

**8th ENII Immunology Summer School,
May/June 2013**

State of the Art in Cytometry

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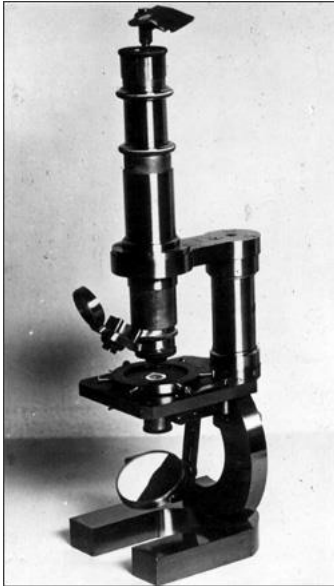
Presentation Outline

- **Technologies for Biology Research**
- **Flow and Image Cytometry Basics**
- **Examples from Genomics and Proteomics**
- **Optimizing Multi-Parameter Experiments**
- **Additional Considerations for Imaging**
- **Intra-vital Microscopy**
- **In-vivo Flow Cytometry**
- **Evolving Technologies**

Biology Research Targets and Tools

Organism	NMR	Contrast agents
	X-ray imaging	Affinity reagents
Organ	Ultrasound	- antibodies
	2-photon imaging	- probes
Tissue	In-vivo cytometry	Enzyme substrates
	Light microscopy	Labels
Single Cell	Electron microscopy	- absorbance
	Flow cytometry	- fluorescence
Organelle	Cell imaging	- element tags
	NA sequencing	
Macromolecule	Mass spectrometry	
	TIRF microscopy	
Small molecules	Electrophoresis	Sample prep

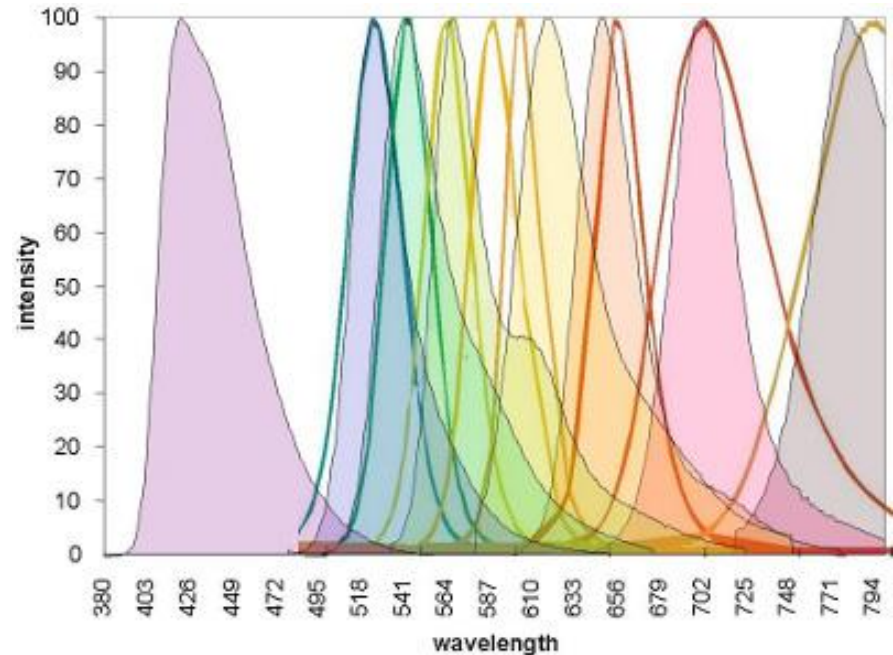
Single Cell Analysis Microscopy and Flow Cytometry



Cytometry Basics

Physical parameters

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

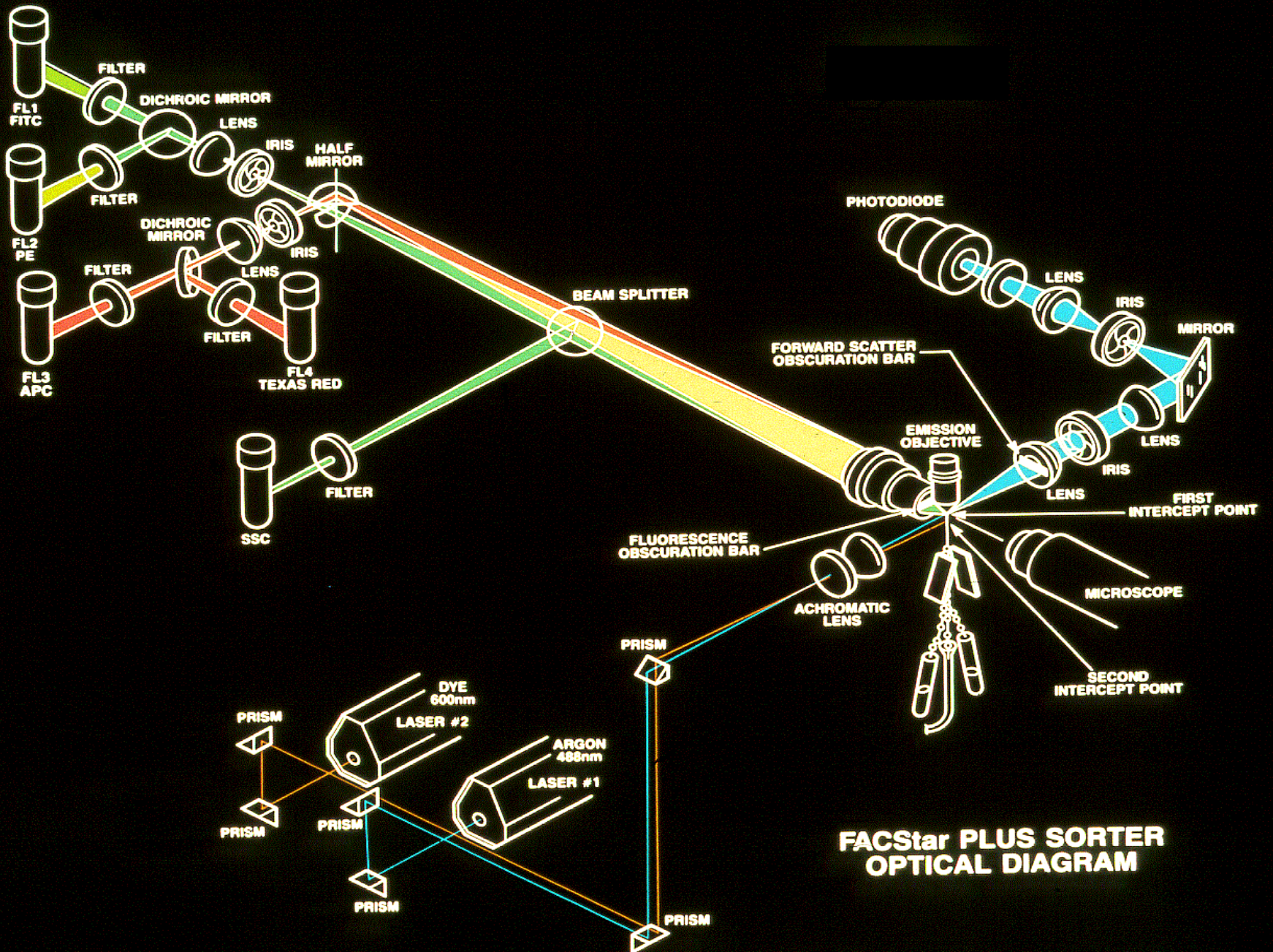


<http://www.dvssciences.com/technical.html>

Flow and Imaging Cytometry Features

Single particle (cell) analysis with

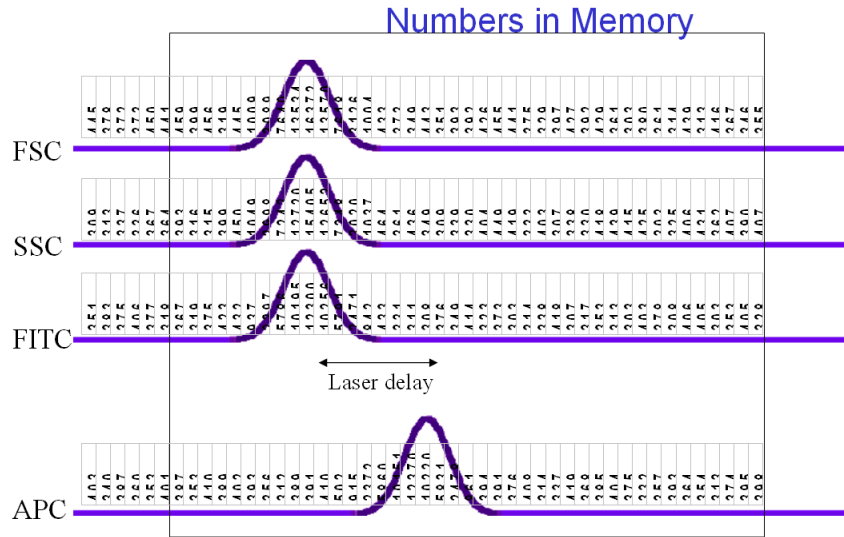
- High sensitivity (single molecule sensitivity by fluorescence) I,F
- Wide dynamic count range (10^3 to 10^7 cells mL^{-1}) F
- Particle sizes from 0.2 to 20 microns I,F
- High analysis rates to $\sim 10^5$ particles sec^{-1} F
- Direct size and 3D spatial information I
- Multi-color fluorescence, multi-parameter analysis F,I
- Wide dynamic range for fluorescence (10^5) F
- Direct kinetic measurements I
- Viable cells can be re-covered F,(I)
- Measurement of adherent cells I
- Good ease-of-use F,(I)



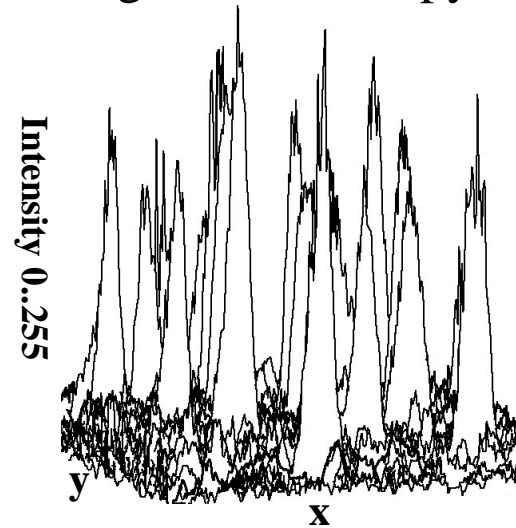
**FACStar PLUS SORTER
OPTICAL DIAGRAM**

Basic Data Processing

Flow Cytometry

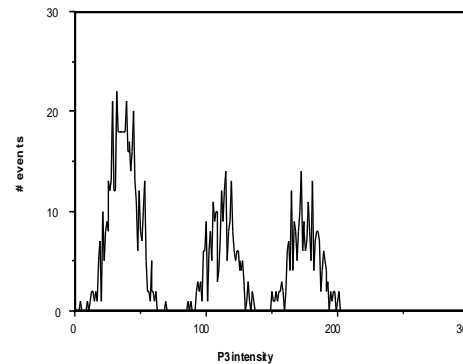


Digital microscopy

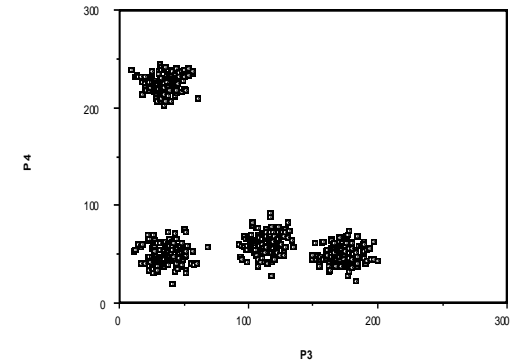


Cell	P1	P2	P3	P4	P5	Pop#
1	242	135	704	175	612	1
2	146	132	690	178	566	1
3	269	147	89	206	580	3
4	442	143	399	250	255	4
5	212	167	155	926	526	2
6	269	2	659	207	575	1
7	204	232	112	171	679	3
8	152	74	160	828	532	2
			...			
9997	215	119	138	936	662	2
9998	244	50	72	261	543	3
9999	214	137	174	1014	597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

