discovery Assays (linked to underlying data)



# Mass Cytometry: 30+ parameters & no compensation



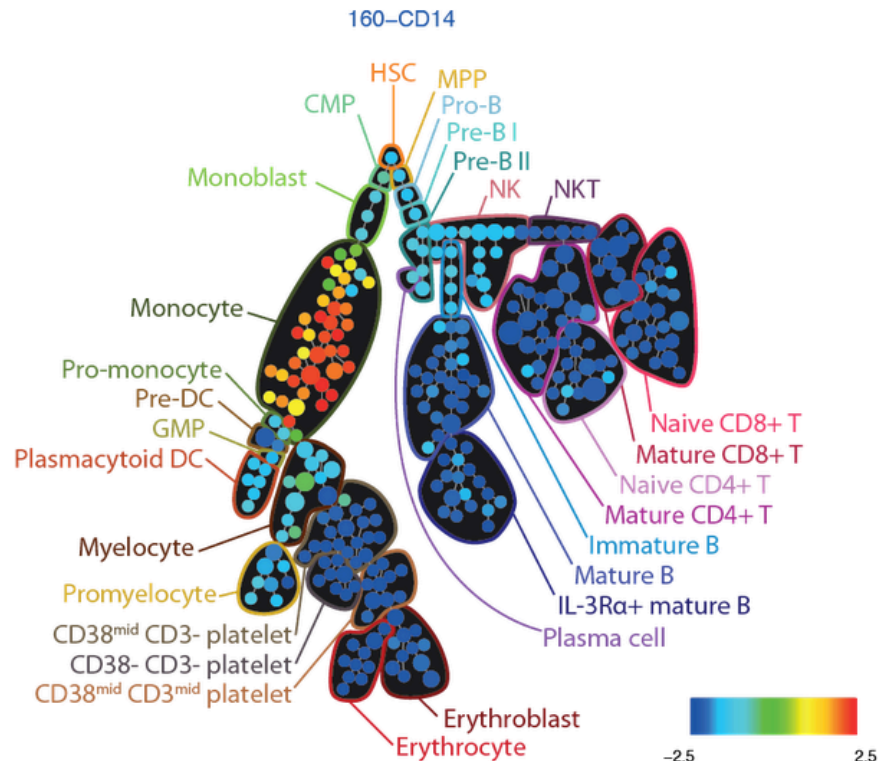


# SPADE allows for analysis of high dimensional cytometry

Figure S4b (Surface-only tube)

Expression of immunophenotype surface markers overlaid onto the SPADE plots of healthy human bone marrow: The expression of an additional 18 surface markers from the 31 surface marker analysis of the same sample was overlaid on the SPADE plot from 13 core surface markers. These 18 surface markers were not used in the SPADE plot and their localized expression is based solely on the shared expression patterns of the 13 core surface markers.

Note: This figure was created in a flow cytometry analysis package called SPADE (not yet published) using data gated and exported from Cytobank. The link below will bring you to an illustration from which you can export the exact gated data used in this figure.



[View Gated Data In Cytobank](#)

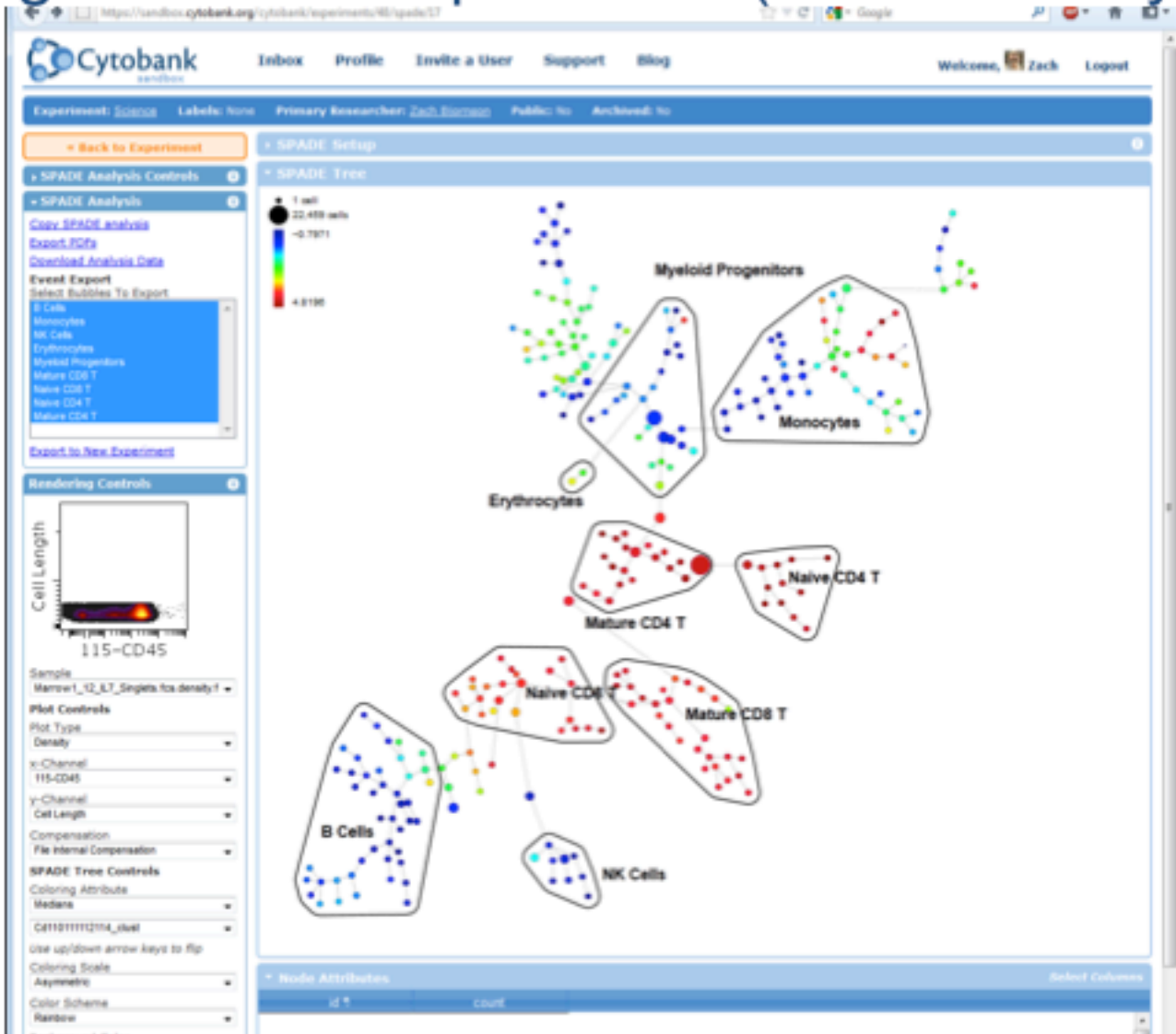
[Back to Top](#)

