



Intravital Microscopy, Flow Cytometry and Cell Sorting

7 – 13 July 2013 | Berlin, Germany

Cytometry Basics and New Developments

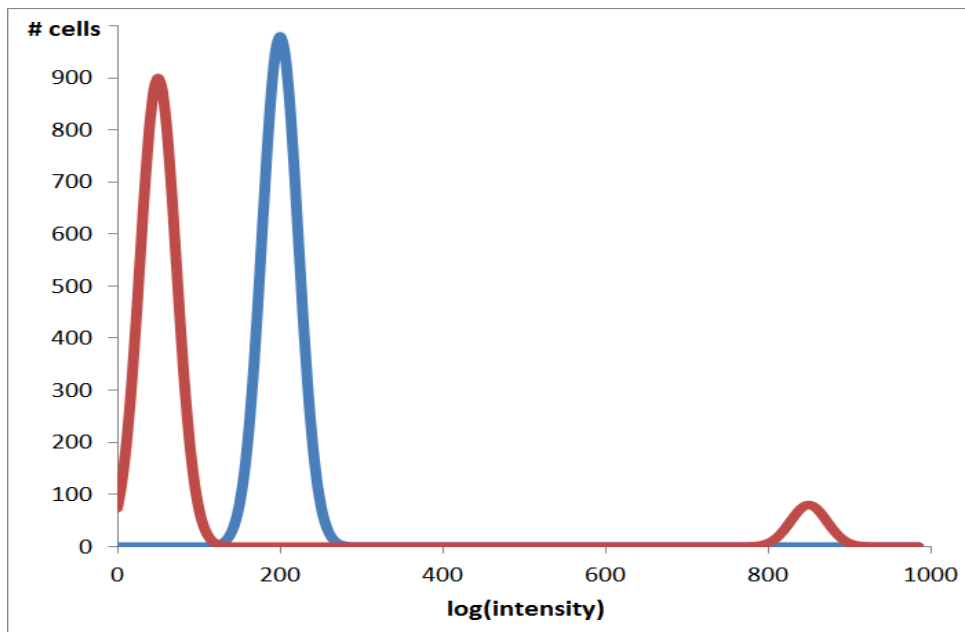
Diether Recktenwald

Desatoya LLC, Reno NV, USA

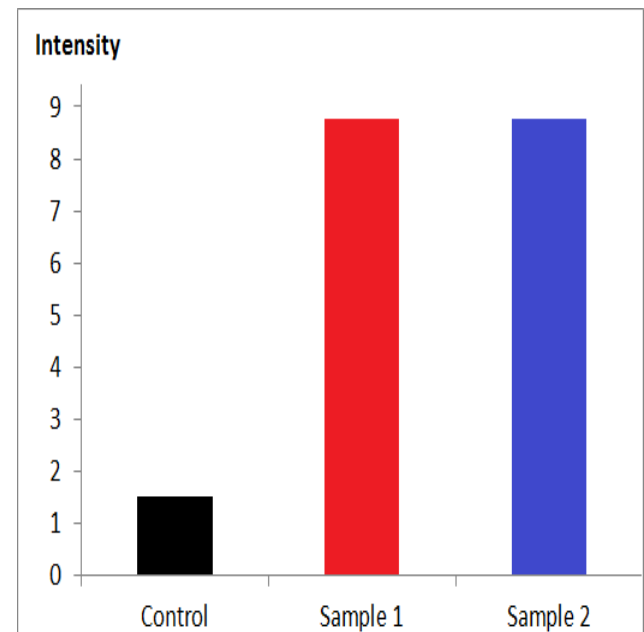
<http://www.desatoya.com>

Why Single Cell Analysis

Intensity Histogram for Single Particles

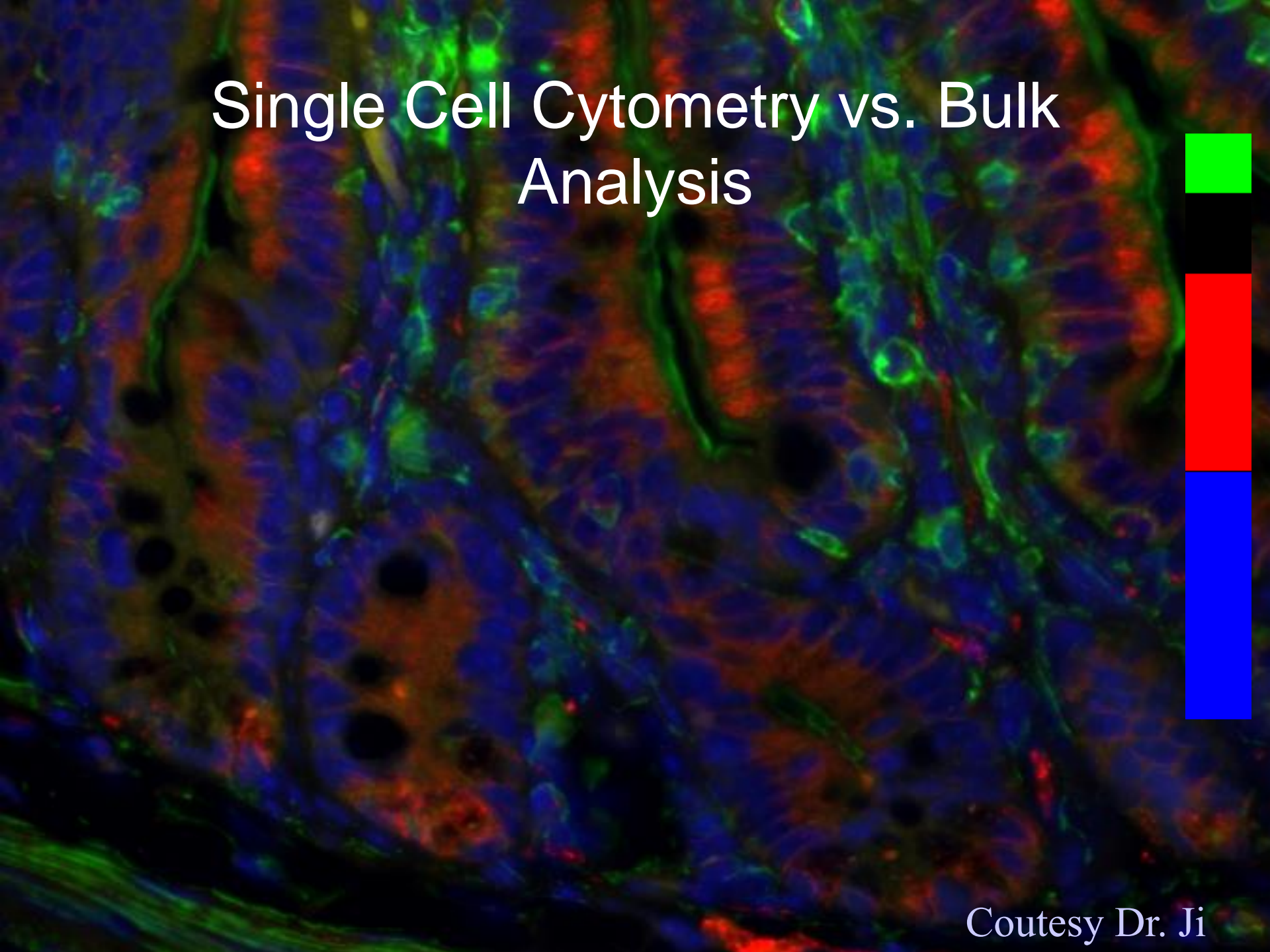


Intensity per Sample



Cell by cell intensity analysis detects population heterogeneity.

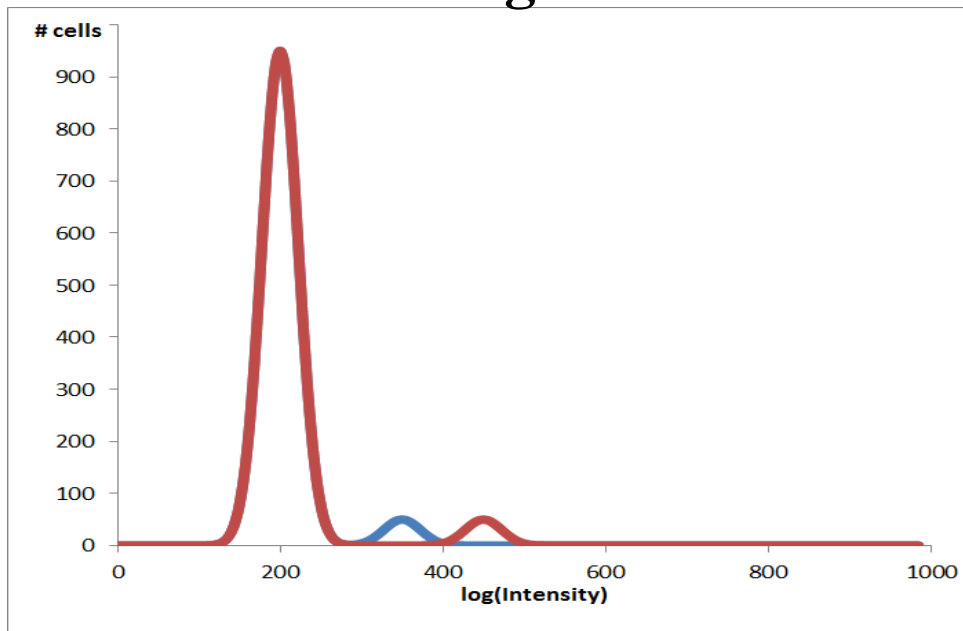
Single Cell Cytometry vs. Bulk Analysis



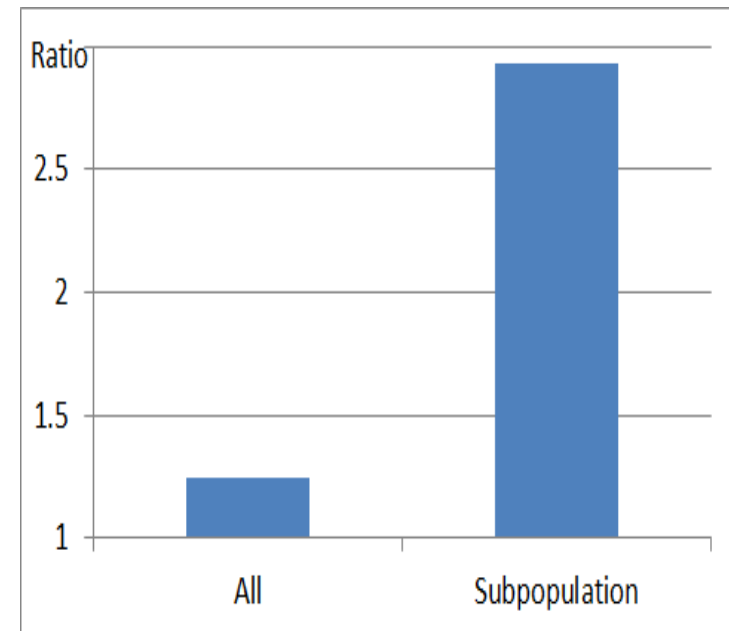
Courtesy Dr. Ji

Benefits of Subset Analysis

Intensity Histogram

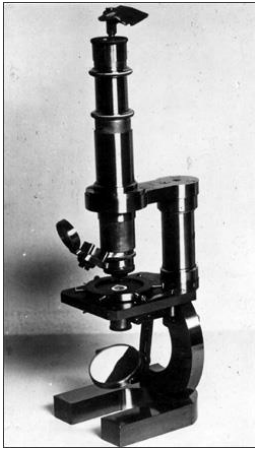


Intensity Ratios



Subpopulation analysis detects changes better, especially for rare subpopulations.