Event histogram

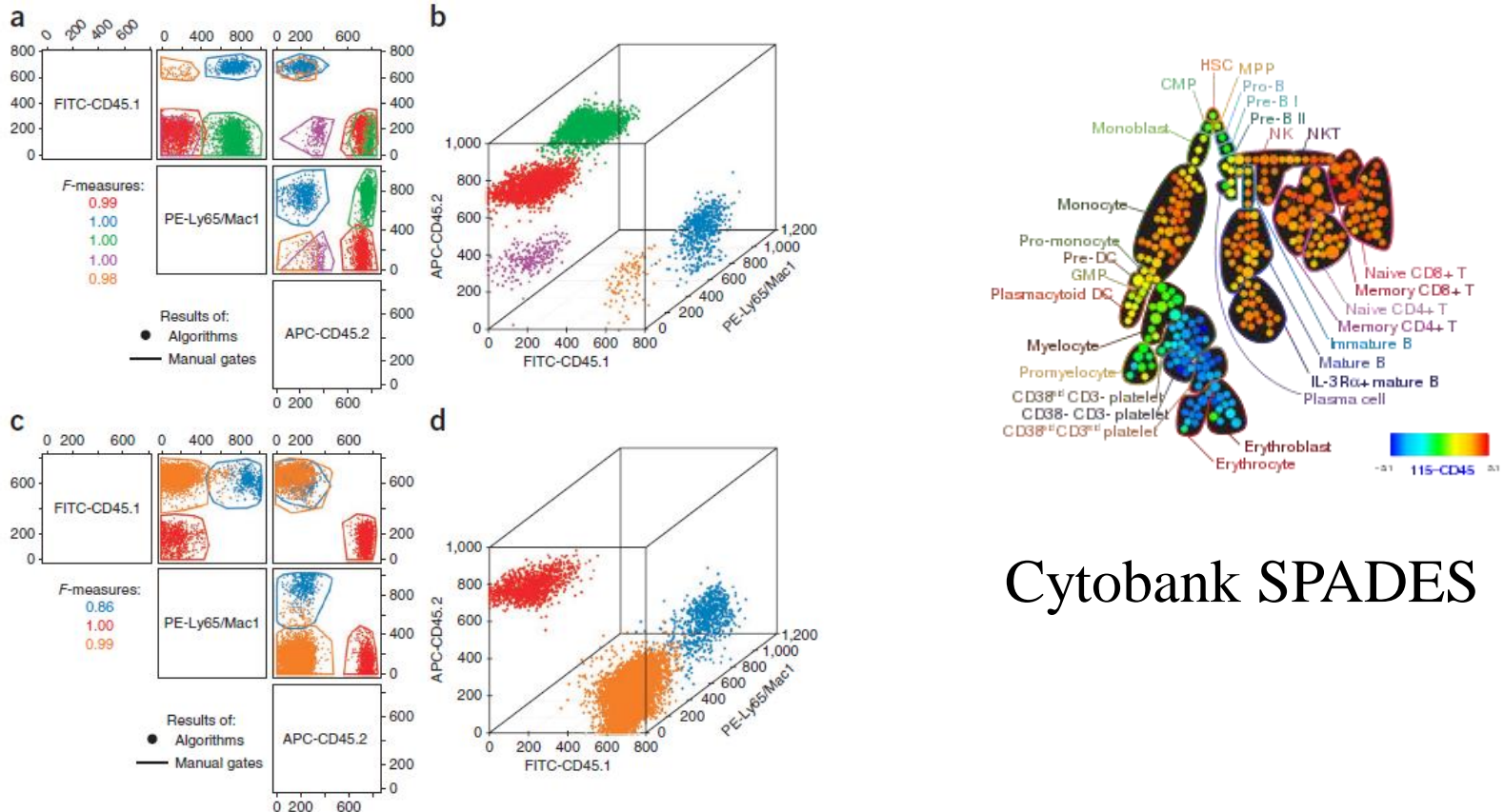


"Dotplot"



for >2 parameters: gating, cluster analysis, ...
For many samples and parameters: bioinformatics

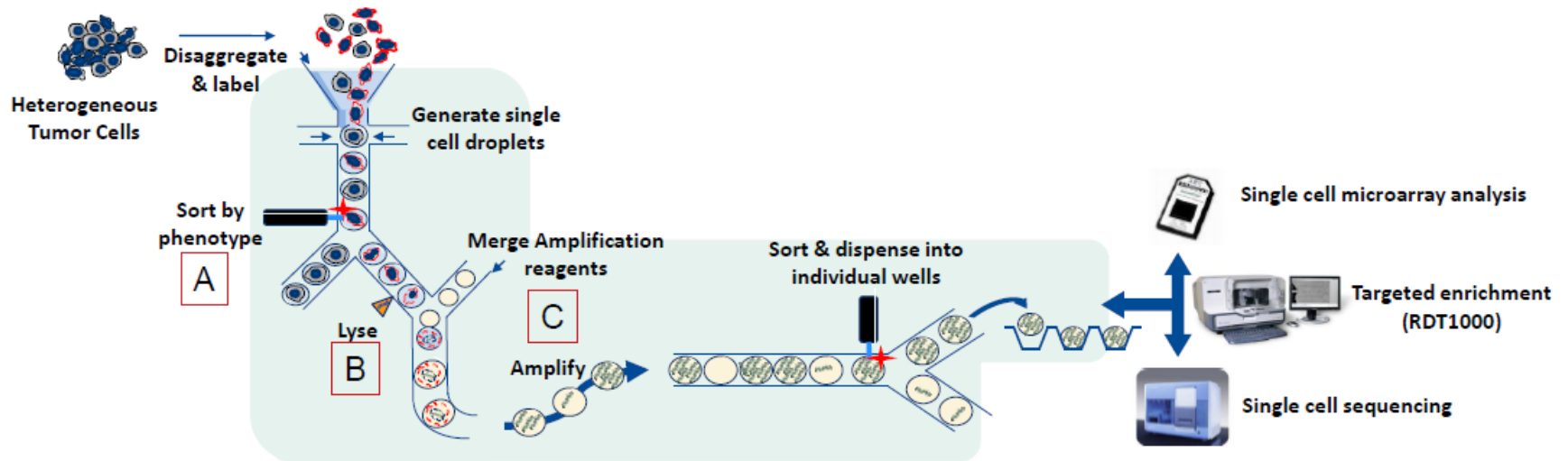
Bioinformatics



Cytobank SPADES

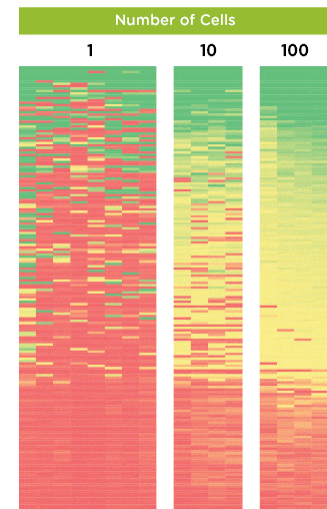
FlowCAP Consortium;
 Nature Methods 2013;10, 228ff

Single Cell Genomics



Source: Raindance Technologies

Single cell analysis reveals heterogeneity, which is masked by averaging, when analyzing groups of cells.

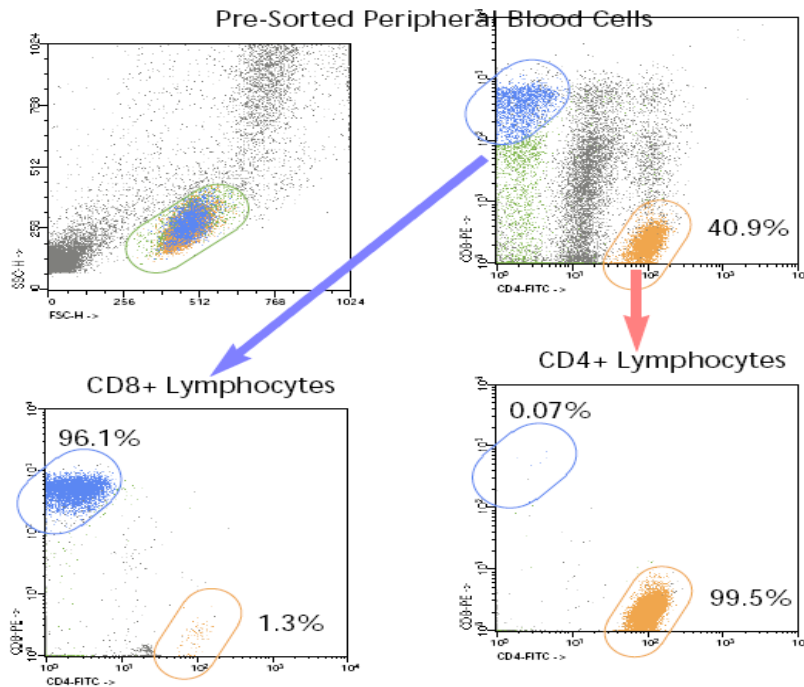


Source:
<http://www.nanostring.com>

Sorting for Cell Surface Proteomics

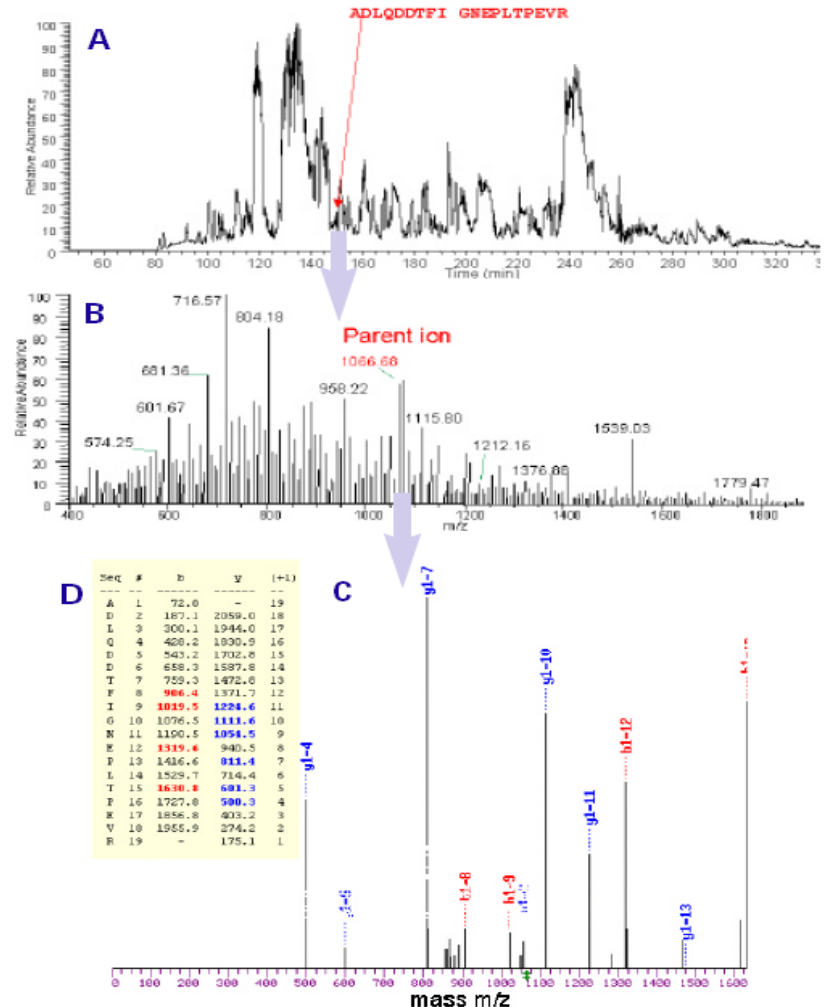
Cell surface proteome by FACS sorting, followed by LC MS

(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.

