



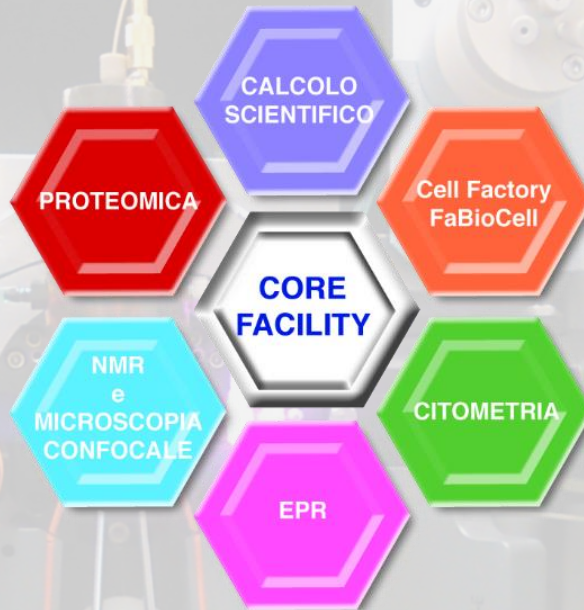
## *Servizio Grandi Strumentazioni e Core Facilities*



# Evoluzione della Citometria a flusso: dall'era policromatica alla massa

Valentina Tirelli

Luca Pasquini



# La citometria a flusso: principi ed applicazioni

*Valentina Tirelli*

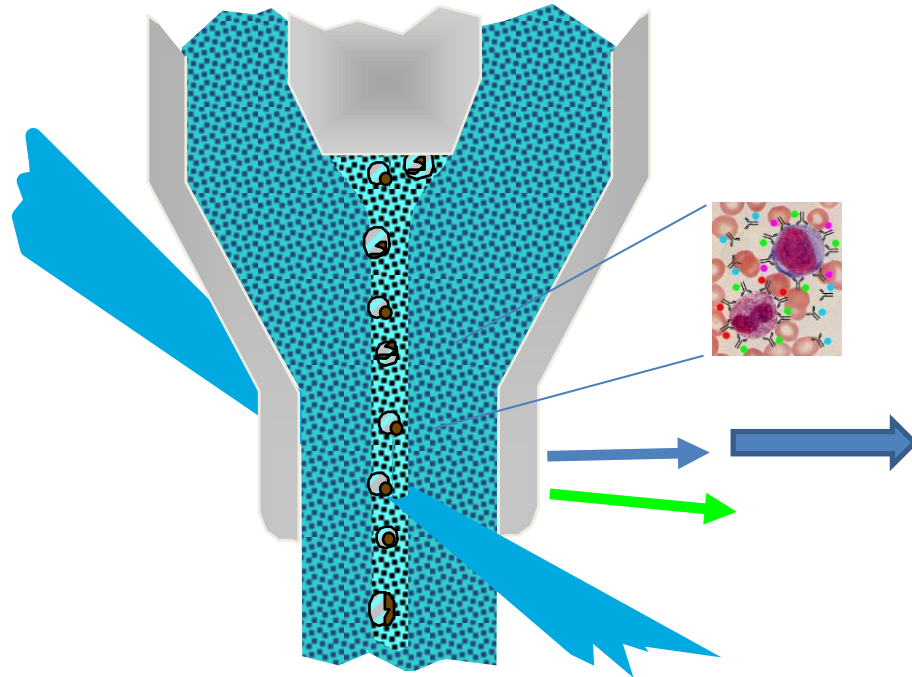




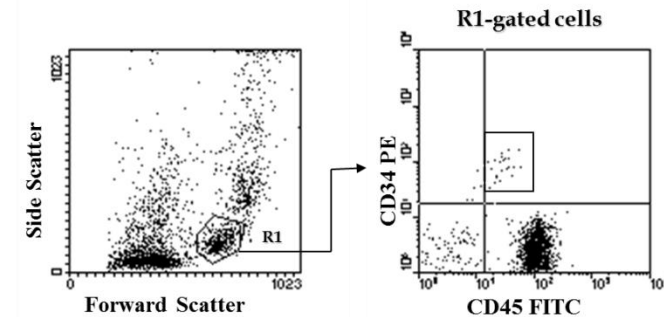
# Flow Cytometry: principle



Flow cytometry measures multiple characteristics of individual particles flowing in single file in a stream of fluid.



Light scattering at different angles can distinguish differences in size and internal complexity, whereas light emitted from fluorescently labeled antibodies can identify a wide array of cell surface and cytoplasmic antigens

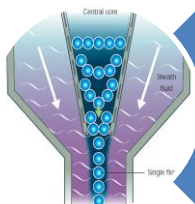


**The basic principle of flow cytometry is the passage of cells in single file in front of a laser so they can be detected, counted and sorted**

detection and enumeration of specific cell types in complex mixtures of cells, such as whole peripheral blood

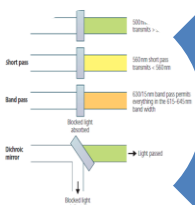


# How the flow cytometer works



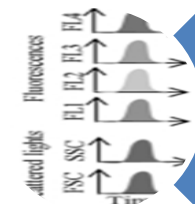
## Fluidic system:

Cells pass one by one through one or more laser beam in a flow cytometer....



## Optic system:

...photodiode and photomultiplier convert and amplify light signals (photons) to an electric signals...



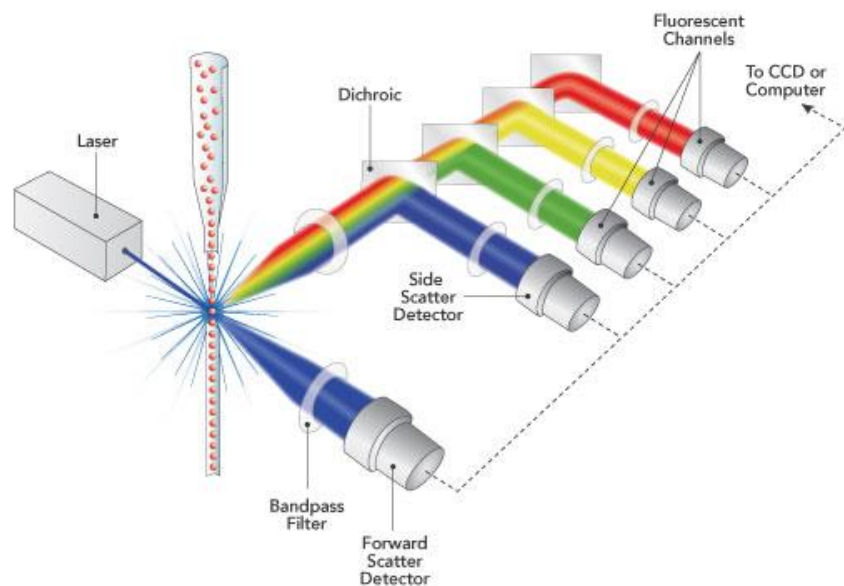
## Electronic system

...this pulse is digitalized, recorded to be analyzed ...



## Data analysis

...with specific computer software





# Advantages



Flow cytometry provides **rapid** analysis of **multiple characteristics** of single cells

Acquisition of **high rate of events** ( $>10^6$  cells)



## Polychromatic flow cytometry

Increasing the number of markers that distinguish rare cells can lead to enhanced assay sensitivity and specificity



## Gating strategy

To maximize detection of the rare event, more than one fluorescence parameter should be used to positively identify the cell of interest (compound gating). Likewise, there should be at least one fluorescence parameter for which the rare event is negative (negative gating)



## Rare-cell populations

stem cells, circulating endothelial cells, circulating tumor cells, and residual disease cells

The information obtained is both **qualitative and quantitative**

Possibility to physically separate sub-populations (**cell sorting**)



# Measurable parameters



Cell surface antigens (CD markers)

DNA (cell cycle analysis, cell kinetics, proliferation etc.)

RNA

Chromosome analysis and sorting (library construction, chromosome paint)

Proteins

Intracellular antigens (cytokines, secondary mediators etc.)

Nuclear antigens

Enzymatic activity

pH, intracellular ionized calcium, magnesium, membrane potential

Membrane fluidity

Apoptosis (measurement of DNA degradation, mitochondrial membrane potential, permeability changes, caspase activity)

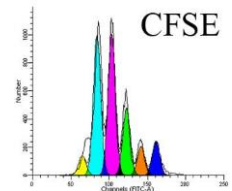
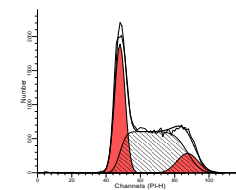
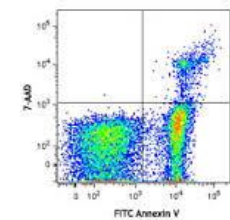
Cell viability

Monitoring electroporation of cells

Oxidative burst

Characterising multi-drug resistance (MDR) in cancer cells

Cellular pigments (Chlorophyll or phycoerythrin)





# MULTIPARAMETRIC Immunophenotyping



+								-							
+				-				+				-			
+		-		+		-		+		-		+		-	
+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-
+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-

Ab:

