

AKADEMIE FÜR IMMUNOLOGIE

SPRING SCHOOL ON IMMUNOLOGY

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Principles of Flow and Image Cytometry & New Technologies for Single Cell Analysis

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Key Reference Abbreviation

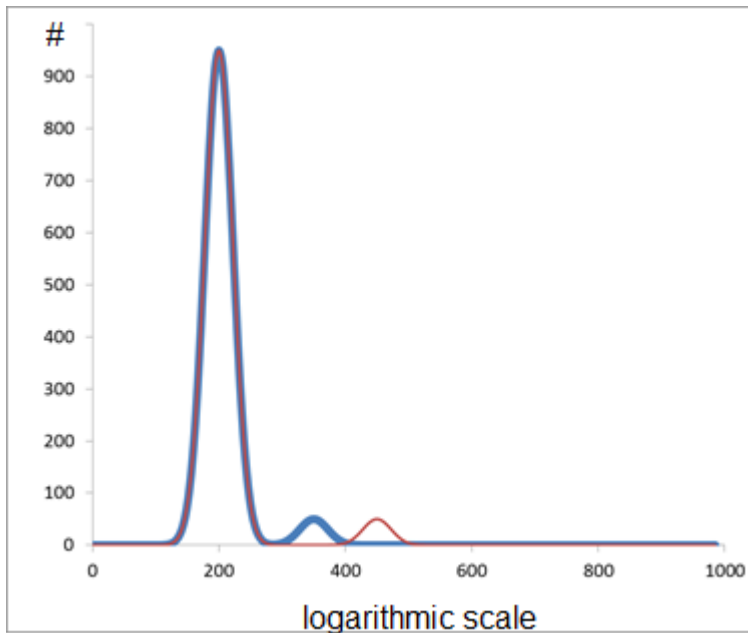
In this document “GUIDELINES” is used for
Cossarizza, Andrea, et al. (2017) "Guidelines for the use of
flow cytometry and cell sorting in immunological studies."
European journal of immunology 47 (10) 1584 - 1797

The GUIDELINES contain contributions from 231 experienced
scientists from 192 institutes worldwide, describing their
recommendations for the optimal use of flow cytometry.

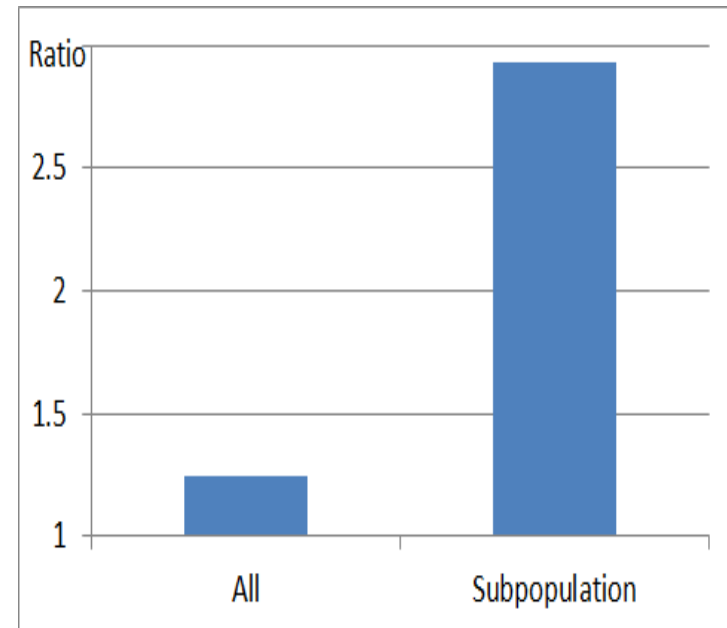
References with a DOI number can be located with an Internet
search.

Why Cell Subset Analysis at the Single Cell Level

Intensity Histogram



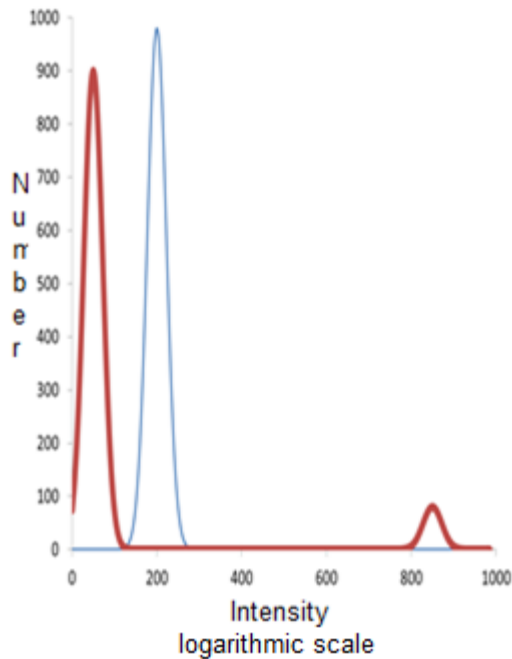
Intensity Ratios



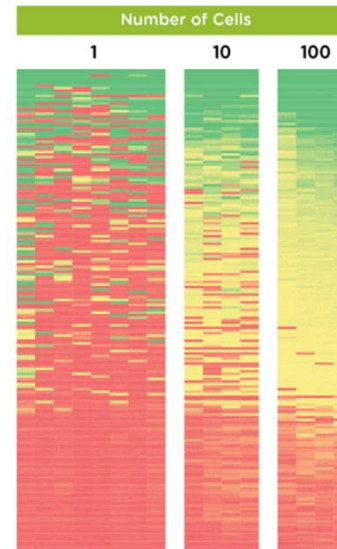
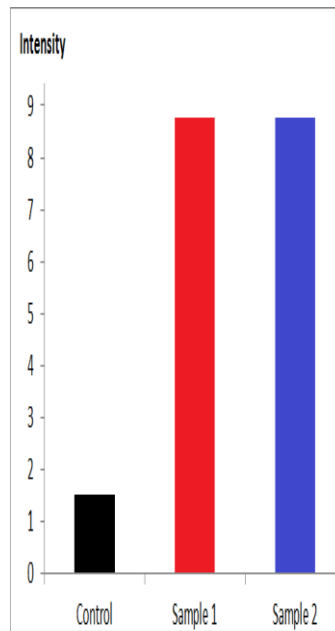
Subpopulation analysis detects changes better, especially for rare subpopulations.

Why Single Cell Analysis

Intensity Histogram for Single Particles



Intensity per Sample



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

Cell by cell intensity analysis detects population heterogeneity.