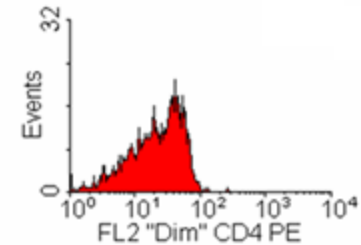
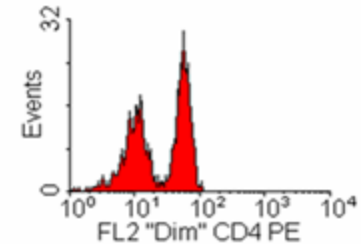
Cell Surface Proteome of Sorted Cells, measured by LC ESI MS



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 tracked 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).

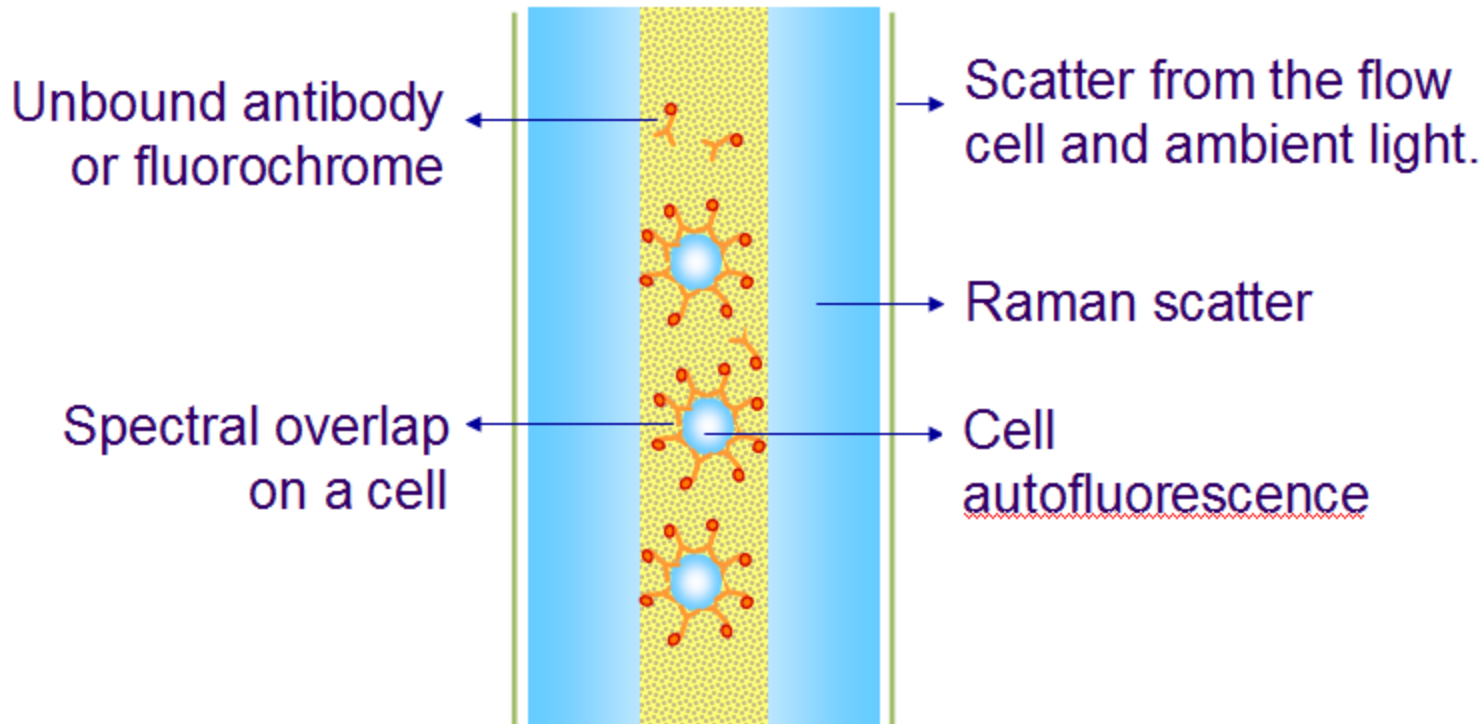
Optimizing Multi-Color Experiments

- Population separation
- Br and Qr
- Data Display
- Controls
- Microscopic Imaging

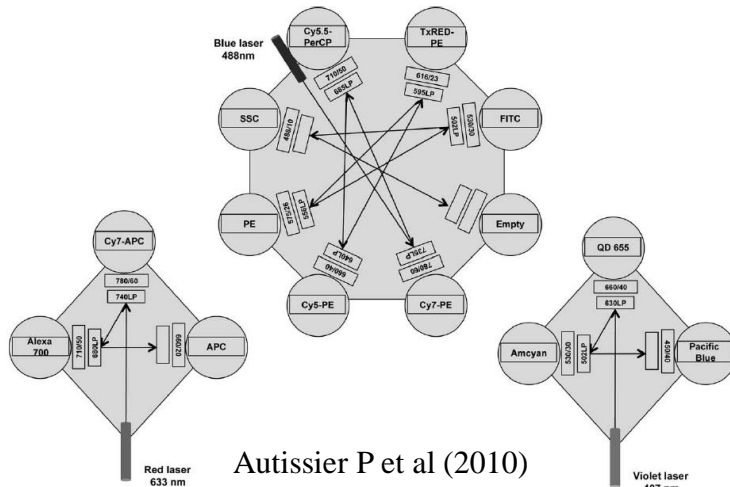
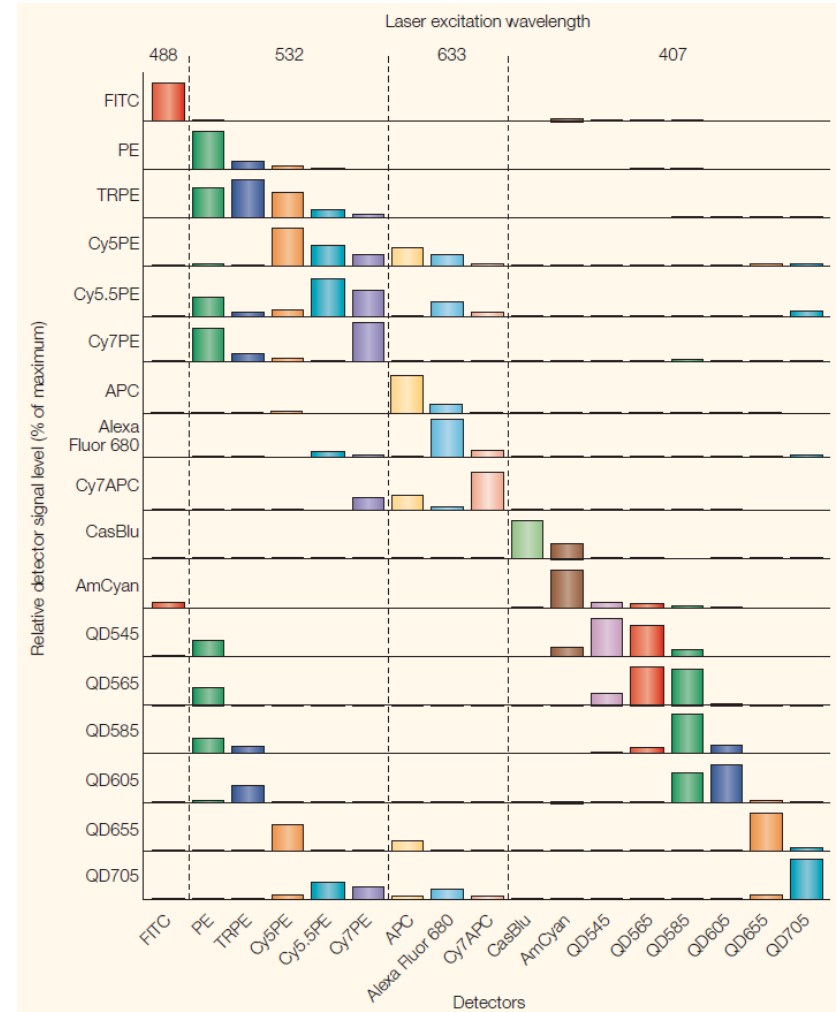
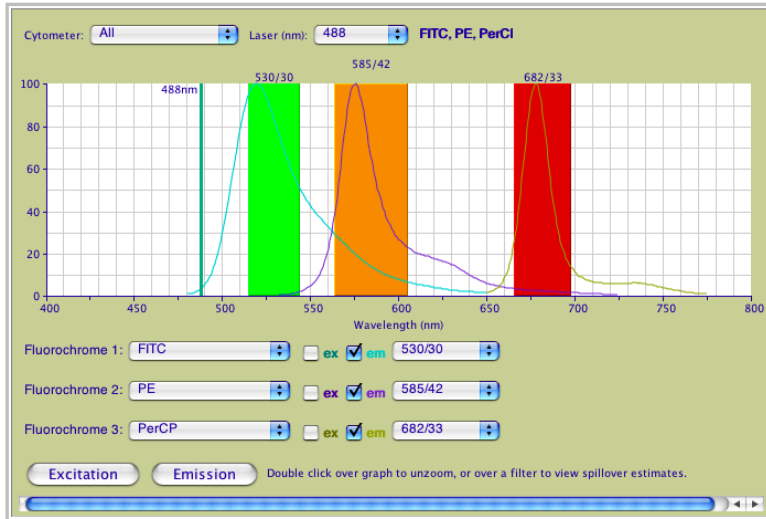


Instrument Evaluation Br

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



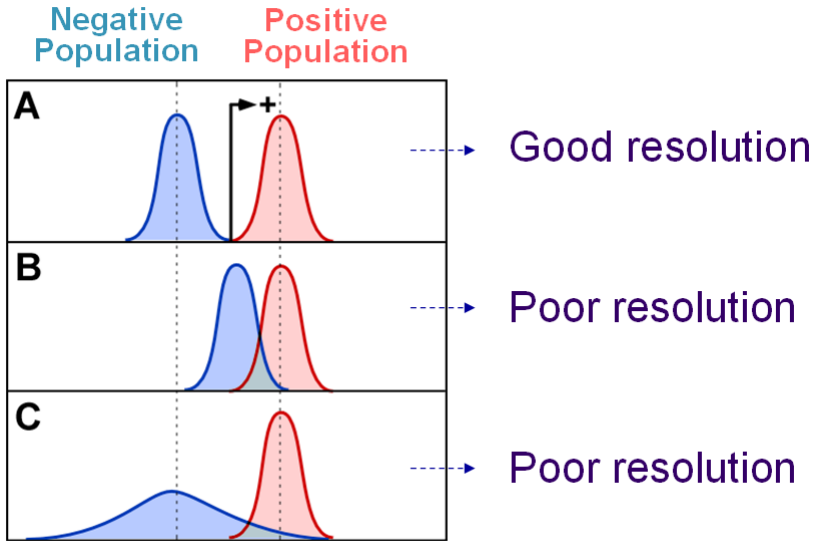
Filter Arrangement and Spectral Overlap (not relevant for element mass cytometry)