From Early to Present Microscopy and Flow Cytometry

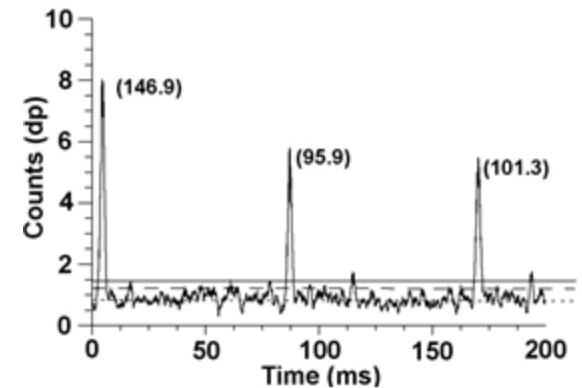


Source: BD Biosciences

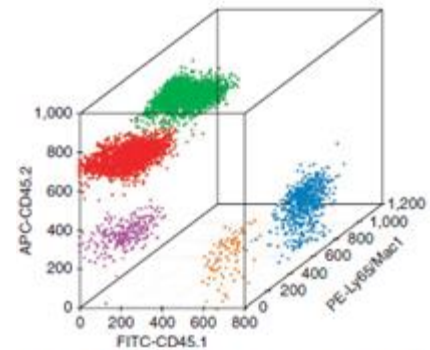
Flow and Imaging Cytometry Features

Single cell analysis with

- High sensitivity (single molecule sensitivity) I,F
- Wide dynamic range (10^0 to 10^7 mL⁻¹) F
- High analysis rates to $\sim 10^5$ sec⁻¹ F
- Light scatter F
- Direct size and 3D information I
- Multi-parameter analysis I,F
- Direct kinetic measurements I
- Live/dead discrimination I,F
- Viable cells can be re-covered F,(I)
- Measurement of adherent cells I
- Good ease-of-use F,(I)



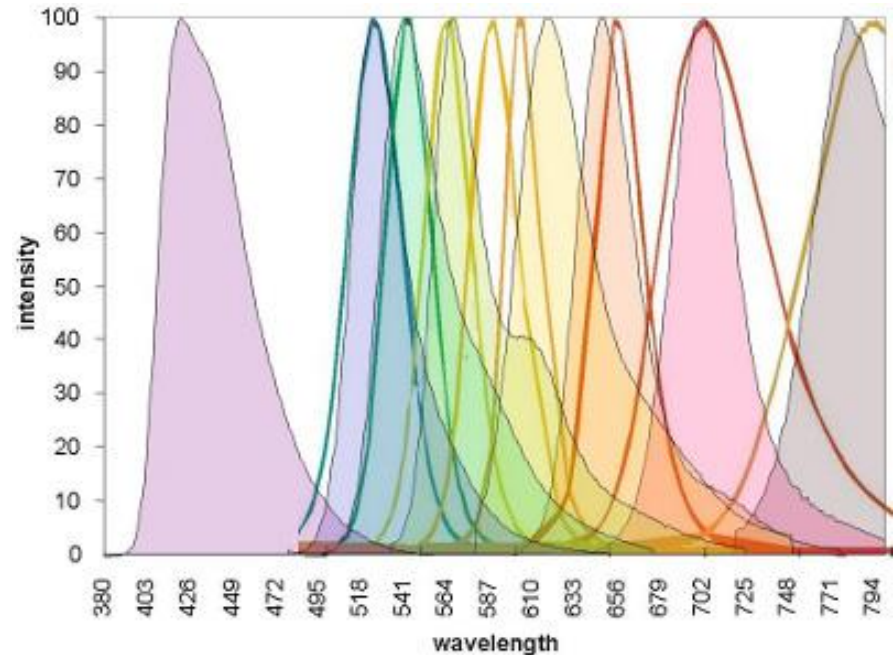
Rob Habbersett & Jim Jett, LANL



FlowCAP Consortium; Nature Methods 2013

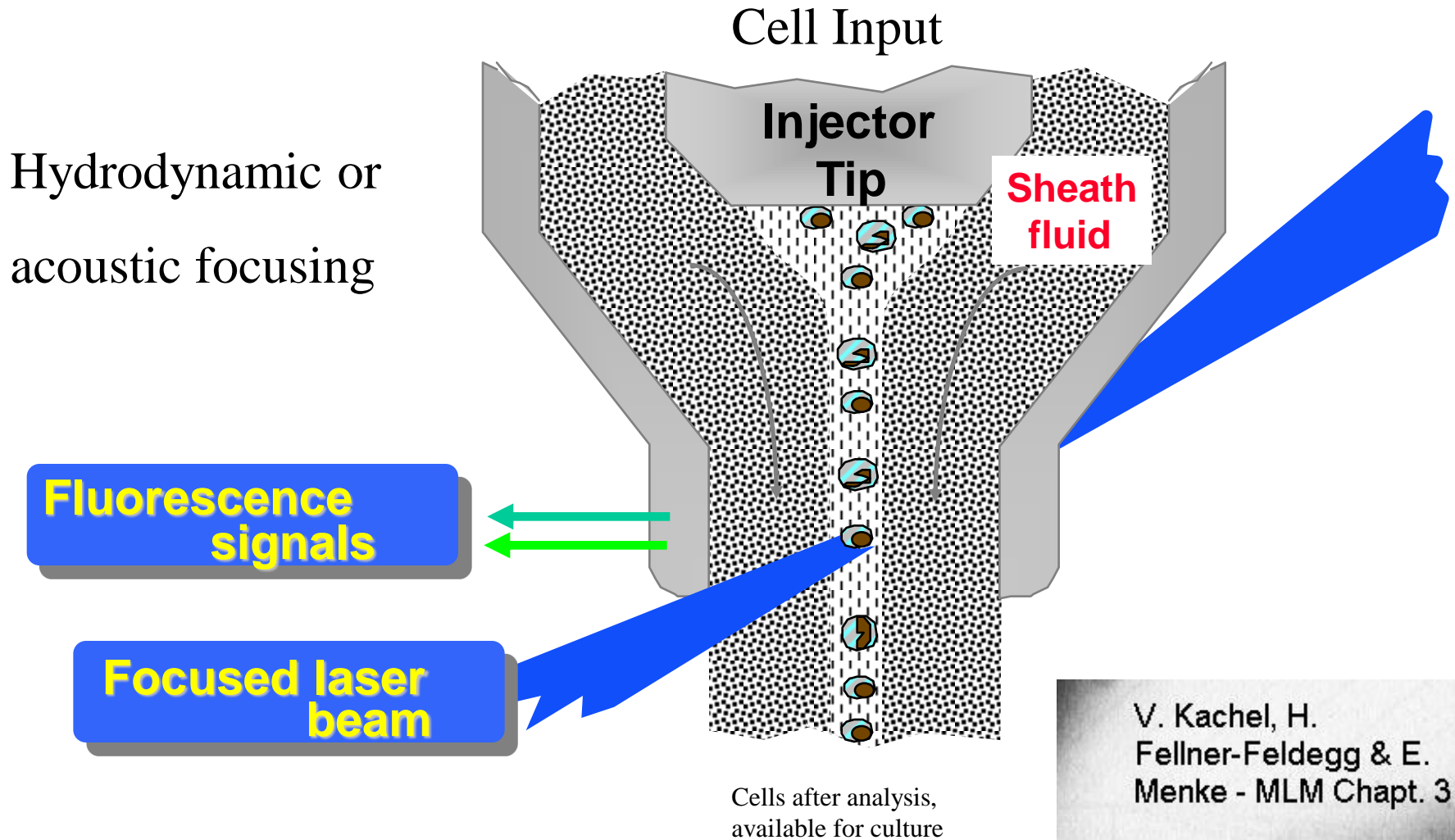
Physical parameters

- Light scatter
- Fluorescence
- Phosphorescence
- Raman
- Element mass
- Electrical properties
e.g. impedance
- ...