A

+

B

+

C

+

D

+

E = 32

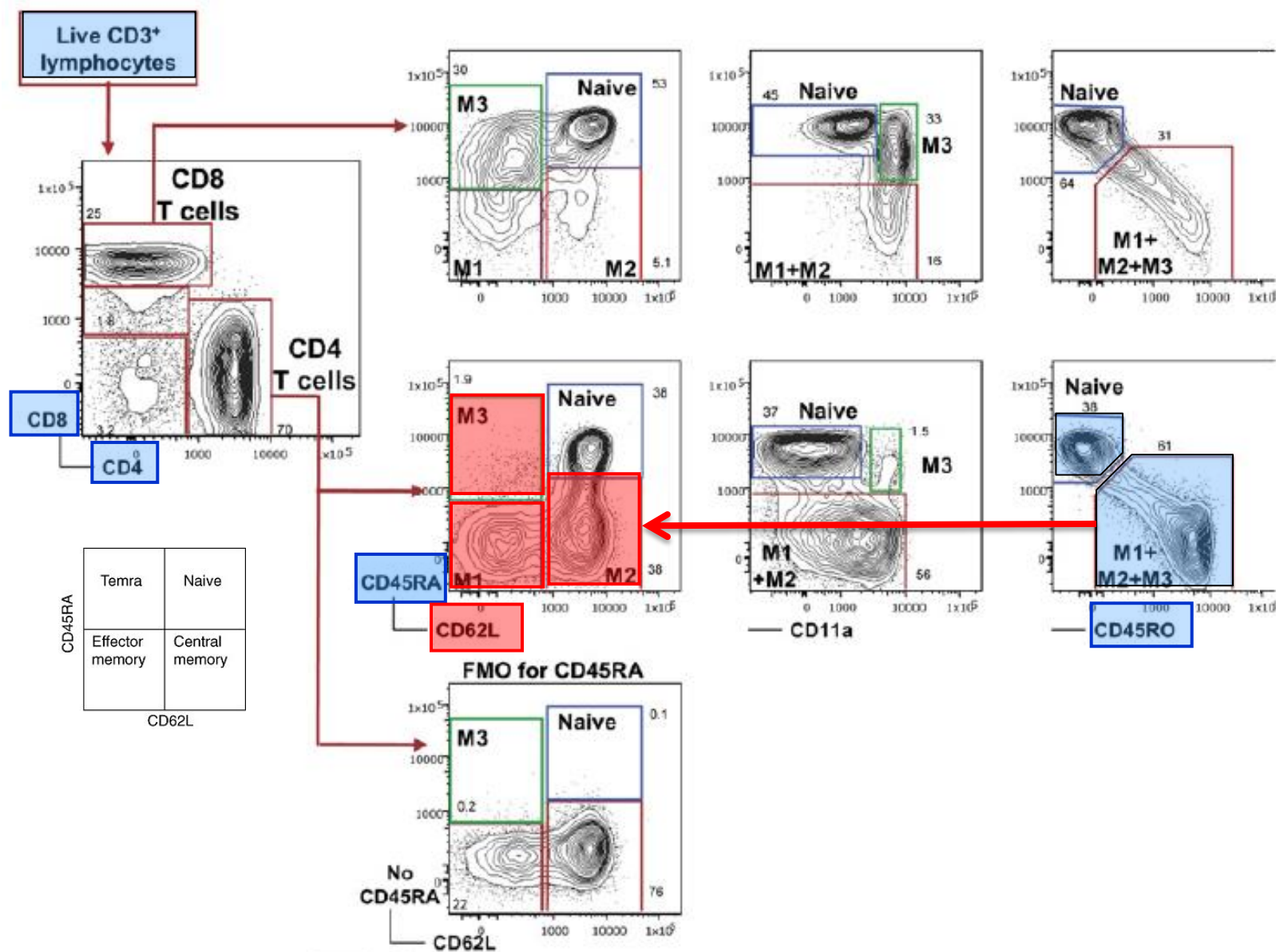
phenotype: A+B-C-D+E+

$a^n$

$a = 2$   $n =$  number of antibodies



# MULTIPARAMETRIC Immunophenotyping







# Gating strategy to detect rare population

Multiparameter flow cytometry (FCM) is ideal for rapid processing of high numbers of cells per second and is commonly utilized to quantify CECs and EPCs

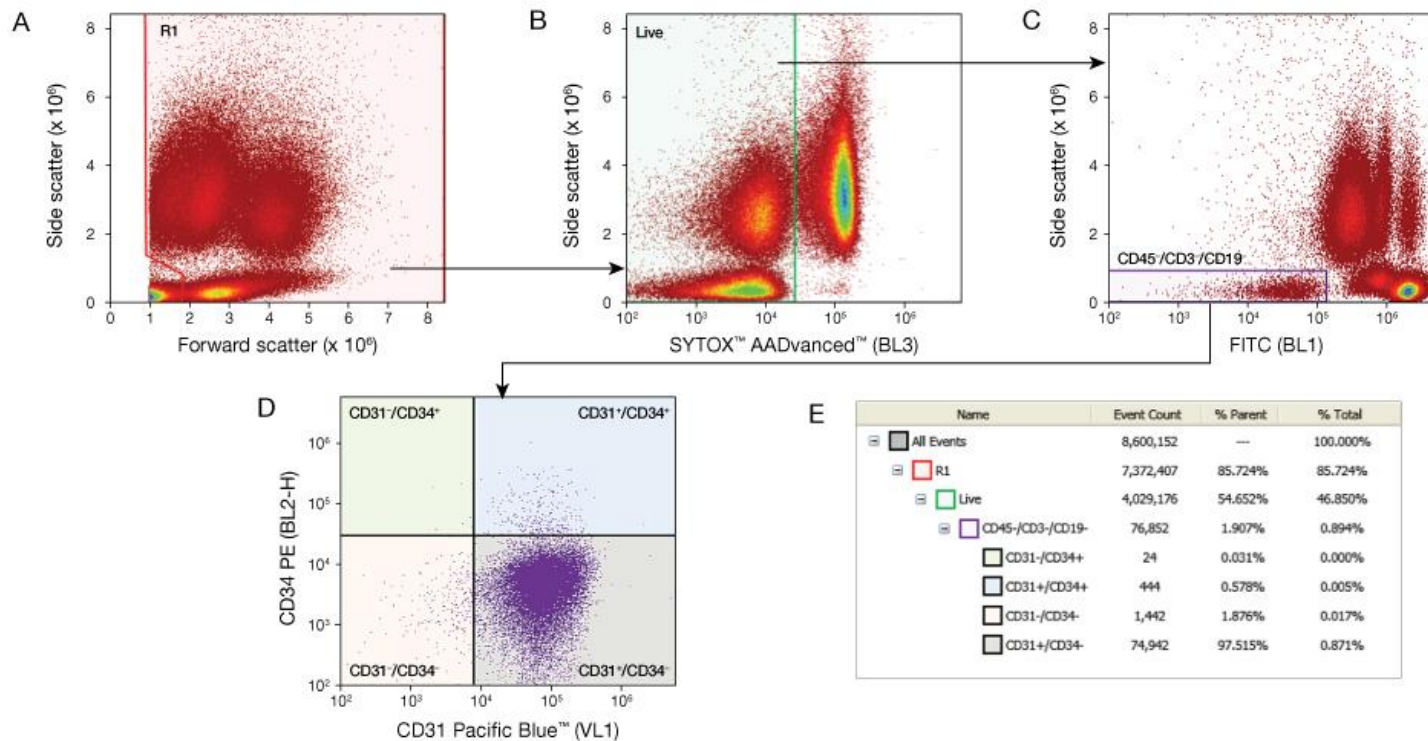
CECs and EPCs are rare in healthy subjects and increase in different tumor



Tool for monitoring clinical outcome and tumor treatment (prognostic marker in specific tumor types)

exclusion of debris and dead cells

Gating the negative population



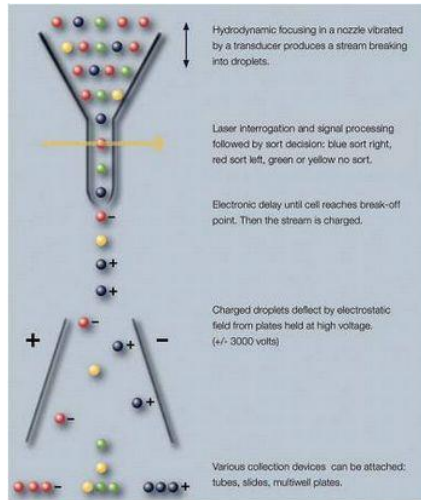
**FITC dump channel**  
Since CECs are negative for all three of these markers, all positive cells can be eliminated from further analysis using only one fluorescence channel



# Cell sorting



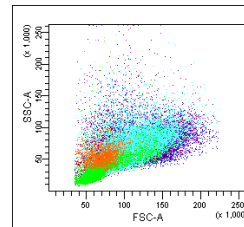
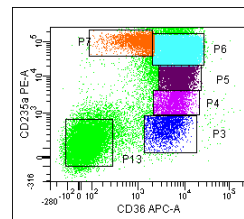
## Cell sorting principle



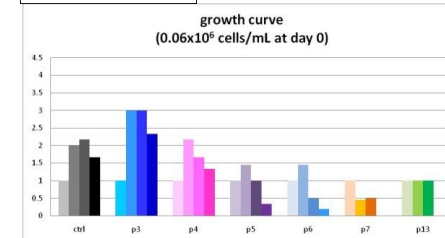
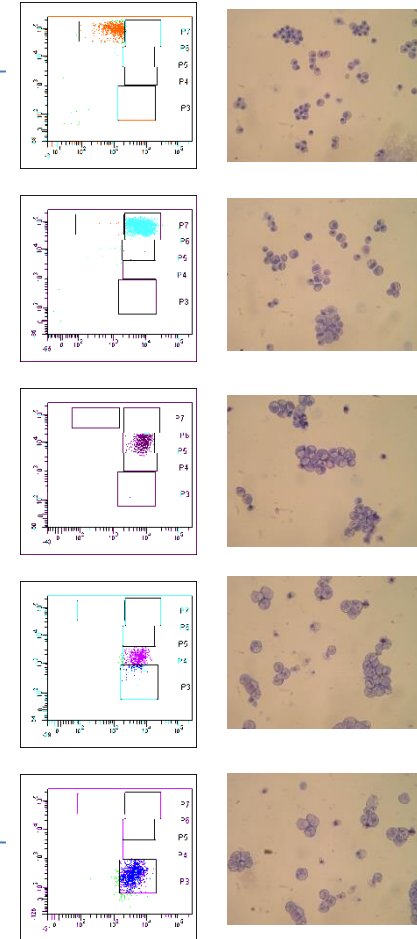
## BD FACS Aria I



Day 10



## Sorted populations Purity >95%



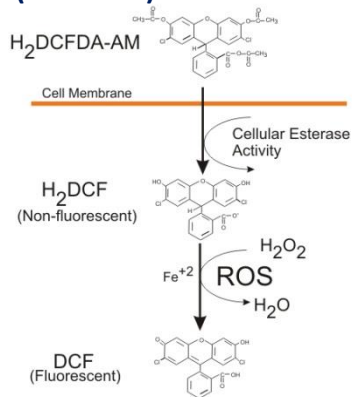


# ROS detection



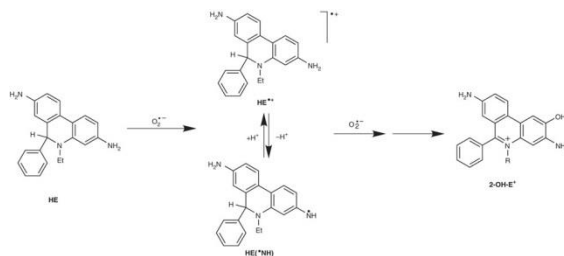
Fluorescent dyes used frequently for measuring hydrogen peroxide, superoxide, and peroxynitrite in biological systems.

## Dichlorodihydrofluorescein (DCFH-DA)



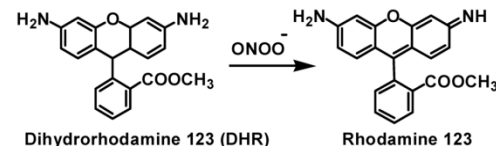
*Free Radic Biol Med.* 2012 Jan 1; 52(1): 1–6.

## Hydroethidine (HE) and Mito-SOX or Mito-HE

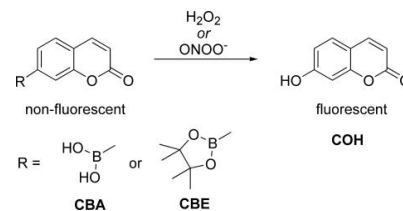


*Nature Protocols* 3, - 8 - 21 (2008)

## dihydrorhodamine (DHR)



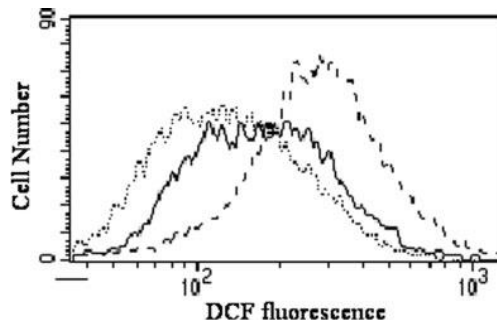
## Boronate-containing fluorophores (e.g., coumarin boronate)



*J Biol Chem.* 2010 May 7; 285(19): 14210–14216.

The generation of reactive oxygen and nitrogen species has been implicated in the onset and progression of several diseases (e.g., atherosclerosis, cancer, diabetes, neurodegeneration)

Representative histograms showing ROS levels in peripheral blood mononuclear cells of euthyroid (—), hypothyroid ( ), and hyperthyroid patients (- - -).