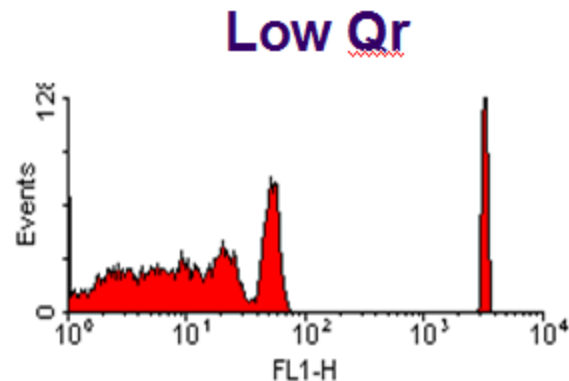
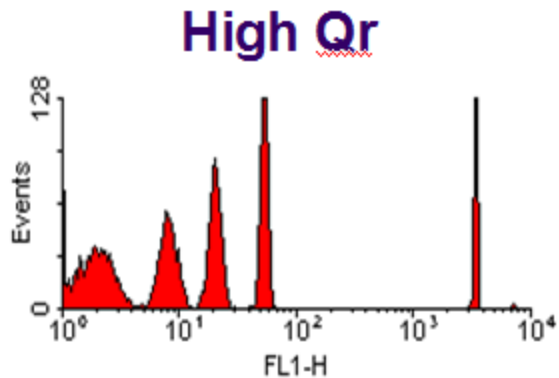
Autissier P et al (2010)
Cytometry 77A, 410ff

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

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

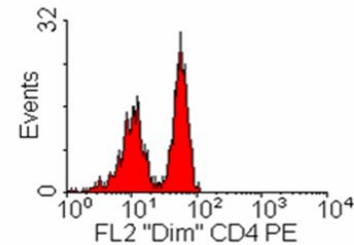
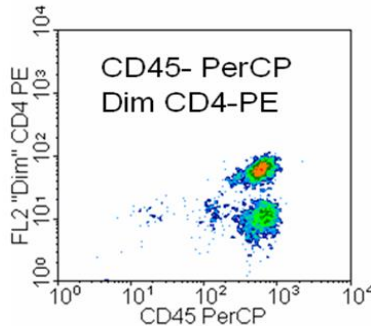
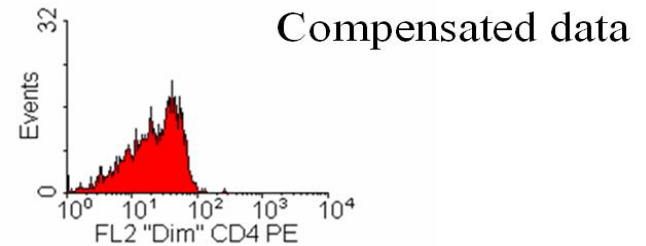
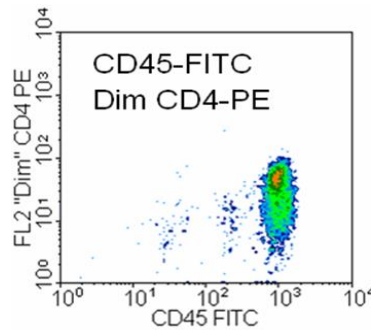
$$\frac{\text{Stain index}}{2 * SD_{neg}}$$

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

- Dye properties (brightness and spectral overlap)

next slide:

- Gain (PMT, CMOS, CCD) settings
- Data Display
- Controls

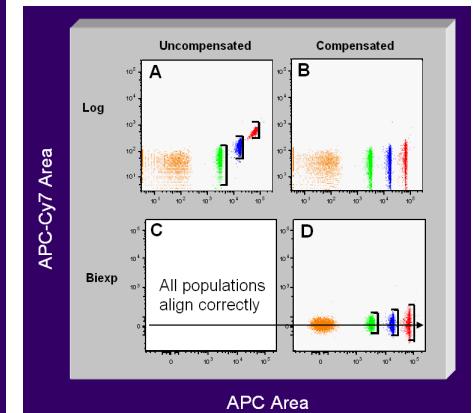
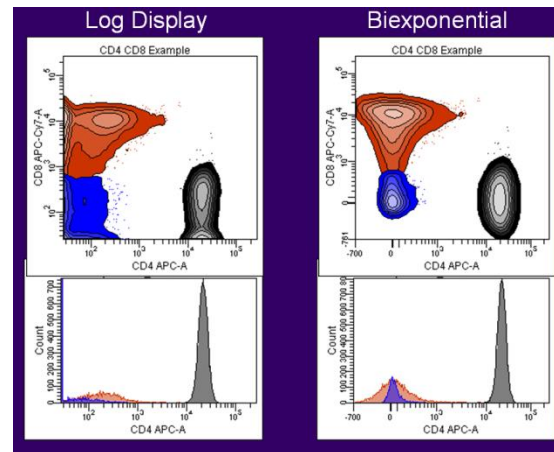
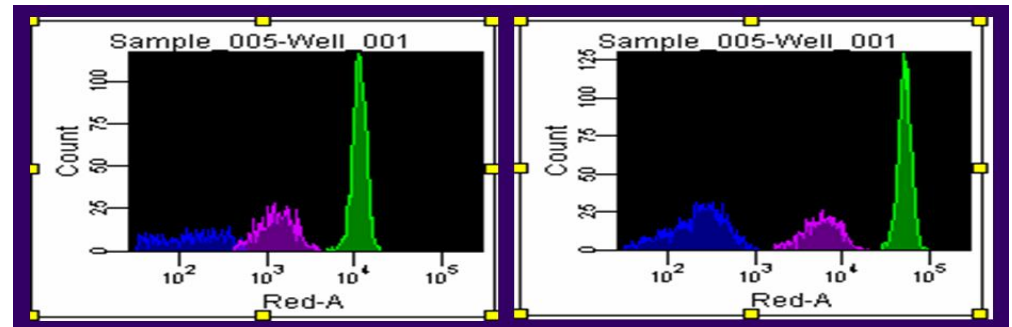


Better separation with less spectral overlap.

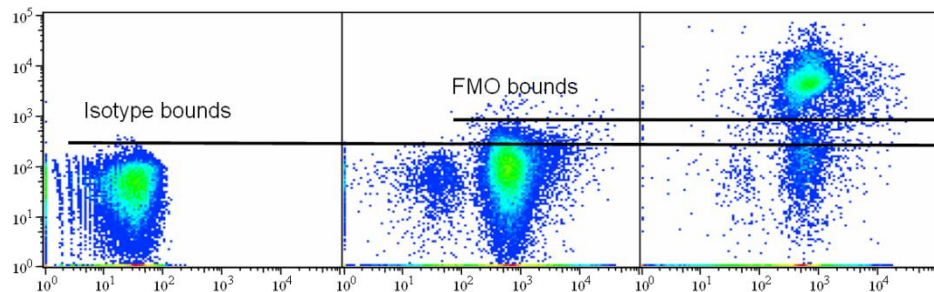
Optimizing cytometry measurements (II)

