

# AML MRD by multiparameter flow cytometry using LAIP and LSC: Methodological aspects in a multicentric study of the French AML Intergroup

Adriana PLESA , Lyon , France

ESCCA, Thessaloniki, Greece, 27 Septembre 2017



*Hôpitaux de Lyon*



Collaborative InterGroup for Acute Leukemia









**Centre Hospitalier Régional  
Universitaire de Lille**

# Map of the centers involve on French AML Clinical Trial



AML Clinical Centers: 54

	Patients inclusion		Nb of center
	45%		9
	14%	}	5
	34%		25
	7%		15
			

# AML MRD flow

## 1) *Why?*

- *Flow Labs and French Clinical Trial organisation*
- *ELN 2017 Guidelines requirements- MRD mandatory (molecular and/or flow)*
- *Harmonisation of Flow Clinical MRD report*

## 2) How?

Methodology multicentric approach:

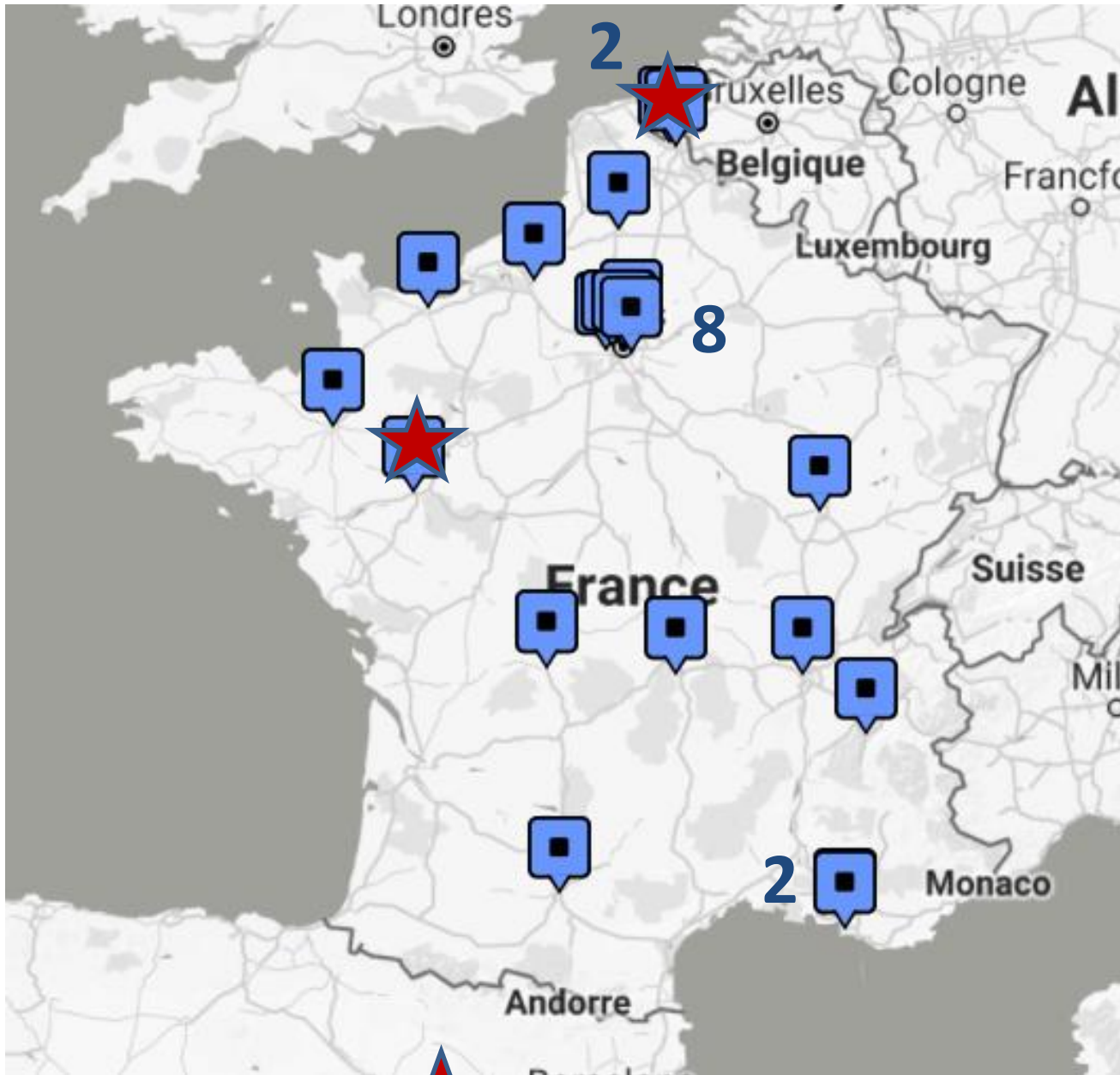
- panel design: simple, reproducible, sensitivity, cut
- CANTO vs NAVIOS « miroir »
- gating strategy

## 3) Ready to start

- CQA: verify gating strategy
- CQE: global check of panel implementation ( n/reg BM)
- Patient follow up



**25 Flow Labs: 13 BD (Canto / Lyric) - 12 BC (Navios)**



**2 Molecular Labs**

**LAM ≥ 18 ans et < 61 ans**

**Induction : Randomisation R1 : R1-DAUNO vs R1-IDA**

**Daunorubicine** : 90 mg/m<sup>2</sup>/j J1 à J3 en 30 mn  
**Cytarabine** : 200 mg/m<sup>2</sup> IV SE 24h J1 à J7  
Myélogramme à J15, G-CSF si < 5% blastes

**Idarubicine** : 9 mg/m<sup>2</sup>/j J1 à J5 en 30 mn  
**Cytarabine** : 200 mg/m<sup>2</sup> IV SE 24h J1 à J7  
Myélogramme à J15, G-CSF si < 5% blastes

**Bilan évaluation entre J28-J35**

(Sang, moelle + MRD1\*\*)

Si RC/RCp/Rci

Si Echec

**Randomisation R2 : R2-IDAC vs R2-HDAC**

**Consolidation**

**Rattrapage**

Selon R2

Selon R2

Selon R2

Selon R2

**IDAC 1**

**Cytarabine** 1.5g/m<sup>2</sup>/12h J1, J3, J5  
G-CSF à partir de J8

**HDAC 1**

**Cytarabine** 3g/m<sup>2</sup>/12h J1, J3, J5  
G-CSF à partir de J8

**IDAC 1 (= conso 1 si RC)**

**Cytarabine** 1.5g/m<sup>2</sup>/12h J1, J3, J5  
G-CSF à partir de J8

**HDAC 1 (= conso 1 si RC)**

**Cytarabine** 3g/m<sup>2</sup>/12h J1, J3, J5  
G-CSF à partir de J8

**Evaluation post IDAC 1 ou HDAC 1**

(Sang, moelle + MRD2\*\*)

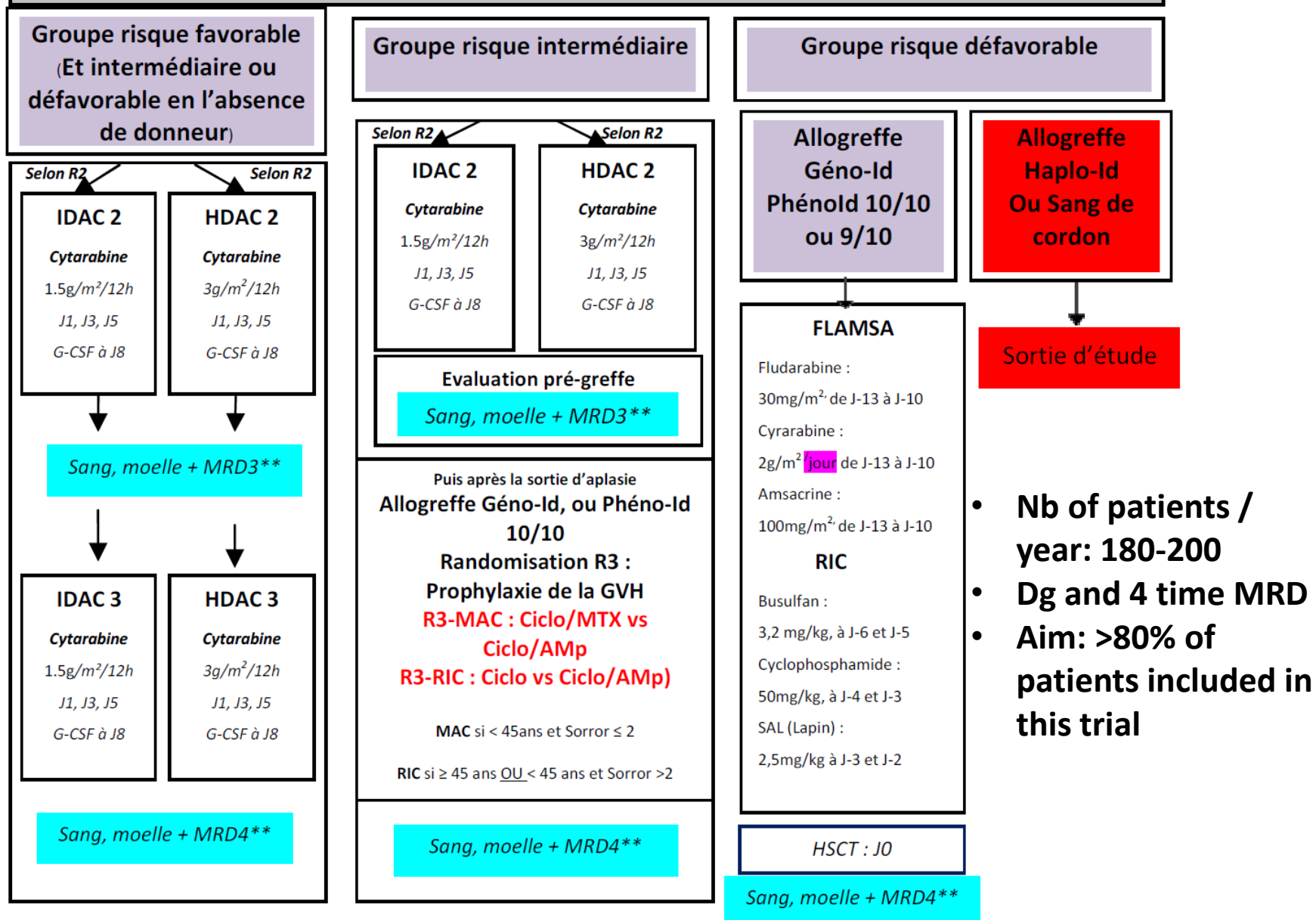
Si RC/RCp/Rci persistante

Si Echec

\*\*MRD2  
décisionnelle pour  
allogreffe

Sortie d'étude

# Traitement post rémission



\*\* : MRD centralisée (ALFA ou GOELAMS) pour NPM1mut (en local pour les autres marqueurs)