Flow Cytometer Components

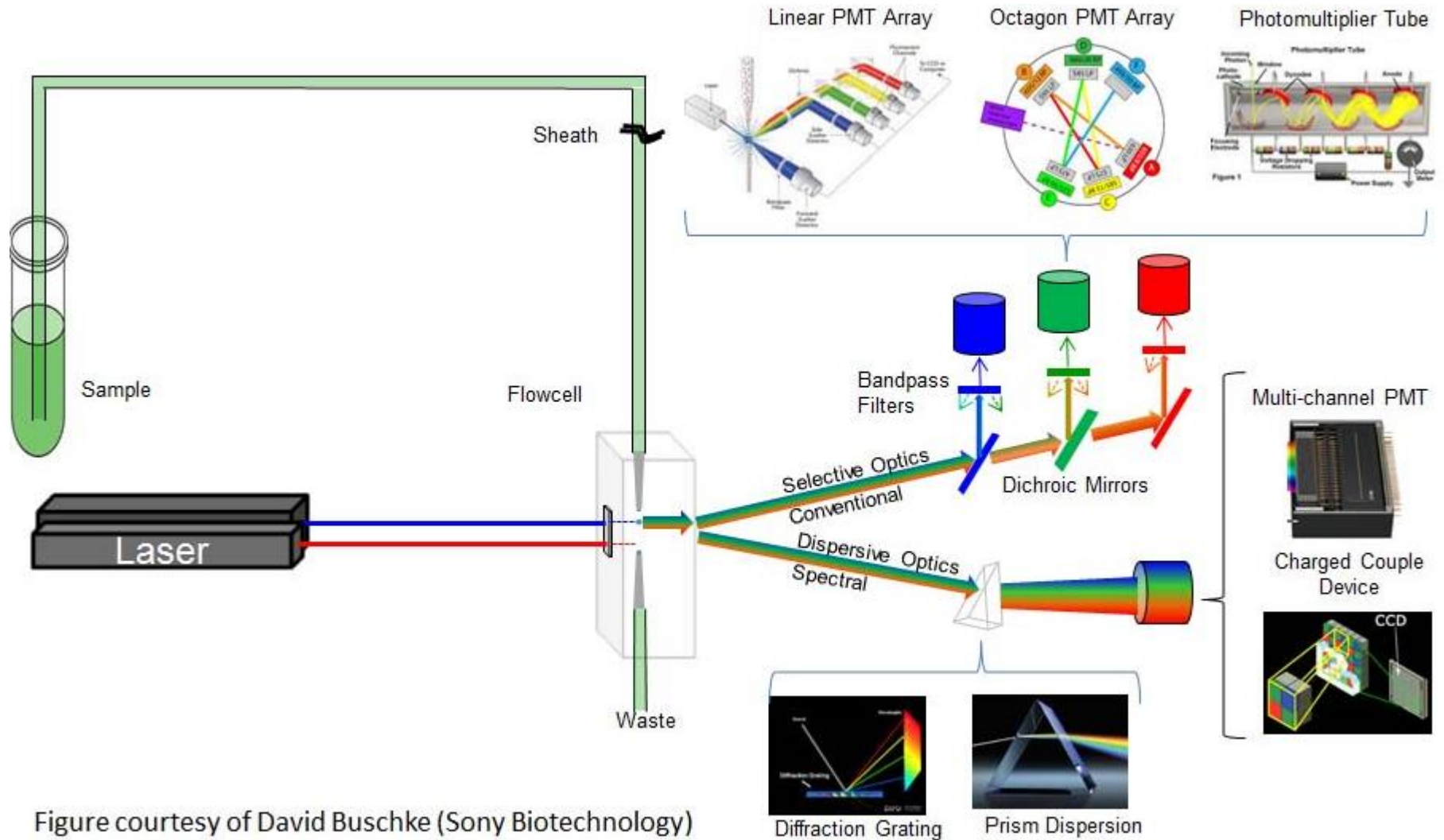


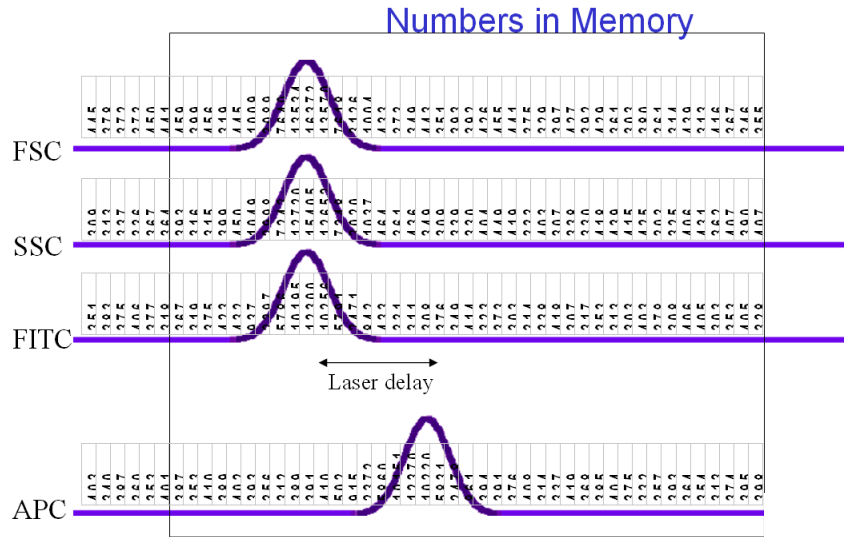
Figure courtesy of David Buschke (Sony Biotechnology)

GUIDELINES Flow cytometers, pages 1596- 1608

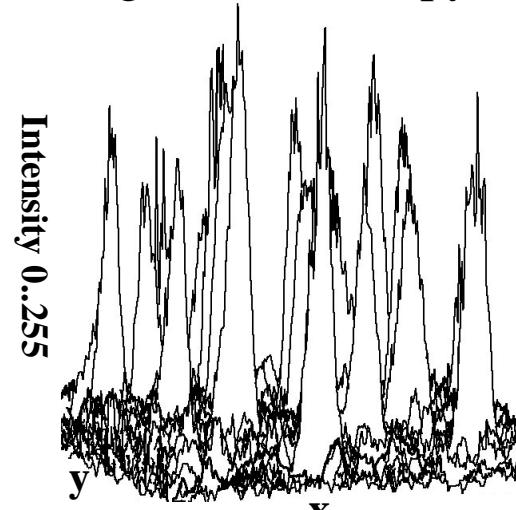
Dichroic filters vs. Multispectral cytometry: Feher K et al.(2016) Cytometry 89A: 681-9

Basic Data Processing

Flow Cytometry

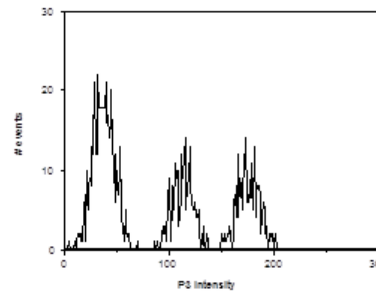


Digital microscopy

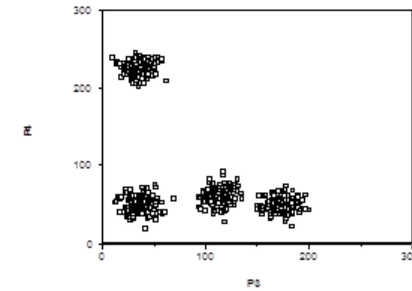


Cell	P1	P2	P3	P4	P5	Pop#
1	242	135	704	175	612	1
2	146	132	690	178	566	1
3	269	147	89	206	580	3
4	442	143	399	250	255	4
5	212	167	155	926	526	2
6	269	2	659	207	575	1
7	204	232	112	171	679	3
8	152	74	160	828	532	2
			...			
9997	215	119	138	936	662	2
9998	244	50	72	261	543	3
9999	214	137	174	1014	597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

Event histogram



"Dotplot"



for >2 parameters: gating, cluster analysis, ...

For many samples and parameters: bioinformatics

N. Aghaeepour et al. (2013) Nature Methods 10:228ff

E.Lugli, M.Roederer, A.Cossarizza (2010) Cytometry 77A:705ff

GUIDELINES Data Analysis, pages 1651-65

Hi-Parameter Data Processing

TOOL	PURPOSE
PhenoGraph	Clustering
X-Shift	Clustering
ACCENSE	Clustering
DensVM	Clustering
FlowSOM	Clustering
SPADE	Clustering
Citrus	Clustering, differential abundance analysis
Cydar	Clustering, differential abundance analysis
ACDC	Cell type assignment
SCAFFoLD	Cell type assignment, cellular trajectory mapping
Statistical SCAFFoLD	Cell type assignment, cellular trajectory mapping, differential abundance analysis
Wanderlust	Cellular trajectory detection
DREMI/DREVI	Cellular trajectory detection
t-SNE	Dimensionality reduction

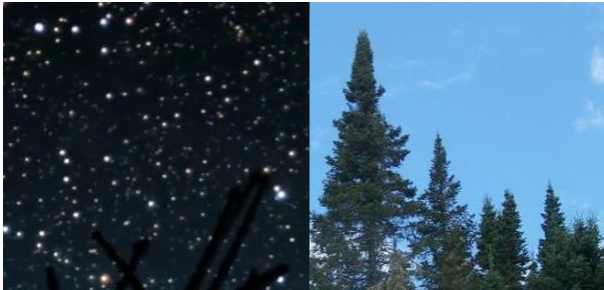
Olsen L R et al. (2019)

DOI: 10.1002/cyto.a.23621

more at: Palit S et al. (2019) doi:<http://dx.doi.org/10.1101/473215>

Mair F et al. (2016) doi: 10.1002/eji.201545774 (mini-review by Monday speaker)

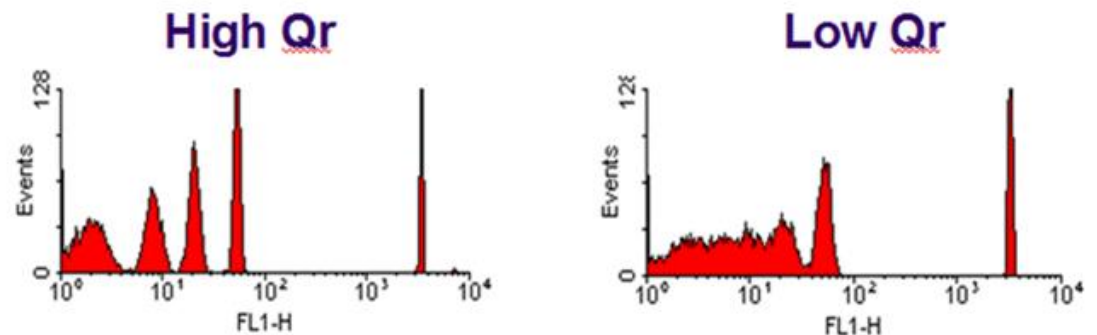
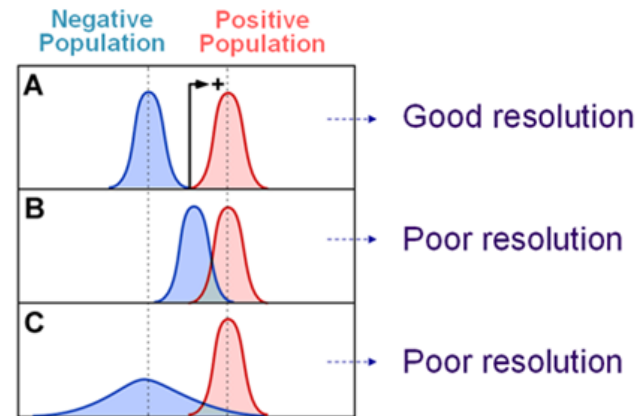
Instrument Evaluation Br, Qr