[www.cytobank.org/nolanlab](http://www.cytobank.org/nolanlab)

Bendall et. al Science 2011

Qiu et al Nature Biotech 2011

## Large Scale Computations (delivered to your browser)



## Cytobank Premium

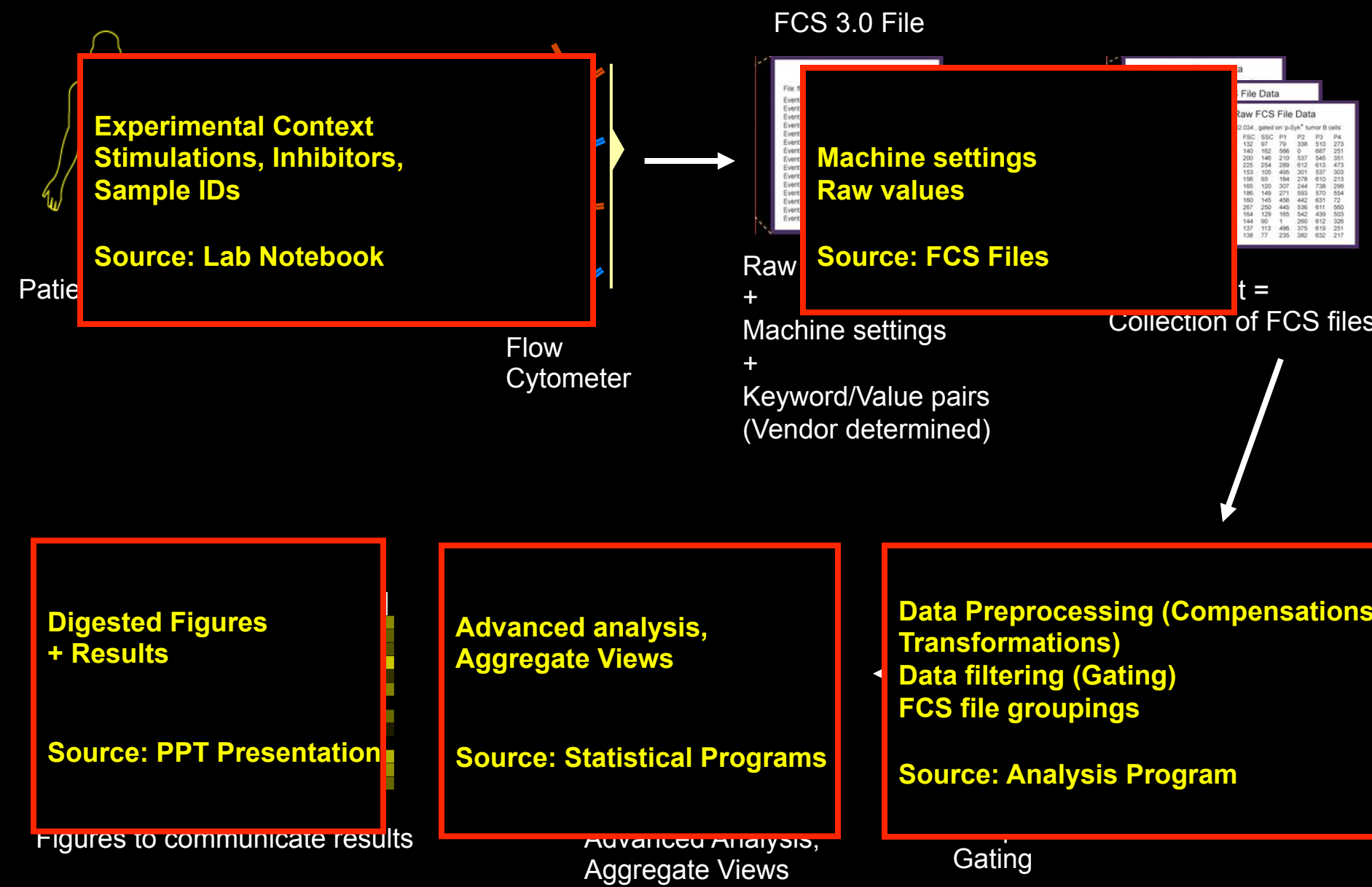
Flow Cytometry  
Examples  
(Lymphoma, Drug  
Discovery, Mass  
Cytometry)