## Diagnosis and management of AML in adults: 2017 ELN recommendations from an international expert panel

Hartmut Döhner,<sup>1</sup> Elihu Estey,<sup>2</sup> David Grimwade,<sup>3</sup> Sergio Amadori,<sup>4</sup> Frederick R. Appelbaum,<sup>2</sup> Thomas Büchner,<sup>5</sup> Hervé Dombret,<sup>6</sup> Benjamin L. Ebert,<sup>7</sup> Pierre Fenaux,<sup>8</sup> Richard A. Larson,<sup>9</sup> Ross L. Levine,<sup>10</sup> Francesco Lo-Coco,<sup>4</sup> Tomoki Naoe,<sup>11</sup> Dietger Niederwieser,<sup>12</sup> Gert J. Ossenkoppele,<sup>13</sup> Miguel Sanz,<sup>14</sup> Jorge Sierra,<sup>15</sup> Martin S. Tallman,<sup>10</sup> Hwei-Fang Tien,<sup>16</sup> Andrew H. Wei,<sup>17,18</sup> Bob Löwenberg,<sup>19</sup> and Clara D. Bloomfield<sup>20</sup>

**Monitoring of MRD.** Two approaches can be used to detect MRD, that is, multiparameter flow cytometry (MFC) and molecular techniques, including real-time quantitative PCR (RT-qPCR), digital PCR, and next-generation sequencing–based technologies. Standardized RT-qPCR assays are now available to detect AML-associated genetic lesions (Table 4). Each methodology

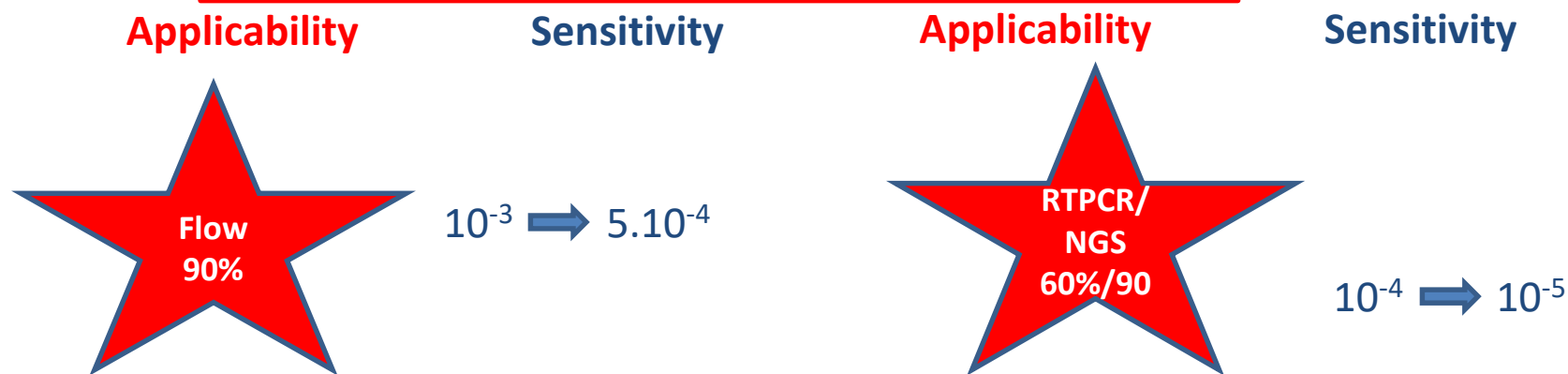


Table 6. Response criteria in AML

Category	Definition	Comment
<b>Response</b>		
CR without minimal residual disease (CR <sub>MRD-</sub> )	If studied pretreatment, CR with negativity for a genetic marker by RT-qPCR, or CR with negativity by MFC	Sensitivities vary by marker tested, and by method used; therefore, test used and sensitivity of the assay should be reported; analyses should be done in experienced laboratories (centralized diagnostics)

# New paradigm in MRD LAM: LSC approach

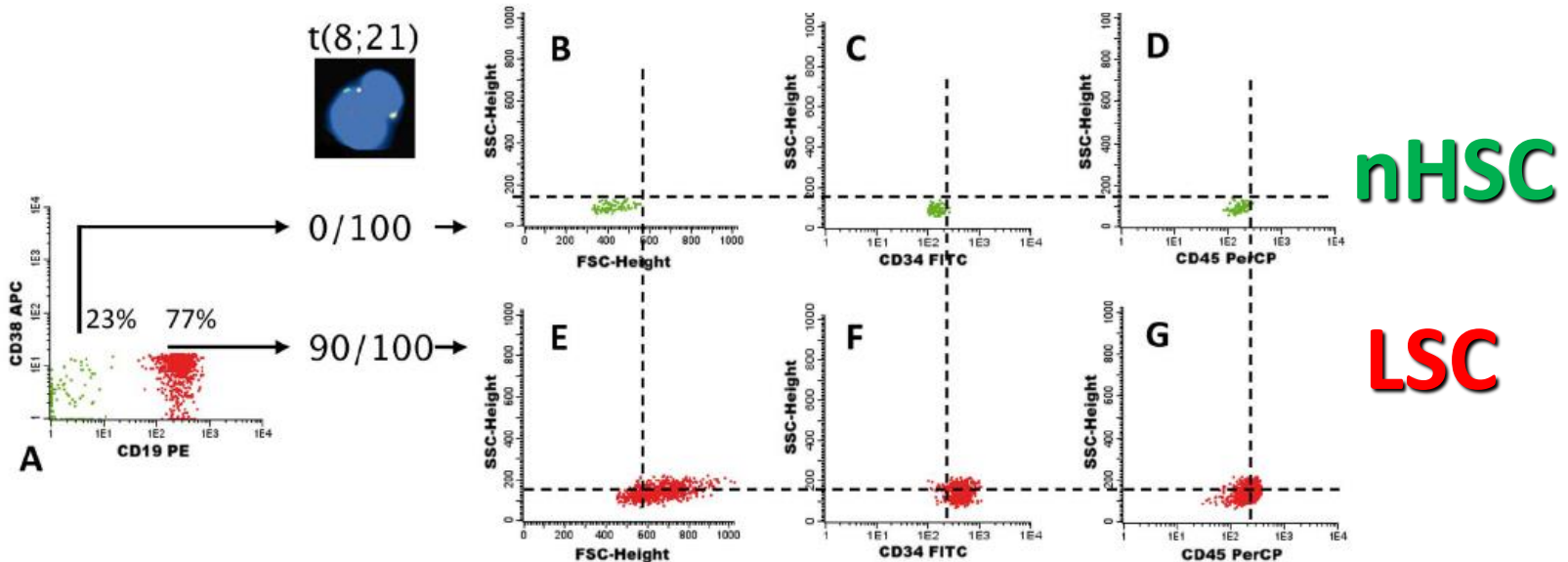
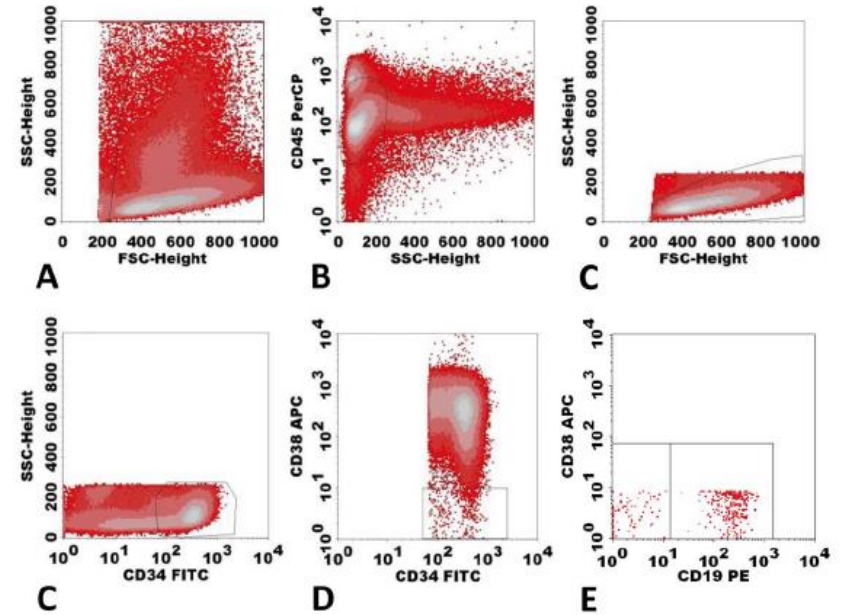
OPEN ACCESS Freely available online