- Gain (PMT, CMOS, CCD) settings

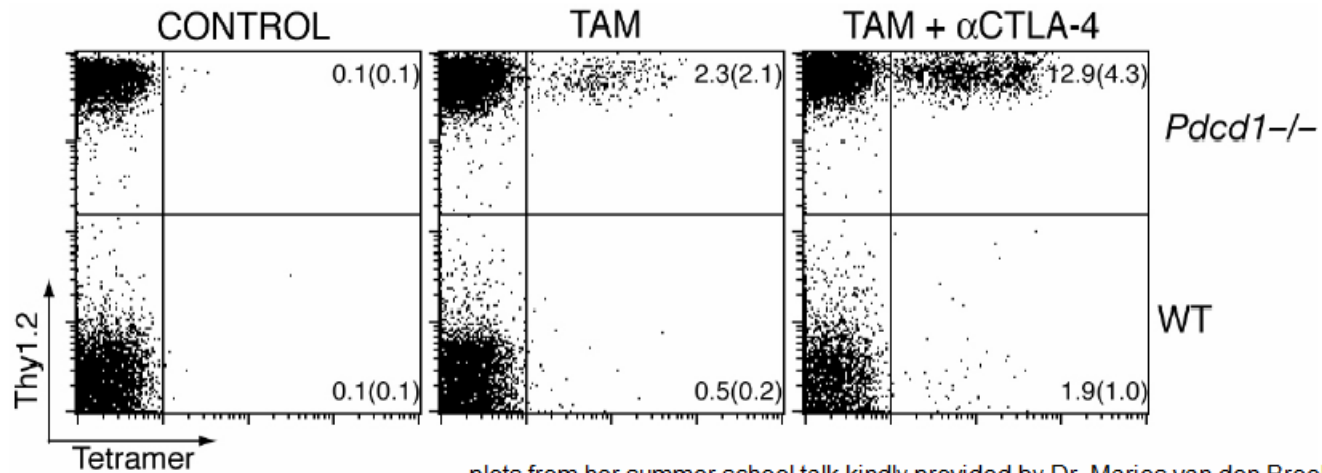
- Data Display



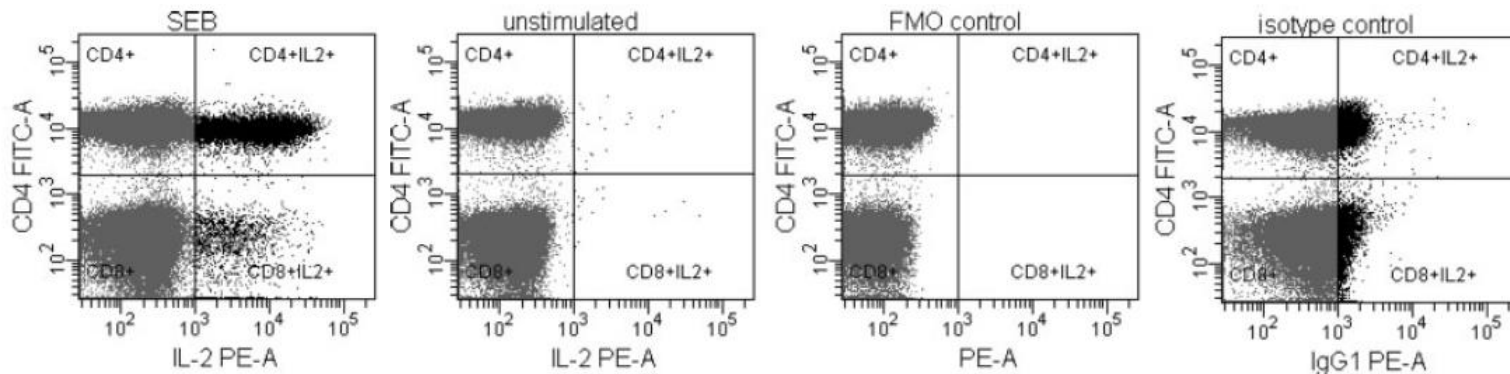
- Controls



Setting Gates and Markers

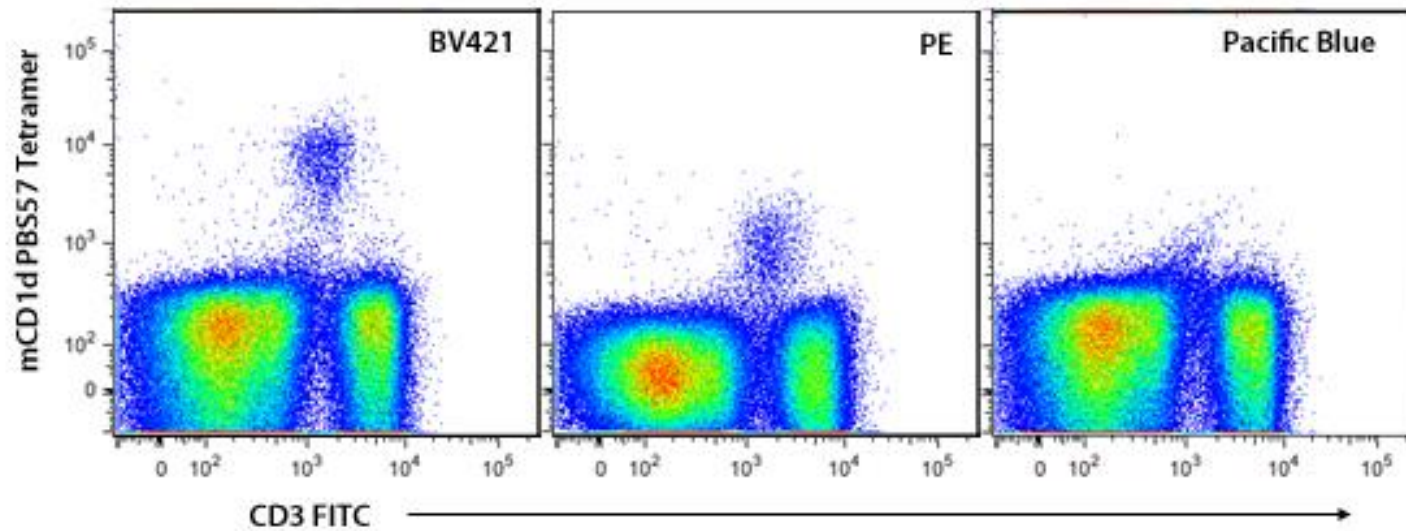


plots from her summer school talk kindly provided by Dr. Maries van den Broek



Adapted from: Maecker HT 2006, Cytometry 69A, 1037ff

Use of Brighter Labels

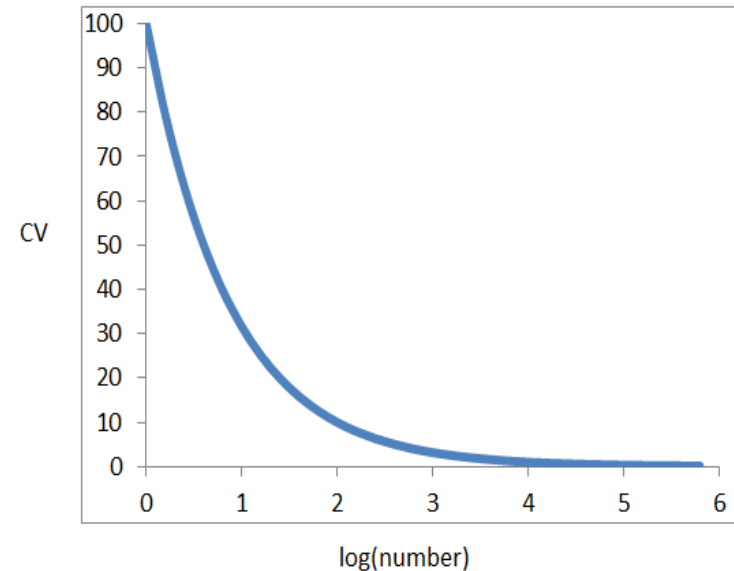


<http://www.biolegend.com/brilliantviolet>

Cell Counting (abs. counts or percentages)

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

Multi-parameter Fluorescence Cytometry Summary

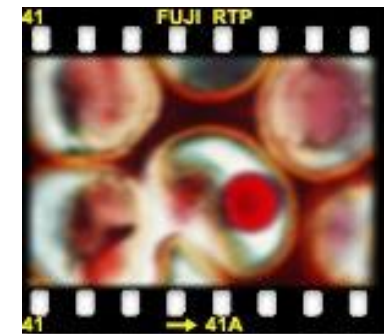
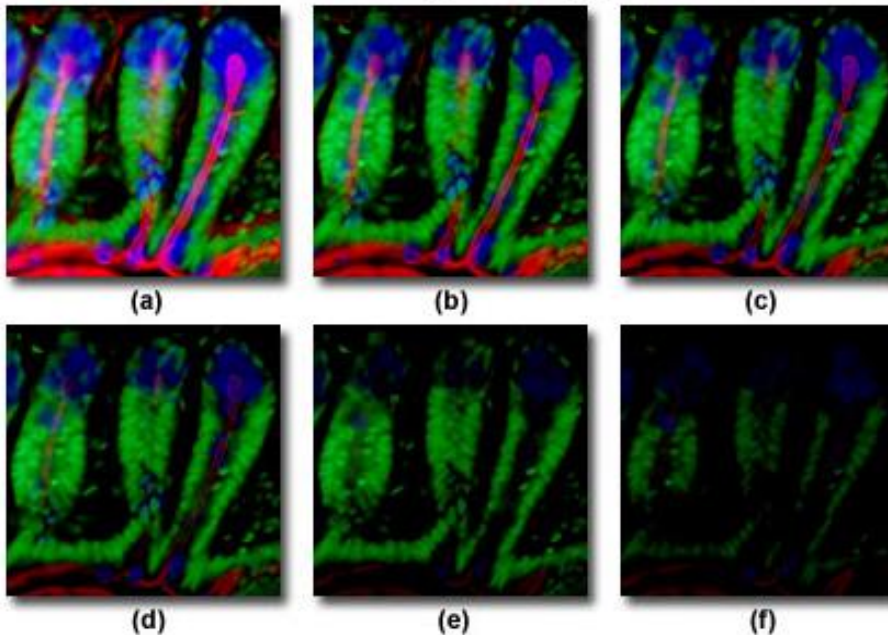
- Know your instrument status e.g. Qr & Br for different channels
- Use high enough gain settings to maximize sensitivity
- An antibody/dye combination that marginally allows discrimination of positives/negatives in a single color assay is unlikely to contribute anything helpful in a multicolor experiment (adequate use of dim and bright labels).
- Avoid spillover from bright cell populations into channels requiring high sensitivity
- Beware of tandem dye degradation
- Internal controls are essential
- Use a gate and marker approach consistent with your experiment objective
- Keep counting statistics limitations in mind

Quantitative Multi-color Microscopy (I)

Additional factors

- Field to field focus
- Photobleaching

Differential Photobleaching in Multiply-Stained Tissues



Out of Focus

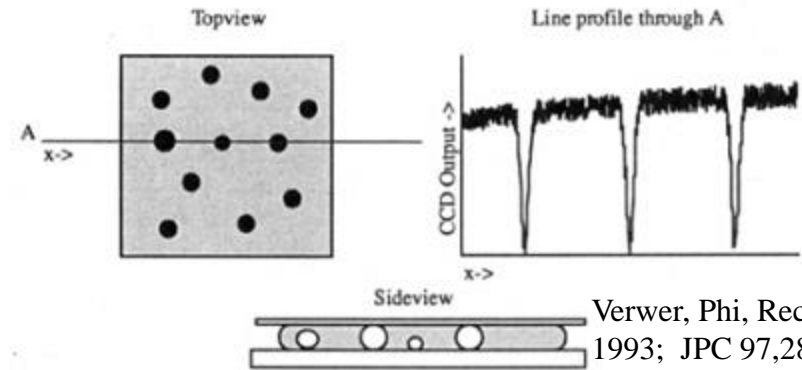
Images from

<http://micro.magnet.fsu.edu/primer/index.html>

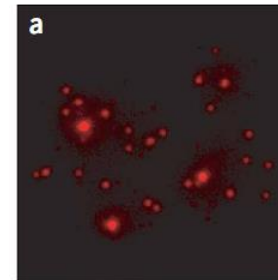
Quantitative Multi-Parameter Microscopy (II)

Selected capabilities

- Intensity calibration by volume exclusion
- Single molecule observation
- Low complexity, low resolution cytometry (Shapiro H, “Cellular Astronomy”)



Verwer, Phi, Recktenwald
1993; JPC 97,2868-70



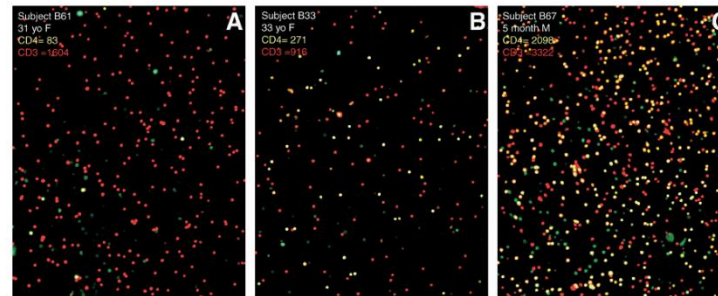
Single m-RNA
molecule analysis.
Robert H Singer's
group, Nature S&MB
2008



+



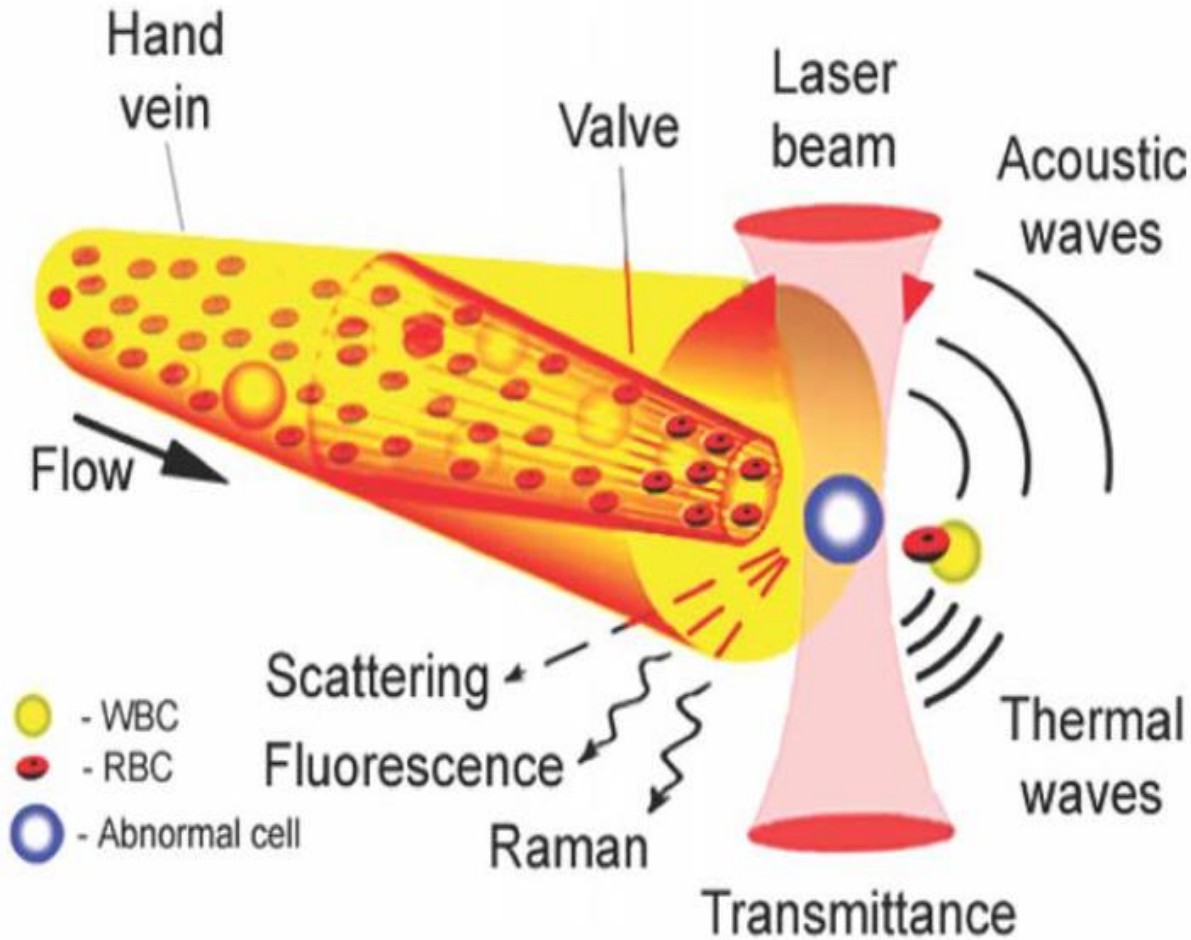
=



Rodriguez WR, McDevitt JT, PLoS Medicine 2005; (CD3+CD4+ yellow, CD3+CD8+ red, monocytes green)