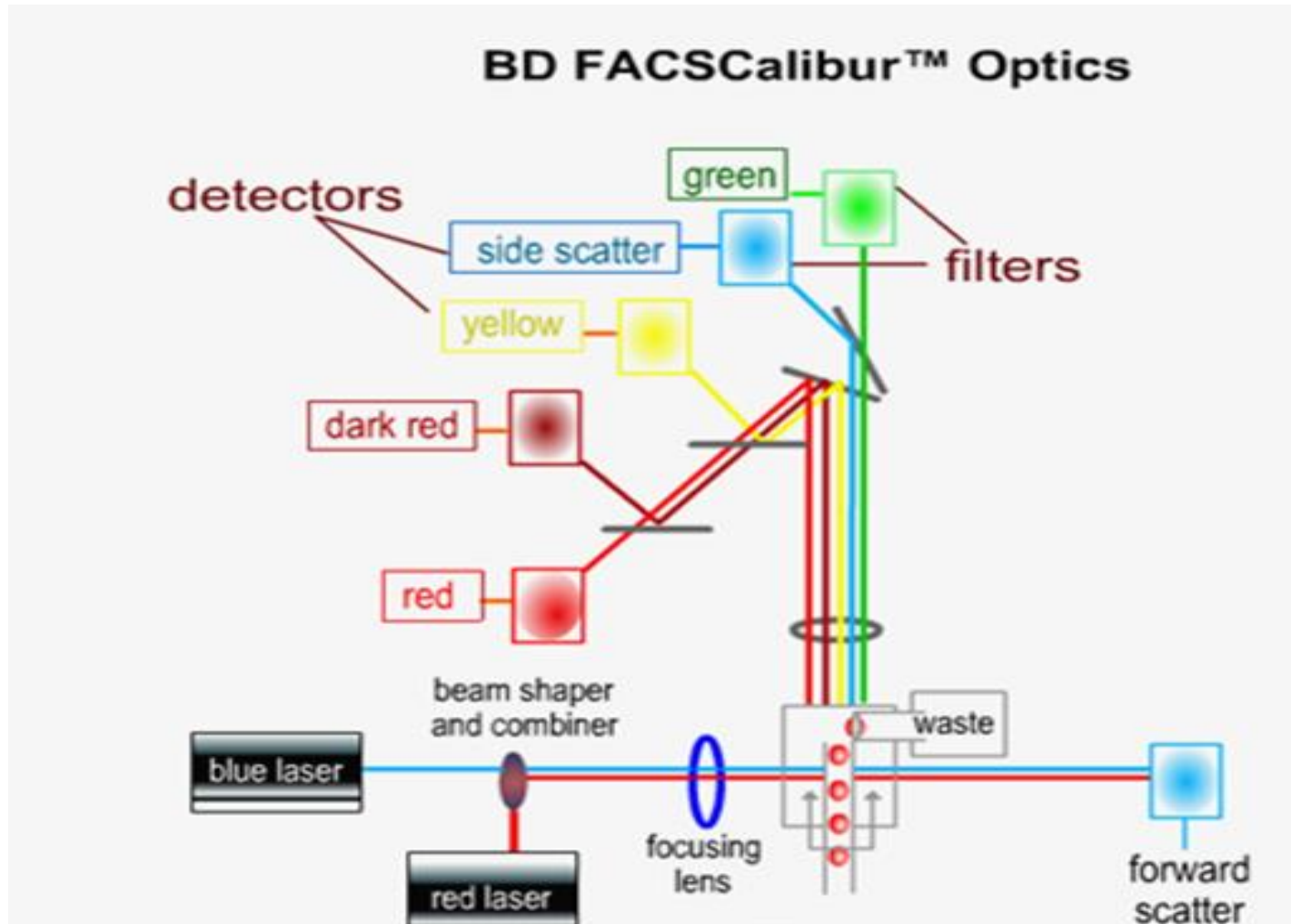


<http://www.dvssciences.com/technical.html>

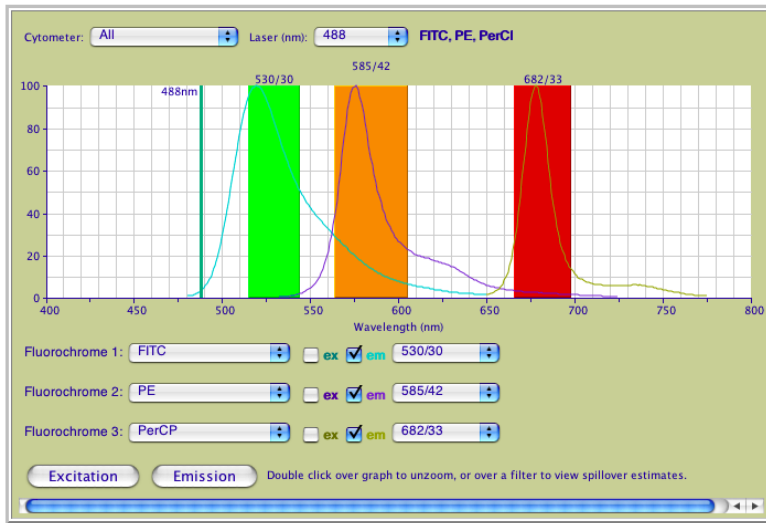
Flow Cytometer Fluidics



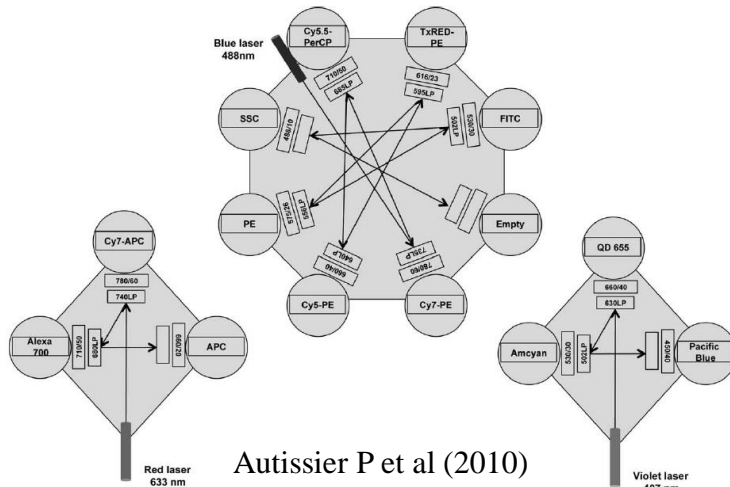
Flow Cytometer Optical Systems



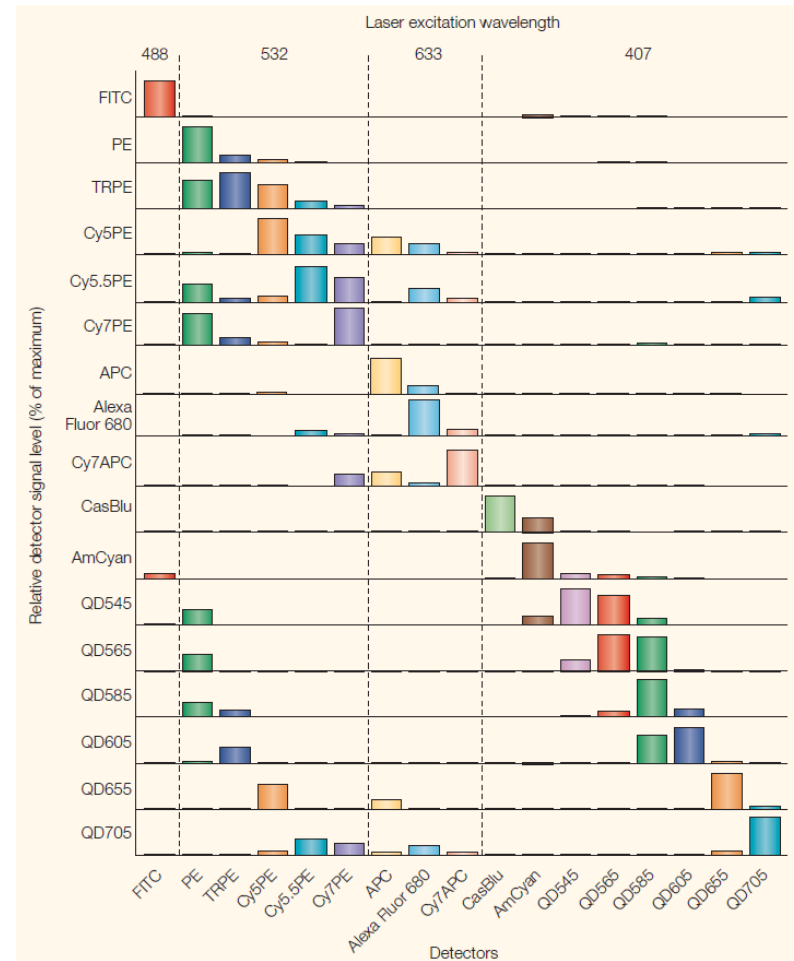
Filter Arrangement and Spectral Overlap



BD Biosciences



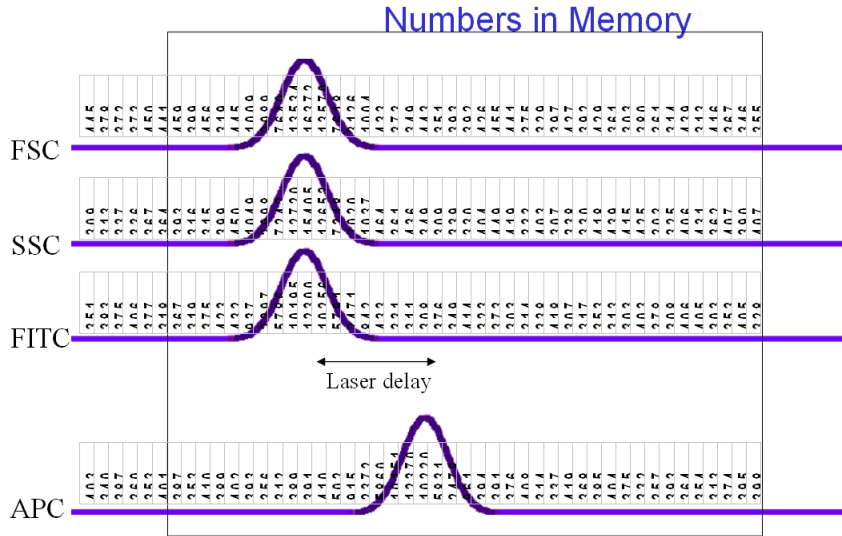
Autissier P et al (2010)
Cytometry 77A, 410ff



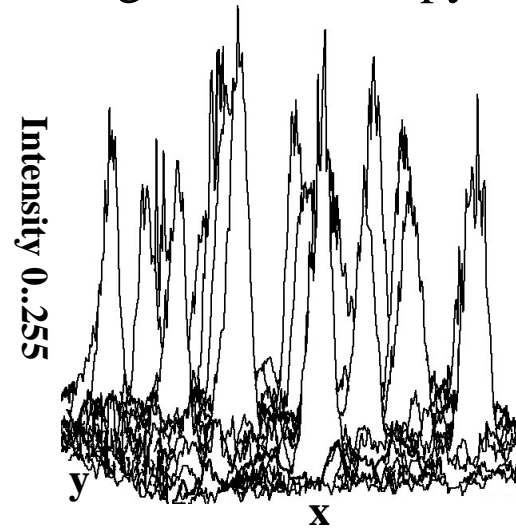
Perfetto SP et al (2004)
Nature Reviews Immunology 4, 648ff