## Leukemic Stem Cell Frequency: A Strong Biomarker for Clinical Outcome in Acute Myeloid Leukemia

Monique Terwijn<sup>1</sup>, Wendelien Zeijlemaker<sup>1</sup>, Angèle Kelder<sup>1</sup>, Arjo P. Rutten<sup>1</sup>, Alexander N. Snel<sup>1</sup>, Willemijn J. Scholten<sup>1</sup>, Thomas Pabst<sup>2</sup>, Gregor Verhoef<sup>3</sup>, Bob Löwenberg<sup>4</sup>, Sonja Zweegman<sup>1</sup>, Gert J. Ossenkoppele<sup>1</sup>, Gerrit J. Schuurhuis<sup>1\*</sup>

<sup>1</sup>Department of Hematology, VU University Medical Center, Amsterdam, The Netherlands, <sup>2</sup>Department of Medical Oncology, Inselspital, Bern University Hospital, University of Bern, Bern, Switzerland, <sup>3</sup>Department of Hematology, University Hospital Leuven, Leuven, Belgium, <sup>4</sup>Department of Hematology, Erasmus University Medical Center, Rotterdam, The Netherlands



I

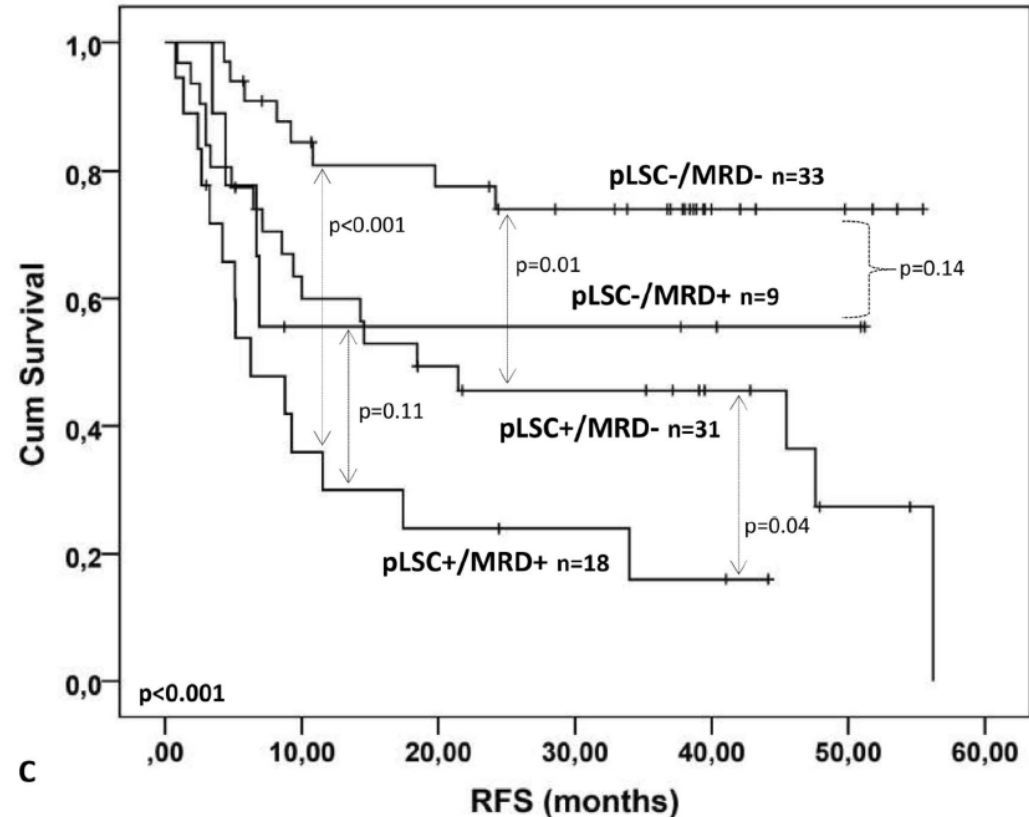
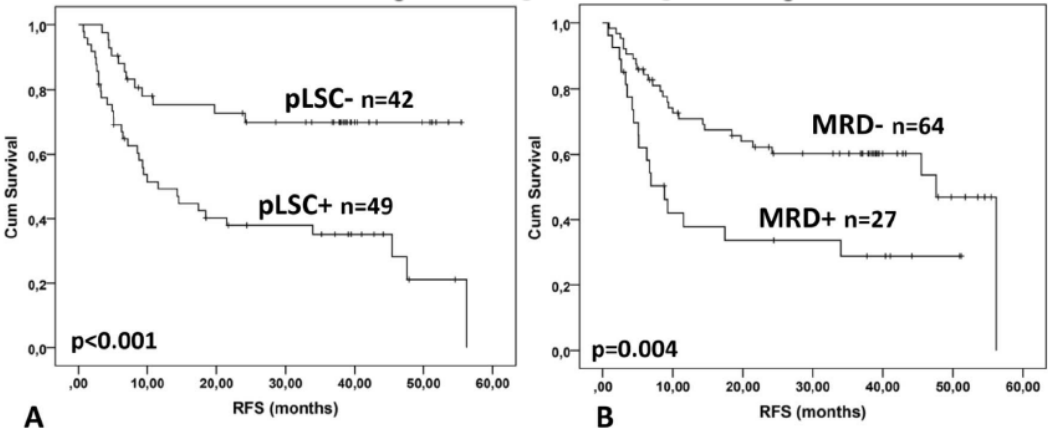
I