*Cytometry Part B (Clinical Cytometry)* 70B:20–23 (2005)



# Proliferation assay



**Table 1: Some Common Cell Proliferation Dyes**

Dye	Cellular Location	Excitation Laser	Emission Peak
CFSE	intracellular	488 nm	~519 nm
CellTrace Violet	intracellular	405 nm	~455 nm
<u>eFluor670</u> proliferation dye	intracellular	633 nm	~670nm
PKH26	membrane	488, 532, 561 nm	~567 nm
PKH67	membrane	488 nm	~502 nm
CellVue Claret	membrane	633 nm	~675 nm

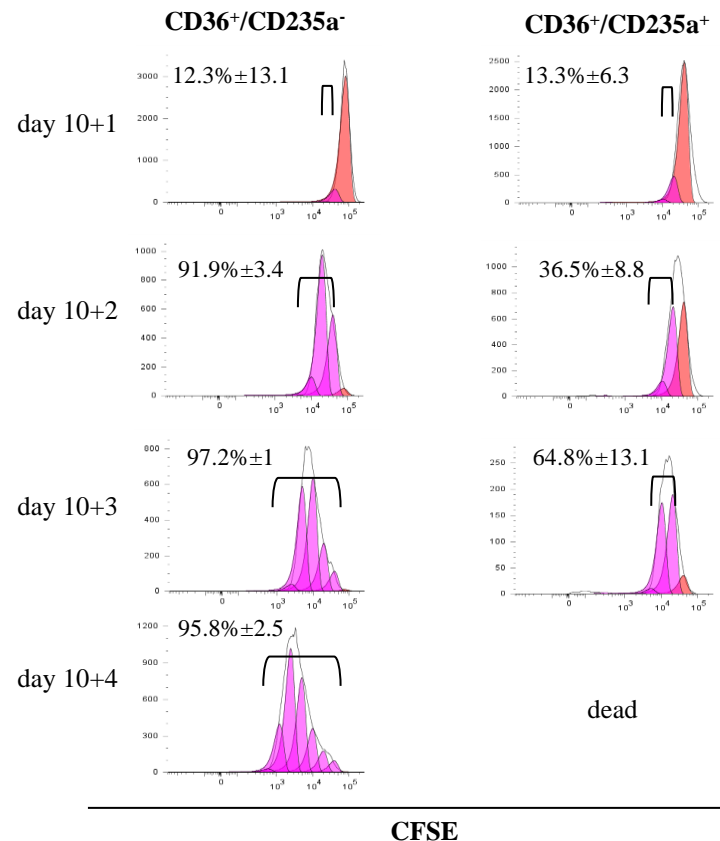
## Possible applications:

- Proliferation of immune cells in response to stimulation
- Self-renewal of stem cells
- Biological homeostasis
- Tumor cell proliferation

Cell proliferation assays are widely used in cell biology to measure cellular metabolic activity in response to stimuli such as growth factors, cytokines and other media components. The development of new treatments for cancer, for example, target the proliferation of the cancer cells,



# Proliferation assay



	Day			
	10+1	10+2	10+3	10+4
<b>Proliferation index</b>				
<b>iEBs</b>	1.02±0.01	1.53±0.03	2.35±0.07	2.76±0.21
<b>mEBs</b>	1.07±0.04	1.12±0.07	1.41±0.34	b.d.
<b>Division index</b>				
<b>iEBs</b>	0.04±0.01	1.32±0.08	2.25±0.07	2.65±0.27
<b>mEBs</b>	0.14±0.07	0.4±0.08	0.8±0.11	b.d.

**Table II. Proliferation and Division indexes values of 10 days sorted iEBs and mEBs populations in Hema culture conditions**

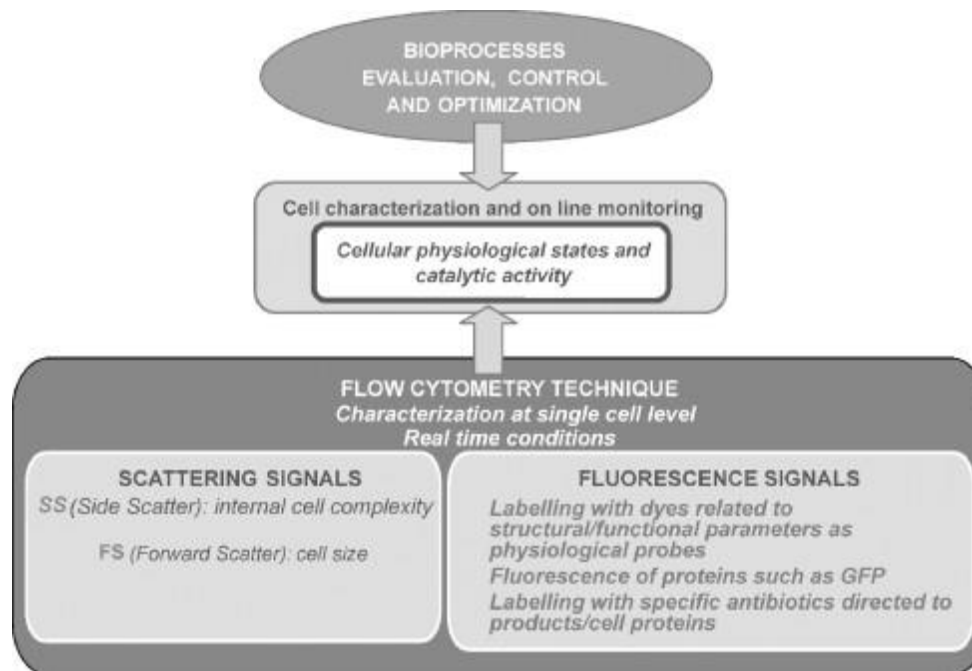
**The Percent Divided** – the measure of the percent of the input cells that entered division.

**The Division Index** – a measure of the average number of divisions which includes the undivided cells.

**The Proliferation Index** – the average number of divisions that exclude the undivided cells.



# Application of flow cytometry to industrial microbial bioprocesses



more efficient and less time-consuming methods to assess the presence of microorganisms, as well as their viability for bioprocesses control and improvement.

Rapid detection of microorganism in sample



More accurate quality control  
Optimization of bioprocess

**Industrial microbial bioprocesses application:  
dairy industry, beverages industry, environmental and water system**



## Industrial microbial bioprocesses application: dairy industry, beverages industry, environmental and water system

**Table 5**  
Some FC applications to water and environmental microbiology.

Application	Microorganism	Functional or structural parameter	Reference
Characterization of freshwater and marine waters	<i>Bacterial cells</i>	DNA content, Membrane integrity and esterase activity	[16]
Cell functionality after peracetic acid treatments	<i>S. Typhimurium</i>	DNA	[43]
Cell enumeration in marine and ballast water samples	<i>Escherichia coli</i> and <i>Bacillus cereus</i>	Nucleic acid content	[92]
Characterization of activated sludge	<i>Comamonas testoteroni</i> , <i>Paracoccus pantotrophus</i> , <i>Escherichia coli</i>	Respiratory activity	[109]
Enumeration of viable cells in water after chlorination	<i>Legionella pneumophila</i>	Esterase activity	[120]
Cell survival in seawater during starvation	<i>Escherichia coli</i> and <i>S. Typhimurium</i>	Membrane integrity and potential	[134]
Effectiveness of disinfection treatments	<i>Escherichia coli</i> , <i>Salmonella Typhimurium</i> , <i>Shigella flexneri</i> , <i>Enterococcus faecalis</i>	Membrane integrity and potential, pump activity	[138,151]
Estimation of total bacteria in live and fixed samples from fresh and saline waters	<i>Salmonella typhimurium</i>	Nucleic acids	[252]
Cell characterization in tropical marine environments	Bacterioplankton	DNA	[257]
Assessment of marine bacterial death	Natural bacterioplankton	Nucleic acids staining	[258]
Cell viability of in different types of water	<i>Aeromonas hydrophila</i>	DNA content and membrane integrity	[259]
Detection and viability assessment of water pathogens	<i>Giardia lamblia</i>	Fluorescent antibodies and membrane integrity	[260]
Chlorination treatment	<i>Nitrosospira</i> spp.	Membrane integrity and esterase activity	[262,263]
Evaluation of different environmental conditions and disinfection processes on VBNC cells in water	<i>Legionella pneumophila</i>	Esterase activity	[264]
Drinking water treatment processes	Heterotrophic bacteria	Nucleic acids detection	[266]
Determination of assimilable organic carbon in drinking water	<i>Pseudomonas fluorescens</i>	Nucleic acids	[267]
Assessment of water quality in papermaking	<i>Bacterial cells</i>	DNA content and membrane integrity	[268]



# FCM in drinking water



► The Federal Council



Schweizerische Eidgenossenschaft  
Confédération suisse  
Confederazione Svizzera  
Confederaziun svizra

The Federal Council  
The portal of the Swiss government

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Start > Documentation > Media releases > Drinking water unexpectedly rich in microbial life

◀ Documentation

◀ Back to overview



## Media releases

Media releases by the Federal Council

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## Drinking water unexpectedly rich in microbial life

Dübendorf, 24.01.2013 - Flow cytometry (FCM) can now be officially used for the quantification of microbial cells in drinking water. The new analytical method – developed at Eawag and extensively tested both in Switzerland and abroad – has been incorporated into the Swiss Food Compendium (SLMB) by the Federal Office of Public Health (FOPH). FCM provides much more realistic results than the conventional method, in which bacterial colonies are grown on agar plates. The results demonstrate that even good-quality drinking water harbours 100 to 10,000 times more living cells than the conventional plate count method would suggest.

**Real time quality programs could lead to substantial industrial cost reductions**



## Naturally fluorescence pigments in phytoplankton

Fluorophore	Exciting Laser	Major Emission Wavelength
Chlorophyll a,b	488	>640 nm
Phycoerythrin	488	575 nm
C-phycoyanin	640	650 nm
R-phycoyanin	640	646 nm
Allophycocyanin	640	660 nm

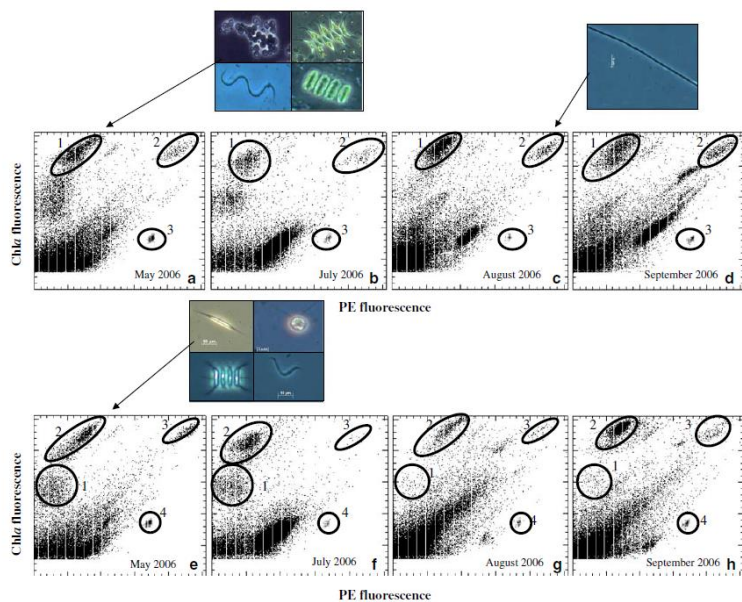


Fig. 2 PE vs. Chl a fluorescence cytograms showing the seasonal evolution of the phytoplankton composition (a–d) for two clusters isolated from station 501 of the Reservoir Mame (1 mixed green algae community, 2 *Pseudanabaena immetica* (picture of a filament), 3 beads) and e–h for three clusters isolated from station 553 of the Reservoir Mame (1 *Chlorella vulgaris*, 2 mixed green algae commu-

nity, 3 mixed green algae community, 4 beads). Pictures of the mixed green algae communities with *Chlorella vulgaris*, *Scenedesmus acuminatus*, *Hyaloraphidium contortum*, *S. linearis* (upper left, from up left to right down) and with *Ankara lanceolata*, *Chlamydomonas* sp., *Scenedesmus spinosus*, *Monoraphidium* sp. (middle, from up left to right down)