Basic Data Processing

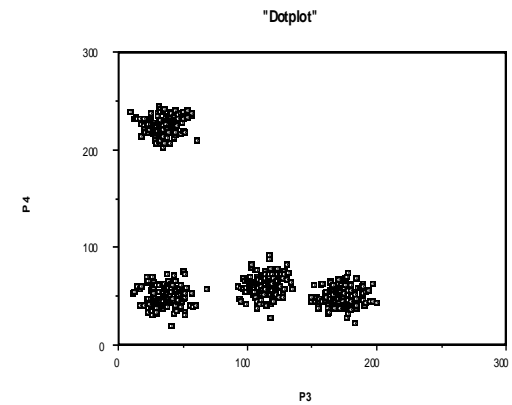
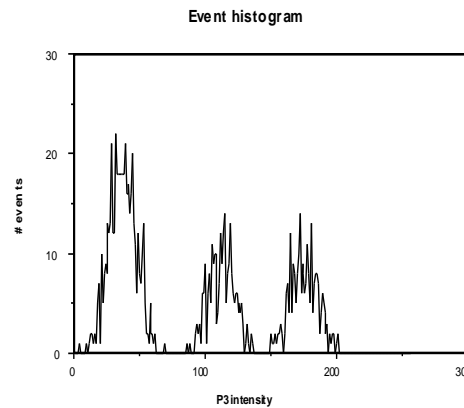
Flow Cytometry



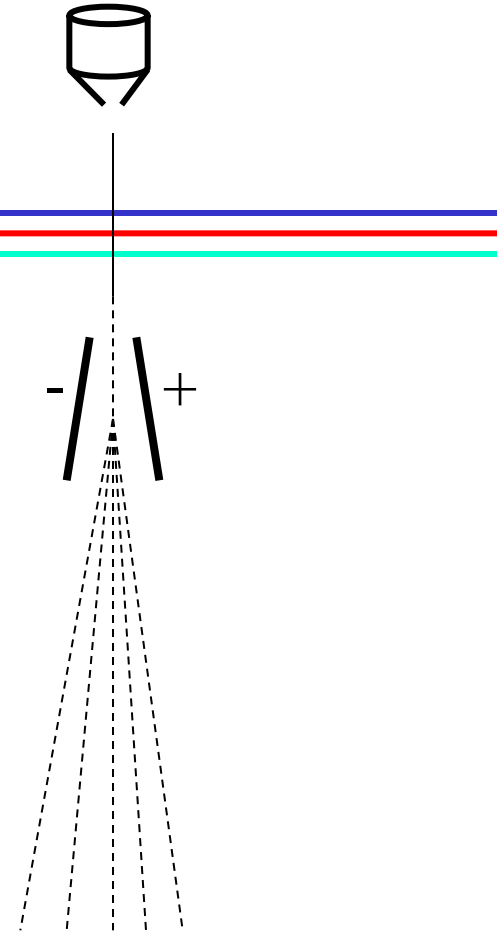
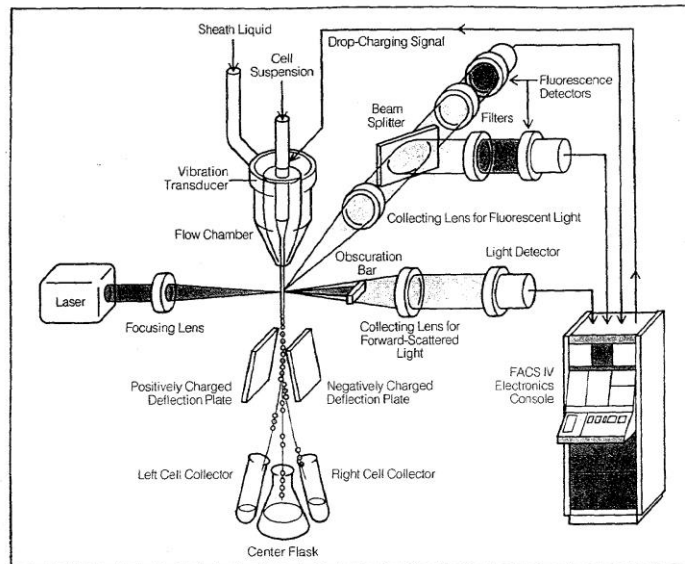
Digital microscopy



Cell	P1	P2	P3	P4	P5	Pop#
1	242	135	704	175	612	1
2	146	132	690	178	566	1
3	269	147	89	206	580	3
4	442	143	399	250	255	4
5	212	167	155	926	526	2
6	269	2	659	207	575	1
7	204	232	112	171	679	3
8	152	74	160	828	532	2
			...			
9997	215	119	138	936	662	2
9998	244	50	72	261	543	3
9999	214	137	174	1014	597	2
10000	312	87	110	904	560	2



“Droplet-based” Sorting



Sorting for gene expression analysis in single cells

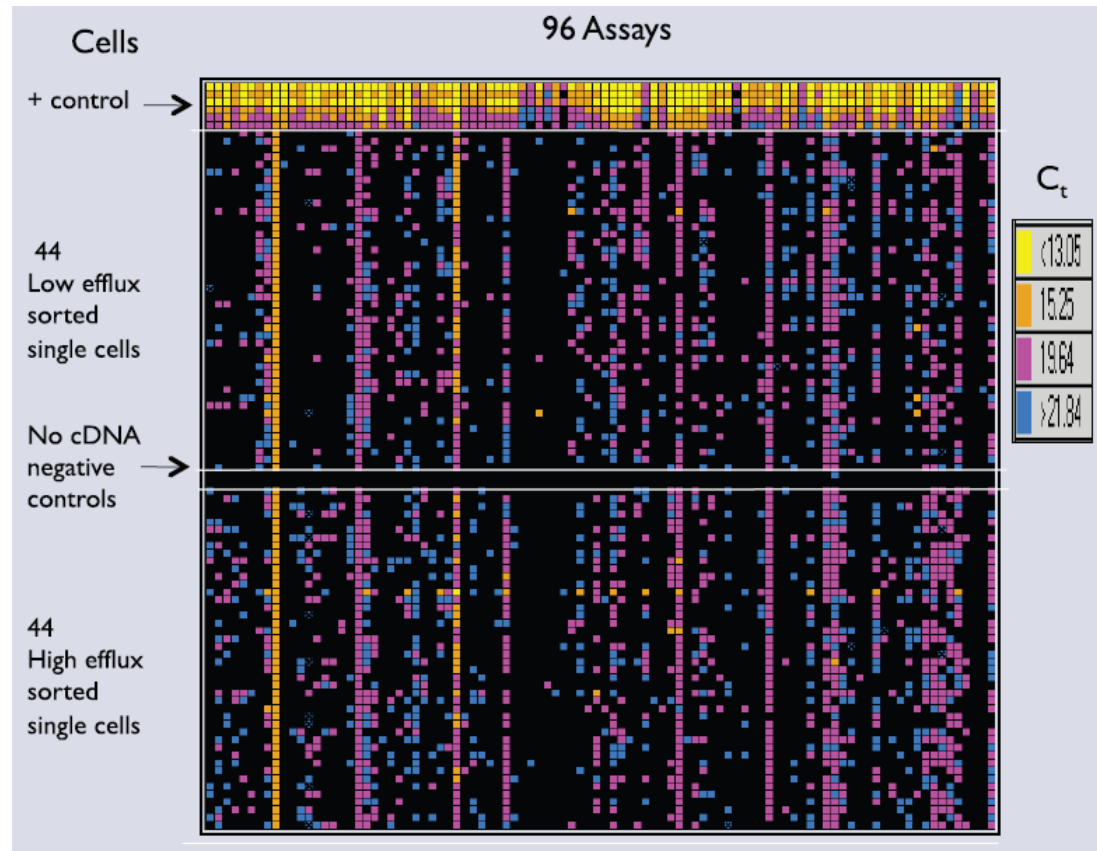
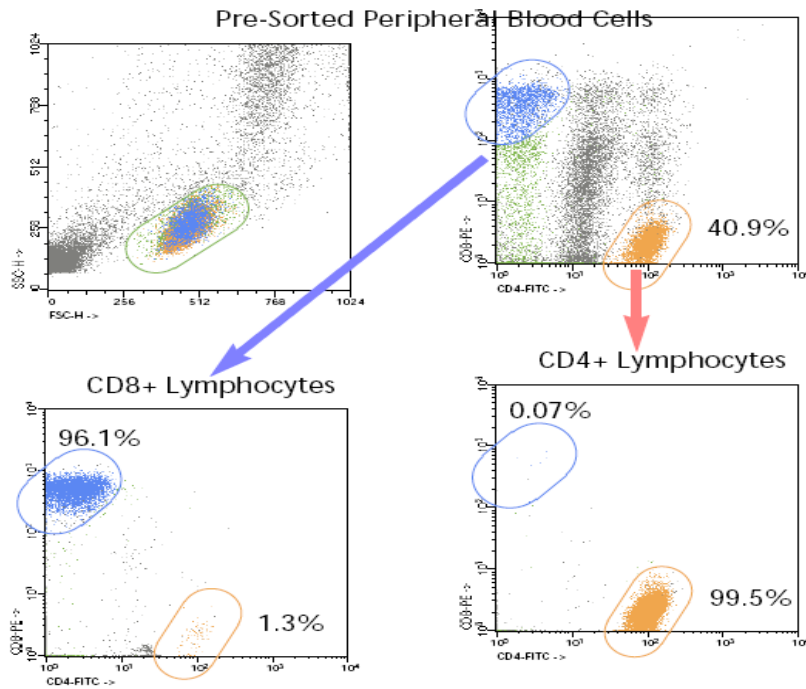


Figure 9: C_t heat map from 96 PCR assays, are shown on a 96.96 chip. Each point represents gene expression within a single cell. Some genes are more highly expressed in SP High than SP Low cell population.