# New concept of AML follow up: Scoring of MRDflow

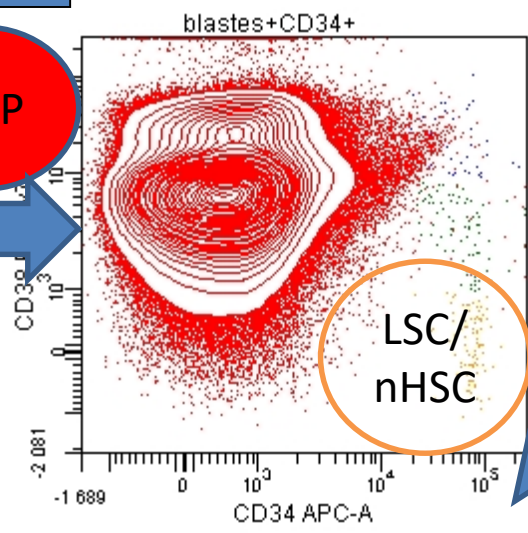
Leukemic Stem Cell Frequency in AML

## (LAIP/DFN/LSC)

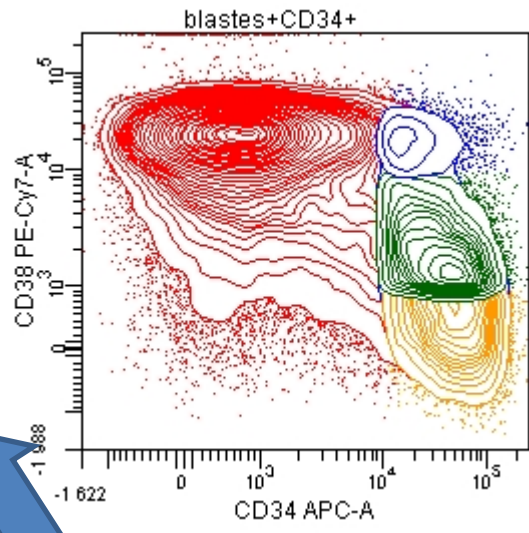


AML dg CN, NPM1+FLT3ITD- CN, NPM1+FLT3ITD+ +8, del20q-, NPM1-FLT3ITD+

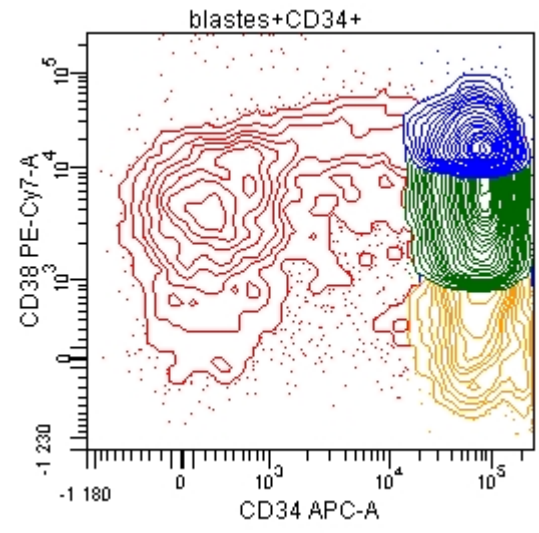
LAIP



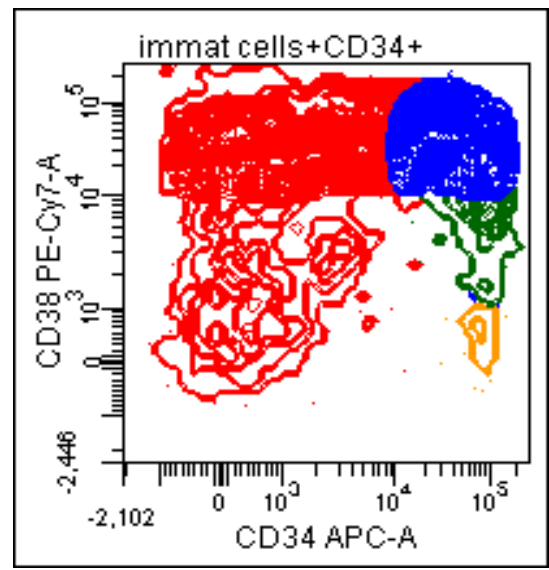
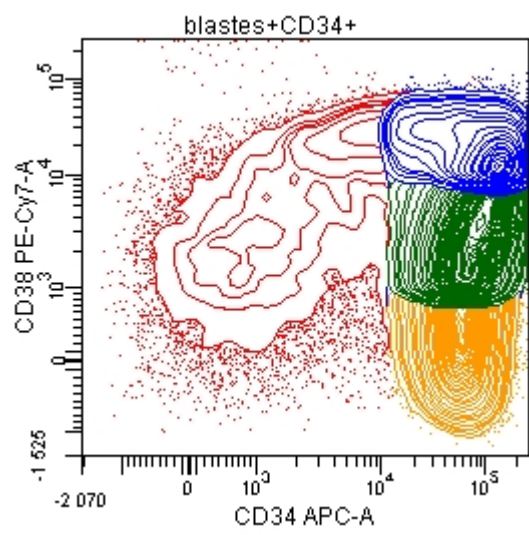
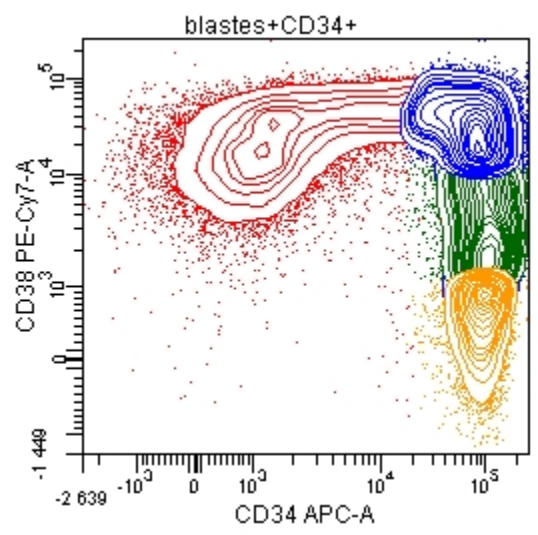
Caryotype CX>5, WT1+, dupMLL



EVI1+, del7q



nBM



# AML MRD flow

## 1) Why?

- Flow Labs and French Clinical Trial organisation
- ELN 2017 Guidelines requirements- MRD mandatory (molecular and/or flow)
- Harmonisation of Flow Clinical MRD report

## 2) *How?*

### ***Methodology multicentric approach:***

- ***panel design: simple, reproducible, sensitivity, cost***
- ***CANTO vs NAVIOS « miroir »***
- ***gating strategy***

## 3) Ready to start

- CQA: verify gating strategy
- CQE: global check of panel implementation ( n/reg BM)
- Patient follow up

# MRD Flow panel design rational

**Aim:** simple, reproducible, sensitivity, cost

- 1) **LAIP approach** (*Leukemia Associated ImmunoPhenotype*)- at diagnosis
- 2) **DFN approach-** *Different of Normal- pattern of normal myeloid differenciation (reactionel vs regeneration BM)*
- 3) **LSC approach** : 34+38-

## **Rational**

- LAIP (Kern, Venditti, Wood, Schuurhuis, Freeman ...)
- Backbone 34/38 CSH/LSC approach (Dick&Bonnet, Nature 2009)
- LMPP-like/GMP-like LSC approach (Goardon, Cancer Cell 2011)
- LSC Flow Signature (Schuurhuis, PlosOne2014): quantification at dg and in MRD follow up

	FL1	FL2	FL3	FL4	FL5	FL6	FL7	FL8
T 1	CD7/56	CD13	CD33	CD34	CD38	CD117	CD19	CD45
T 2	CD90	MIX* TIM3+CLL1	CD123	CD34	CD38	CD117	CD45RA	CD45

\*New markers of LSC prospectively evaluated in addition to TIM3 and CLL1: CD97, GPR56...

Supp Tube	CD36	CD11b	CD33	CD34	HLADR	CD117	CD4	CD45
-----------	------	-------	------	------	-------	-------	-----	------

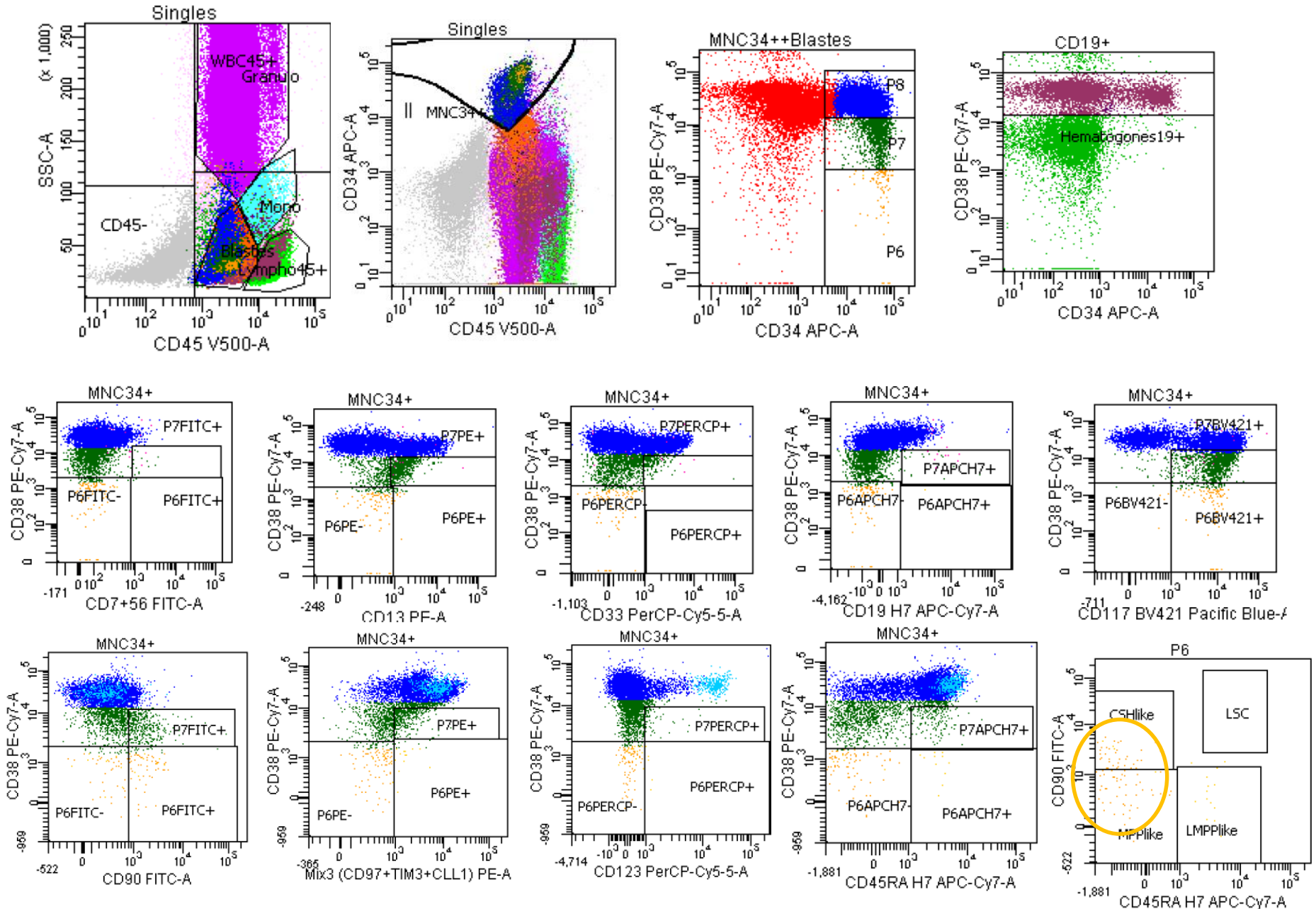
## Leukemic stem cells: identification and clinical application

Diana Hanekamp<sup>1</sup> · Jacqueline Cloos<sup>1,2</sup> · Gerrit Jan Schuurhuis<sup>1</sup>