Cytobank: A cloud-  
computing platform for  
Cytometry

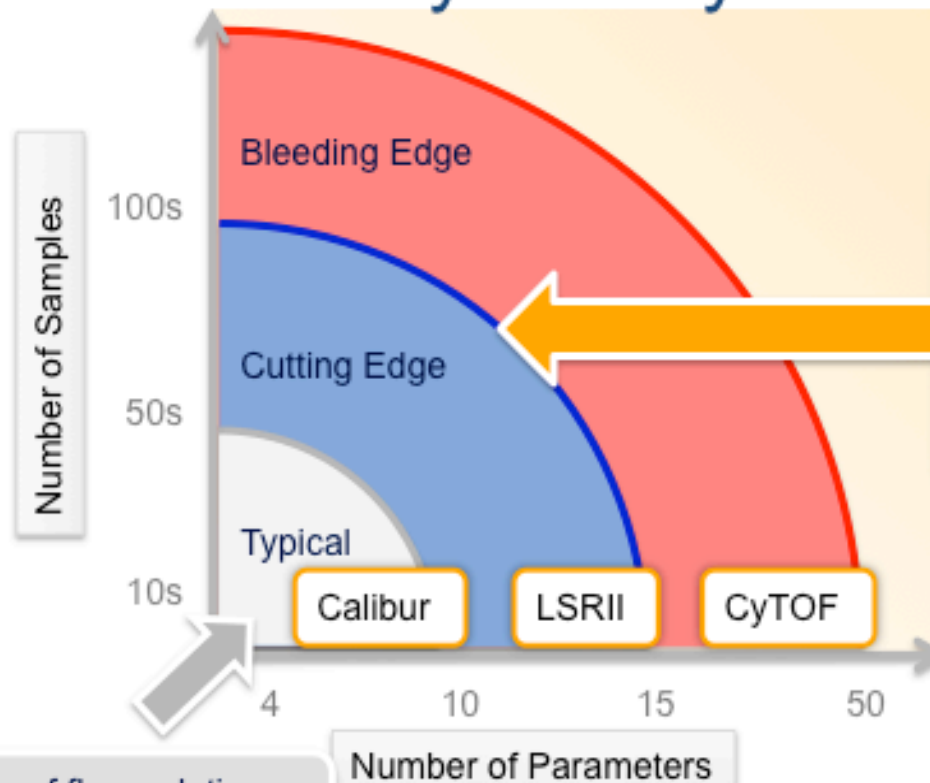
Annotations using NCBO  
BioPortal

Cytobank Reports  
A new way to publish data

# Analysis and annotation of flow cytometry data is fragmented



# Trends in Cytometry & Cell Analysis (I)



- Majority of flow solutions built for this space

## Future of Flow

- New solution required
- Organize/manage experiments
- Communicate & collaborate
- Annotations & analyses linked
- Novel analyses & visualizations
- Scalable compute resources
- Platform to build on top of

# Cytobank is for *managing, sharing & analyzing* flow experiments over the web



Experiment =  
Collection of FCS files



Experiment Analysis



Experiment Details

# Create a new experiment and upload FCS files



Inbox Profile Admin Help

Create a New Experiment

Getting Started!

Credits Available: 3/20  
[Request More Credits](#)

All Experiments (32)

Select Label:

Filter Experiments 0

All experiments (32)

My experiments (30)

Public experiments (2)

Trash (14)

Manage Projects

Manage My Projects

Views (2)

My Labels

Instructions for Banking

Bank New Experiment

Experiment Name

Project

Primary Researcher

Upload Experiment Files

Allow PG/P

Files:

Filename	FCS Version	Upload?
panel001 - 1.01) ELM unstim	FCS2.0	<input checked="" type="checkbox"/>
panel002 - 1.02) ELM 20n Flt3L	FCS2.0	<input checked="" type="checkbox"/>
panel003 - 1.03) ELM 20n GMCSF	FCS2.0	<input checked="" type="checkbox"/>
panel004 - 1.04) ELM 20n G-CSF	FCS2.0	<input checked="" type="checkbox"/>
panel005 - 1.05) ELM 20n IL-3	FCS2.0	<input checked="" type="checkbox"/>
panel006 - 1.06) ELM 20n IFNg	FCS2.0	<input checked="" type="checkbox"/>
panel007 - 2.01) Asta unstim	FCS2.0	<input checked="" type="checkbox"/>
panel008 - 2.02) Asta 20n Flt3L	FCS2.0	<input checked="" type="checkbox"/>
panel009 - 2.03) Asta 20n GMCSF	FCS2.0	<input checked="" type="checkbox"/>
panel010 - 2.04) Asta 20n G-CSF	FCS2.0	<input checked="" type="checkbox"/>
panel011 - 2.05) Asta 20n IL-3	FCS2.0	<input checked="" type="checkbox"/>
panel012 - 2.06) Asta 20n IFNg	FCS2.0	<input checked="" type="checkbox"/>
panel013 - 3.01) Grim unstim	FCS2.0	<input checked="" type="checkbox"/>
panel014 - 3.02) Grim 20n Flt3L	FCS2.0	<input checked="" type="checkbox"/>
panel015 - 3.03) Grim 20n GMCSF	FCS2.0	<input checked="" type="checkbox"/>
panel016 - 3.04) Grim 20n G-CSF	FCS2.0	<input checked="" type="checkbox"/>
panel017 - 3.05) Grim 20n IL-3	FCS2.0	<input checked="" type="checkbox"/>

Upload Progress:

Uploading to <http://staging02.cytobank.org/cytobank/uploadServlet>

Launch web browser &  
Login to Cytobank

Create a new Experiment

Upload FCS Files

# Organize information around samples using Experimental Variables

Variable Types	Experiment Variable Values
Individual Patients:	<ul style="list-style-type: none"> <li>AML Patient 4 (P04)</li> <li>AML Patient 5 (P05)</li> </ul>
Phospho-proteins:	<ul style="list-style-type: none"> <li>Phosphorylated Stat3 (p-Stat3)</li> <li>Phosphorylated Stat5 (p-Stat5)</li> </ul>
Conditions:	<ul style="list-style-type: none"> <li>Unstimulated</li> <li>GM-CSF</li> <li>IL-3</li> <li>G-CSF</li> <li>IFN<math>\gamma</math></li> <li>Flt3 Ligand (FL)</li> </ul>
Cell Populations:	<ul style="list-style-type: none"> <li>Intact Cells</li> </ul>

Figure Dimensions: Conditions > Channels > Populations > Individuals

Available Dimensions - Click to toggle

Conditions Channels Populations Dosages Timepoints Individuals Sample Types Fcs Files

Arrange Dimensions - Drag Left/Right to Reorder

Conditions Edit

Columns

Channels Edit

Rows

Populations Edit

Table 1

Individuals Edit

Table 2

## Select a Condition

All Conditions

Unstim FL GM-CSF G-CSF IL-3 IFN $\gamma$

All Conditions

Untagged

Filter

Move to... Move file(s)

Unstim Tagged Files

1.01) ELM unstim  
AML-P01 Unstim Panel1.Fcs

1.07) ELM unstim  
AML-P01 Unstim Panel2.Fcs

1.13) ELM unstim  
AML-P01 Unstim Panel3.Fcs

2.01) Acla unstim  
AML-P02 Unstim Panel1.Fcs

2.07) Acla unstim  
AML-P02 Unstim Panel2.Fcs

FL Tagged Files

1.02) ELM 20n Flt3L  
AML-P01 FL Panel1.Fcs

1.08) ELM 20n Flt3L  
AML-P01 FL Panel2.Fcs

1.14) ELM 20n Flt3L  
AML-P01 FL Panel3.Fcs

2.02) Acla 20n Flt3L  
AML-P02 FL Panel1.Fcs

2.08) Acla 20n Flt3L  
AML-P02 FL Panel2.Fcs

Assign samples to experimental variables

Draw Gates

Save & Return Select Rectangle Ellipse Polygon Quadrant Split Merge Reset

File AML-P01 IFN $\gamma$  Panel2.Fcs - 5.52 ELM 20n IFN $\gamma$

View

Active Population: Live Cells

Active Compensation: File Compensation

Plot Settings

Populations Manage View

Y: Stat5-Au647 Log

X: Stat1-Au647 Log

Population signaling ... was created. Double-click the gate to select it.

List of gates: Live Cells, signaling gate

Selected gate: Name: signaling gate

Global Stencil Alone Lock

Remove Check Gate

Identify populations of interest (gating)

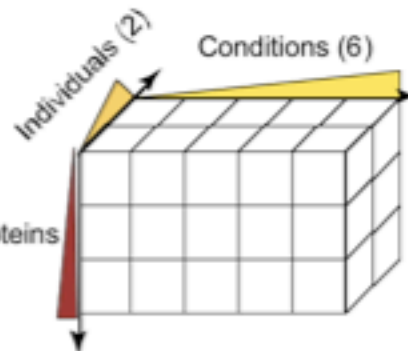
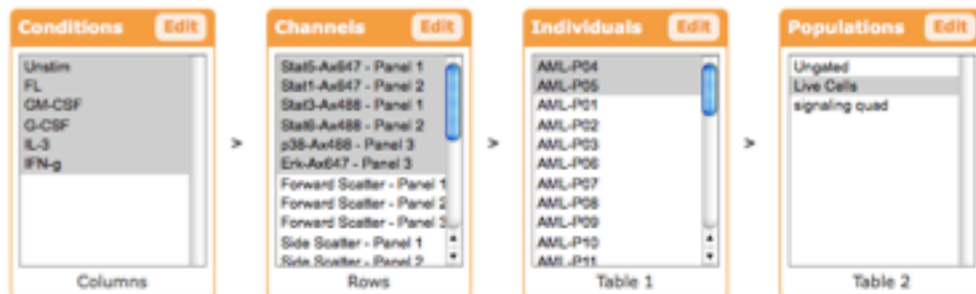


# Use Experimental Variables to create and pivot figures

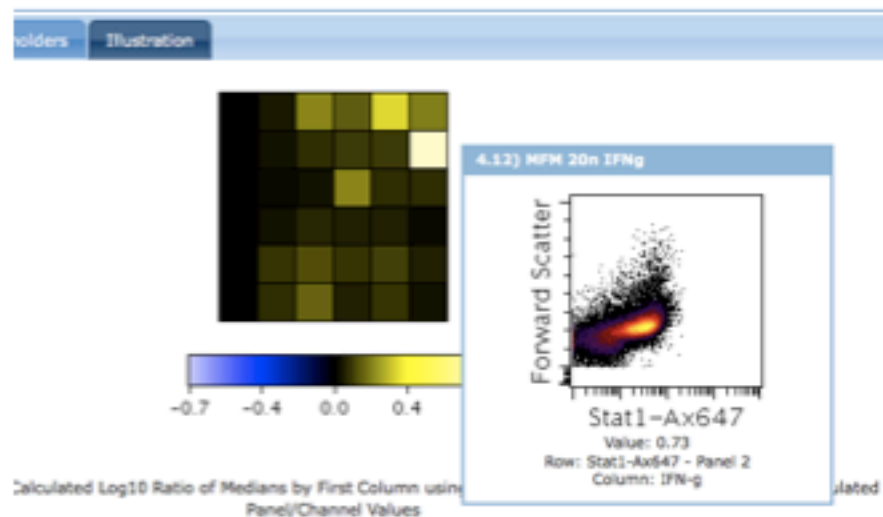
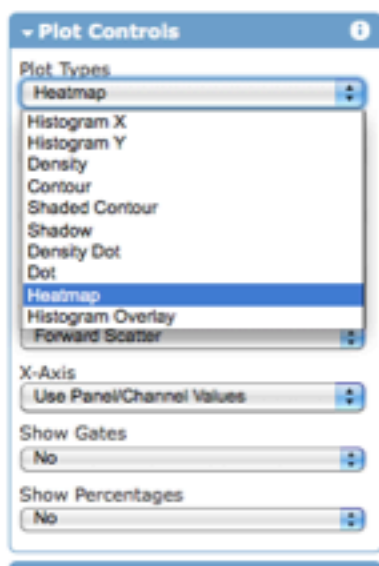
Available Dimensions - Click to toggle