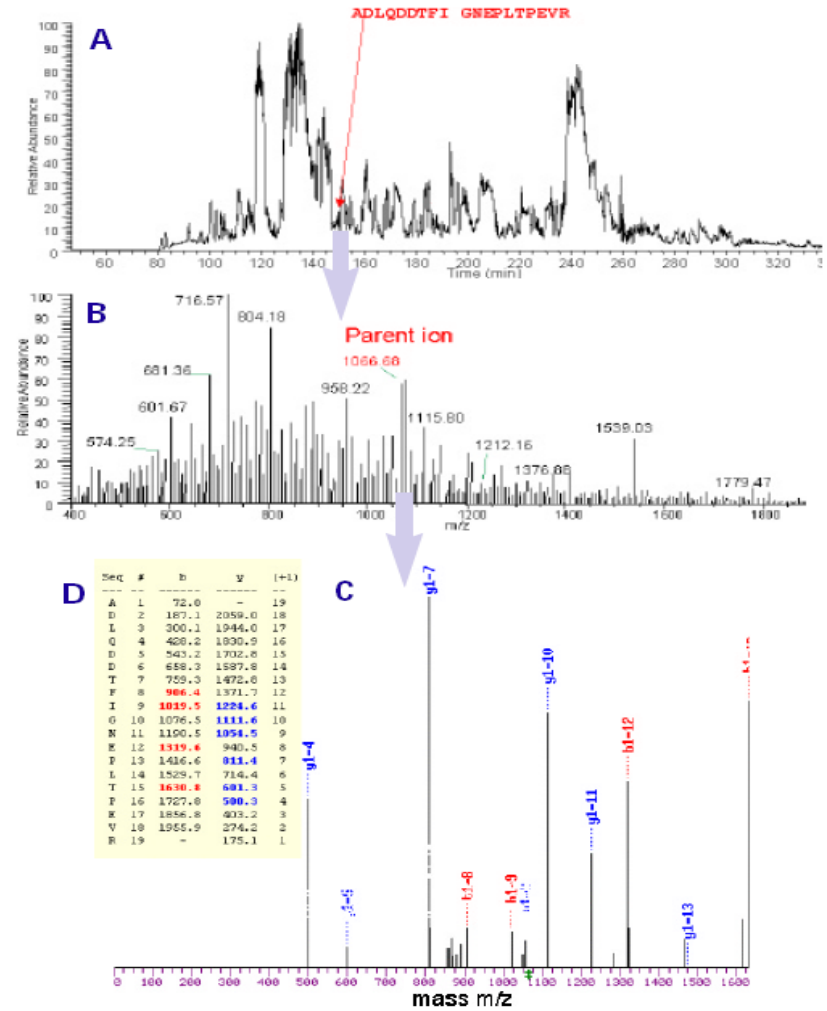
Sorting for Cell Surface Proteomics

Cell surface proteome by FACS sorting, followed by LC MS

(in collaboration with Thermo Finnigan, San Jose, CA)

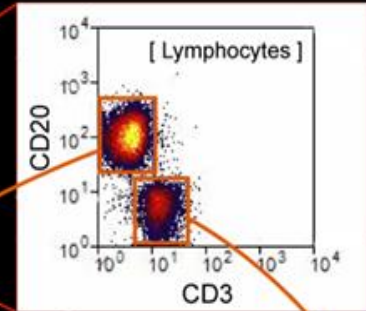
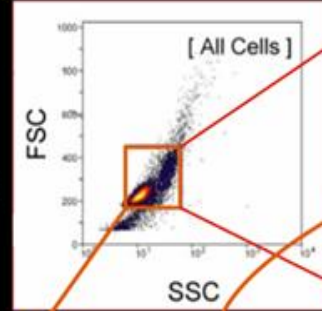
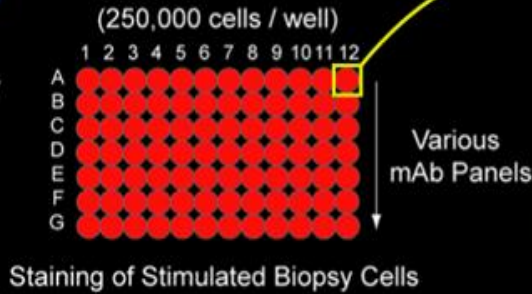
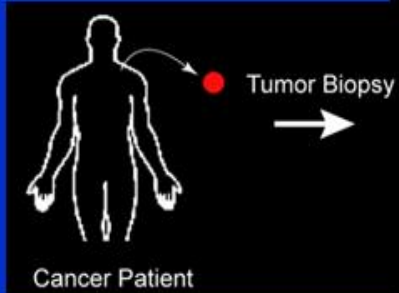


The dot plots show the sorting strategy used for stained peripheral blood cells and population purity after sorting for CD4- and CD8-positive cells. CD4 cells were gated on scatter and FITC fluorescence; CD8 bright cells were gated on scatter and RPE fluorescence. Sorted populations showed >95% purity.

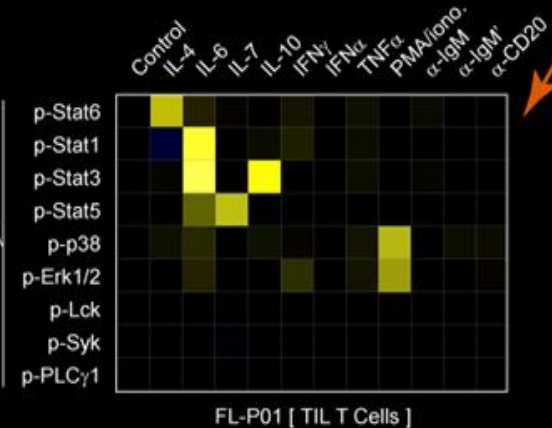
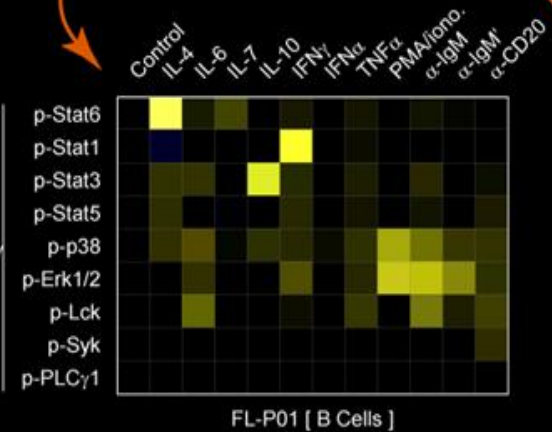
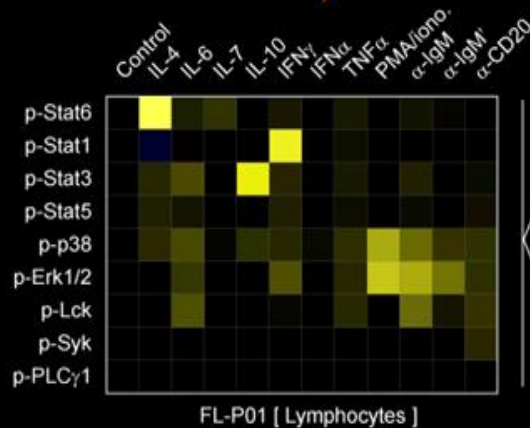
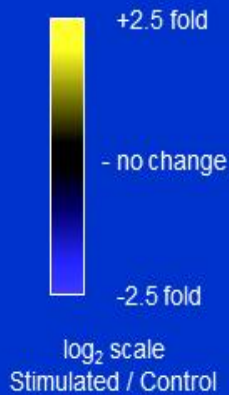


Peptide mixtures were separated by reverse phase HPLC (A) as described in Methods. Eluted peptides were subjected to electrospray injection into the mass spectrometer and analyzed for their mass/charge ratio (m/z value) (B). Selected ions were collected in the ion trap. These parent ions were cracked by collision ion dissociation to produce a range of fragment sizes (C) that were compared to predicted peptide sequences in the human database using TurboSequest (D).