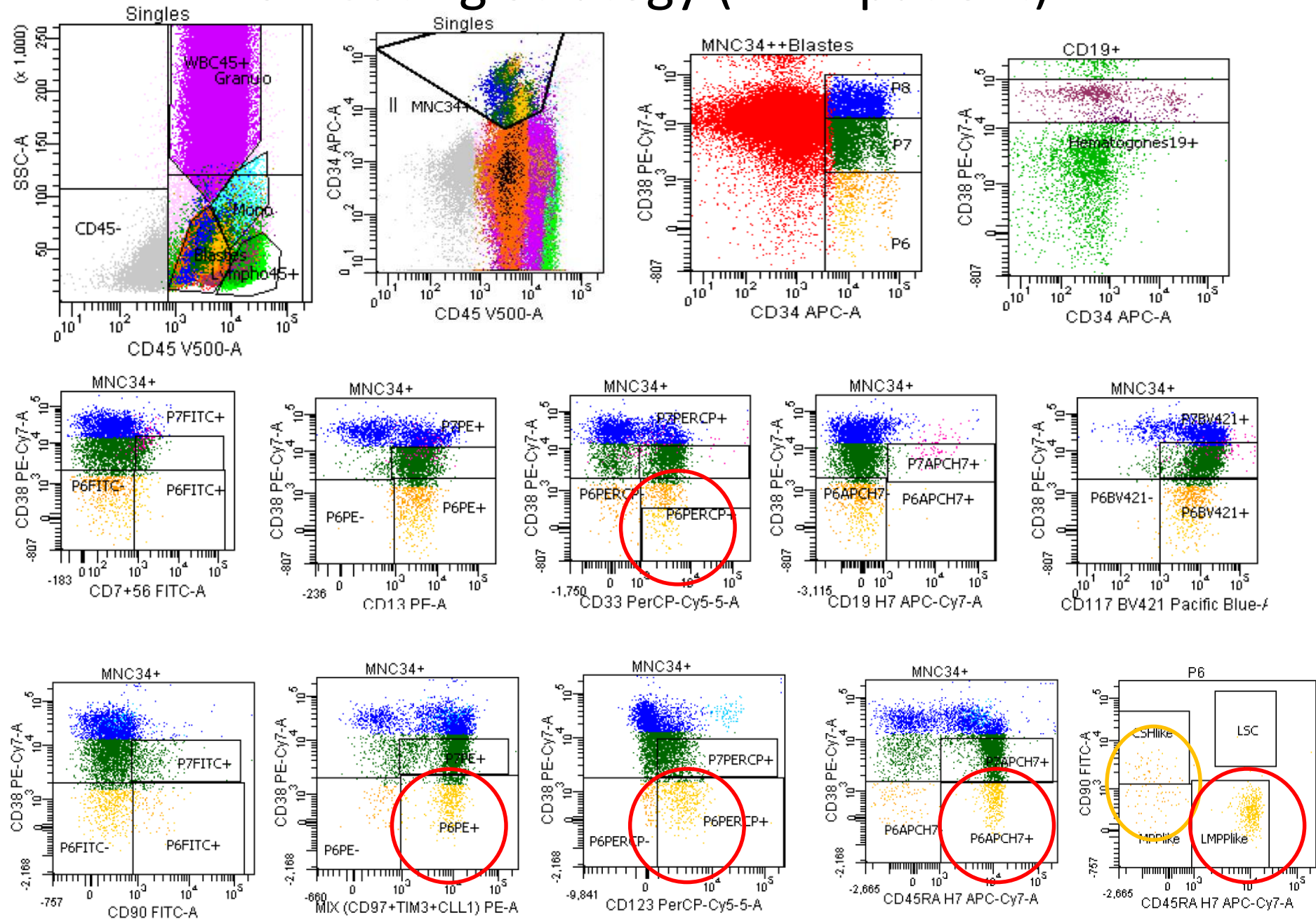
**Table 1** Distinct leukemic stem cell markers

Marker	Identified as	Expression	References				
			Normal	In AML (%)	HSC	CD34+ CD38– LSC	
★ IL1RAP	IL1R3	T cells		79	–	+	[70–72]
★ CLL-1	CLEC12A, MICL, DCAL-2	Myeloid cells		70	–	+	[6]
★ TIM-3	T-cell Ig Mucin 3	Activated T cells, NK cells		91	–	+	[73]
★ CD2	SRBC, LFA2, T11	T cells, NK cells		87	–	+	[14]
★ CD7	GP40, TP41, LEU-9	T cells		43	–	+	[6]
CD11b	Integrin alpha M, Mac-1	Myeloid cells		55	–	+	[6]
CD22	BL-CAM, Siglec-2	B cells		51	–	+	[6]
★ CD25	IL2RA, TAC	Activated B and T cells		25	–	+	[74]
★ CD33	P67, Siglec-3	Myeloid cells, NK cells		82	+	++	[6] [75]
★ CD44	Adhesion molecule	Ubiquitously		100	+	++	[6]
★ CD45RA	Tyrosine phosphatase receptor type C	T cells, myeloid cells		65	–	+	[76]
★ CD47	Integrin-associated protein (IAP)	Ubiquitously		100	+	++	[77]
★ CD56	N-CAM, MSK39	NK cells, activated T cells		32	–	+	[6]
★ CD96	TACTILE	Activated T cells		33	–	+	[6]
★ CD99	MIC2, single-chain type-1 glycoprotein	Myeloid cells		83	–	+	[78]
★ CD123	IL3R	Myeloid cells		82	+	++	[6] [48] [79]

# LSC Gating Strategy (nBM)



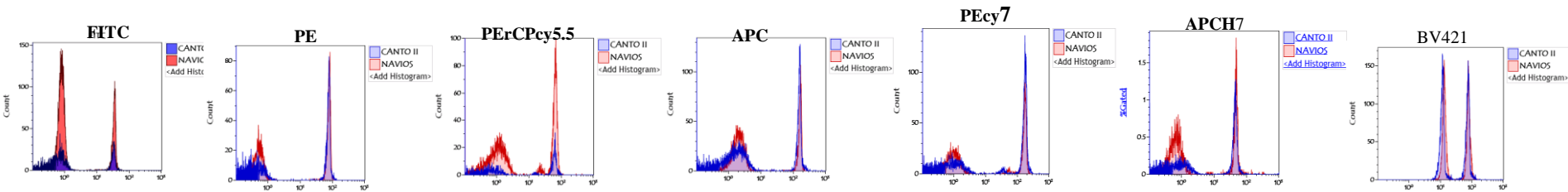
# LSC Gating Strategy (AML patient)



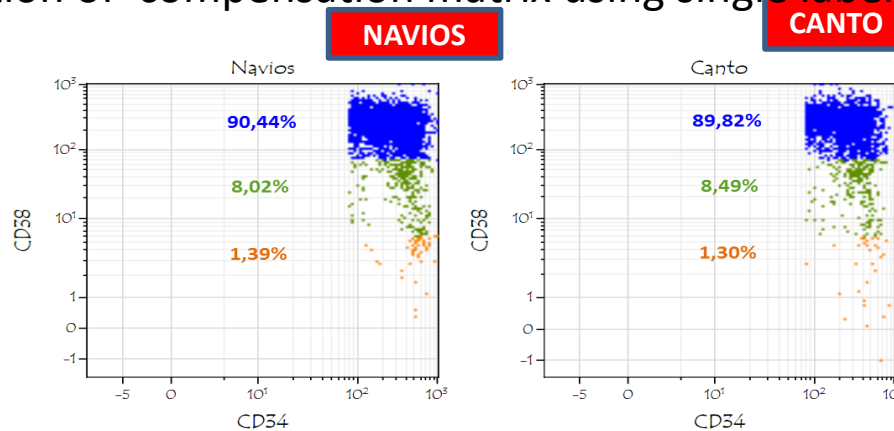
# Challenge : to achieve harmonisation of MRD results between 25 FlowLabs

## CANTO vs NAVIOS « miroir »

- 1) Harmonisation of preanalytical treatment of the samples: BM dilution ctr, Bulk Lysis, nb of events acquisition (**mandatory >500 000/tube**)
- 2) « Mirroring » CANTO vs NAVIOS platforms
  - Setting of voltage for the chosen channel trough acquisition of rainbow beads without compensation to reach target of MFI values Multicentrique on the CANTO platform;
  - transposition to NAVIOS platform by:  $\text{New MFI Target} = \text{CANTO target} / 256$



- 3) Evaluation of the homogeneity of the results using normal BM cells labeled with the panel after elaboration of compensation matrix using single labels Ab



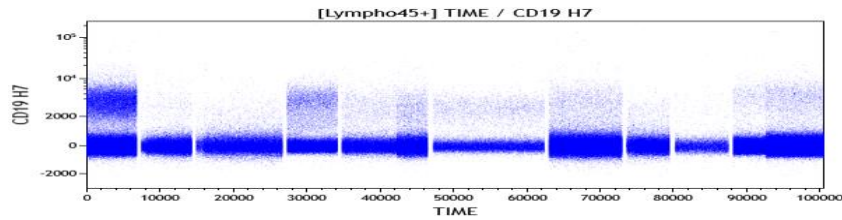
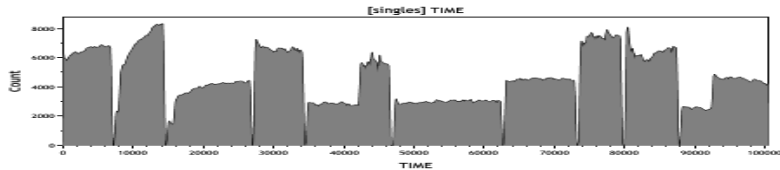
Comparable flow cytometry data can be obtained with two types of instruments, Canto II, and Navios. A GEIL study.

Solly F, Rigollet L...Lacombe F, Béné MC. *Cytometry A*. 2013 Dec;83(12):1066-72.