In-vivo Multi-parameter Cytometry

Single cell analysis in living animals



Issues:

- tissue optics
- object motion
- flow rate
- Labelling
- ...

Intravital Microscopy

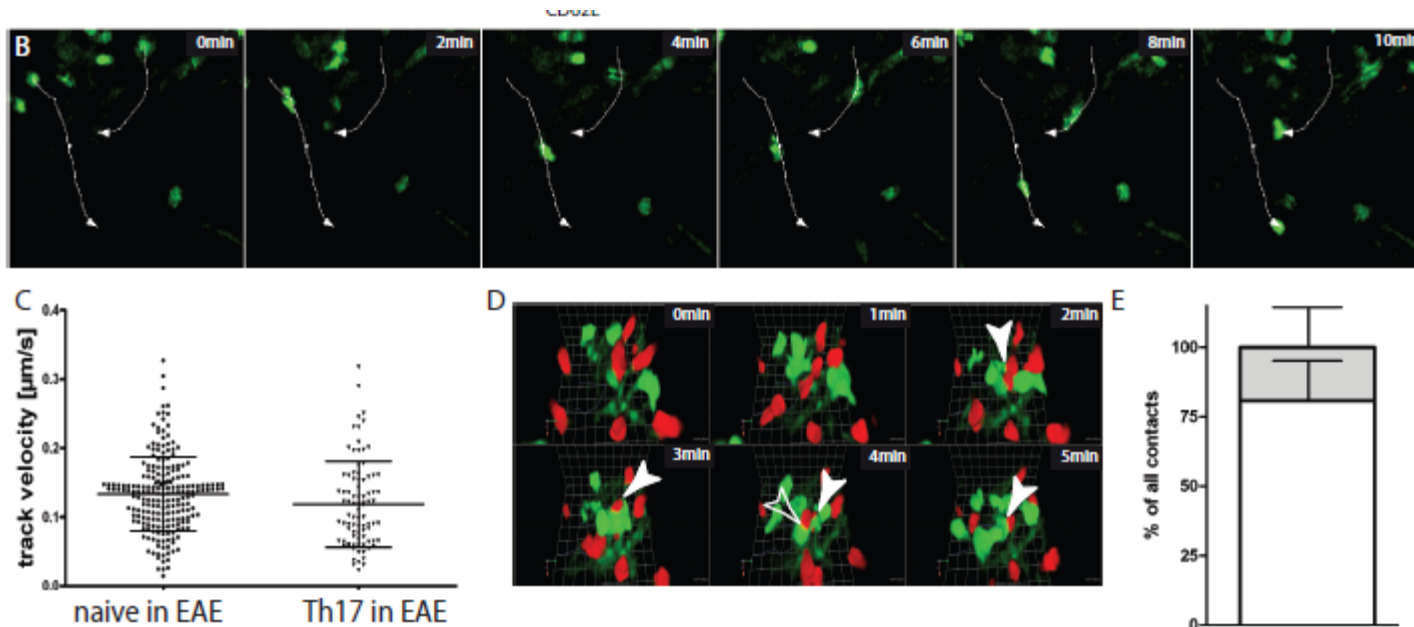


Figure 1 Intravital imaging reveals high motility of naïve CD4⁺ T cells in the inflamed CNS. Naïve OT2 EGFP T cells were intravenously injected into EAE affected mice at the peak of disease (clinical score 2.5) or locally applied onto the imaging field. Intravital TPLSM on the brain stem of these mice was performed 12 - 24 hours after naïve T cell injection or 30 minutes after local application. Adoptive EAE was induced by transfer of *in vitro* generated encephalitogenic 2d2 Th17 T cells into *C57BL/6 RAG1*^{-/-} mice. **(A)** T cell phenotype of MACS isolated naïve T cells was confirmed by FACS analysis prior to experiments. Surface antigen expression of CD62L, CD25, CD69 and CD44 was determined on CD4⁺ lymphocytes. **(B)** A representative time lapse series derived from intravital TPLSM demonstrates rapid movement of naïve T cells deep in CNS tissue (100-150 μm). Two cell tracks are shown exemplarily by white arrows. For further details see also Additional File 1 (Scale bar: 10 μm). **(C)** Cell track velocities of naïve OT2 (N = 212) and effector 2d2 Th17 (N = 87) cells at the peak of disease in the inflamed CNS were quantified. The mean track velocities from 4 independent experiments are shown (\pm SD). **(D)** Contacts (arrowheads) between encephalitogenic 2d2 Th17 effector T cells (EGFP,green) and naïve OT2 (tdRFP red) could be observed during intravital TPLSM, as revealed by the reconstructed 3D time lapse series (80-110 μm). These interactions were mainly short and random like (open arrowhead) although some static long-lasting contacts (filled arrowhead) could be also detected. **(E)** To quantify effector-naïve T cell interactions we analyse the co-localisation area of EGFP and tdRFP as previously described [19]. We discriminated short (random) contacts (< 5 min) from long-lasting (most probably non-random) interactions (\geq 5 min) and observed that $81\% \pm 14$ formed short interactions with effector T cells (white bar) and $19\% \pm 14$ formed long-lasting interactions. Data are shown as percentage of all contacts from two independent experiments (\pm SD).

Conclusions

Multi-parameter cytometry

Optimized flow and imaging single cell cytometry with adequate bio-informatics tools provide quantitative molecular measurements of biological processes at organism, cellular and sub-cellular level.

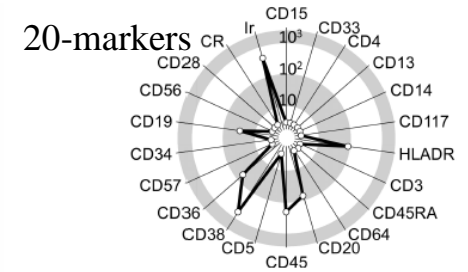
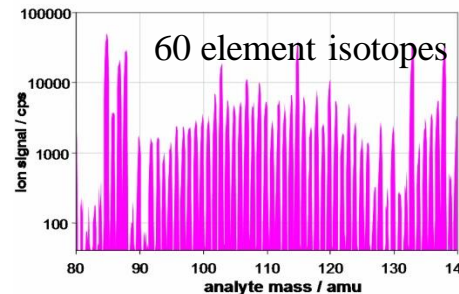
New developments in many areas have simplified the tools for the biologist.

Evolving Technologies for Cytometry

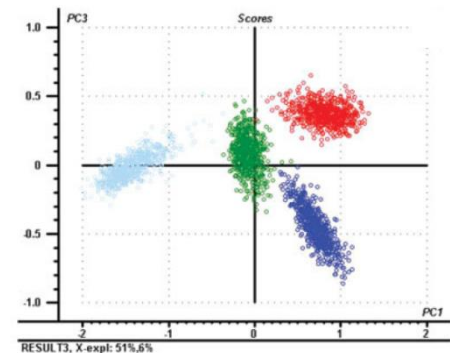
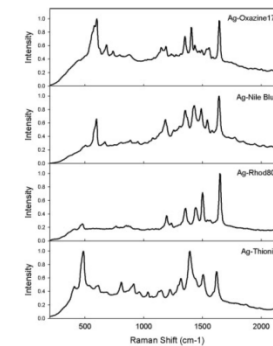
- Multi-parameter cytometry approaches
- High speed imaging in flow
- Single cell sorting
- Low complexity cytometers for cell (subset) counting
- “Label-free” cell analysis
- Sample preparation micro-fluidics
- Fluorescent polymers as labels for high sensitivity
- Affinity reagents (antibodies)

New Developments for in-vitro Multi-parameter Cytometry

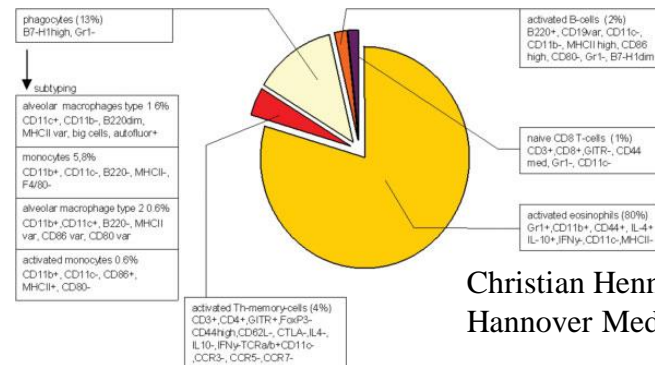
- **Element-Label Flow Cytometry (CyTOF, addresses fluorescence spectral overlap issue by using elements as labels, Anal. Chem., 2009, 81 (16), pp 6813–6822)**
- **Sequential Stain De-stain Cytometry (Cytometry, 2009, 75A(4), pp 362-370)**
- **SERS-Label Flow Cytometry (uses spectral fine-structure to distinguish labels, Cytometry, 2008, 73A(2), pp 119-128)**
- **SONY spectral analysis**