Subset Analysis Example



profiling a lymphoma patient

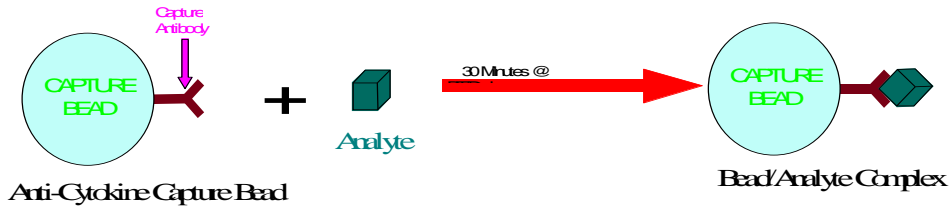


Slide provided by Nikesh Kotecha,
 CytoBank Inc, Mountain View CA

Data:
 J. Irish,
 Stanford

BeadArray Assays

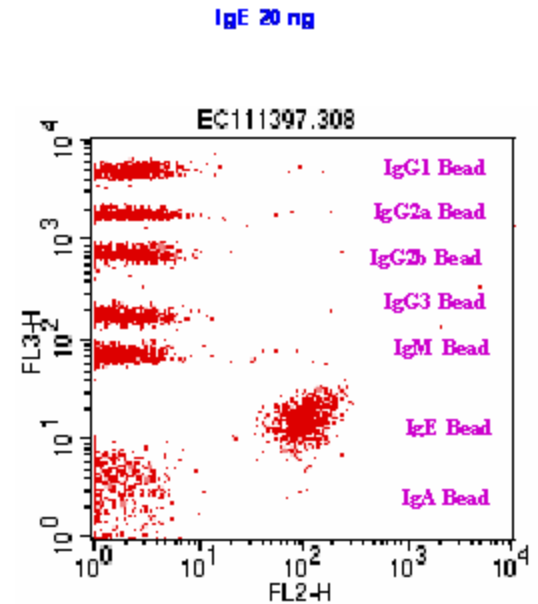
Step 1: Analyte



Step 2: Detector Antibody

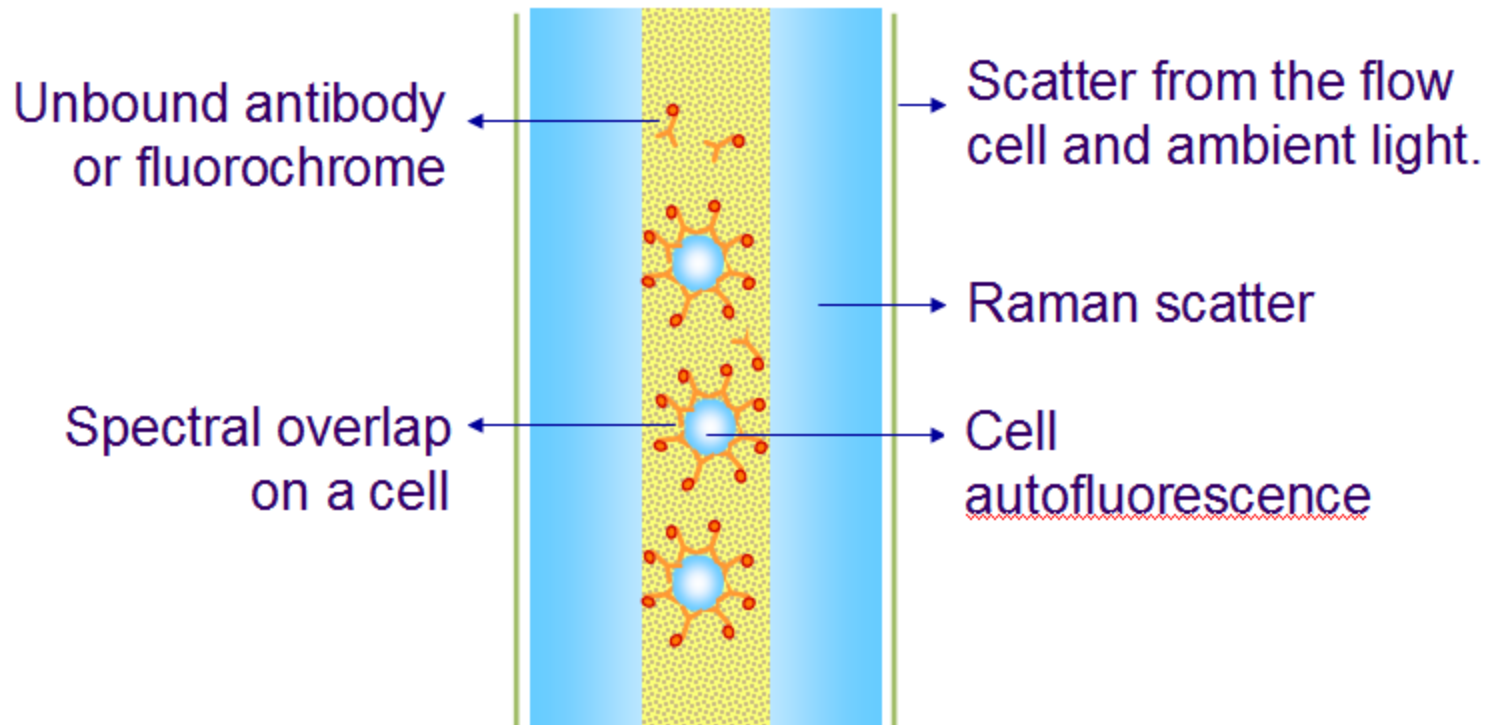


Step 3: Reporter Molecule

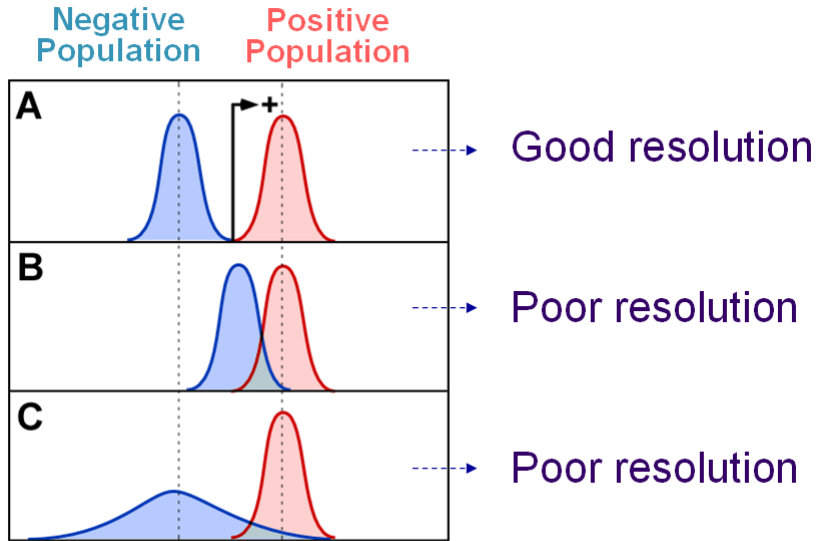


Instrument Evaluation Br

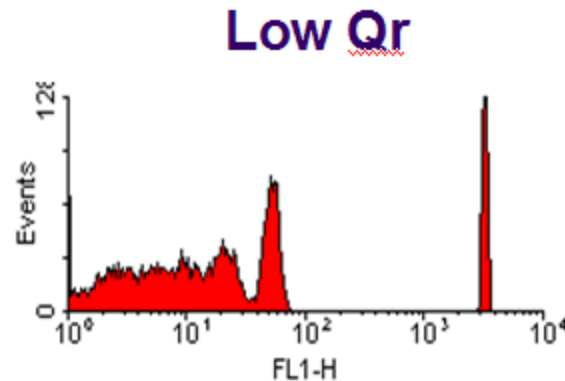
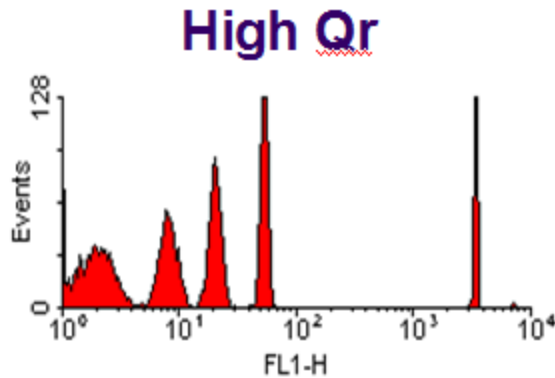
Relative B (Br) is a measure of true optical background in the fluorescence detector.



Instrument Evaluation Qr



$$Qr = \frac{\# \text{ photoelectrons}}{\# \text{ fluorescence molecules}}$$



Optimizing cytometry measurements (I)

- Background light

The total measurement SD is the sum of the error contributions from all sources:

$$SD = \sqrt{SD_{optical}^2 + SD_{intrinsic}^2 + SD_{illumination}^2 + SD_{noise}^2}$$

When two dyes (PE and FITC) are measured by a single detector the SD is additive:

$$SD_{optical} = \sqrt{SD_{PE}^2 + SD_{FITC}^2 + SD_{background}^2}$$

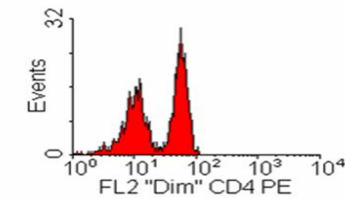
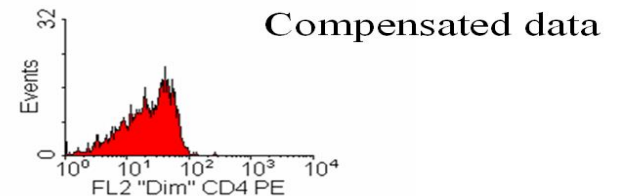
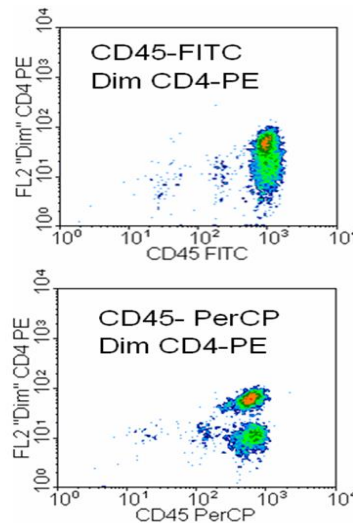
Reagent performance $\frac{\text{Stain index}}{2 * SD_{neg}}$

$$\frac{Medium_{pos} - Medium_{neg}}{2 * SD_{neg}}$$

- Dye properties (brightness and spectral overlap)

next slides:

- Multi-color dye overlap
- Gain (PMT, CMOS, CCD) settings
- Data Display
- Controls

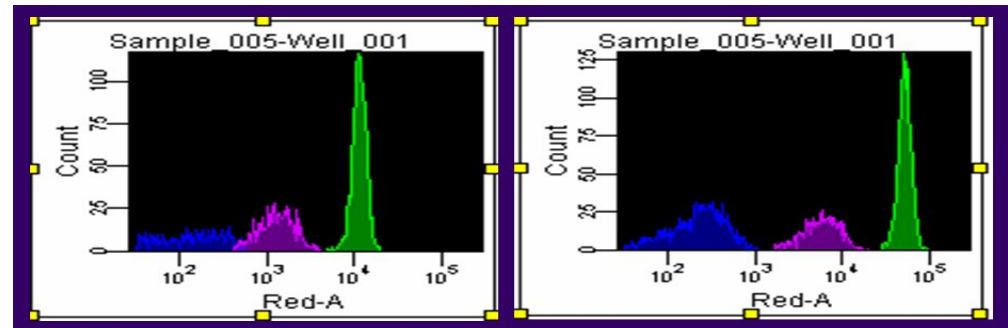


Better separation with less spectral overlap.

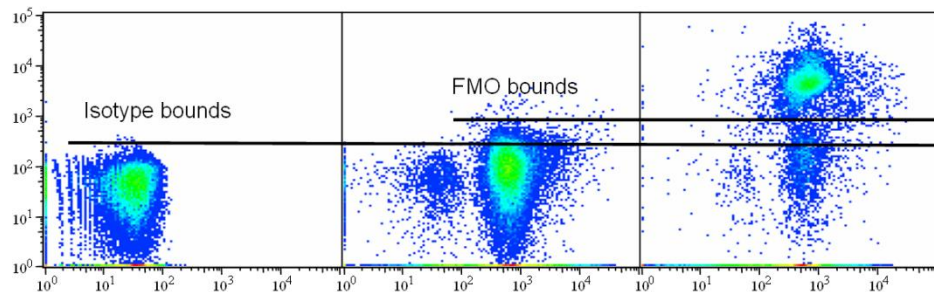
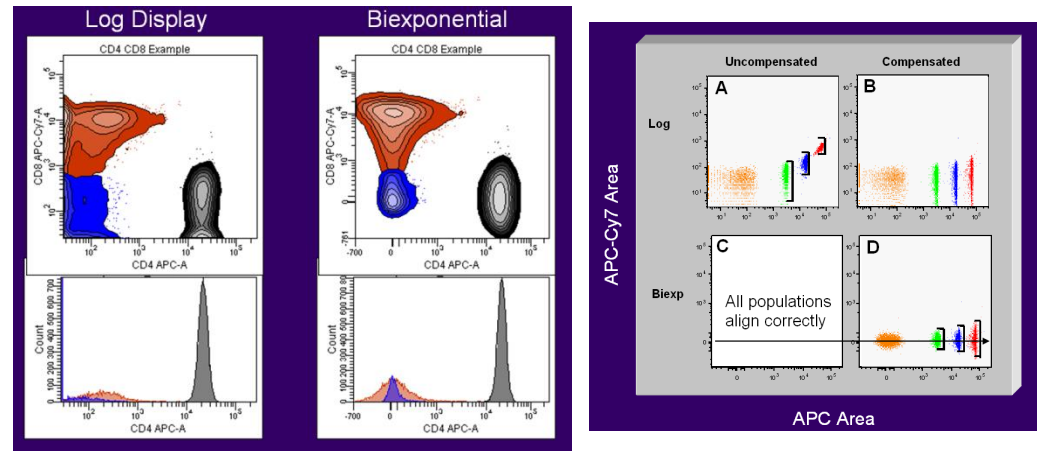
Optimizing cytometry measurements (II)