Cellamare M., et al. (2009) *J Appl Phycol*

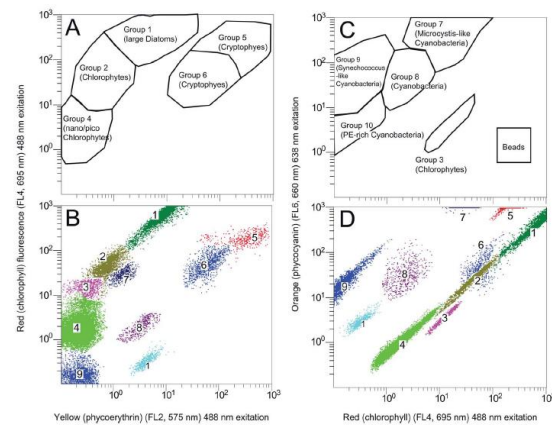


Fig. 1 Flow cytometry panels showing gating profiles for phycoerythrin (FL2 – 575 nm) against chlorophyll (FL4 – 695 nm) fluorescence (a) and chlorophyll (FL4 – 695 nm) against phycoerythrin (FL6 – 660 nm) (c). Composite plots (data from different samples consolidated) showing example data for phytoplankton groups from the River Thames at Wallingford, UK (b and d).

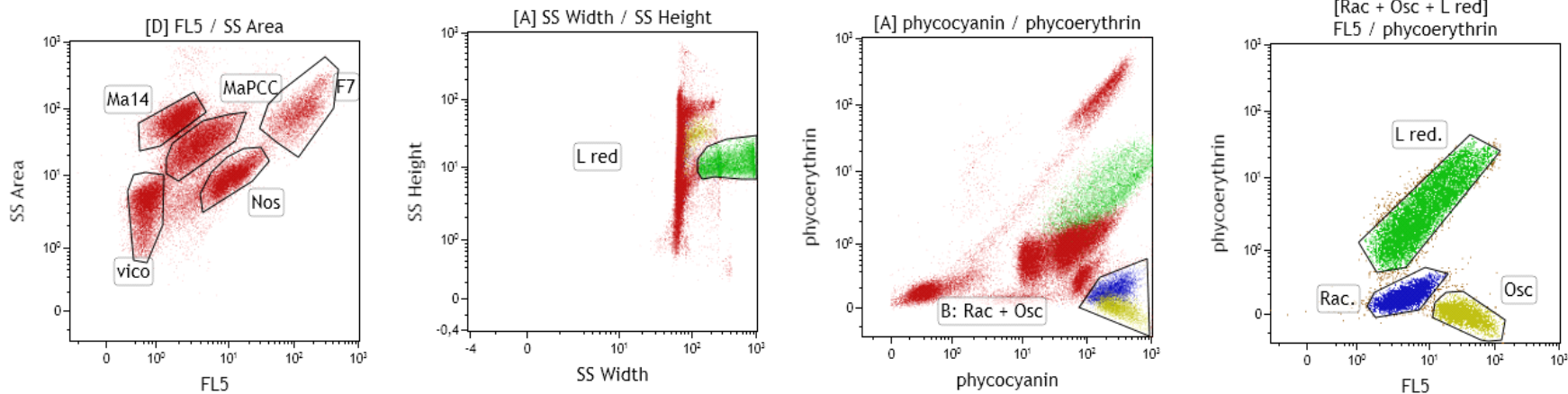
*Environ. Sci.: Processes Impacts*, 2014, 16, 594–603



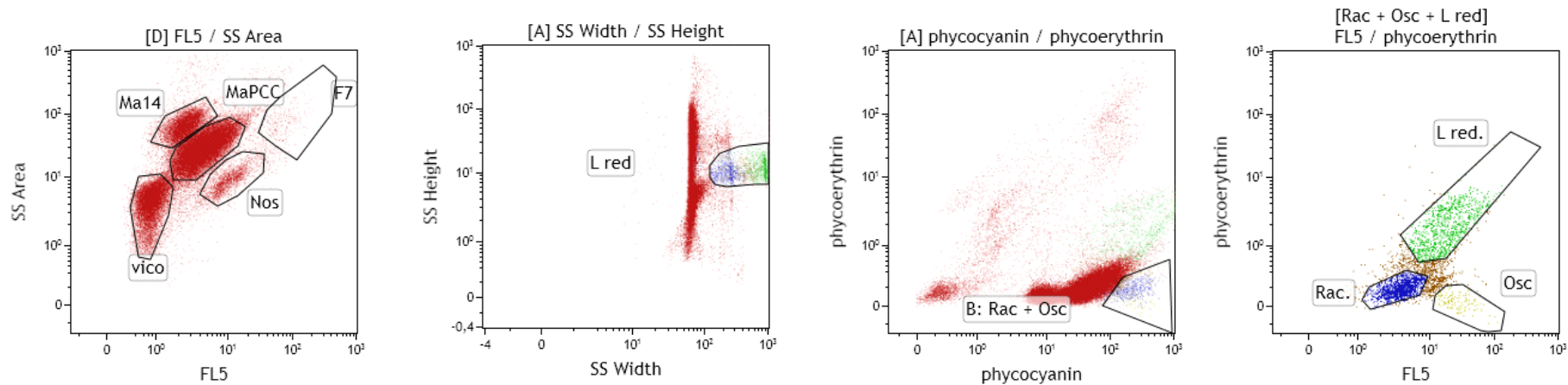
# FCM in Aquatic microbiology



## Merge campioni acquisiti



## Mix culture





# Multicolor flow cytometry: choice of fluorochromes



Multicolor flow cytometry rapidly reveals a large amount of biological information from a single sample. Over the past few years, the number of parameters (and consequently colors) simultaneously analyzed in typical flow cytometry experiments has increased

One consideration when performing multicolor fluorescence studies is the possibility of **spectral overlap between fluorophores** and when making decisions about which fluorochromes to use in your experiments, you'll want to know their relative **emission spectra**

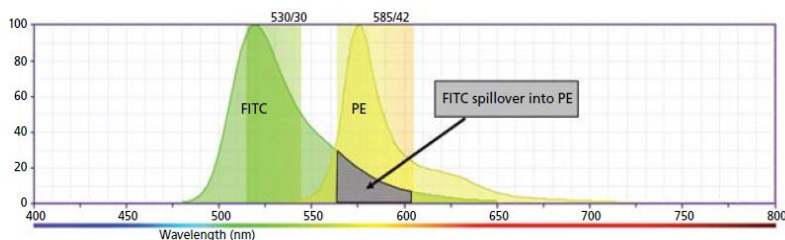
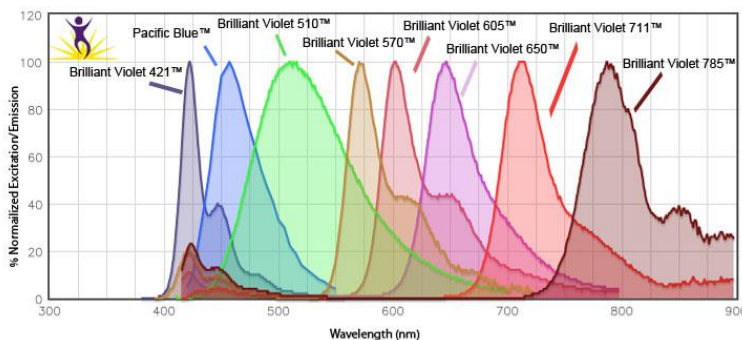


Figure 1. Example of FITC spillover into the PE channel.

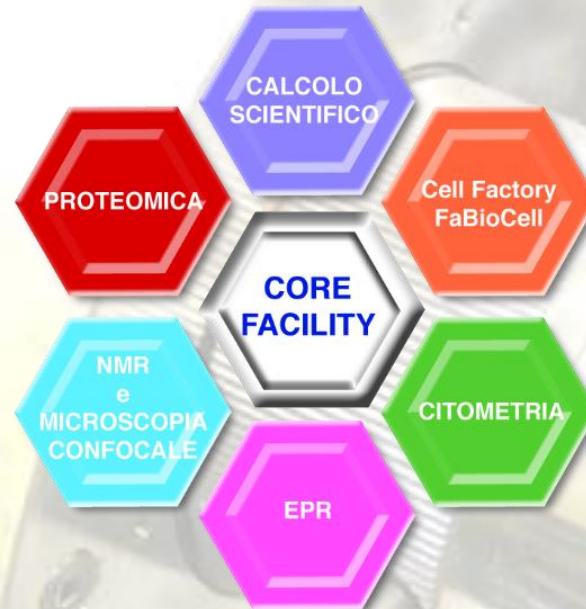
➔ need for compensation

Alternatively you can combine fluorophores that can only be activated by specific individual laser lines, (providing the lasers are spatially separated), but as you increase the number of fluorophores this becomes very difficult



## Brightness of various fluorochrome conjugates

Relative Brightness	Reagent	Filter
BRIGHTEST	Brilliant Violet™ 421	450/50
	PE	575/26
	Brilliant Violet 605	610/20
	BD Horizon PE-CF594	610/20
	PE-Cy5	670/14
BRIGHT	APC	660/20
	PE-Cy7	780/60
	Alexa Fluor® 647	660/20
MODERATE	PerCP-Cy5.5	695/40
	Alexa Fluor® 488	530/30
	FITC	530/30
	BD Horizon V450	450/50
DIM	Pacific Blue™	450/50
	Alexa Fluor® 700	730/45
	PerCP	695/40
	APC-Cy7	780/60
	AmCyan	525/20
	BD Horizon V500	525/20
	BD APC-H7	780/60



# Citometria di Massa: principi ed applicazioni

*Luca Pasquini*



**Citometria a flusso**

**Spettrometria di Massa**

Anticorpi coniugati  
(con metalli rari)