Scott Tanner, DVS Sciences Inc



John Nolan, La Jolla Bioengineering Institute

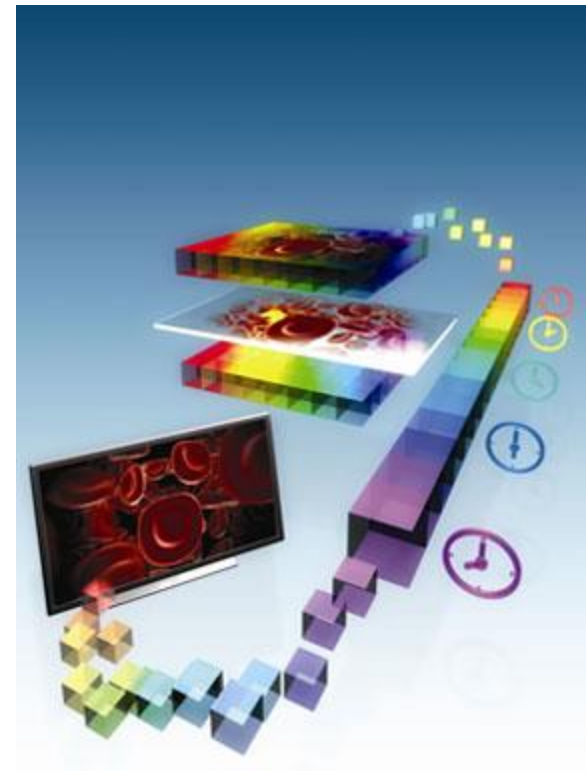


31-marker analysis

Christian Hennig, ChipCytometry
Hannover Medical School

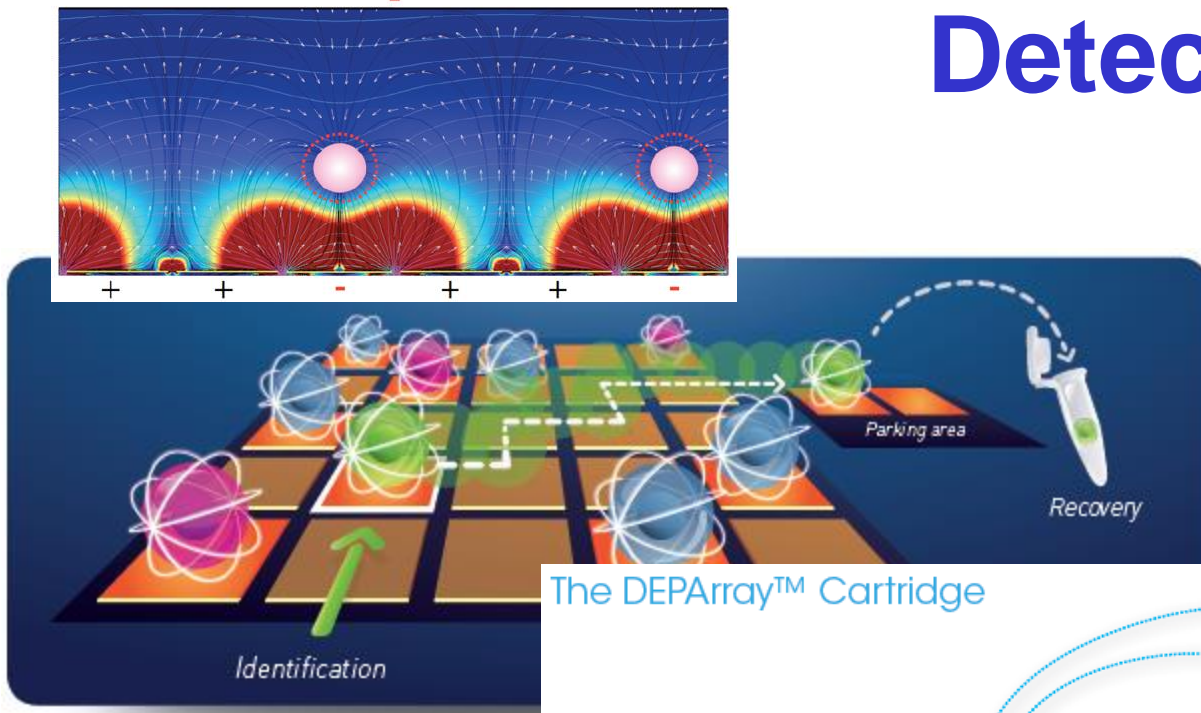
High speed imaging in flow

- ImageStream (EM Merck)
- Bahram Jalali group, UCLA
- ...



<http://www1.ee.ucla.edu/Research-highlights-jalali-4.htm>

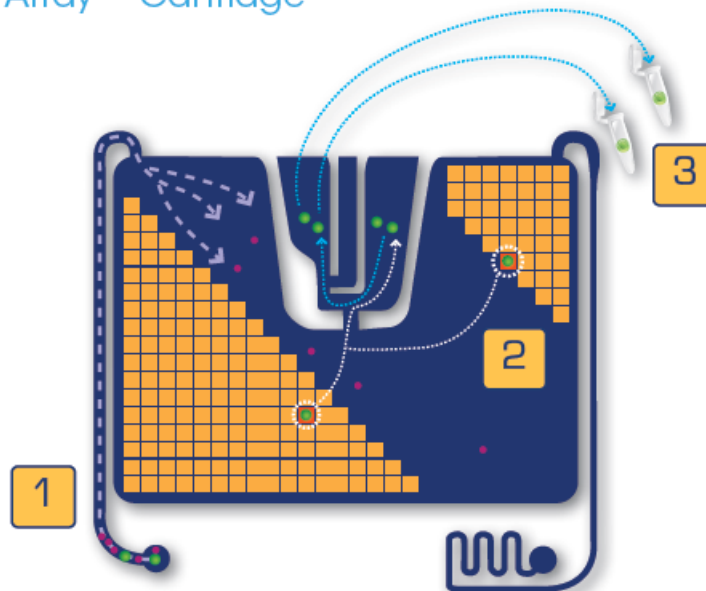
Single Cell Sorter with Microscopic Detection



The DEPArray™ Cartridge

Cells are transferred to a special slide with 40,000 “cages”. Cells of interest are identified by fluorescence microscopy and sorted by the instrument.

Cell movement with dielectric forces.
DEPArray
Silicon Biosystems,
Bologna, IT



1

Inject, trap and image cells

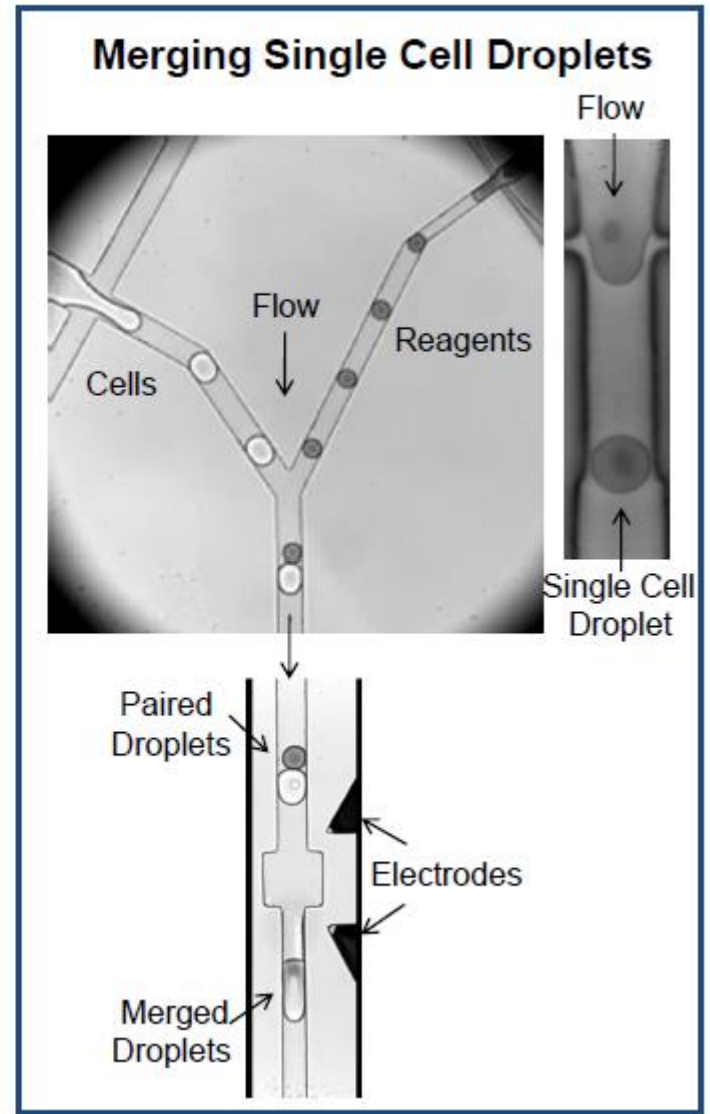
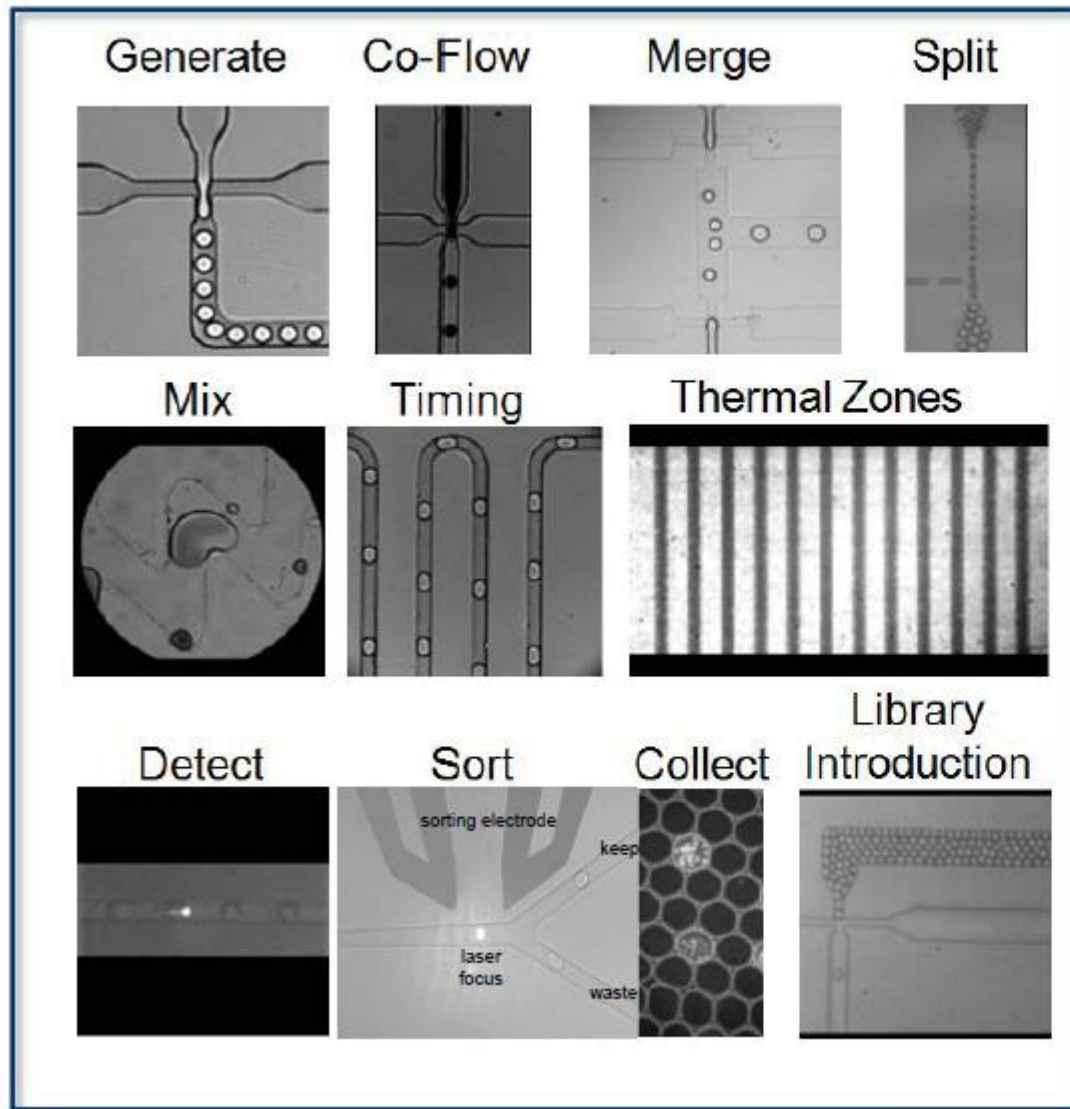
2