Conditions Channels Populations Dosages Timepoints Individuals Sample Types Fcs Files

Arrange Dimensions - Drag Left/Right to Reorder



Arrange figure using  
Experimental Variables



Plot controls to specify  
plot type and display

# Share results and analyses with collaborators and the community

[Irish 2004 AML Dataset](#) > Nikesh's Working Illustration  
Illustration Author: Nikesh Kotecha  
Primary Researcher: Jonathan Irish  
Created: Tue Mar 30 11:03:00 -0500 2010 | Updated: Tue Mar 30 11:16:00 -0500 2010

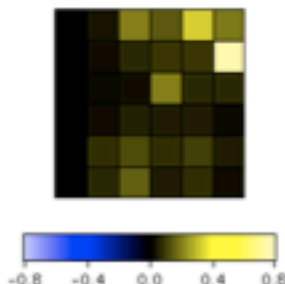


[Display as PDF](#)

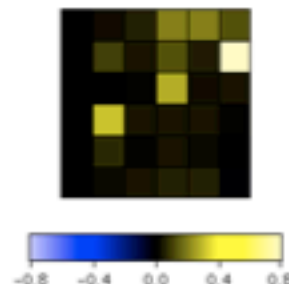
[Jump to Gating Hierarchy](#)

[Share Illustration](#)

AML-P04, Live Cells



AML-P14, Live Cells



Individual and project level  
sharing for collaborators



[Inbox](#) [Profile](#) [Admin](#) [Help](#)

[Create a New Experiment](#)

[Getting Started!](#)

Credits Available: 4/79

[Request More Credits](#)

[Filter Experiments](#)

[All experiments](#) (131)

[My experiments](#) (92)

[Public experiments](#) (51)

[Trash](#) (17)

[Manage Projects](#)

All Experiments (131)

Select Label:  [Apply Label](#) [Delete Experiments](#)

Experiment Name

☐ P Healthy PBMC Mini Profile - Comp Tester

☐ P Irish 2004 AML Dataset

Make experiments public  
to share with community

## Cytobank Benefits

- Easy way to manage, share and backup your fcs files
- Share analyses and experiments with colleagues and collaborators
- Managers and PIs – eliminate fear about “losing your data” (when people leave their lab)
- Core facilities/Service labs can use Cytobank for value added services
  - *Data management and backup*
  - *Remote analysis and support*
- Capture experiment information while creating figures
  - *“What did I measure in that experiment from 6 months ago again?”*

# The Cytobank Platform

Data Collection Hooks  
Integration with Core  
Facilities  
(e.g. Stanford)

Translational  
Assays  
Drug Screening  
(Clinical) Reports

High Dimensional  
Cytometry,  
Mass Cytometry,  
(SPADE, Large Scale  
Computations)

Community Resources  
(BD FACSSelect)  
Public Cytometry  
Collections



Accessible from any web-enabled device  
anywhere

Flow Cytometry  
Examples  
(Lymphoma, Drug  
Discovery, Mass  
Cytometry)

Cytobank: A cloud-  
computing platform for  
Cytometry

Annotations using NCBO  
BioPortal

Cytobank Reports  
A new way to publish data

## NCBO: Key activities - <http://www.bioontology.org/>

- **We create and maintain a library of biomedical ontologies.**
- **We build tools and Web services to enable the use of ontologies and their derivatives.**
- **We collaborate with scientific communities that develop and use ontologies.**

Ontology  
Services

Views



- Download
- Traverse
- Search
- Comment

## Mapping Services



- Create
- Download
- Upload

## Widgets



- Tree-view
- Auto-complete
- Graph-view

## Annotation



## Term recognition

## Data Access



Fetch “data”  
annotated with a  
given term

<http://bioportal.bioontology.org>



Jump To:

[Legend](#)

- Malignant **melanoma** (synonym)
- Amelanotic **melanoma** (preferred name)
- Excision of **melanoma** (preferred name)
- Melanoma** in situ (preferred name)
- Melanoma** vaccine (preferred name)

Expression, Expression of bladder, bladder, smooth, bladder muscle, muscle, smooth muscle, cells, mechanical, mechanical stimulation, stimulation, Chronic, results, bladder overdistension, associated, associated with, with, loss, genes, altered





# Integration with NCBO BioPortal

Experiment: [BCR Signaling in Follicular Lymphoma \(Blood 2006\)](#)

Labels: None

Primary Researcher: [Jennifer Davis](#)

Public: No

Archived: No

« Back to Inbox

My Working Illustration »

Actions

Experiments

[Edit Experiment Details](#)

[Reset Experiment](#)

Reagents

[Edit reagents](#)

FCS Files

[Download FCS Files](#)

[Upload More FCS Files](#)

Cloning/Copying

[Clone Experiment](#)


Ontology Annotations


Search for Ontology Annotations


Experiment

Sharing Permissions

Full Access Users

 [Jennifer Davis](#) [PR]

 [Angela Landrigan](#) [x]

 [Nikesh Kotecha](#) [x]

[Invite a new user](#)

Share with a User (Full Access)

Getting Started: Experiment Details

60-Second Overview

This Experiment Details Page contains information about the experiment listed above.