Br, optical background from

- Cell autofluorescence
- Flow cell reflections
- Ambient light
- Free fluorochrome
- Raman scatter
- Spectral overlap

Qr, photon detection efficiency



Signal Overlap and “Compensation”

Calculation of concentrations from optical/mass intensities

$$I_1 = a_{11} * c_1 + a_{12} * c_2 + a_{13} * c_3$$

$$I_2 = a_{21} * c_1 + a_{22} * c_2 + a_{23} * c_3$$

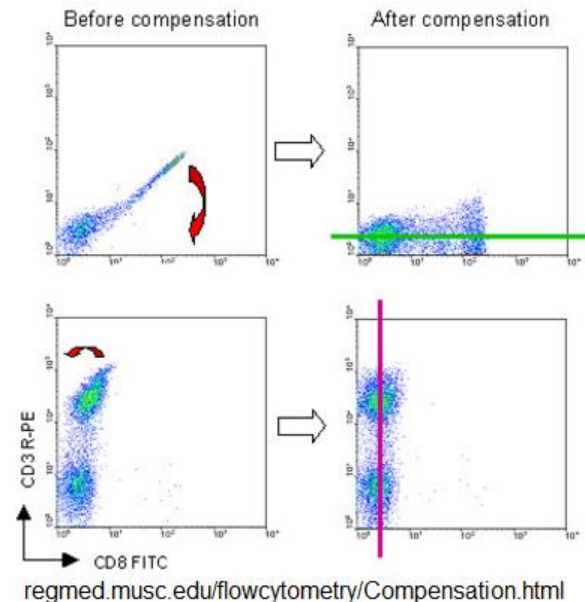
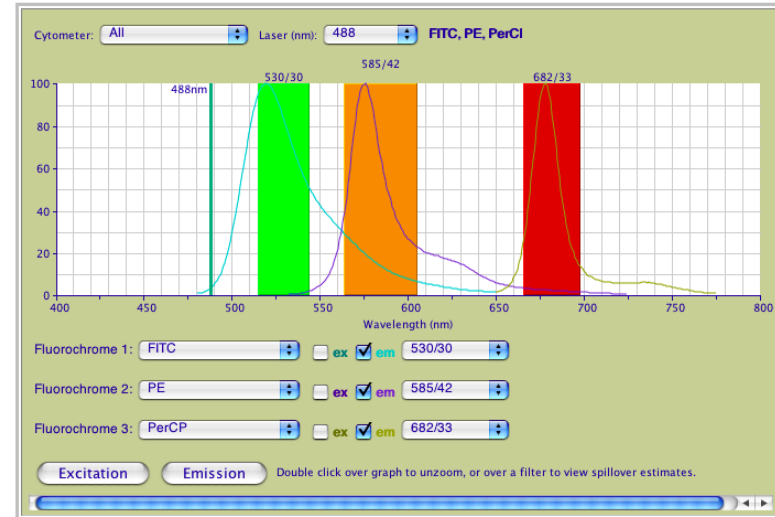
$$I_3 = a_{31} * c_1 + a_{32} * c_2 + a_{33} * c_3$$

a_{ik} : “compensation” matrix numbers

I_i : measured intensities

c_k : label concentrations

Solve n equations with n unknowns
(in spectral cytometry generally many more equations than unknowns)



Cytometer Measurements

Single Cell Analysis

- * Cell-concentration
- * Cell size
- * Cell arrangement in clusters
- * Mass of multiple cellular components per cell
- * Distribution of component mass in subsets
- * Temporal change of the above parameters
- * Subset fractions
- * Cell shape

Non-cell applications

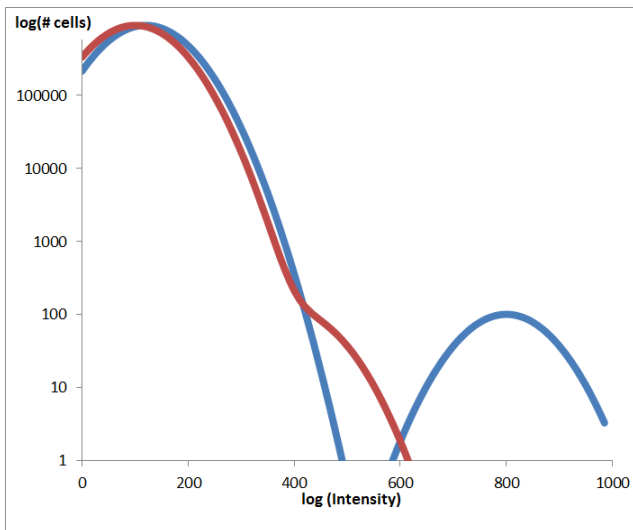
- * Highly multiplexed bead-based immunoassays
- * Single molecule counting

Label Selection

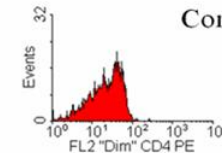
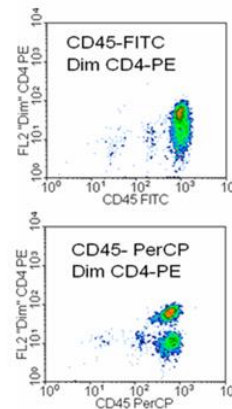
- Detection System
- Brightness
- Spectral Overlap
- Application (surface vs. internal)

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

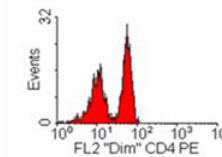
$$\frac{\text{Medium}_{pos} - \text{Medium}_{neg}}{2 * SD_{neg}}$$



Brightness and Separation



Compensated data



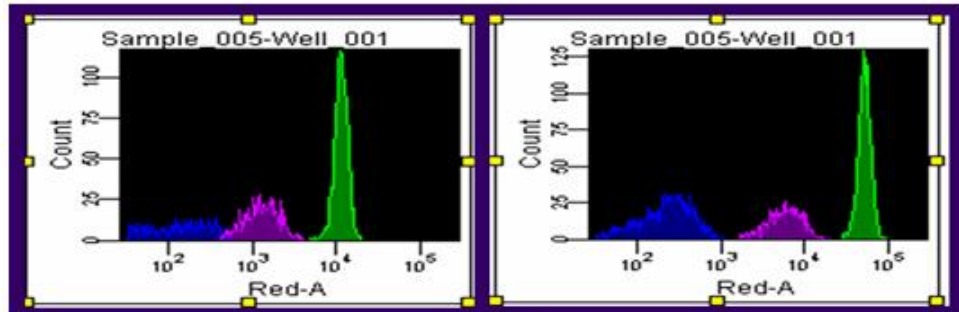
Better separation with less spectral overlap.

Spectral Overlap and Separation

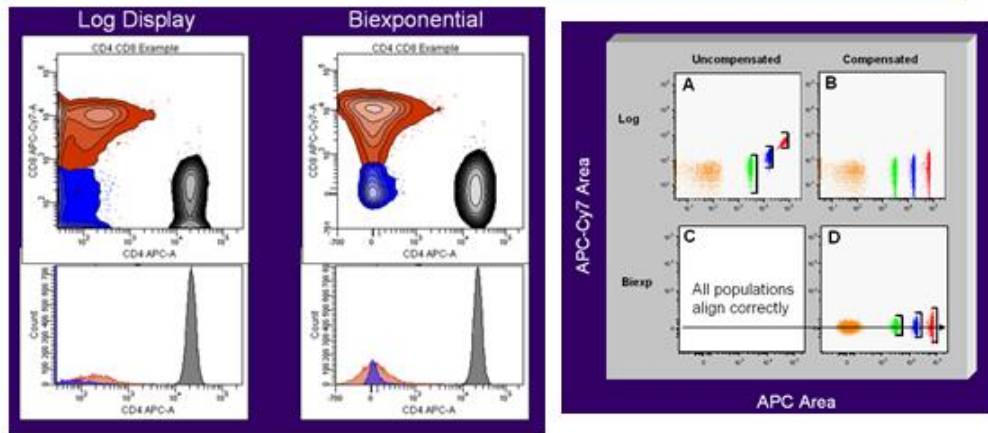
More info: Maecker HT et al. (2004) Cytometry 62A:169-173