Risoluzione TOF

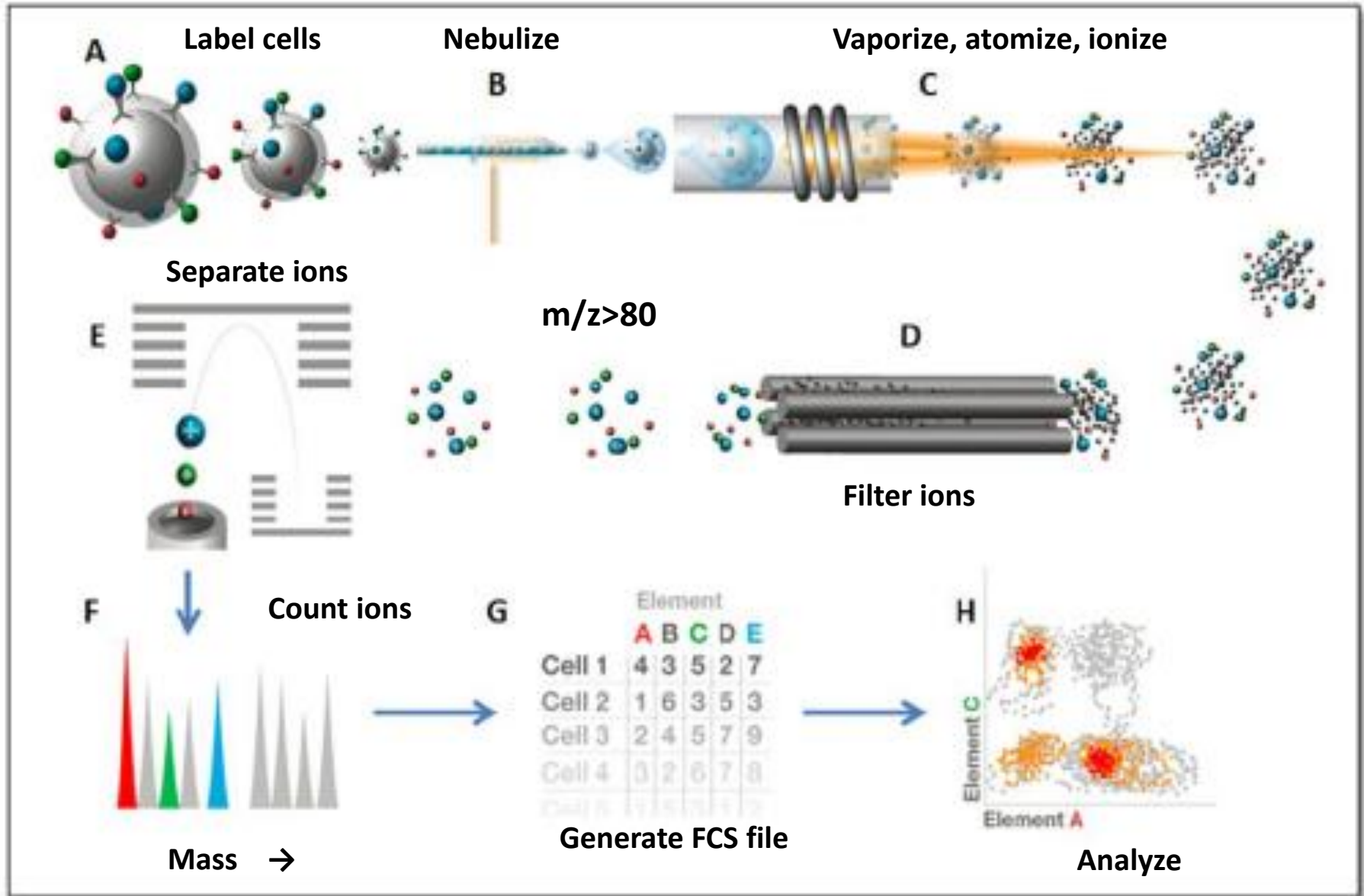
**Citometria di Massa  
(*CyTOF*)**



***Notevole incremento del numero  
di parametri valutabili!!!***



# Citometria di Massa (workflow)





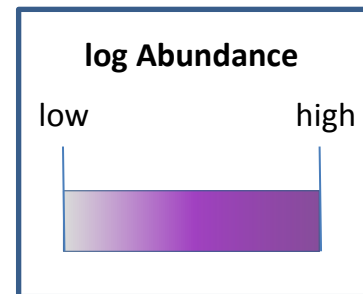
# Lantanidi



## log Abundance in Earth's crust

1. Drag cursor around plot area to show information.  
2. Click on element within plot area to go to that element.

H																				He
Li	Be						B	C	N	O	F									Ne
Na	Mg						Al	Si	P	S	Cl									Ar
K	Ca	Sc	Ti	V	Cr	Mn	Fe	Co	Ni	Cu	Zn	Ga	Ge	As	Se	Br				Kr
Rb	Sr	Y	Zr	Nb	Mo	Tc	Ru	Rh	Pd	Ag	Cd	In	Sn	Sb	Te	I				Xe
Cs	Ba	Lu	Hf	Ta	W	Re	Os	Ir	Pt	Au	Hg	Tl	Pb	Bi	Po	At				Rn
Fr	Ra	Lr	Rf	Db	Sg	Bh	Hs	Mt	Uun	Uuu	Uub	Uut	Uuq	Uup	Uuh	Uus				Uuo



La	Ce	Pr	Nd	Pm	Sm	Eu	Gd	Tb	Dy	Ho	Er	Tm	Yb
Ac	Th	Pa	U	Np	Pu	Am	Cm	Bk	Cf	Es	Fm	Md	No

## LANTANIDI

## log Abundances in humans

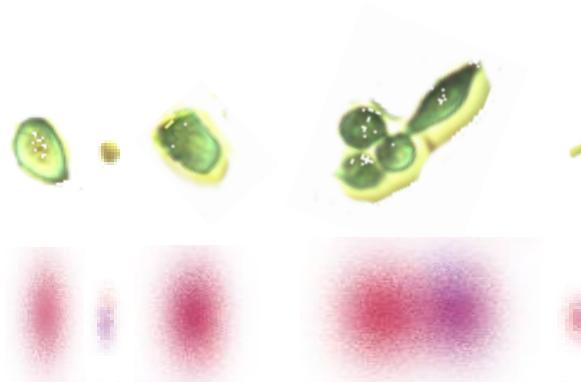
1. Drag cursor around plot area to show information.  
2. Click on element within plot area to go to that element.

H																				He
Li	Be						B	C	N	O	F									Ne
Na	Mg						Al	Si	P	S	Cl									Ar
K	Ca	Sc	Ti	V	Cr	Mn	Fe	Co	Ni	Cu	Zn	Ga	Ge	As	Se	Br				Kr
Rb	Sr	Y	Zr	Nb	Mo	Tc	Ru	Rh	Pd	Ag	Cd	In	Sn	Sb	Te	I				Xe
Cs	Ba	Lu	Hf	Ta	W	Re	Os	Ir	Pt	Au	Hg	Tl	Pb	Bi	Po	At				Rn
Fr	Ra	Lr	Rf	Db	Sg	Bh	Hs	Mt	Uun	Uuu	Uub	Uut	Uuq	Uup	Uuh	Uus				Uuo

La	Ce	Pr	Nd	Pm	Sm	Eu	Gd	Tb	Dy	Ho	Er	Tm	Yb
Ac	Th	Pa	U	Np	Pu	Am	Cm	Bk	Cf	Es	Fm	Md	No



# Identificazione degli eventi



Campione iniettato

Torch

TOF

Mass detection

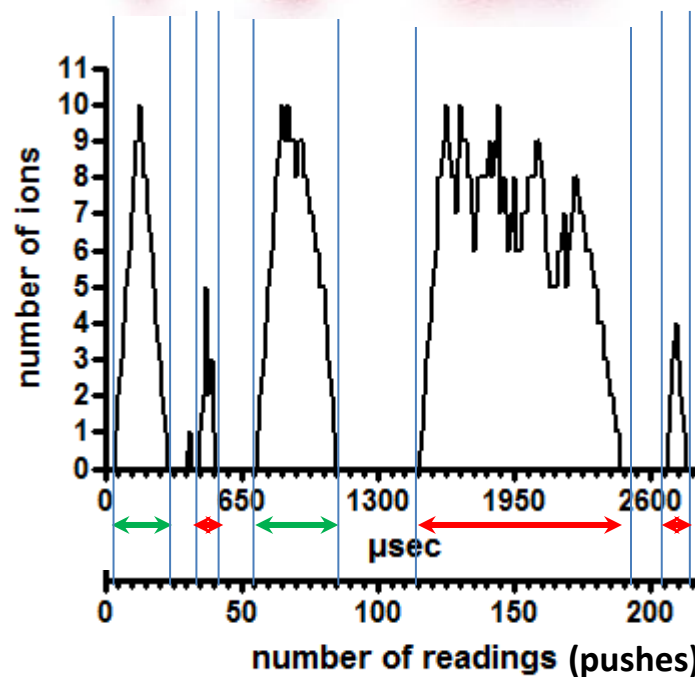
(76800 spectra/s)



1 spectra/13 $\mu$ s = 1 push

Cutoff (cell event):