[Analyze in Cytobank](#)

Ontology Annotation Search

Select Text to Annotate:

☒ Experiment Name

☒ Purpose

☒ Experiment Variables

☒ Channel Names

BCR Signaling in Follicular Lymphoma (Blood 2006) An example of BCR signaling in human follicular lymphoma samples. When looking at the B cells within these human follicular lymphoma (FL) tumors, BCL2 marks the tumor cells (it is overexpressed due to the t14;19 translocation, a signature of FL). So in the illustrations, BCL2+ B cells are lymphoma cells, whereas BCL2- B cells are non-malignant B cells within the tumor. You can see that their signaling differs in response to BCR engagement. Also present within the tumor are T cells, which provide another control for these experiments comparing the kinetics of BCR signaling in tumor and non-malignant B cells within the same tumor sample. a-BCR, a-BCR + H2O2, 0m, 4m, 16m, 60m, 90m, FL Patient 10, FL Patient 9 bcl2-pe, cd20c-percpcy5.5, fl1-a, forward scatter, lambda-fitc, p-erk12-ax647, side scatter, p-syk-ax647

Submit



# Ontologies (Initial Set) Used in Cytobank

## Keywords

- Gene Ontology, Gene Ontology Extension,
- ICD10, ICD10CM,
- Logical Observations Identifier Names and Codes (LNC),
- MedDRA (MDR),
- NCI Thesaurus (NCIt),
- RadLex (RID),
- SNOMED Clinical Terms (SNOMEDCT),
- Medical Subject Headings (MSH),
- Online Mendelian Inheritance in Man (OMIM)
- Molecule Role (INOH Protein name/family name ontology) (IMR)

# Add Keywords To a Cytobank Experiment

tumor and non-malignant B cells within the same tumor sample. a-bcr, a-bcr + H2O2, 0m, 4m, 16m, 60m, 90m, FL Patient 10, FL Patient 9 bcl2-pe, cd20c-percp5.5, fll-a, forward scatter, lambda-fitc, p-erk12-ax647, side scatter, p-syk-ax647"

## Keywords (74)

(lymphatic tissue carcinoma) or (lymphoma)	cell	lymphoma cell count	provide (2)
90m	cells (3)	lymphoma cells (2)	response (3)
answer	chromosomal translocation	lymphoma, follicular	routine signature
anterior	process	malignant lymphoma	sample
apoptosis regulator bcl-2	control (4)	mass	scientific control
b-cell cll/lymphoma 2	control group	mouse lymphoma	signal transduction
b-lymphocyte	due to (2)	murine t-	signaling (2)
b-lymphocytes	example	lymphocytes	signature
bcl2	follicle	murine tumor cells	specimen
bcl2 gene	follicular	neoplasm (3)	t-lymphocyte
bcl2_human	follicular lymphoma (3)	neoplasms	t-lymphocytes
bcl2_mouse	forward (2)	non-malignant	translocation
bcl2_rat	h2o2	patient (4)	tumor (3)
bcr (4)	homo sapiens (2)	patients	tumor cells
bcr gene	human (2)	pharmacokinetics	tumor cells, uncertain whether benign or
bcr protein	human - origin	positive	malignant
biospecimen	humans	present (3)	tumor tissue
biospecimen core resource	hydrogen peroxide	presentation	veterinary patient
breakpoint cluster region	kinetics (2)	prevention & control	
	lymphoma (5)	protein domain	

## Annotations (107)

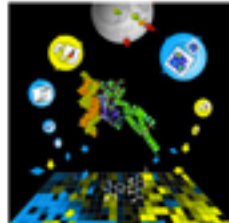
Preferred Name	Term ID	Ontology	Semantic Type(s)
Cells	<a href="#">D002477</a>	<a href="#">Medical Subject Headings</a>	T025: Cell
Cells	<a href="#">MTHU001933</a>	<a href="#">Logical Observation Identifier Names and Codes</a>	T025: Cell
Cells	<a href="#">LP14738-6</a>	<a href="#">Logical Observation Identifier Names and Codes</a>	T025: Cell
Cell	<a href="#">Cell</a>	<a href="#">NCI Thesaurus</a>	T999: NCBO BioPortal concept
tumor	<a href="#">npo:NPO_1573</a>	<a href="#">NanoParticle Ontology</a>	T999: NCBO BioPortal concept
Tumor	<a href="#">LP7664-8</a>	<a href="#">Logical Observation Identifier Names and Codes</a>	T191: Neoplastic Process

Flow Cytometry  
Examples  
(Lymphoma, Drug  
Discovery, Mass  
Cytometry)

Cytobank: A cloud-  
computing platform for  
Cytometry

Annotations using NCBO  
BioPortal

Cytobank Reports  
A new way to publish data



## Nolan Lab

**Principal Investigator:** Garry P. Nolan, Ph.D.  
**Affiliation:** Stanford University, Stanford, CA  
**Curator:** Garry Nolan  
**Website:** <http://www.stanford.edu/group/nolan>  
**Research:** cancer, cell signaling, immunology, leukemia, intracellular cytometry, mass cytometry, systems biology

**Synopsis** We analyze cell signaling directly by next-generation mass cytometry and traditional flow cytometry, focusing on following multiple phosphoproteins in complex populations of primary cells such as mouse cells and human clinical samples. Using mass cytometry, up to 34 simultaneous protein parameters can be measured in single cells including multiple kinases, phosphoproteins, cell cycle proteins, and other parameters, enabling resolution of cellular activation states.