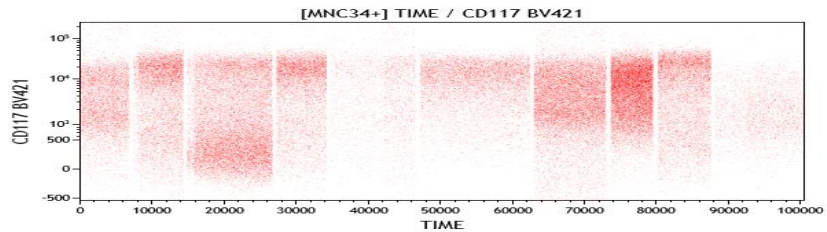


# Merged dataset obtained from 10 BM samples shared between Lyon and Lille Labs

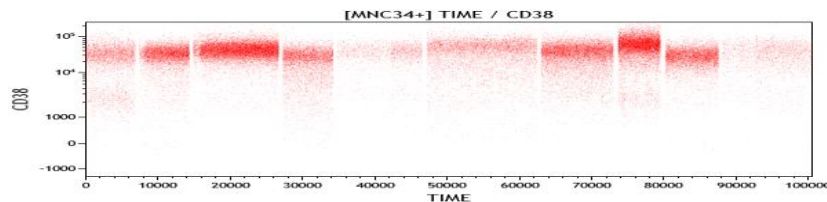
## T1 CANTO (Lyon)



Gate Ly CD45



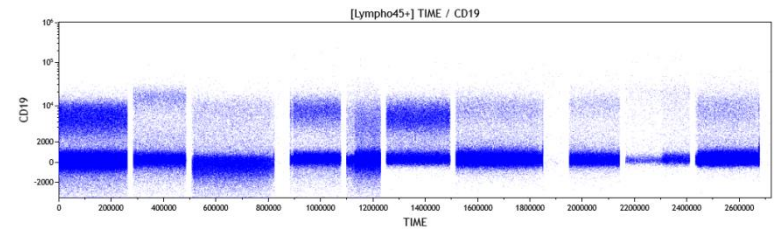
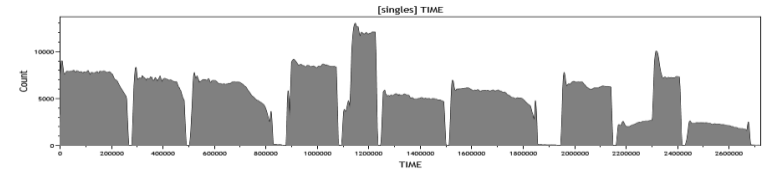
CD117



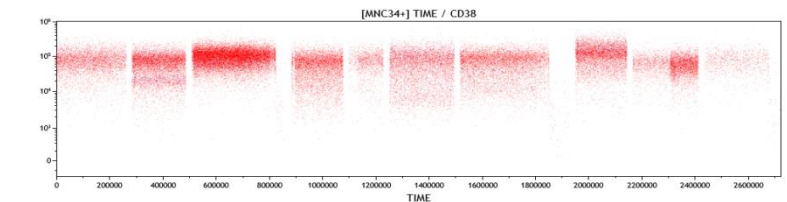
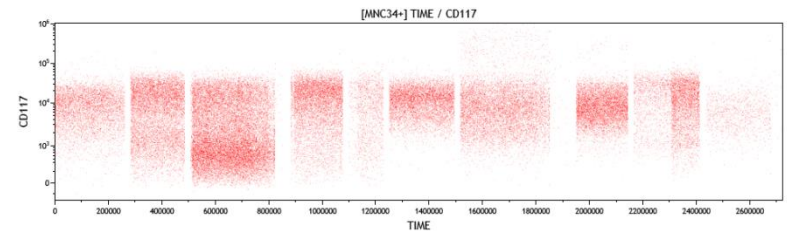
CD38

Gate MNC CD34+

## T1 NAVIOS (Lille)



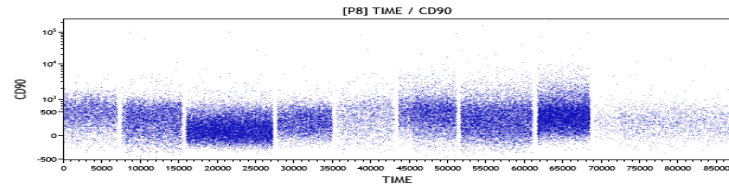
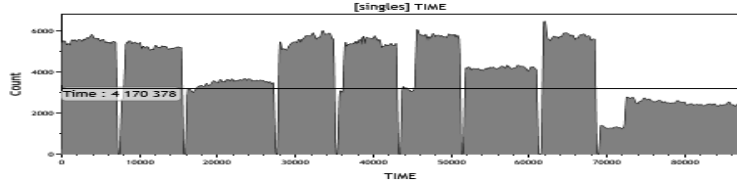
Gate Ly CD45



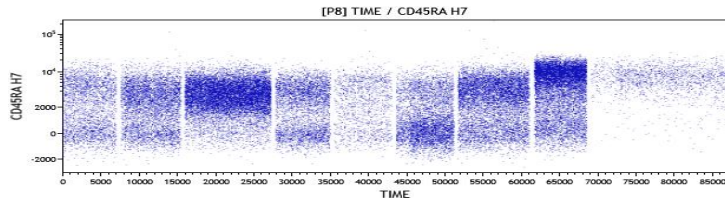
Gate MNC CD34+

# Merged dataset obtained from 9 BM samples shared between Lyon and Lille Labs

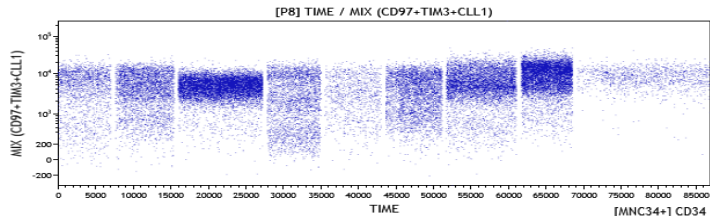
## T2 Canto (LSC markers)



CD90

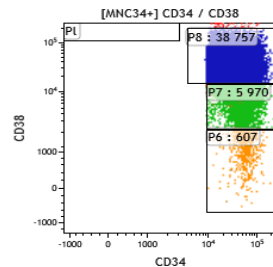


CD45RA

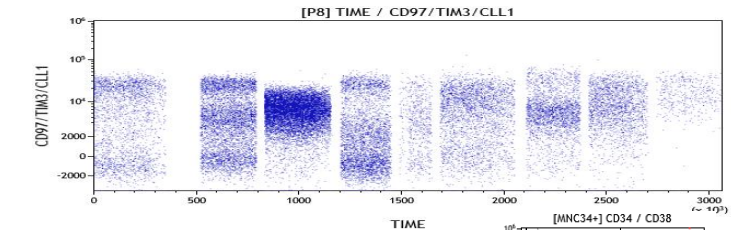
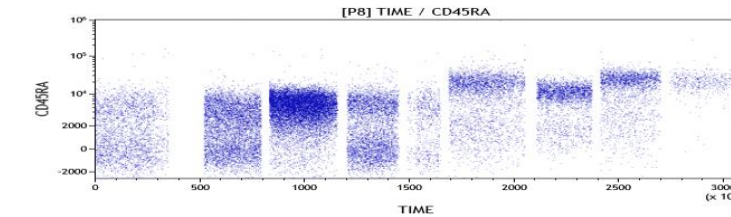
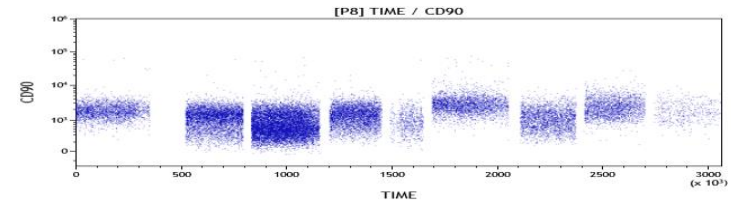
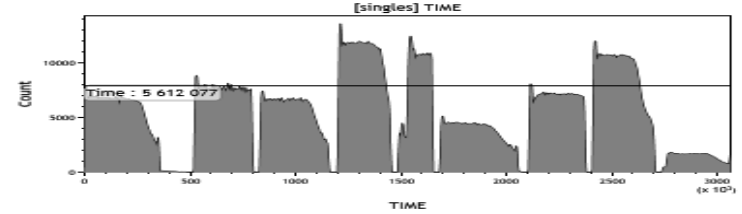