Optimizing cytometry measurements

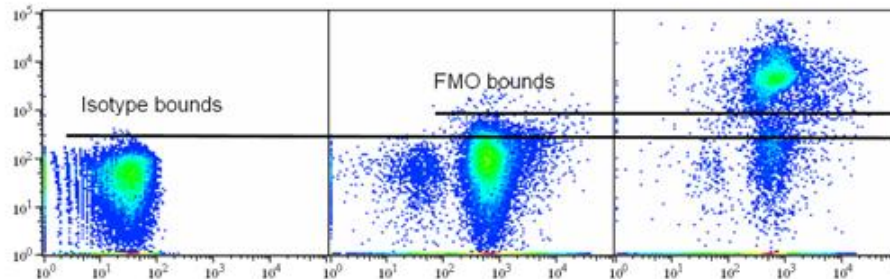
- Gain (PMT, CMOS, CCD) settings



- Data Display



- Controls

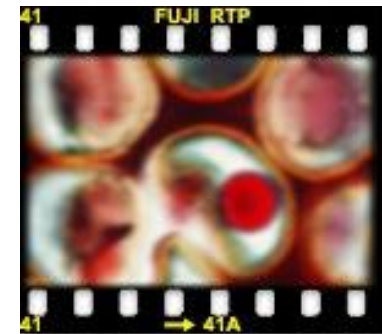
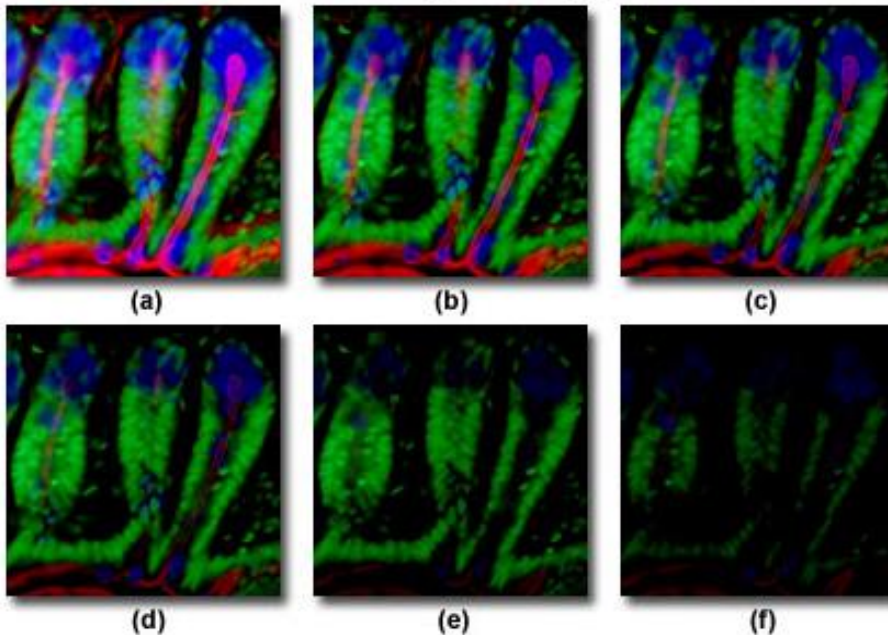


Quantitative Multi-color Microscopy

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>

Multiparameter Cell Analysis

- Imaging
 - Immunofluorescence (IF)
 - Sequential Stain-Destain IF
 - ...
- Flow Cytometry
 - IF, SERS labels
 - CyTOF
 - Sequence coding
 - Direct Raman
 - Label-free Impedance
 - Imaging e.g. Imagestream
- In-vivo Imaging and Flow Cytometry

Fluorophore	Laser Line, nm	Max Ex, nm	Max Em, nm	Relative Brightness
BD Horizon Brilliant Ultraviolet 395	355	348	395	
Alexa Fluor 350	355	340	440	
BD Horizon Brilliant Ultraviolet 496	355	348	496	
BD Horizon Brilliant Ultraviolet 563	355	348	563	
BD Horizon Brilliant Ultraviolet 661	355	348	661	
BD Horizon Brilliant Ultraviolet 737	355	348	737	
BD Horizon Brilliant Ultraviolet 805	355	348	805	
DyLight 405	405	400	420	
Alexa Fluor 405	405	401	420	
BD Horizon Brilliant Violet 421	405	407	421	
eFluor 450	405	405	445	
BD Horizon V450	405	404	448	
Super Bright 436	405	414	436	
Pacific Blue	405	401	452	
BD Horizon Brilliant Violet 480	405	436	478	
BD Horizon V500	405	415	500	
BD Horizon Brilliant Violet 510	405	405	510	
BD Horizon Brilliant Violet 570	405	407	574	
Super Bright 600	405	414	600	
BD Horizon Brilliant Violet 605	405	407	602	
Super Bright 645	405	414	645	
BD Horizon Brilliant Violet 650	405	407	650	
Super Bright 702	405	414	702	
BD Horizon Brilliant Violet 711	405	407	711	
BD Horizon Brilliant Violet 786	405	407	786	
BD Horizon Brilliant Blue 515	488	490	515	
DyLight 488	488	493	518	
Alexa Fluor 488	488	495	519	
FITC	488	490	525	
PerCP	488	490	675	
BD Horizon Brilliant Blue 700	488	485	693	
PerCP-Cy5.5	488	490	695	
DyLight 550	561	562	576	
PE	488	561	496/546	578
PE-eFluor 610	488	561	496/546	607
PE/Dazzle 594	488	561	496/546	610
PE-Alexa Fluor 647	488	561	496/546	667
PE-Cy5	488	561	496/546	667
PE-Cy5.5	488	561	496/546	695
PE-Alexa Fluor 750	488	561	496/546	779
PE-Cy7	488	561	496/546	785
eFluor 680	640	633	680	
APC	640	650	661	
Alexa Fluor 647	640	650	665	
Cy5	640	649	670	
DyLight 650	640	654	673	
Alexa Fluor 700	640	702	723	
APC-eFluor 780	640	650	780	
APC-Cy7	640	650	785	
APC/Fire 750	640	650	787	

bio-rad-antibodies.com/flow

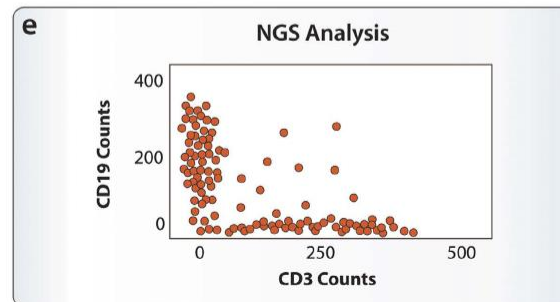
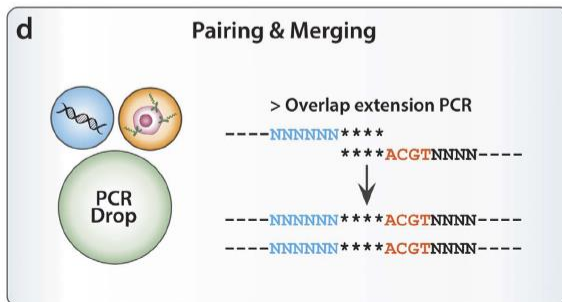
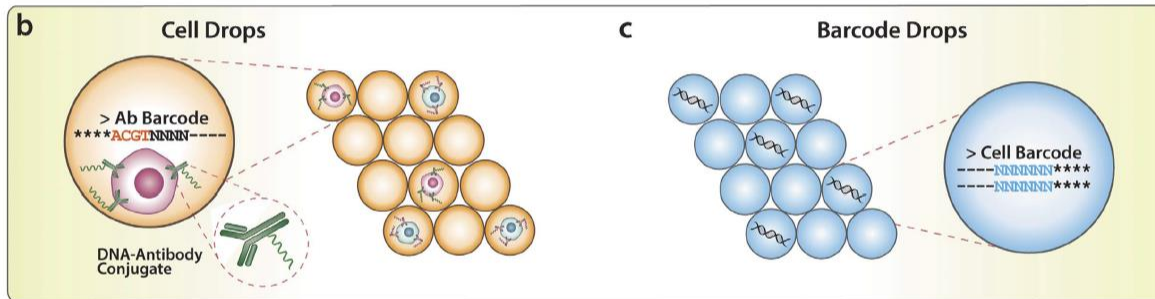
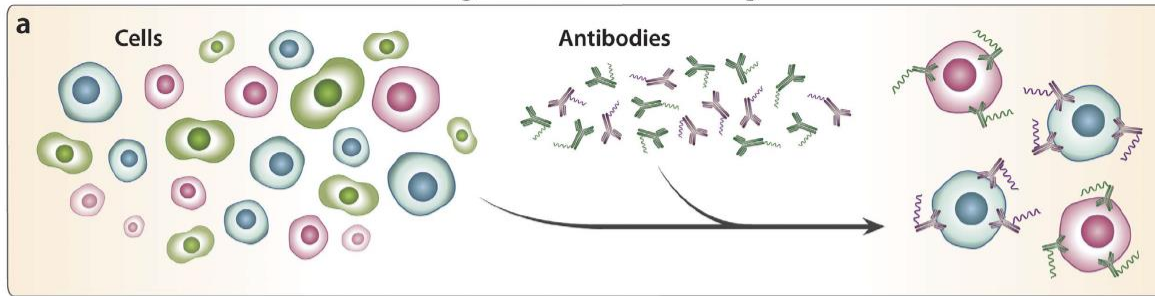
More info: Bendall SC et al. (2012) Nature Biotech. 30:639-47

Maecker H, Trotter J (2011) Multicolor Flow Cytometry Application Note

<http://www.desatoya.com/ScienceTechnology/CytometryWithSorting.htm>

Multiparameter Single Cell Analysis using Sequence Barcodes

Single-Cell Protein Profiling



Conceived to measure unlimited number of markers with single molecule sensitivity (PCR).