Move cells of interest into parking chamber

3

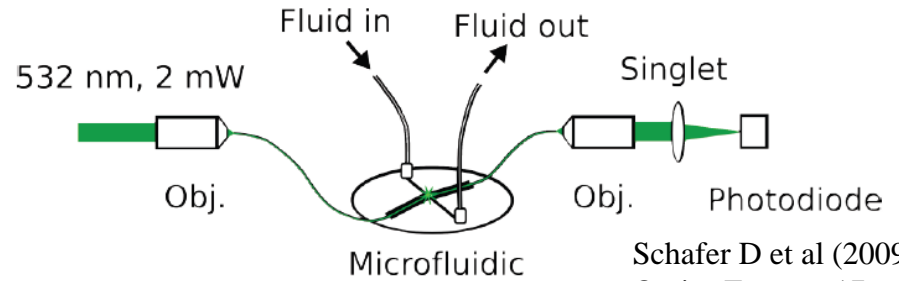
Move individual or multiple cells into recovery chamber and flush

Advanced Single Cell Analysis in Droplets



Low-complexity Cytometers for Cell Counting

Low-end cell counters



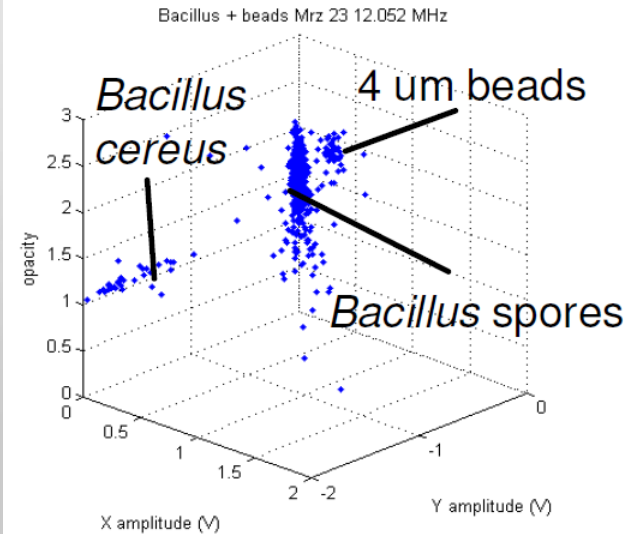
Schafer D et al (2009)
Optics Express 17, 6068ff



Merck Millipore MUSE™

Label-free Cell Analysis

LEISTER : Axetris Impedance flow cytometry



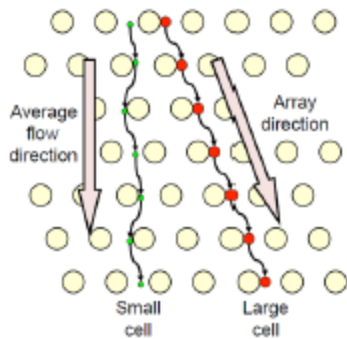
Marco DiBerardino, Leister Axetris

Electrical parameters of living cells (no label required).

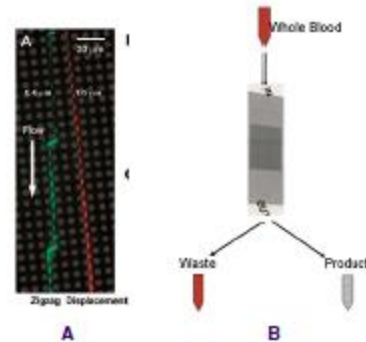
Other parameters: fluorescence polarization, fluorescence lifetime, compressibility, ...

Innovative Sample Preparation

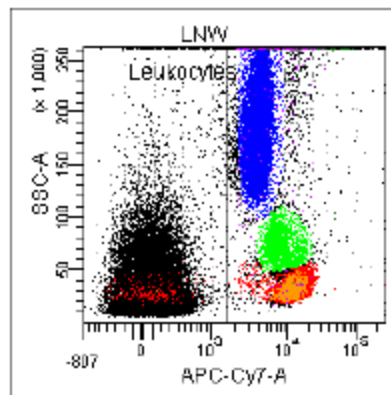
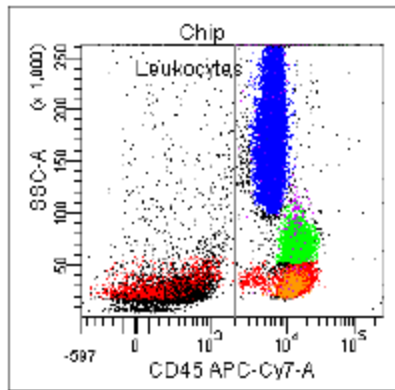
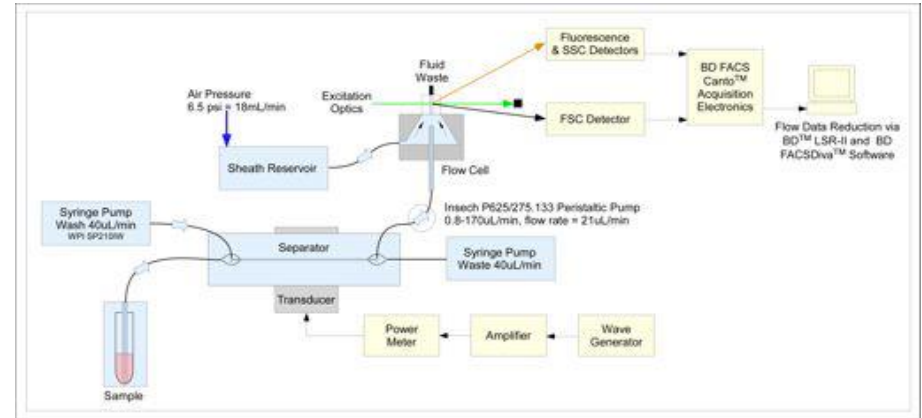
Microfluidic system
for leukocyte isolation
(deterministic lateral displacement)



Chip and new blood separation process



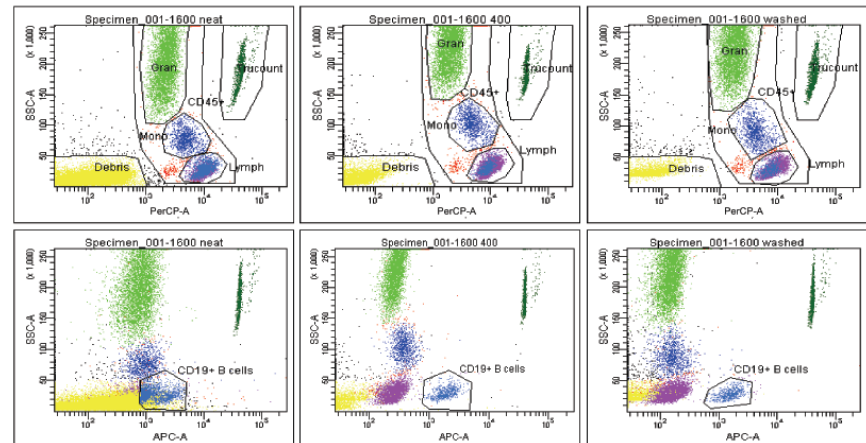
Acoustic particle focusing
for cell washing



Lyse no wash

Chip wash

Centr. wash



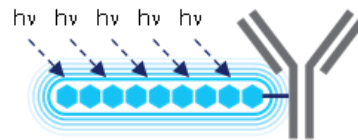
Cyto 2012 poster, Liping Yu et al, GPB and BD Biosciences

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

Bright Fluorescent Polymer Dyes

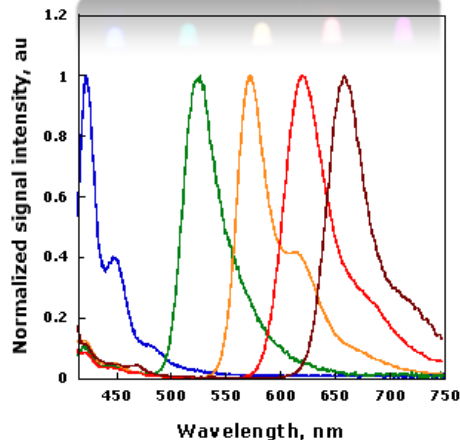
Polymer Based Fluorochromes

- Well defined synthetic organic polymer structures
 - Single conjugation site, defined size, etc.
- Backbone comprised of π -conjugated repeat units
 - Affords massive light harvesting ($\epsilon > 10^6$) materials with high quantum yields
- Tunable architecture adapted for low NSB, high aqueous solubility and spectral performance



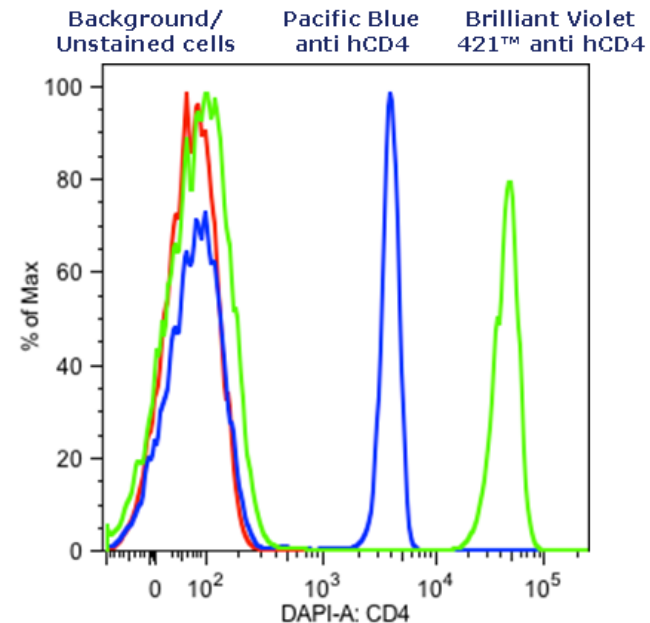
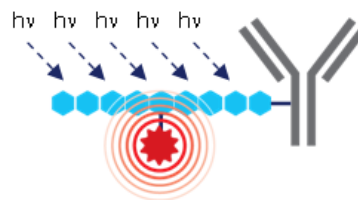
Brilliant Violet 421™

- PE level performance w/ 405nm Laser
- >10x the Stain Index of Pacific Blue
- Enables detection of low abundance targets in multicolor assay panels (e.g. CD56, CD127, etc.)
- Wide range of Ab clones validated



Brilliant Violet Tandems

- Provides a wider range of colors spanning the visible spectrum
 - >6 unique colors validated
- Chemically controlled ratio of donor/acceptor provides:
 - Reproducible performance
 - Low (<5%) compensation at 450nm



Novel Affinity Reagents

- **Antibodies**
 - Antibodies from different species (e.g. Llama 15 kDalton fragments with 10^{-9} M Kd and high stability, potential for intracellular use)
 - Recombinant antibody fragments
 - ...
- **Synthetic affinity reagents**
 - Aptamers
 - Protein scaffolds
 - Molecular Imprinted Polymers

Recent review: Fodey T et al; Trends in Anal. Chem. 30(2011) 254ff

Conclusions

Evolving Technologies

Technology developments in the physical sciences provide tools for deeper molecular understanding of living systems.

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 - Brent Gaylord, Sirigen > BD
 - Collette Rudd, Thermo
- above all BD

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