CLL1/Tim3

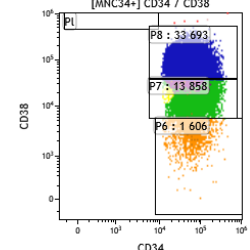
Gate CD34+CD38+hi

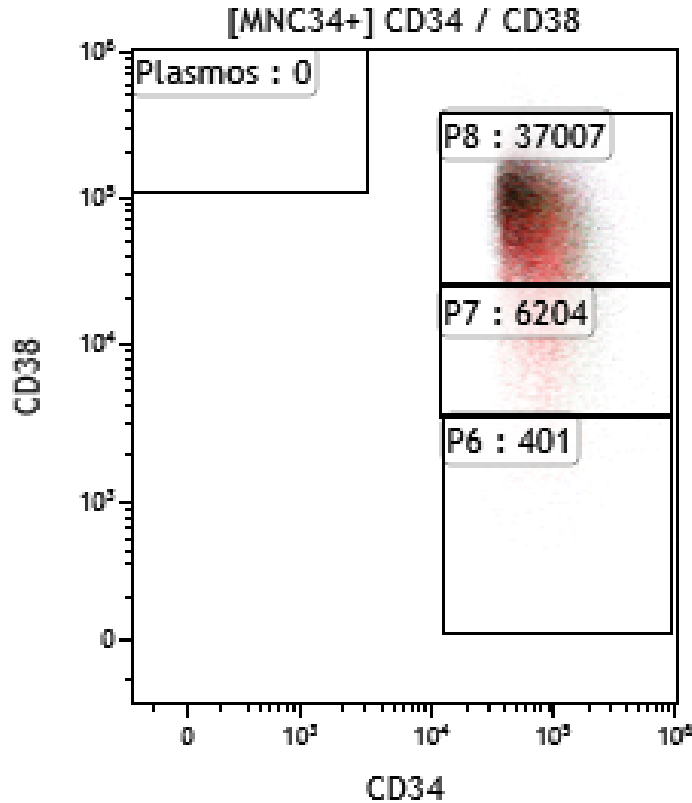


## T2 Navios (LSC markers)

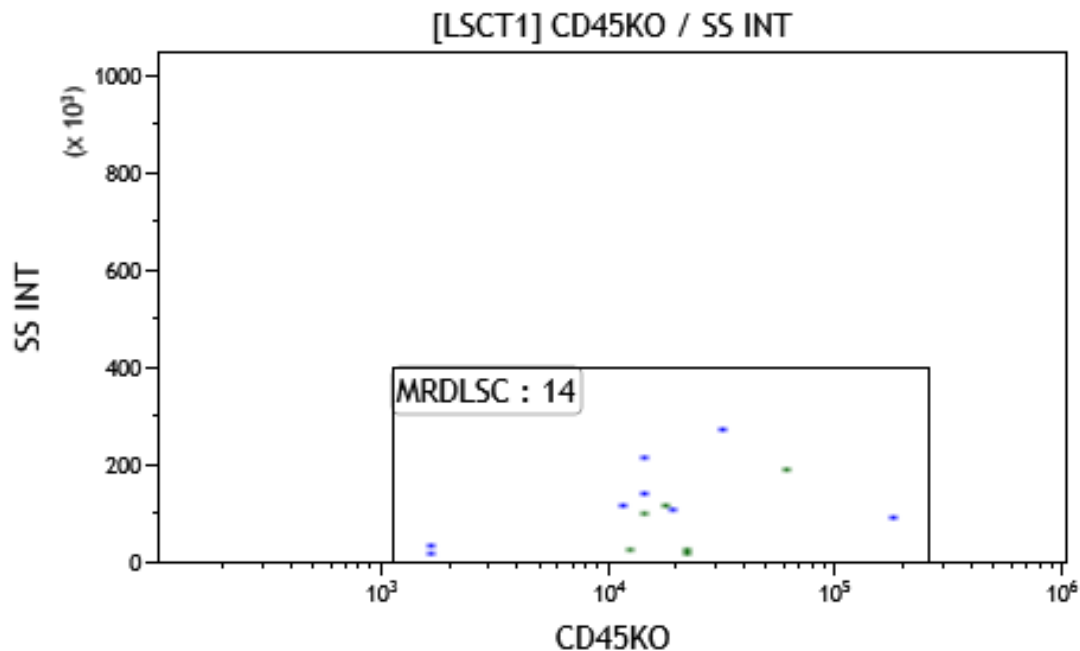


Gate CD34+CD38+hi

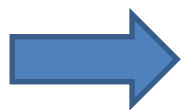




LAIP-P6(34+38-) Boleene MRD  
 LOD<0,0003%(3x10<sup>-6</sup>)  
 5X10<sup>6</sup> ev



MRD LSC LOD



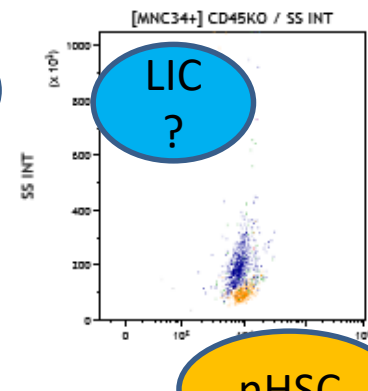
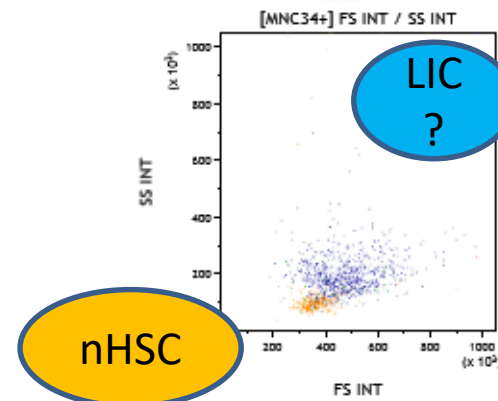
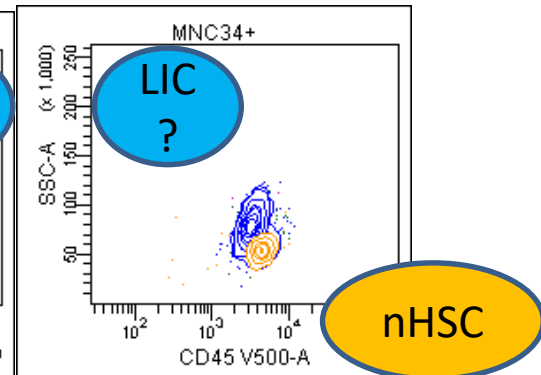
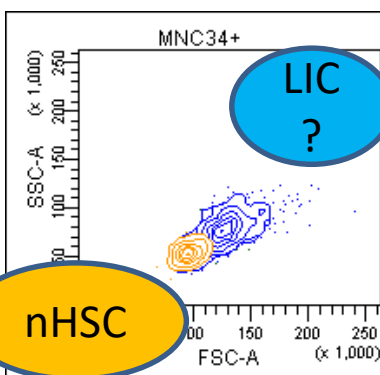
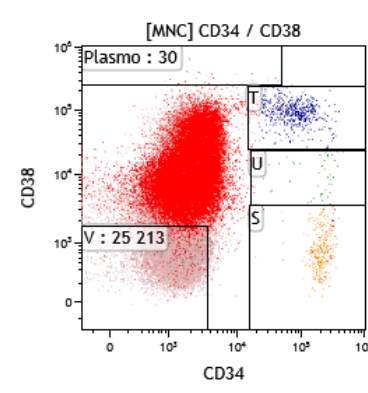
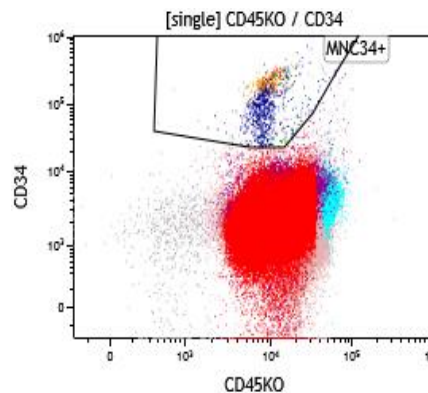
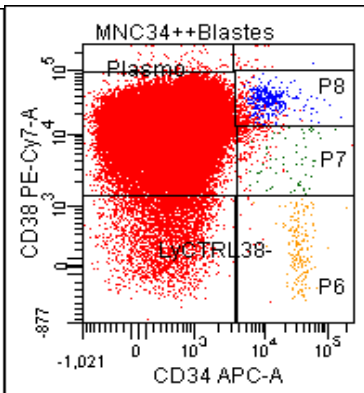
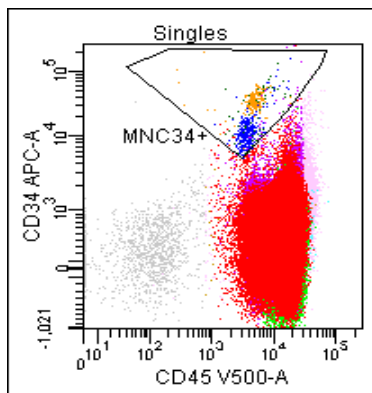
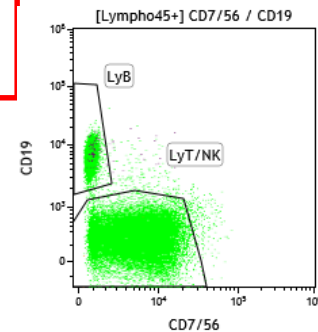
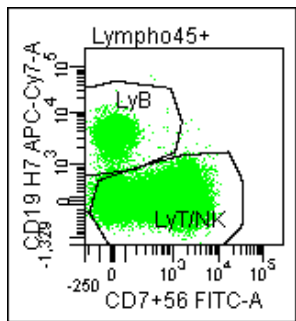
14 ev  
 0,0003%

Gate	Number	%Total
All	14	0,0002
MRDLSC	14	0,0002

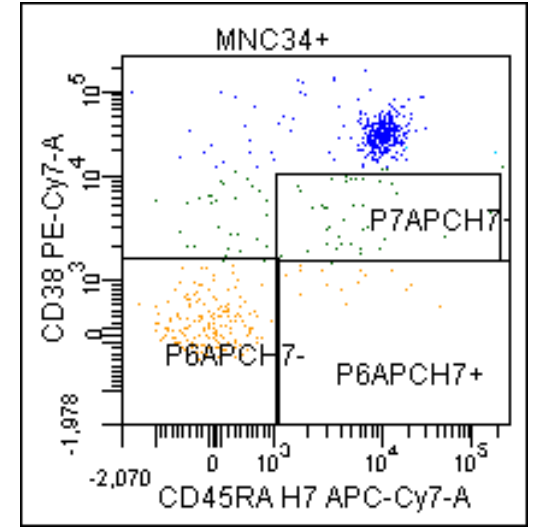
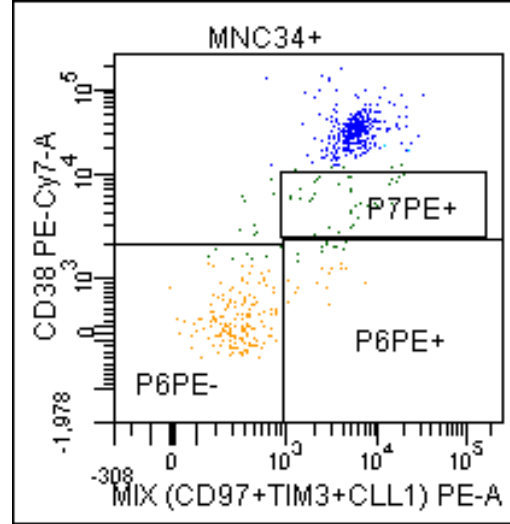
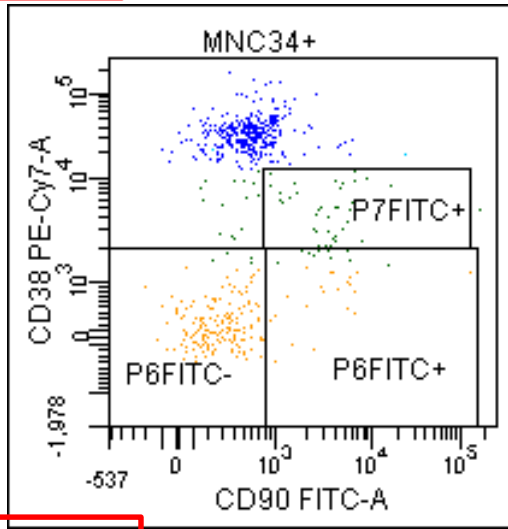
CANTO/  
DIVA