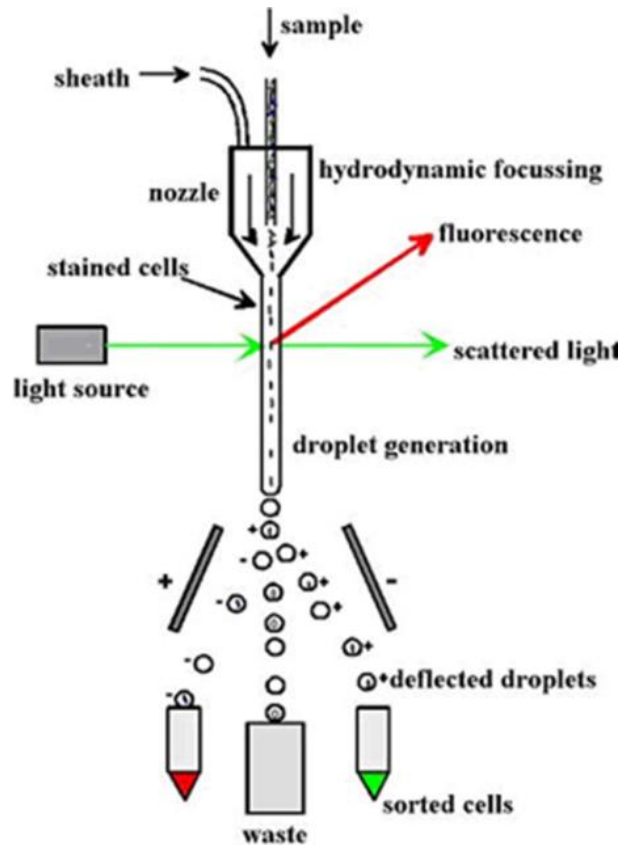
Analysis of the details reveals important limitations e.g. limited dynamic range, issues combining low and high expression markers in addition to common staining artefacts.

Multi-marker Cytometry Measurements

Points To Consider

- Know your instrument status e.g. Qr & Br for different channels
- Use optimal detector settings e.g. high gain to maximize sensitivity (check to avoid off-scale events)
- An antibody/dye combination with poor separation for a single marker assay will not work for a multi-marker measurement
- Use high sensitivity labels for low expression markers and vice versa
- For energy transfer fluorophors beware of spectral drifts by photo-degradation
- Internal controls are essential

Cell Sorting (FACS droplet sorter)



www.lifesciencesfoundation.org/events-The_FACS.html

Application Examples

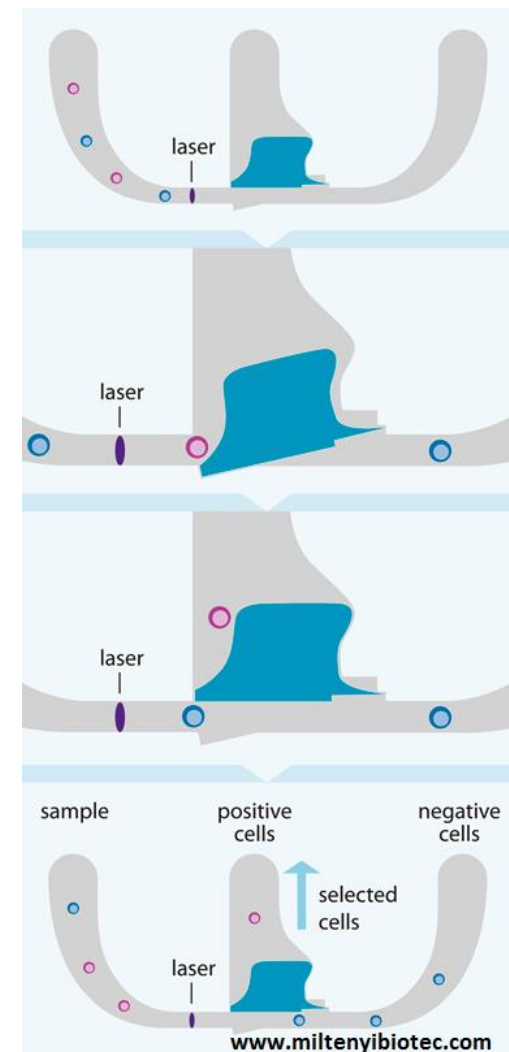
- Chromosomes
- Cloning, Strain Improvement
- Genomics, Proteomics
- Comprehensive single cell analyses (heterogeneity)

Other Cell Sorting Technologies

- Single Cell dispensers
- Tyto/OWL
- DEP sorter
- NanoCollect
- BulkSorting
(Magnetic, Gravity, Acoustic, ...)



DEPArray™ System



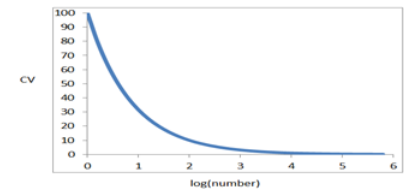
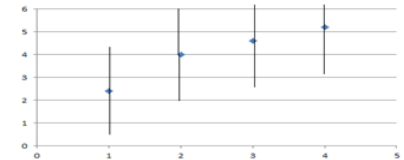
MACSQuant®Tyto™

Rare Cell Analysis and Sorting

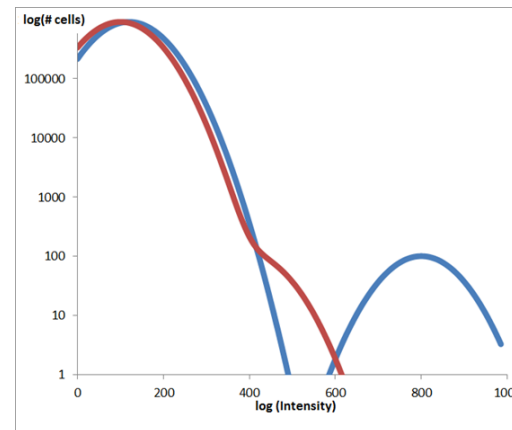
Examples CD34, AC133, antigen specific cells, CTCs

- Poisson count statistics
- Population Separation
- Bulk pre-enrichment or enrichment sorts

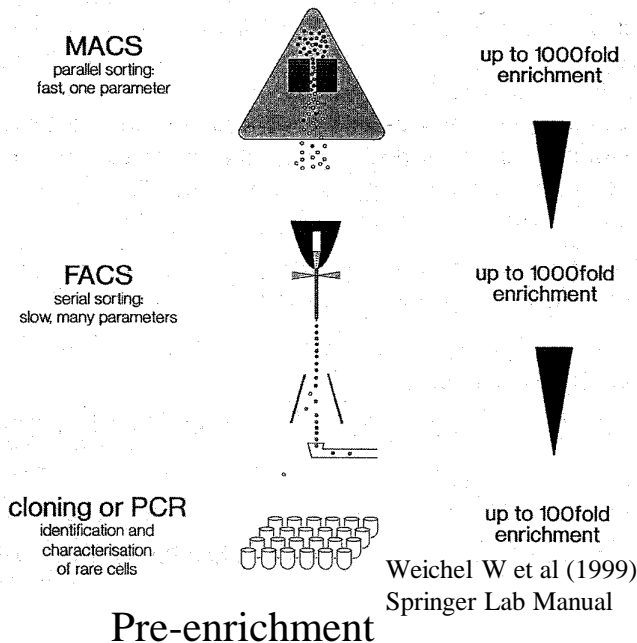
	Sample 1	Sample 2	Sample 3	Sample 4
	6	2	6	8
	3	7	1	6
	1	3	5	3
	1	4	5	6
	1	4	6	3
Mean	2.4	4	4.6	5.2
St.Dev	2.2	1.9	2.1	2.2
		Overall	Mean	4.1
			St.Dev	2.2