10-75 pushes



Signal digitization

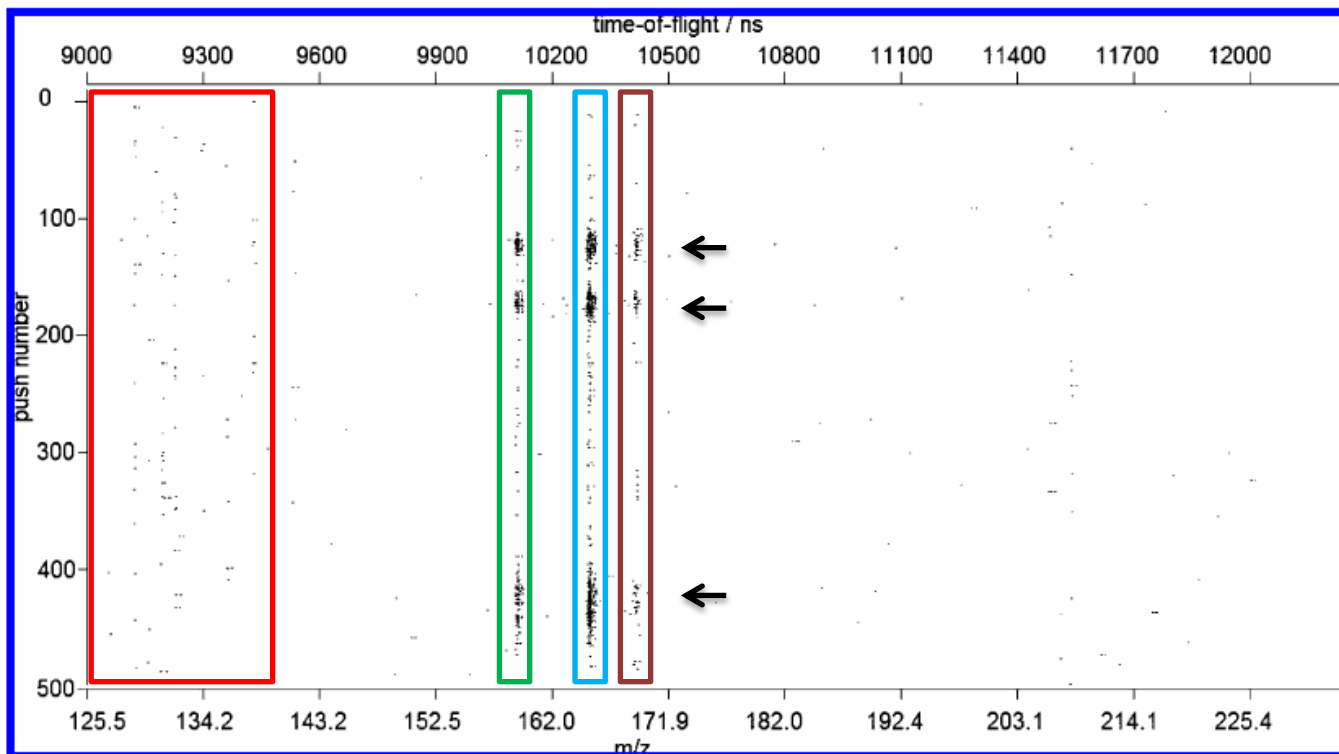
- 103-193 m/z

- 1ns resolution



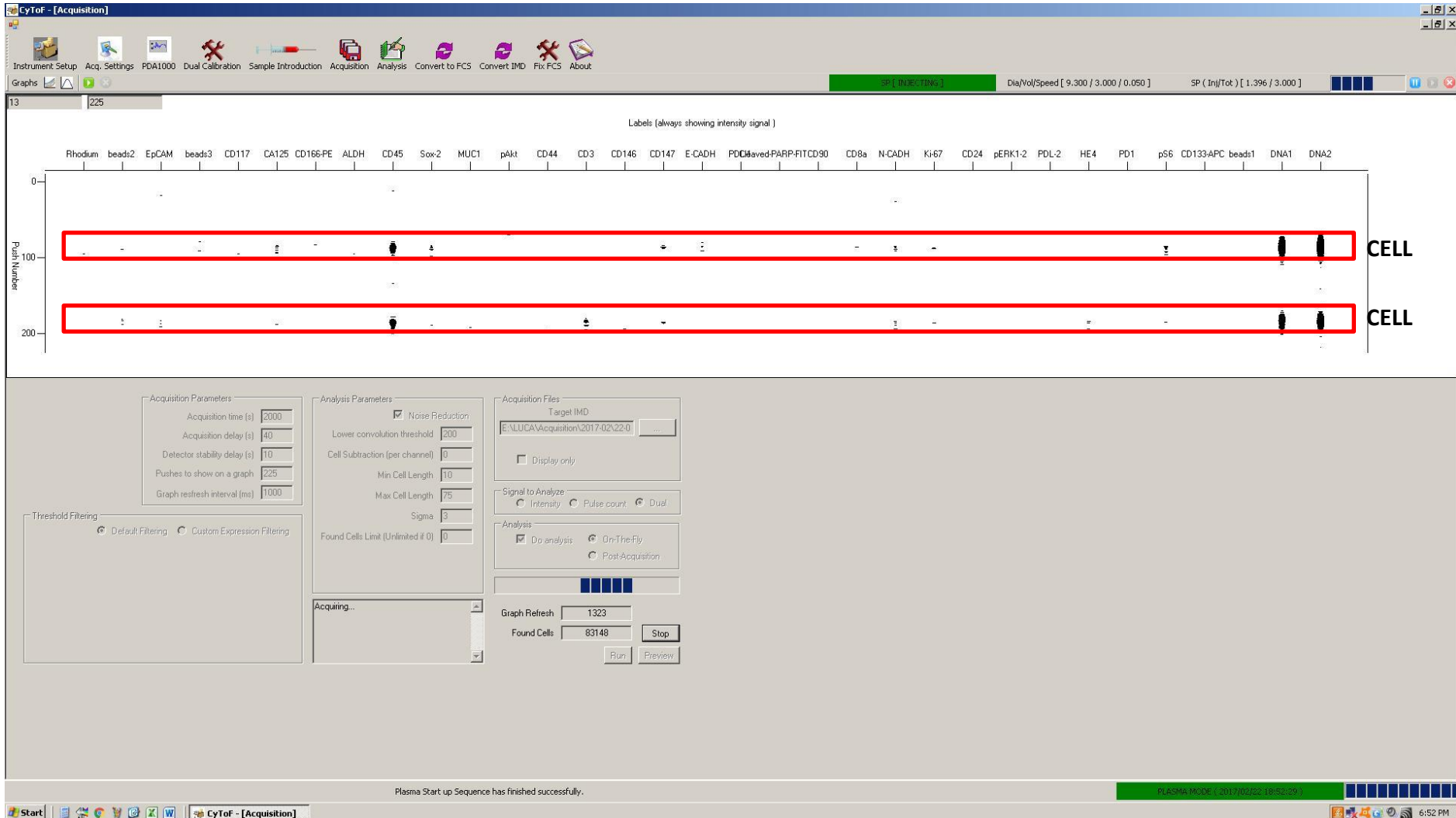


# Identificazione degli eventi

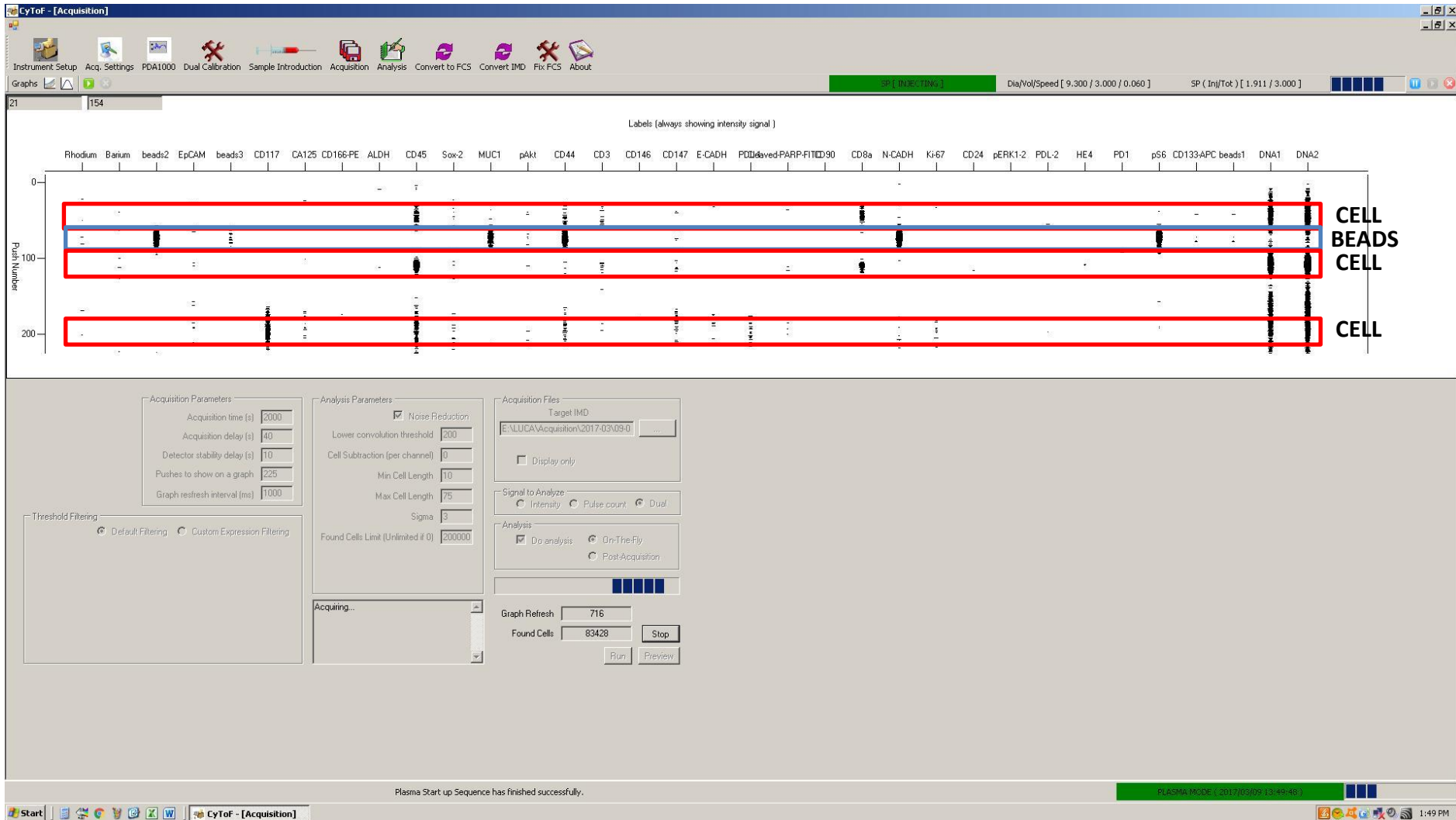


**Figure 4.** Screenshot of 500 consecutive spectra (total of 6.5 ms) for a sample of beads containing **Tb**, **Ho** and **Tm** at 1:2:1 concentration.

# Identificazione degli eventi (densità del campione)



# Identificazione degli eventi (densità del campione)

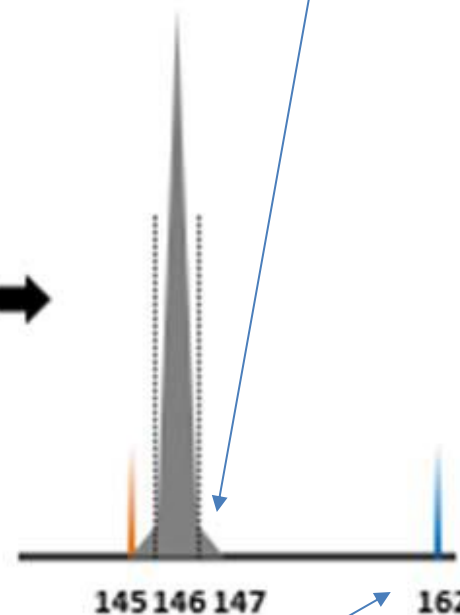
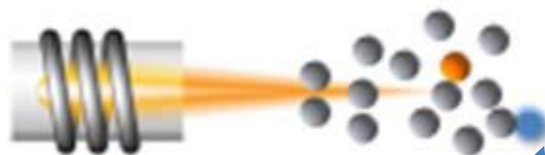
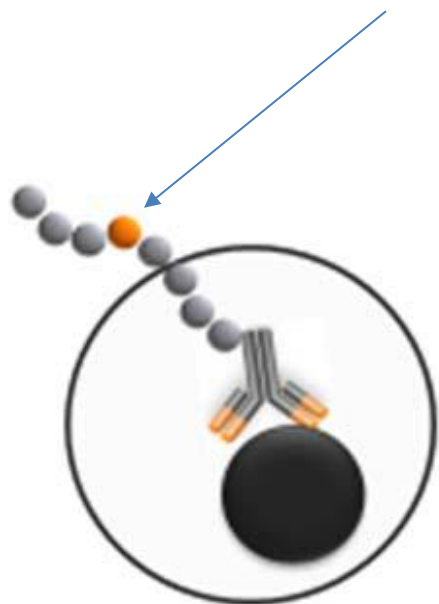