We are using these techniques to study healthy biochemical signaling in the immune system and dysfunctional signaling in hematological malignancies including AML, ALL, JMML, MDS, and follicular lymphoma. We have used this approach to distinguish predictive patterns of intracellular signaling to classify patient responses to chemotherapies and to determine how their signaling systems are altered in disease states. We are also using the technique for drug screening in primary cells to truly select for drugs with efficacies in certain cell subsets but not others.

Autoimmune diseases in which we have particular interest include rheumatoid arthritis and systemic lupus erythematosus. In these diseases, we focus on understanding how the immune system becomes dysregulated as disease comes and goes. We can measure and determine the cellular network states in multiple cell subsets. In cancer, we are working in follicular lymphoma as well as acute myelogenous leukemia where we can look at disease progression as a measure of changes in disease states correlated to particular genetic changes in the genome of human cancer cells. Also we have made determined efforts in understanding how the cancerous microenvironment modulates immune signaling.

### Published Experiments

Article	Journal	Date	Cytobank Report
<a href="#">Single-Cell Mass Cytometry of Differential Immune and Drug Responses Across a Human Hematopoietic Continuum</a> Bendall SC, Simonds EF, Qiu P, Amir ED, Krutzik PO, Finck R, Bruggner RV, Melamed R, Trejo A, Ornatsky OI, Balderas RS, Plevritis SK, Sachs K, Pe'er D, Tanner SD, Nolan GP	Science	May 2011	<a href="#">View Data</a>

Providing published data to the computational biology and cytometry communities



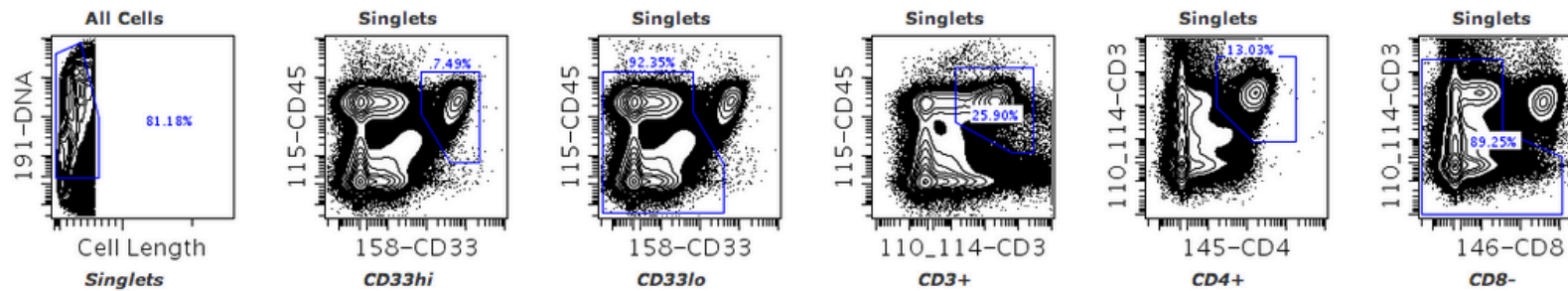
IL7 --> pStat5 in T cells

IL-7 is a canonical activator of T cell proliferation. Here, IL-7 mediated activation of pSTAT5 in T cells is shown as an example of the cell-type specific signaling responses that can be detected by minutes was detected using an antibody against STAT5 phosphorylated at the Y694 residue.

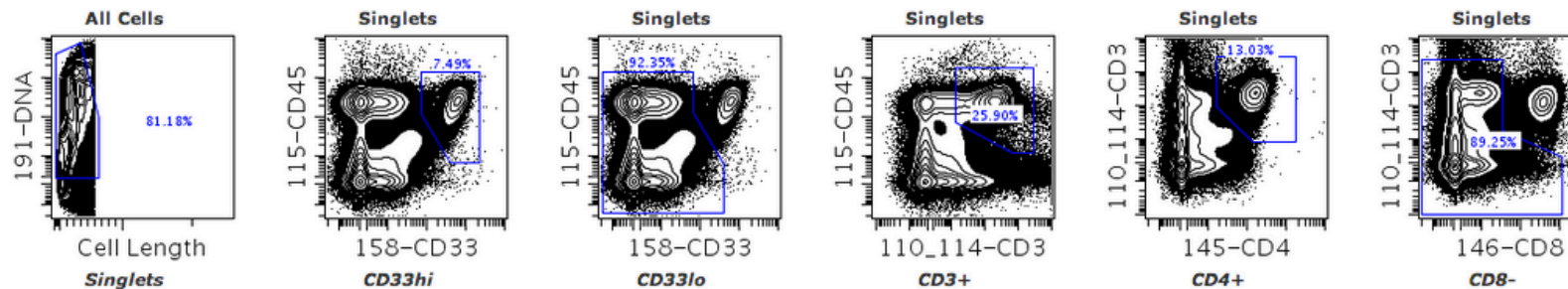
This same data is summarized in 4 squares of the heatmap highlighted in Figure 3B of the paper.