Ignoring Counting Statistics Can Lead to Erroneous Conclusions



Population Separation



Small Particle Analysis

Microorganisms, Organelles, Exosomes

Flow Cytometry

Light microscopy

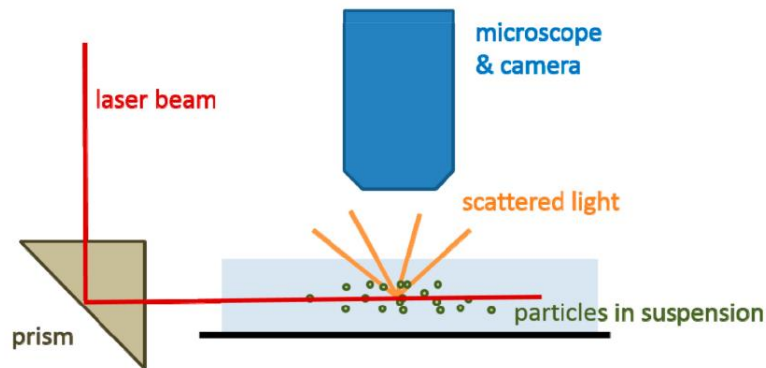
Electron microscopy

Atomic force microscopy

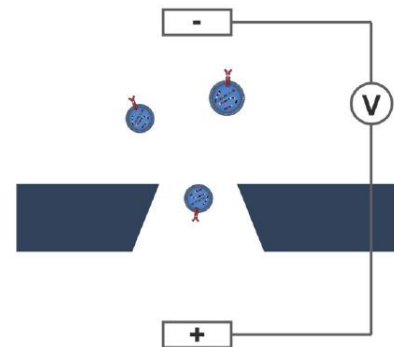
Nanoparticle tracking analysis

Tunable resistive pulse sensing

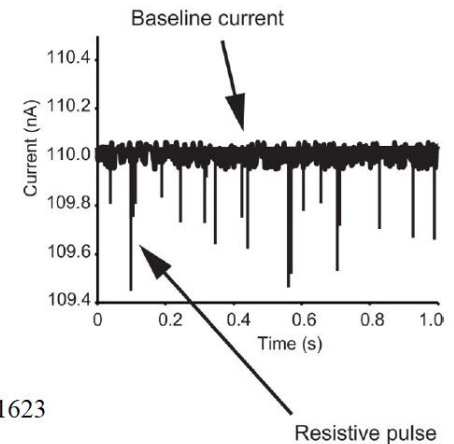
Dynamic light scattering



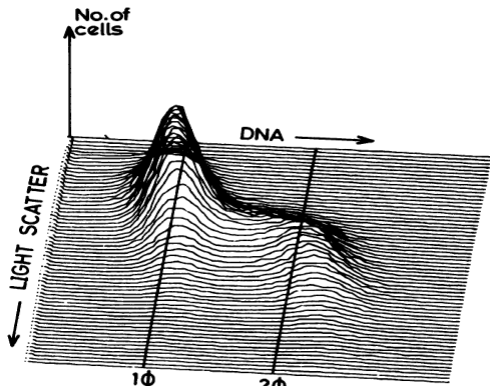
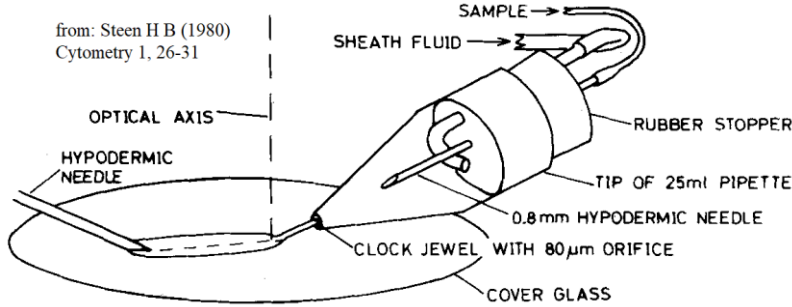
SzatanekR2017 doi:10.3390/ijms18061153



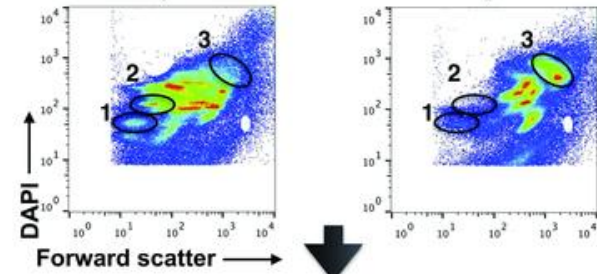
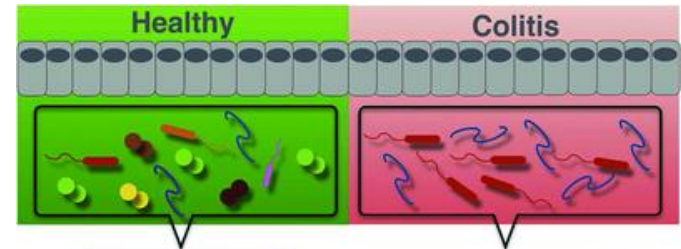
Maas S L N et al (2014) doi:10.3791/51623



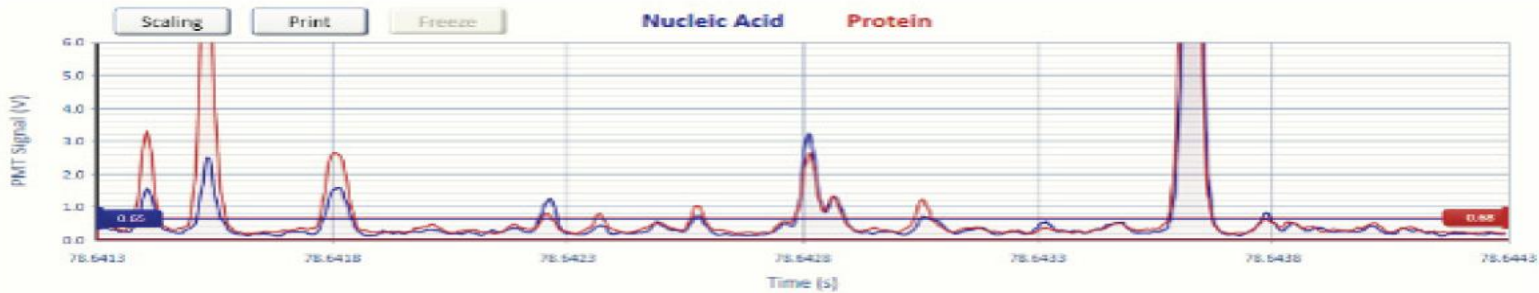
Bacteria & Virus Flow Cytometry



E.coli cell cycle analysis
Skarstad K (1983) J Bacteriol 154(2) 656-662



Zimmermann J et al (2016) doi.org/10.1002/eji.201646297



Photomultiplier Signals from a ViroCyt Virus Counter (Sartorius Stedim)

Organelles

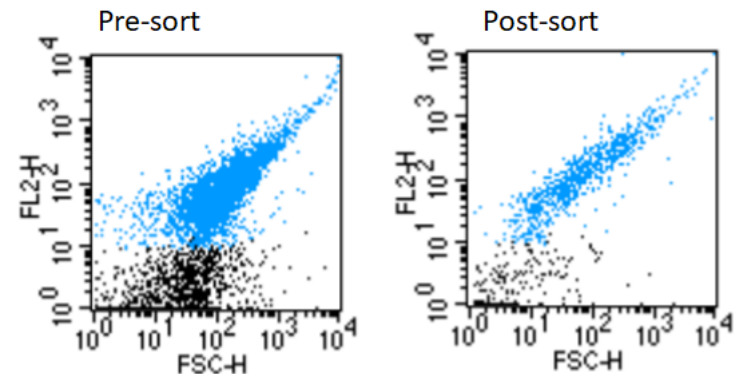
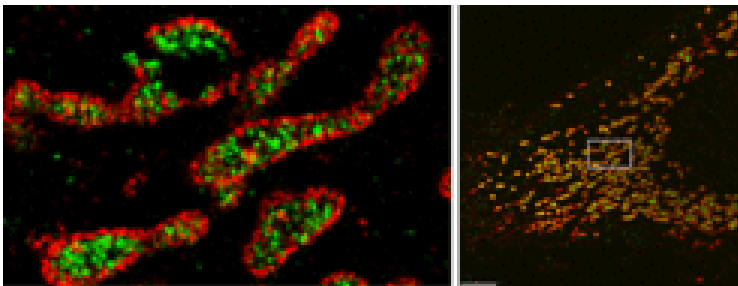
Observation

Fluorescence microscopy
Superresolution microscopy
Electron microscopy
Atomic force microscopy
Flow Cytometry

Purification

Centrifugation
Optical trapping
Field flow fractionation
Inertial flow microfluidics
Affinity binding to beads
FAOS (fluorescence assisted organelle sorting)

www.biochem.mpg.de/en/20171219-schueder-jungmann



Sorting of metaxin-stained liver mitochondria

Data from BD Biosciences

Conclusions / Caveats

- **For optimal results use an adequate technology**

(flow cytometry has enormous capabilities, but is not always the adequate technology to use e.g. single cell kinetics)

- **Understand the limitations of the system**

(complexity, limits of detection, non-specific binding of reagents, ...)