- Gain (PMT, CMOS, CCD) settings

- Data Display

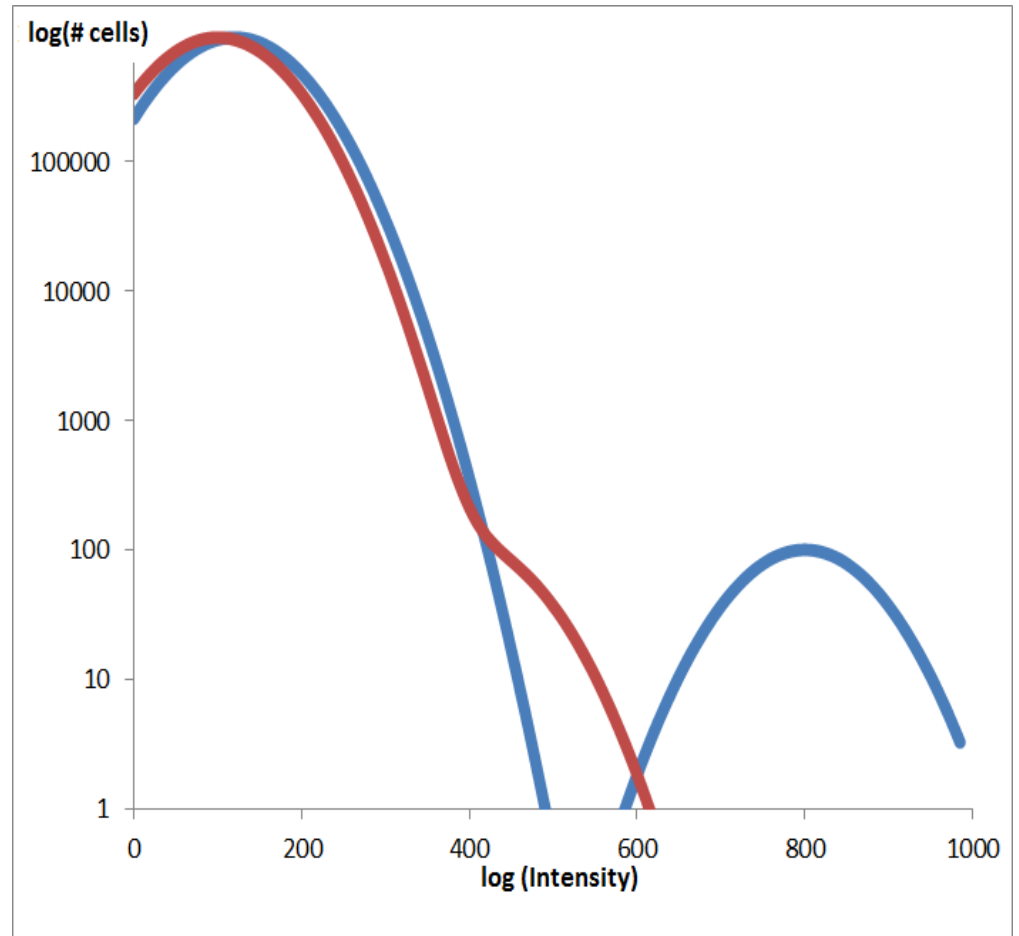


- Controls



Marker Brightness

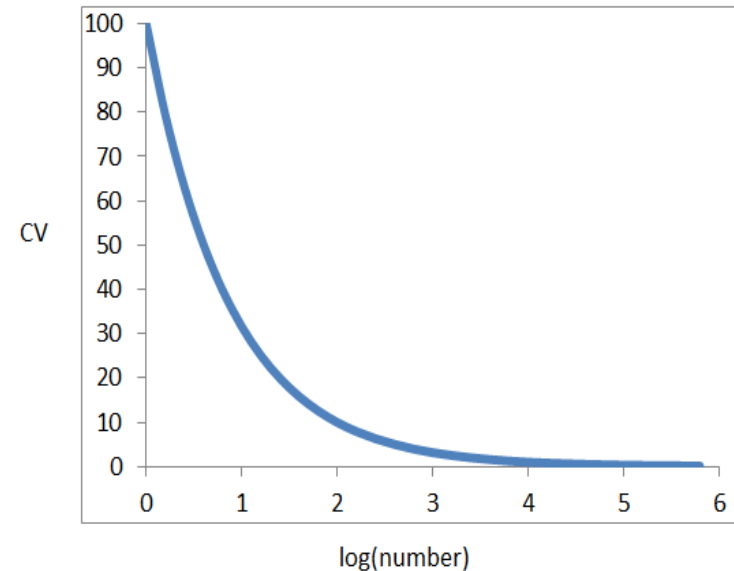
Brighter
markers resolve
rare populations
better



Rare Subpopulation Analysis

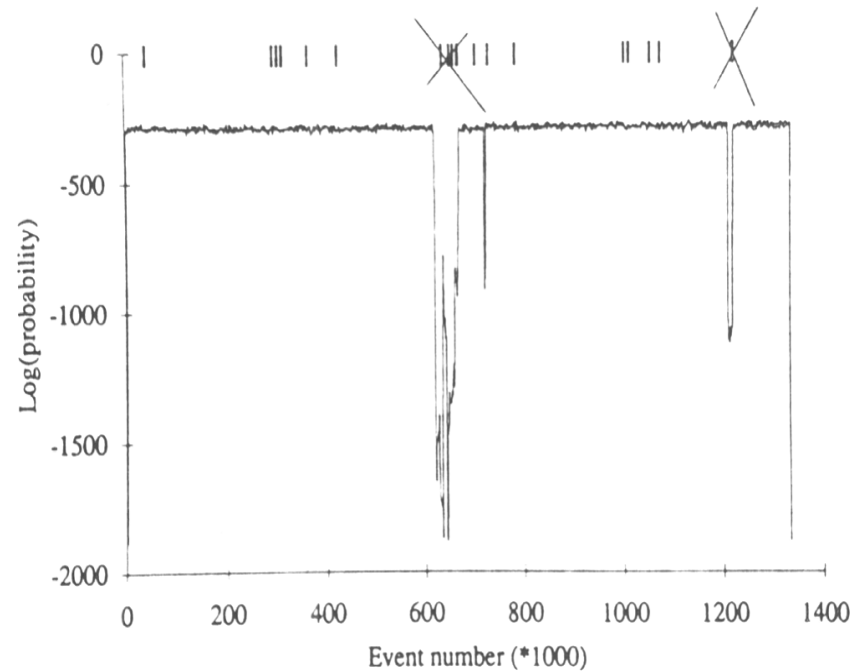
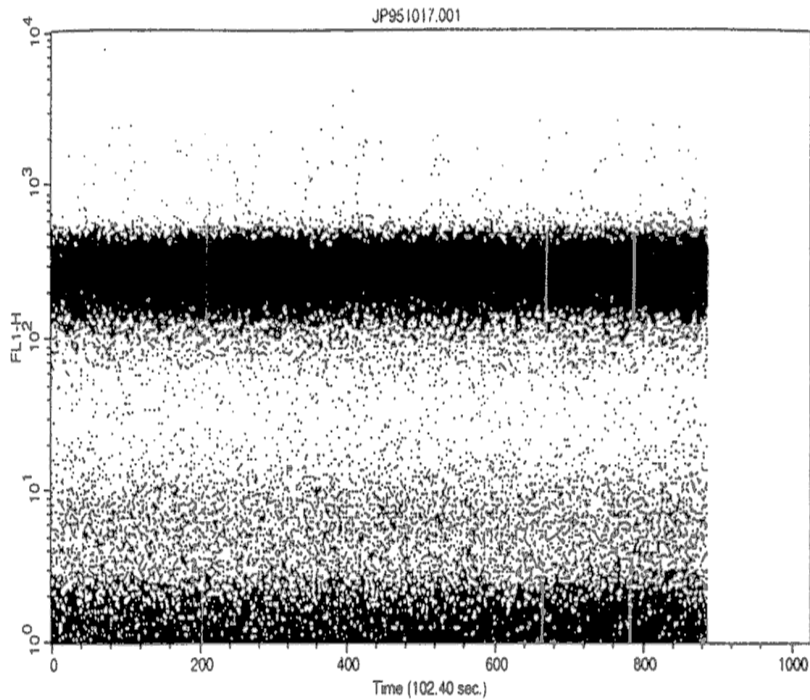
Counting Statistics

	Sample 1	Sample 2	Sample 3	Sample 4
	6	2	6	8
	3	7	1	6
	1	3	5	3
	1	4	5	6
	1	4	6	3
Mean	2.4	4	4.6	5.2
St.Dev	2.2	1.9	2.1	2.2
		Overall	Mean	4.1
			St.Dev	2.2



Ignoring Counting Statistics Can Lead to
Erroneous Conclusions

System Noise



Gross HJ et al, Cytometry 1993

Eliminating system noise from fluidics perturbations and other sources improves the limit of detection for rare cell analysis.

Particle Carry-Over

Carry-over specification examples

(from manufacturer's info)

Beckman Coulter FC500	<1%	
Beckman Coulter	<0.1%	
BD FACSCanto II	<0.1%	loader <1%
BD FACSVerser	<0.5%	
Guava EasyCyte	<0.2%	
Life Technologies Attune		loader <0.5%
Miltenyi MACSQuant	<0.01%	

Eliminating fluidic system particle carry-over
is vital for reliable rare cell analysis.