IL7 --> pStat5 in T cells (Gating Hierarchy)

Mature CD4+ T



Naive CD4+ T



## Targeting the Ideal Reagent Quickly with FACSelect

[illegible]

[www.cytobank.org/facselect](http://www.cytobank.org/facselect)



## CD3e (145-2C11)

## About CD3e (145-2C11)

## Conjugates Shown

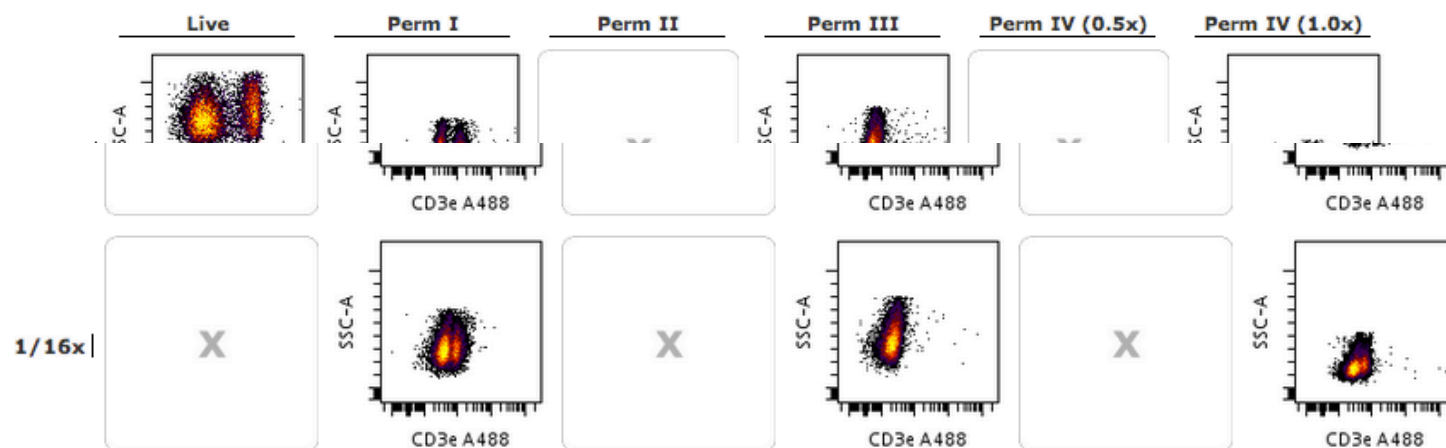


[Back to FACSelect](#)

**Protein Name:** CD3e  
**Clone:** 145-2C11  
**Isotype:** IgG1, k  
**Reactive species:** Mouse  
**Host species:** Armenian Hamster  
**Protocol:** [details](#)  
**Experiment Cell Source:** BALB/c Spleen  
**Cytometer used:** FACSCantoII

[Alexa Fluor@ 488](#) [BD]  
[FITC](#) [BD]  
[PE](#) [BD]  
[PerCP](#) [BD]  
[PerCP-Cy5.5](#) [BD]  
[PE-Cy7](#) [BD]  
[APC](#) [BD]  
[APC-Cy7](#) [BD]

## CD3e A488



Calculated Raw values of statistic using X-Axis channel(s): Use  
 Panel/Channel Values

	Live	Perm I	Perm II	Perm III	Perm IV (0.5x)	Perm IV (1.0x)
1x	4.92	2.07	X	1.18	X	1.66
1/4x	X	2.14	X	1.25	X	1.82
				1.36	X	1.46

[View in Cytobank](#)

[Gating Hierarchy](#)

[Back to Top](#)

## CD3e FITC

Live

Perm I

Perm II

Perm III

Perm IV (0.5x)

Perm IV (1.0x)



## Conclusions: Analysis & Presentation

**Annotate at the cytometer** – it will save time and help find files later

For figures based on large datasets or new statistics / analysis tools, always **show representative primary data**

Organization, experiment design, and **annotation** are critical

- computational analysis (e.g. SPADE)
- collaborations & long term projects
- sharing data with publications

# Cytobank Links/Emails

- Cytobank Documentation – <http://support.cytobank.org>
  - *Documentation, Tutorials and Walkthroughs for Cytobank*
  - *Actively updated. Send comments!!!*
- Cytobank Blog – <http://blog.cytobank.org>
- Support, questions, feedback – [helpdesk@cytobank.org](mailto:helpdesk@cytobank.org)
  - *Quickest way to ask any questions/get support on Cytobank*
  - *Can also fill out a support ticket via Help in Cytobank*
- Cytobank website – [www.cytobank.org](http://www.cytobank.org)
- Cytobank Inc. – [www.cytobankinc.com](http://www.cytobankinc.com)
- Referencing Cytobank:
  - *Kotecha et al. **Web based analysis and publication of flow cytometry experiments**. Current Protocols in Cytometry. July 2010.*

Thank you!

[nikesh@cytobank.org](mailto:nikesh@cytobank.org)

Extras

# Data Security & Privacy

- All connections to the Cytobank app through SSL / HTTPS
  - The same that financial institutions use
- Cytobank's servers are firewalled and hardened to restrict access and prevent attacks.
  - Only members of Cytobank's operations team have root access to the servers.
- All data on any Cytobank is *\*always\** private by default until ***you choose*** to share it with someone
- Each Cytobank installation is self-contained and independent from the others -- no accounts are shared and no data from one instance is visible in another.
  - Customer administrators can choose to require validation of all accounts before they can be used.

# Data Backup & Server Maintenance

- All servers are kept up to date with the latest security patches according to common industry practice
- Cytobank's servers are monitored 24x7.
  - Using services like AlertSite and Nagios
- Cytobank maintains local and remote backups of each user's data.
- All access to the servers and applications are logged.