- **Use appropriate statistical methods**

(understand variance of very low counts during rare cell analysis)

- **Validate results with appropriate controls**

Conclusions

Multi-parameter cytometry

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

Many systems isolate selected single cells.

New developments in many areas provide more tools for cytometry.

More info: Bendall SC et al. (2012) Nature Biotech. 30:639-47

Maecker H, Trotter J (2011) Multicolor Flow Cytometry Application Note

Key Applications

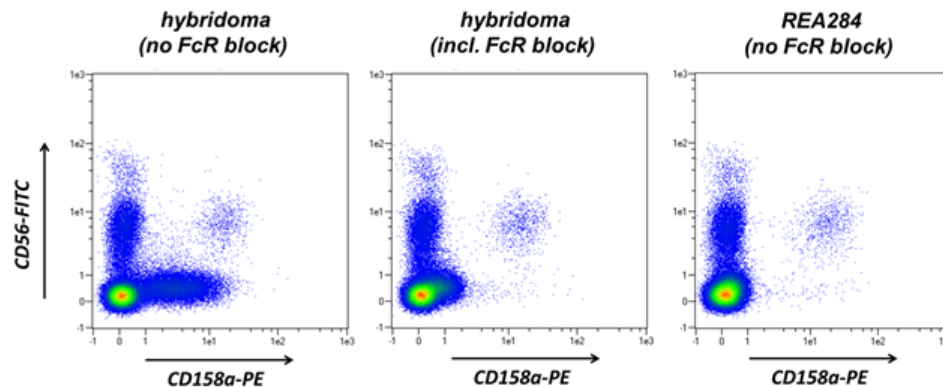
- Multi-parameter immunofluorescence (antibodies)
- Multi-parameter gene expression analysis (NA probes)
- Exosome analysis and sorting
- Single cell sorting for analysis with other technologies
- Cell cycle analysis (high resolution FCM, imaging, BrDU)
- Molecular clustering (fluorescence energy transfer FRET)
- Kinetics (population-based flow cytometry, single cell by imaging; Ca⁺⁺ flux, enzyme activity, cell proliferation)
- Receptor ligand binding (by quantitative fluorescence)
- Single Cell Sequencing (single cell sorting, PCR amplif.)
- Particle-based assays (Luminex-type multiplexed assays)

Reagents

- Nucleic acid Probes
- Antibodies (Human, Mouse, Rabbit; Camel, Llama, Shark)
- Recombinant Antibodies
- Aptamers
- Molecular Imprinted Polymers
- Environment sensitive dyes (DNA dyes, pH probes)
- Enzyme reaction probes (fluorogenic substrates)
- Labels (fluorescent, Raman, mass, sequence “barcodes”)

Llama: 15 kDalton antibodies
 10^{-9} M Kd, high stability

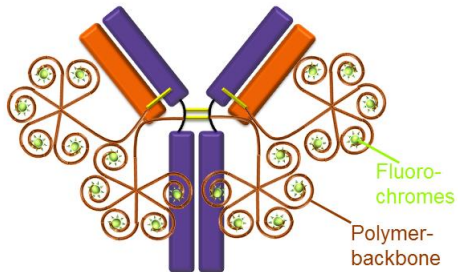
Fodey T et al; Trends in Anal.Chem. 30(2011) 254ff



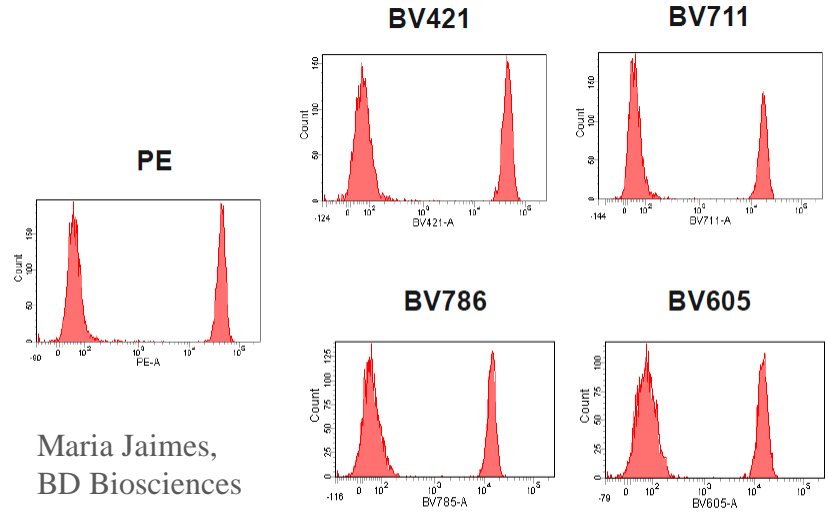
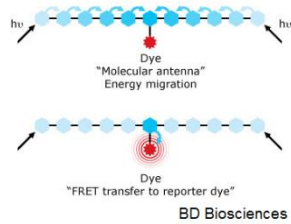
Fc-receptor binding:
CD158a-PE on PBMC

Source: Dr. Christian Dose,
Miltenyi Biotec

New Bright Dyes



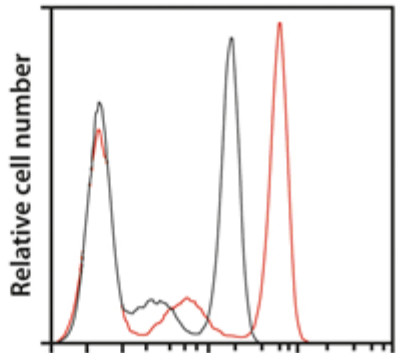
VioBright™ dyes, Christian Dose, Miltenyi Biotec



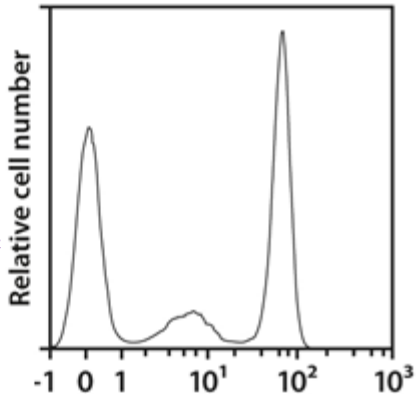
Maria Jaimes,
BD Biosciences

Maria Jaimes, page 17 in https://www.bdbiosciences.com/documents/webinar_071713_multicolor-bv.pdf

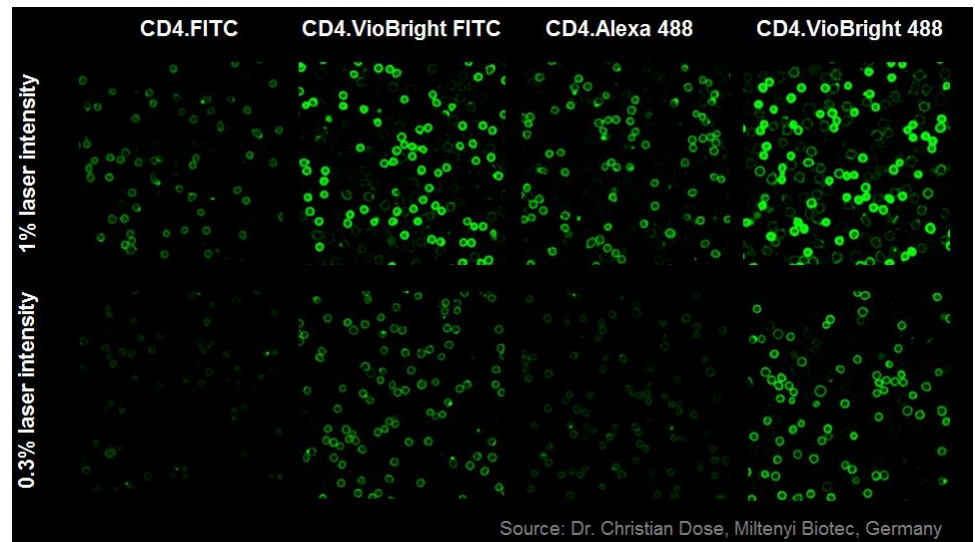
CD4-FITC
CD4-VioBright FITC
(clone Vit 4.3)



CD4-PE
(clone Vit 4.3)