# "Signal Overlap"

Purezza degli isotopi

Sensibilità dello strumento

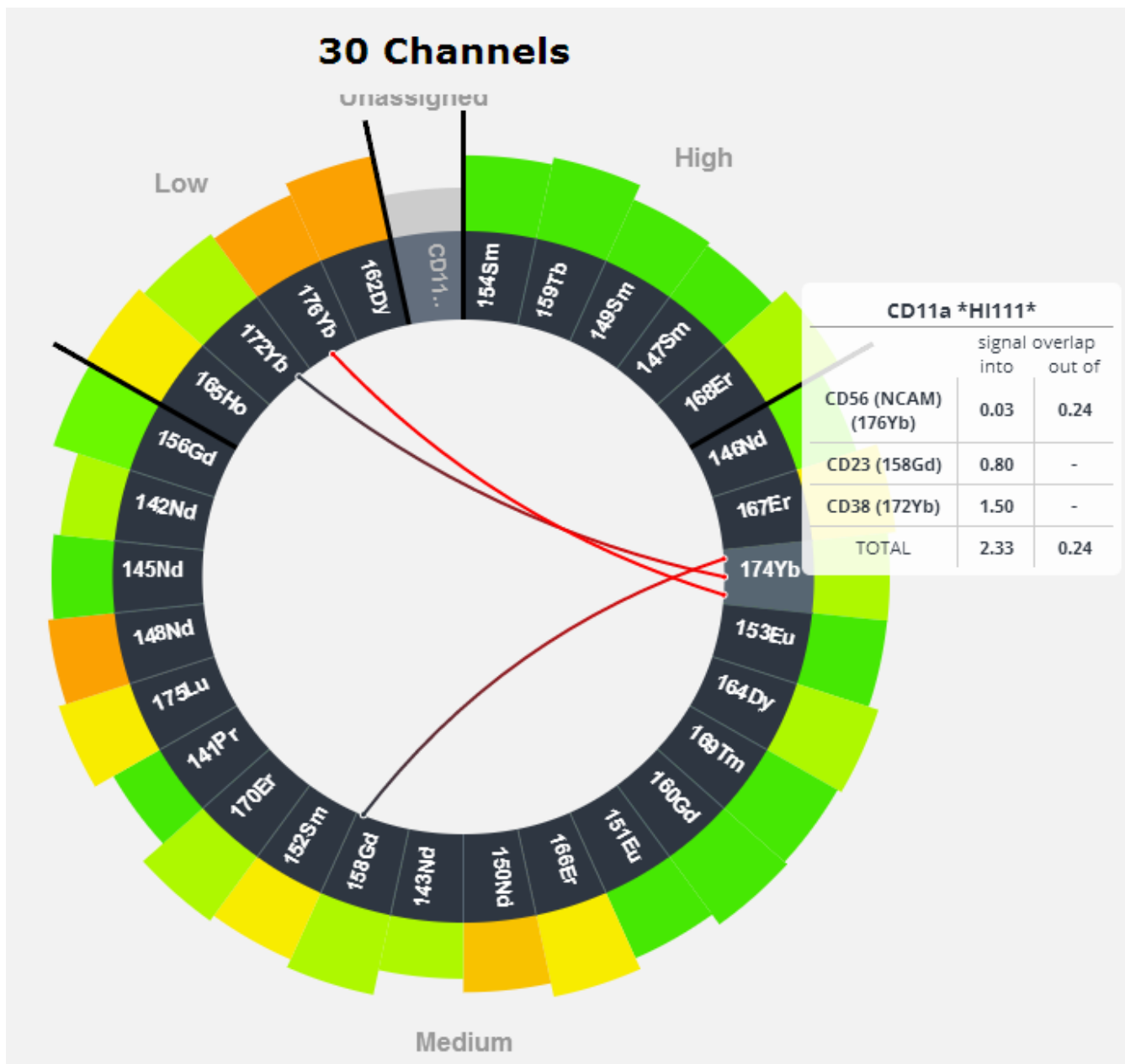


Formazione degli ossidi  
(M+16)



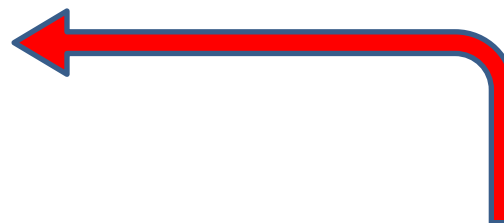
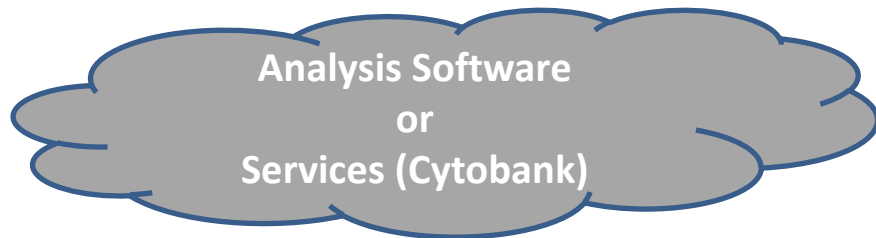


# Signal Overlap (Panel Designer)



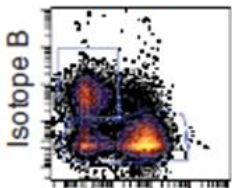
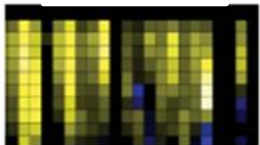

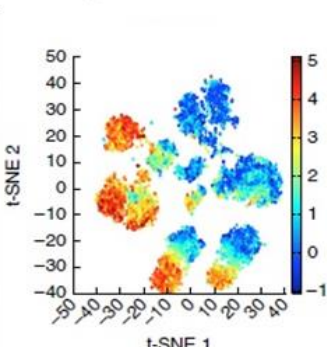
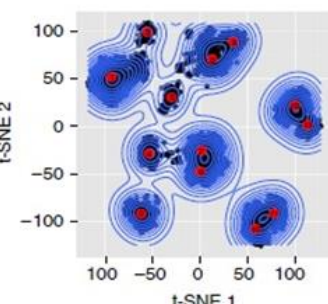


# Analisi dei dati



.FCS files  
(Mass Cytometry data)



2D Plots (FlowJo)  Isotope B Isotope A	Expression & Fold-Changes 	SPADE 
 t-SNE 2 t-SNE 1 viSNE	 t-SNE 2 t-SNE 1 ACCENSE	



# Citometria di massa – CyTOF



## Principali Vantaggi:

- Analisi multiparametrica >35-40 parametri;  
[attualmente in ISS compresa tra i 25 e 30 parametri, Progetto HERCULES (H2020)]
- Analisi contemporanea di antigeni di superficie, intracellulari e fosfoproteine a livello di singola cellula;
- Notevole riduzione del fenomeno di sovrapposizione tra i segnali (non è richiesta compensazione);
- Il costo degli anticorpi (già coniugati con il metallo) è paragonabile a quelli per citofluorimetria;
- Mediante l'utilizzo di kit specifici di coniugazione è possibile produrre Ab-Met "*in-house*";

## Limiti:

- Velocità di acquisizione molto inferiore alla citofluorimetria (~500 cell/s);
- Non è possibile effettuare "cell sorting";
- Solo il 30-40% del campione iniettato viene analizzato;

## Principali Applicazioni:

- Immunofenotipo (Caratterizzazione di popolazioni cellulari e definizione del signaling a livello delle singole sottopopolazioni cellulari)
- Biomarcatori (Identificazione di biomarcatori predittivi della risposta alle terapie anche in specifiche sottopopolazioni)
- Farmacodinamica (Screening farmaci ed identificazione del meccanismo di azione)

**Esperimenti ben costruiti forniscono un numero enorme di informazioni!**



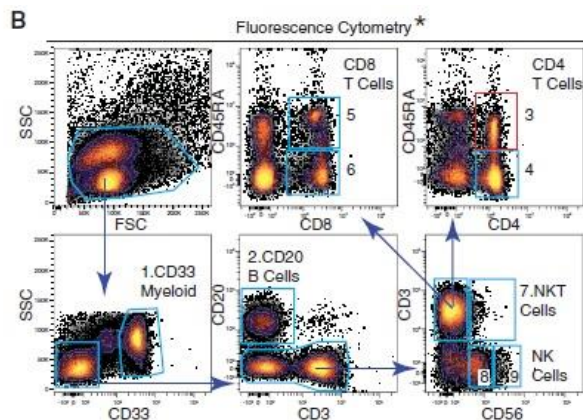




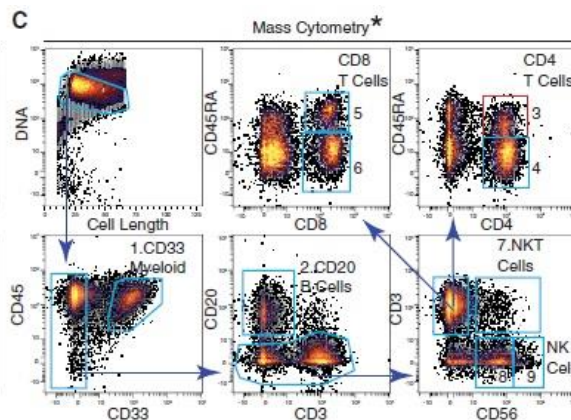
# “Single-Cell Mass Cytometry of Differential Immune and Drug Responses Across a Human Hematopoietic Continuum - Bendall SC et al. Science. 2011”



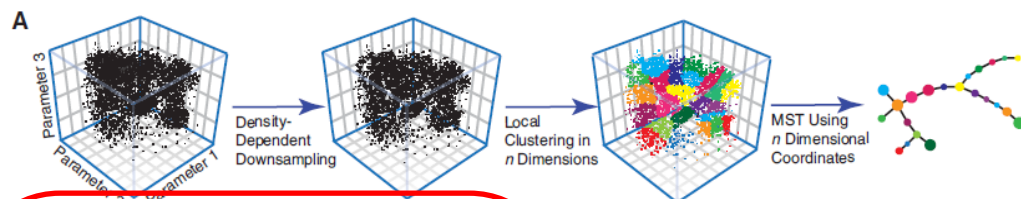
Flow Cytometry



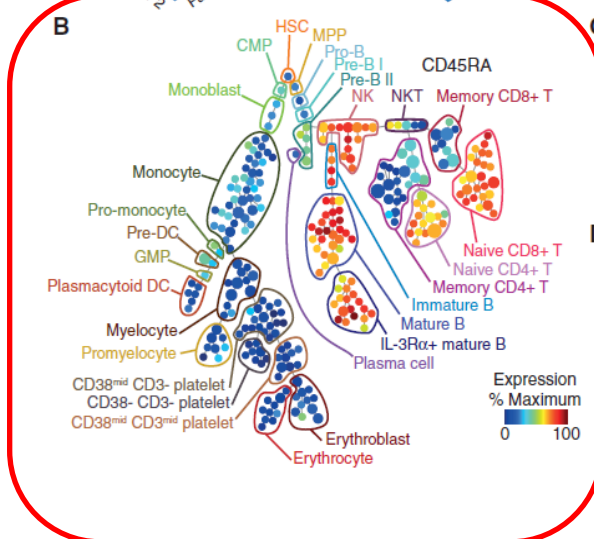
Mass Cytometry



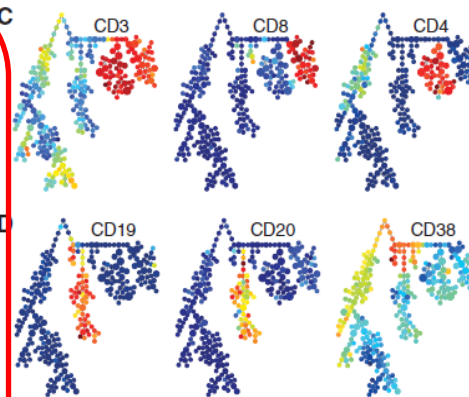
SPADE analysis



13 marcatori di superficie  
+ 18 intracellulari  
(comprese fosfoproteine)