Multi-parameter Fluorescence Cytometry

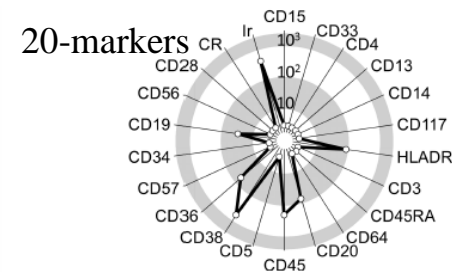
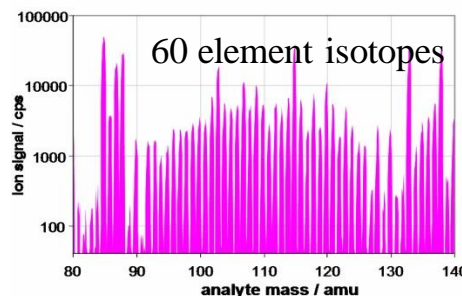
Points To Consider

Summary

- Know your instrument e.g. Qr & Br for different channels, carover, ...
- Use high enough gain settings to maximize sensitivity
- An antibody/dye combination that marginally allows discrimination of positives/negatives in a single color assay is unlikely to contribute anything helpful in a multicolor experiment.
- Avoid spillover from bright cell populations into channels requiring high sensitivity
- Beware of tandem dye degradation
- Internal controls are essential
- Take advantage of bright markers for “dim markers”

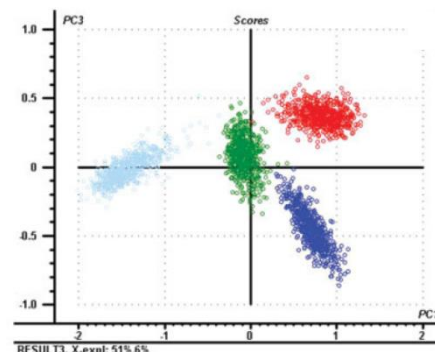
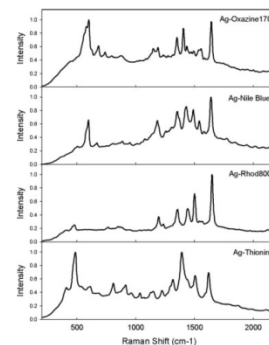
New Developments for Multi-parameter Cytometry

- Element-Label Flow Cytometry (CyTOF, addresses fluorescence spectral overlap issue by using elements as labels, *Anal. Chem.*, 2009, 81 (16), pp 6813–6822)



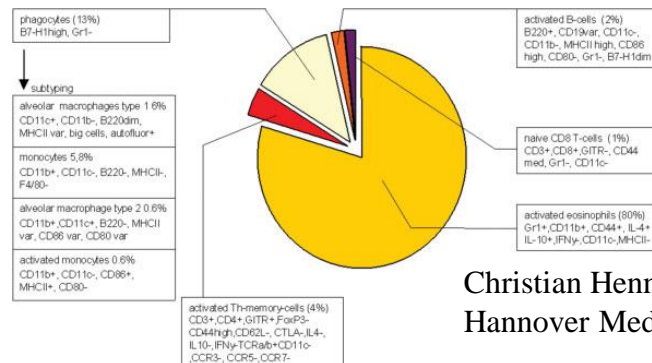
Scott Tanner, DVS Sciences Inc

- SERS-Label Flow Cytometry (uses spectral fine-structure to distinguish labels, *Cytometry*, 2008, 73A(2), pp 119-128)



John Nolan, La Jolla Bioengineering Institute

- Sequential Stain De-stain Cytometry (*Cytometry*, 2009, 75A(4), pp 362-370)
- SONY spectral analysis



31-marker analysis

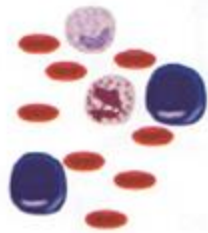
Christian Hennig, ChipCytometry Hannover Medical School

New Developments for Sample Preparation

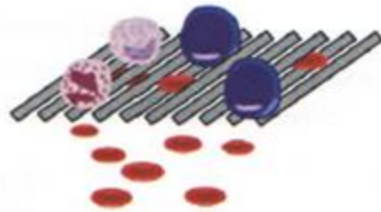
Removing erythrocytes and/or free reagent without centrifugation

Silicon filter for leukocyte/ erythrocyte separation

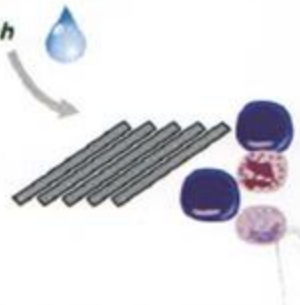
RBC depletion



Cell separation



Wash

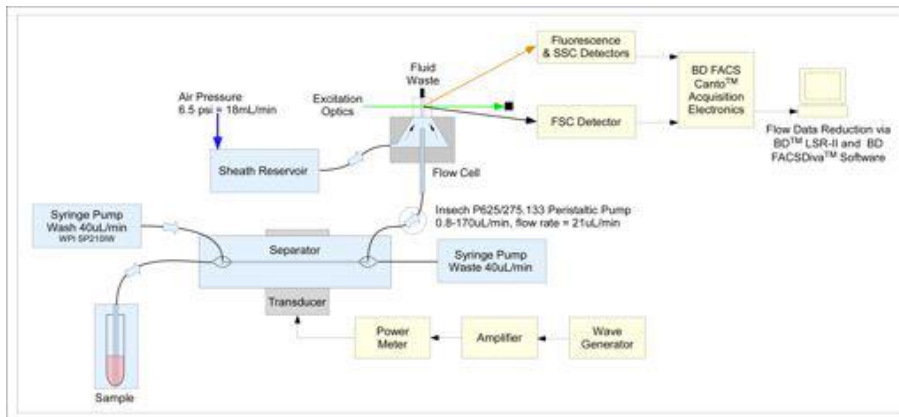


Leukocyte recovery

Flow rate	0.18ml/min	0.33ml/min
Cell load		
10ul (51.1±7.5)×1000 cells	0.98±0.04	0.92±0.07
50ul (350±14.1)×1000 cells	0.75±0.18	0.35±0.15

2011 Liping Yu et al, BD Biosciences and Aviva Inc.

Acoustic particle focusing

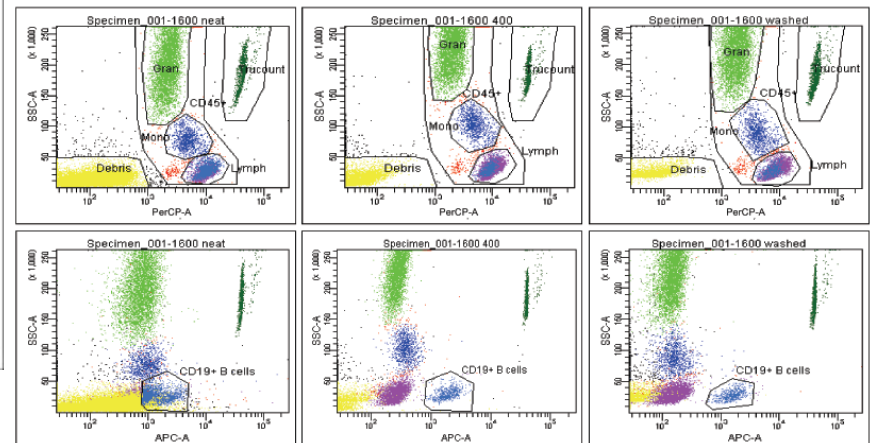


2010, Laurell group, Lund University
& Brian Warner, BD Biosciences

Lyse no wash

Chip wash

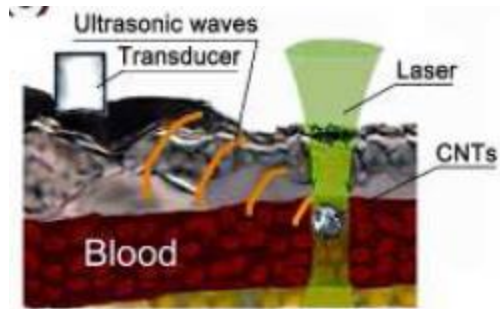
Centr. wash



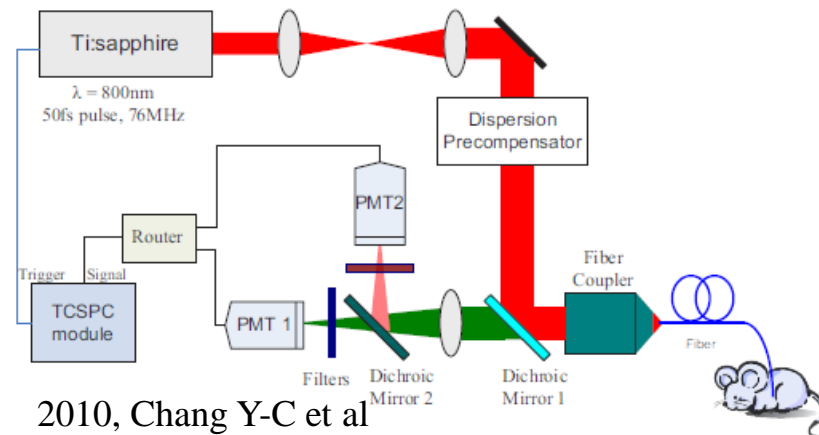
Intra-vital Cytometry

Single cell analysis in living animals

Flow cytometry in blood vessels

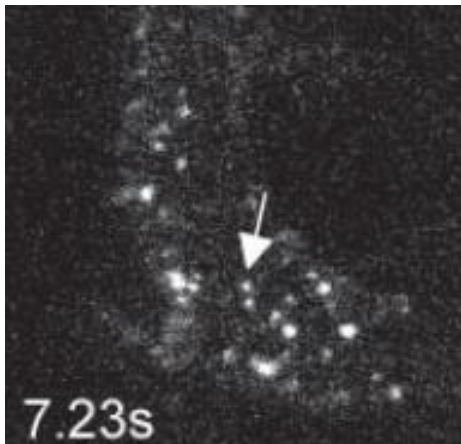


2010, Zharov VP and coworkers



2010, Chang Y-C et al

Microscopy



2011, Runnels JM et al; homing of multiple myeloma cells in bone marrow

Signals from

- 2-photon fluorescence
- bioluminescence
- photo-acoustic effect
- ...

Review paper:

Niesner RA, Cytometry 79A (2011)

Conclusions

New technologies from many fields are rapidly generating new opportunities for cytometry.

Acknowledgements

- Joe Trotter
- Ming Yan
- Ben Verwer
- Maria Jaimes
- Brian Warner
- Ed Goldberg
- Hrair Kirakossian
- Liping Yu
(all BD)
- BD
- Holden Maecker, Stanford
- Bob Hoffman, independent
- Ken Davis, retired
- Bill Godfrey, Danaher
- Brent Gaylord, Sirigen
- Collette Rudd, Thermo

Contact

Diether.Recktenwald@cytometrygroup.org,
Phone: +1-408-658-6074

Diether@desatoya.com
<http://www.desatoya.com>