Source:
Christian Dose
Miltenyi Biotec



Source: Dr. Christian Dose, Miltenyi Biotec, Germany

Technologies for Cell Analysis

	LOD	Mult	Envi	Id	Morph	Res	vivo	lbl-free
Flow cytometry	+++	+	+	+	-	-	+	+
Digital microscopy	+++	++	++	+	+++	++	-	++
2-photon imaging	+++	+	+	+	++	++	+	-
Electron microscopy	+	-	-	+	+++	+++	---	-
NGS	++	+++	---	+++	---	---	---	+++
ELISA	++	+	--	+++	---	---	---	---
Electrophoresis	++	++	+	+	---	---	---	+++
Mass spectrometry	+	+++	---	+++	+	-	---	+++
NMR MRI MRM	--	++	++++	++	++	---	+++	+++
Acoustic imaging	--	--	--	--	++	-	+++	+++
X-ray imaging	--	-	-	-	+++	++	+++	+++
...	na	na	na	na	na	na	na	na

Details at: <http://www.desatoya.com/ScienceAndTechnology.htm>

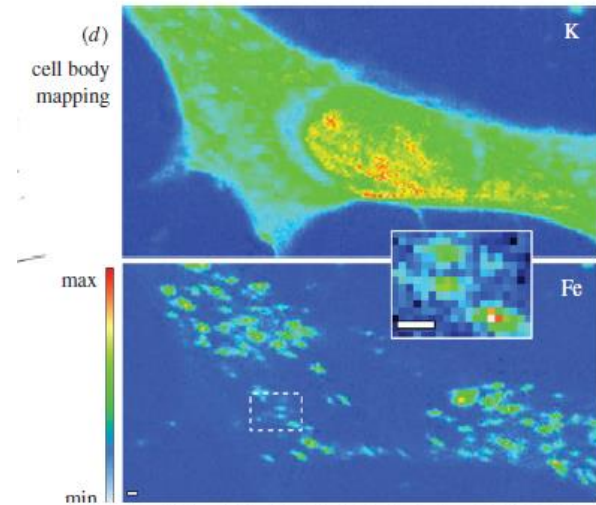
Microfluidics for WBC Isolation From Blood

Method		Dilution factor	WBC Separation efficiency	RBC depletion	Purity		WBC Enrichment	Sample throughput**
					WBC	MNC		
Cross-Flow filter	8	Undiluted	~98%	~99.975%	~70.5%	~28%*	~2000	0.06 µl/min
	12	Undiluted	~97.2%	NA	~96.9%	~39%*	NA	0.33 µl/min
Hydrodynamic filtration	13	10x	NA	NA	~3.6%	~1.4%*	~29	2 µl/min
Hydrophoretic filtration	15	20x (rat blood)	NA	NA	~58%	~23.2%*	~210	0.05 µl/min
Deterministic lateral displacement	9	Undiluted	~96% (WBC) ~95% (MNC)	~99.1%	~9%	~5.5%	~110	0.018 µl/min
Microfiltration using rarchets	26	Undiluted	~98% (WBC)	~100%	~100%	~40%*	NA	0.083 µl/min
Inertial focusing	16	500x	~95%	~94%	NA	NA	NA	3.6 µl/min
	18	400x	~89.7%	~99.8%	~91%	~36.4%*	NA	0.375 µl/min
	19	20x	NA	NA	~48%	~19.2%*	NA	240 µl/min (30 µl/min per channel)
Dielectrophoresis	21	5x	~92.1%	~87%	NA	NA	NA	0.16 µl/min
Leukocyte margination	22	Undiluted	NA	NA	NA	NA	~34	NA
Continuous erythrocyte lysis	23	Undiluted	~100%	>99.5%	NA	NA	NA	0.5 µl/min
	24	10x	~99%	NA	NA	NA	NA	100 µl/min
Slanted hydrodynamic filtration	27	20x	~85%	NA	~80%	~32%*	NA	2 µl/min
Acoustophoresis	This work	20x	>43% (WBC) >87% (MNC)	>99.95%	~54%	~53%	~1000 (WBC) ~2800 (MNC)	5 µl/min

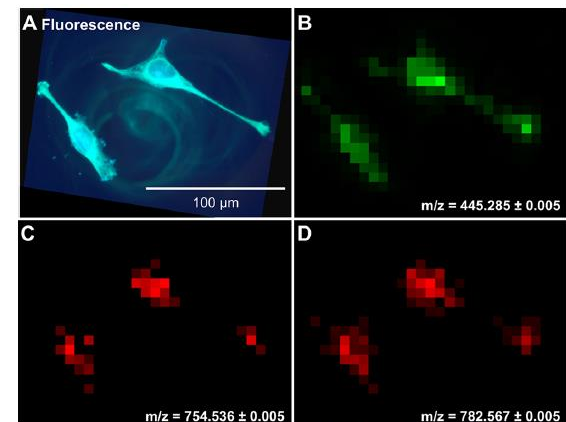
Label-free, continuous separation of WBC from blood using microfluidics. Urbansky A et al (2017) DOI:10.1038/s41598-017-17200-9

Examples of New Detection Technologies

- High spatial resolution and multi-parameter capability with X-ray / synchrotron radiation fluorescence
(super-high resolution with element labels or direct element imaging)
- Medium resolution, multi-parameter mass spectrometric imaging
(CyTOF like element labels, direct metabolite or structural component detection)
- Label-free imaging with Raman
(measuring cellular components by their Raman spectra)
- Label-free medium resolution NMR imaging
(chemical environment sensing)



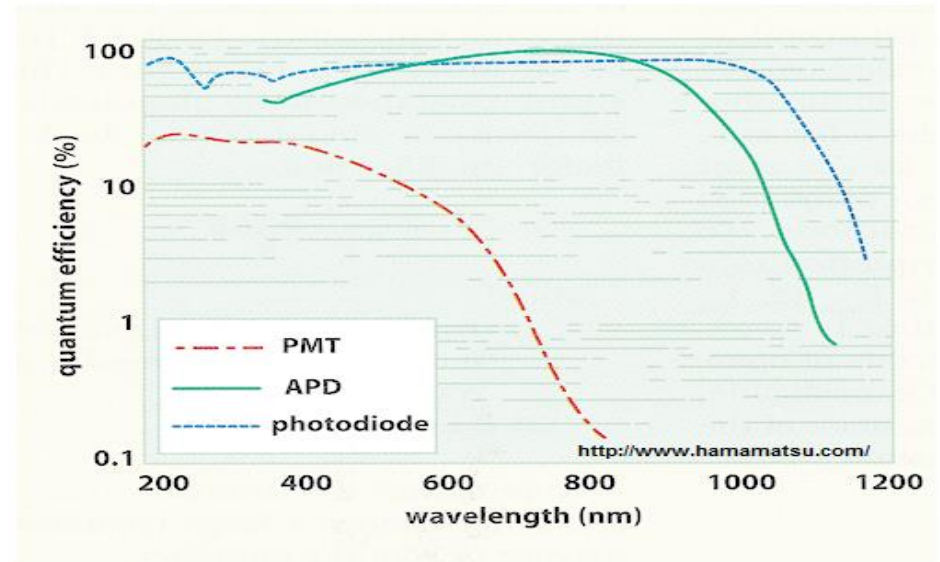
Ortega R et al (2009) J.R.Soc Interface 6: S649-S658



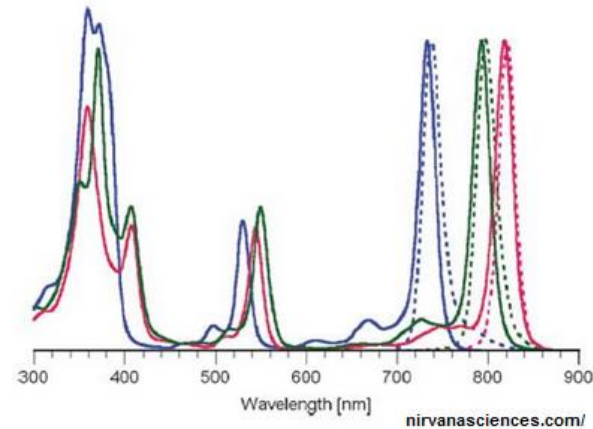
Schober Y et al. (2012) Anal.Chem. 84, 6293ff

New Detector-Label Combinations

- New photodetectors extend the available spectrum
(Si avalanche photodiodes extend detection into the far infrared)



- New dyes add excitation in the UV, some detection in the IR
(Fluorescent polymers, bacteriochlorins, ...)



Conclusions

Evolving Technologies

Technology developments in algorithms, computing, detectors, electronics, nanotechnology, microfluidics, organic chemistry, and recombinant protein technology create the basis for new reliable analytical approaches for a deeper molecular understanding of living systems.

There is substantial value in working with other scientific disciplines.

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More science detail and references: <http://www.desatoya.com>