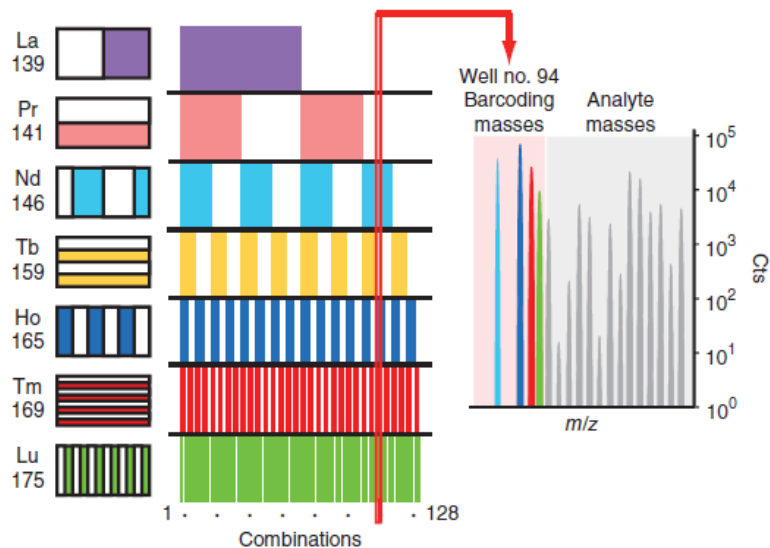
T cell lineage



B cell lineage



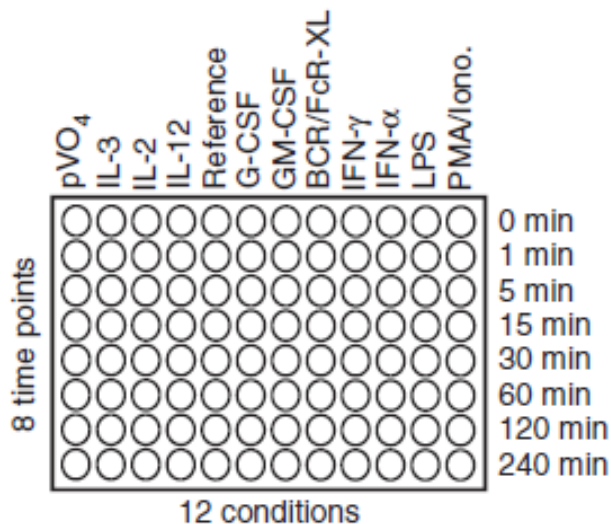
## "CELL BARCODING"



Le cellule presenti in ogni pozzetto sono etichettate con una combinazione unica di 7 Lantanidi ( $2^n$  possibili combinazioni generate)

### Vantaggi del Barcoding:

- Riduzione dell'errore di meccanico (pipetta)
- Variazioni nello staining
- Consumo di anticorpi
- Riduzione tempo di acquisizione

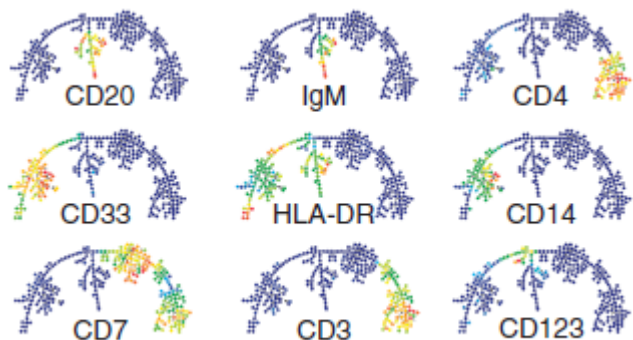


12 differenti stimoli - 8 differenti incubazioni

Cellule raccolte e colorate con 9 marcatori di superficie e 14 marcatori del signaling

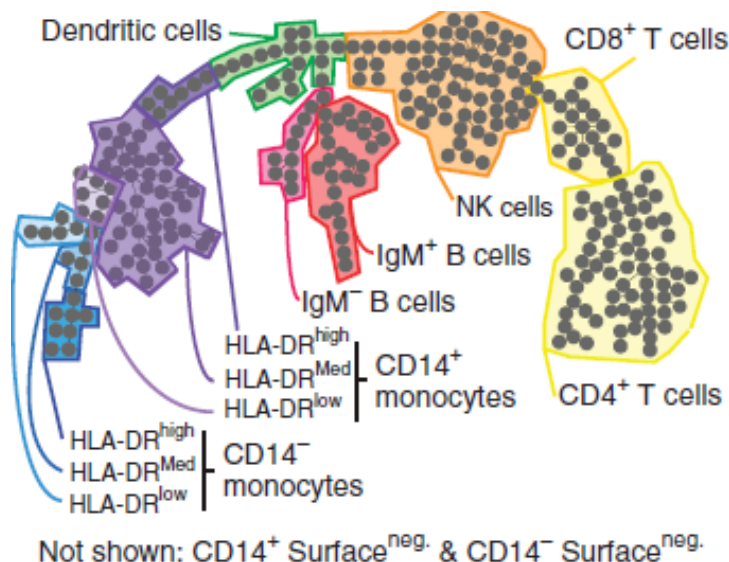


# Multiplexed mass cytometry profiling of cellular states perturbed by small-molecule regulators - Bodenmiller et al. *Nature Biotechnology* 30 (9) 2012

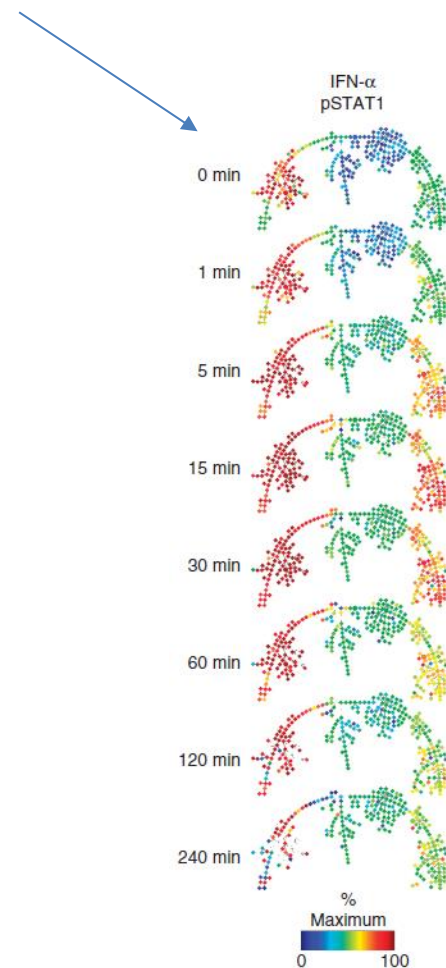


Espressione  
dei 9 marcatori di  
superficie

Identificazione  
delle  
sottopopolazioni



Valutazione dell'attivazione del signaling  
a livello di ogni sottopopolazione ed in  
base al tempo di incubazione per ogni  
stimolo

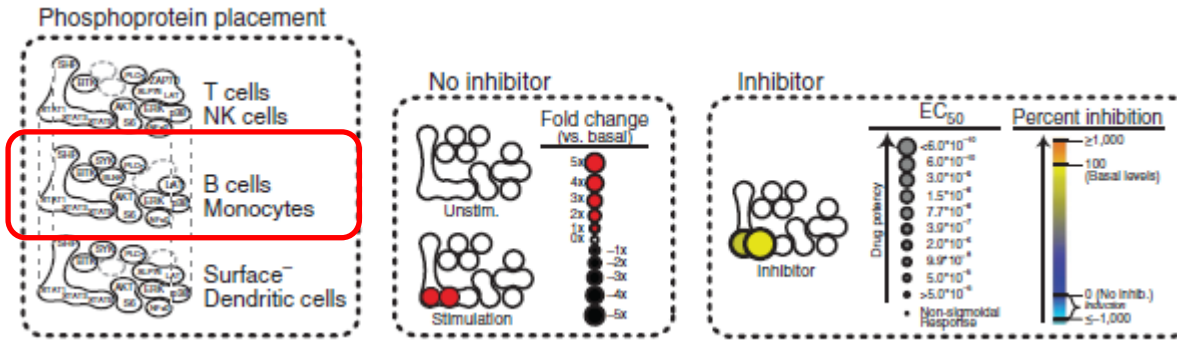




# Multiplexed mass cytometry profiling of cellular states perturbed by small-molecule regulators - Bodenmiller et al. *Nature Biotechnology* 30 (9) 2012

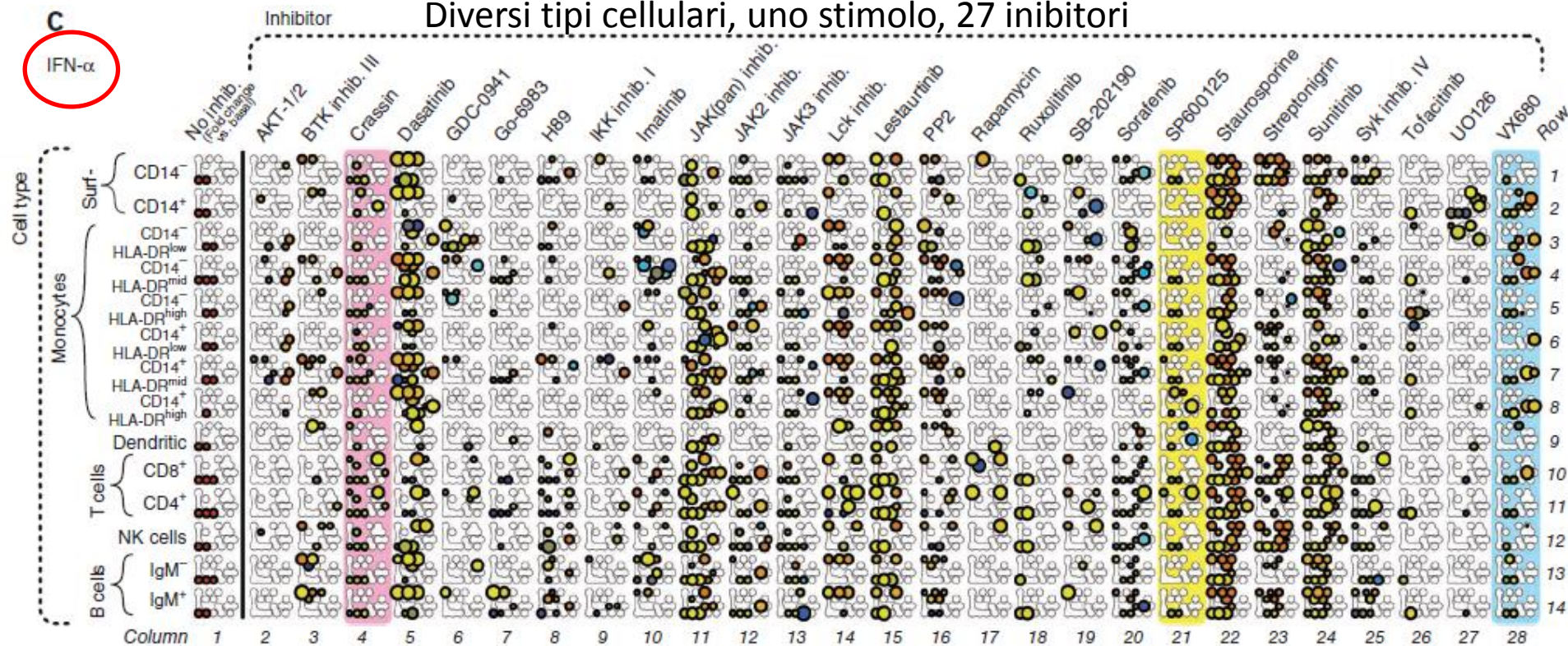


Signaling network



C

Diversi tipi cellulari, uno stimolo, 27 inibitori



# *Thank you!*



Valentina Tirelli - Ed. 15, piano A, st. 12 - t. 2576

Luca Pasquini - Ed.1, piano G, st.16-19 - t. 2422-2632

Prenotazioni Analisi Citometria:

<http://gscf.iss.it/dw/doku.php?id=aree:citometria:prenotazioni>