Harmonisation of detection of LSC using gating strategy  
adaptated to CANTO and NAVIOS softwares  
in an AML CD34- patient  
(BM sample shared between Lyon and Lille Labs)

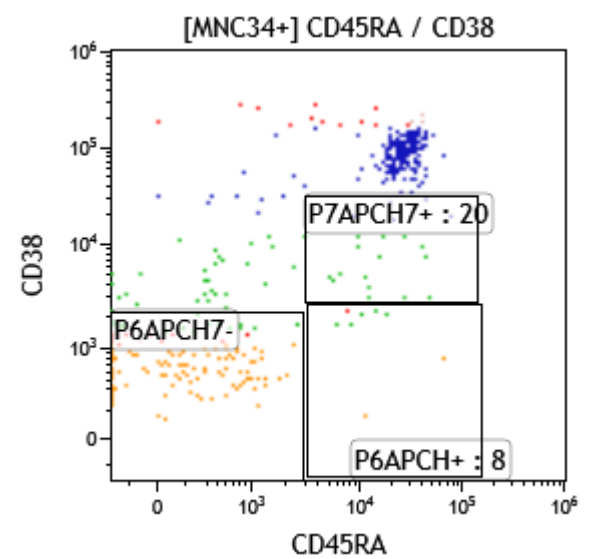
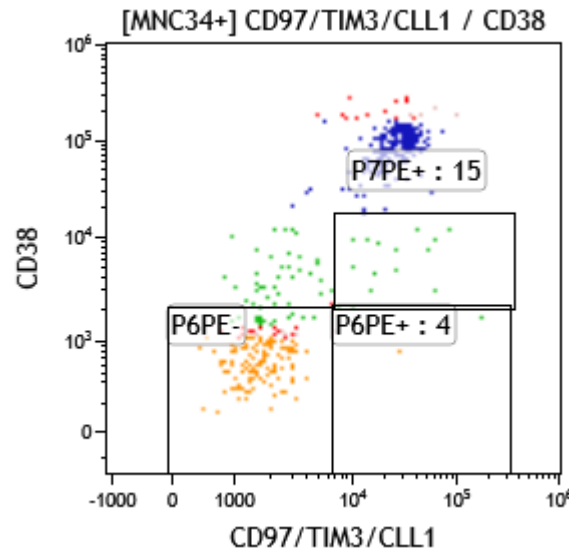
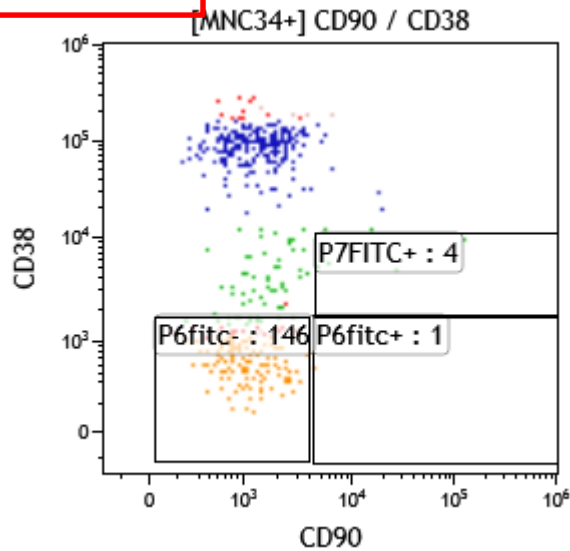
NAVIOS/  
KALUZA



CANTO/  
DIVA



NAVIOS/  
KALUZA



# AML MRD flow

## 1) Why?

- Flow Labs and French Clinical Trial organisation
- ELN 2017 Guidelines requirements- MRD mandatory (molecular and/or flow)
- Harmonisation of Flow Clinical MRD report

## 2) How?

Methodology multicentric approach:

- panel design: simple, reproducible, sensitivity, cut
- CANTO vs NAVIOS « miroir »
- gating strategy

## 3) *Ready to start*

- ***CQA: verify gating strategy***
- ***CQE: global check of panel implementation ( n/reg BM)***
- ***Patient follow up***

## Original Article

## A QA Program for MRD Testing Demonstrates That Systematic Education Can Reduce Discordance Among Experienced Interpreters

Michael Keeney,<sup>1\*</sup> Brent L. Wood,<sup>2,3</sup> Benjamin D. Hedley,<sup>1</sup>  
Joseph A. DiGiuseppe,<sup>4</sup> Maryalice Stetler-Stevenson,<sup>5</sup> Elisabeth Paietta,<sup>6</sup>  
Gerard Lozanski,<sup>7</sup> Adam C. Seegmiller,<sup>8</sup> Bruce W. Greig,<sup>8</sup> Aaron C. Shaver,<sup>8</sup>  
Lata Mukundan,<sup>9</sup> Howard R. Higley,<sup>9</sup> Caroline C. Sigman,<sup>9</sup> Gary Kelloff,<sup>10</sup>  
J. Milburn Jessup,<sup>11</sup> and Michael J. Borowitz<sup>12</sup>

<sup>1</sup>Pathology and Laboratory Medicine, London Health Sciences Centre, London, Ontario, Canada

<sup>2</sup>Seattle Cancer Care Alliance, Seattle, Washington

<sup>3</sup>University of Washington, Seattle, Washington

<sup>4</sup>Department of Pathology, Hartford Hospital, Hartford, Connecticut

<sup>5</sup>National Cancer Institute, National Institutes of Health, Bethesda, Maryland

<sup>6</sup>Oncology, Montefiore Medical Center, Bronx, New York

<sup>7</sup>Department of Pathology, Ohio State University, Columbus, Ohio

<sup>8</sup>Department of Pathology, Microbiology, and Immunology, Vanderbilt University Medical Center, Nashville, Tennessee

Table 2  
Dry Challenges 1-3

	15	15	15 <sup>a</sup>	15	15	15 <sup>a</sup>
# Attempted	15	15	15	15	15	15
% Attempted	100%	100%	100%	100%	100%	100%
# Positive	7	9	7	11	12	9
# Negative	8	6	8	4	3	6
False positive	0	1	2	2	4	0
False negative	2	1	4	1	2	2
Overall concordance (%)	87%	87%	60%	80%	60%	87%
Outside 1/2 log	2	4	6	4	7	3
1/2 log Concordance (%)	87%	73%	60%	73%	53%	80%

<sup>a</sup>The 3rd round of results for two laboratories were submitted after the deadline results were due.

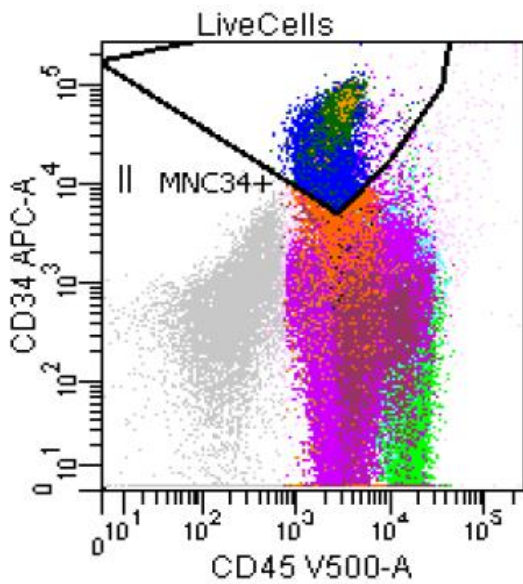
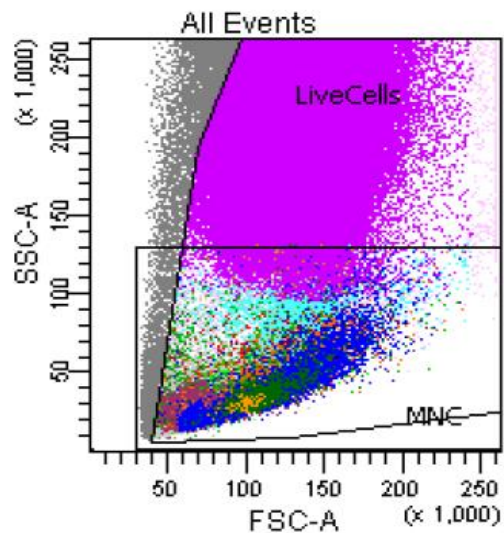
Table 3  
Dry Challenges 4-5

	Statistics for 7 participating laboratories						
# Attempted	10	10	10	10	10	10	10
% Attempted	100%	100%	100%	100%	100%	100%	100%
# Positive	5	5	7	6	5	6	5
# Negative	5	5	3	4	5	4	5
False POS	0	0	3	1	0	1	0
False NEC	0	0	1	0	0	0	0
Overall Concordance (%)	100%	100%	60%	90%	100%	90%	100%
Outside 1/2 log	0	0	4	1	0	1	0
1/2 log Concordance (%)	100%	100%	60%	90%	100%	90%	100%

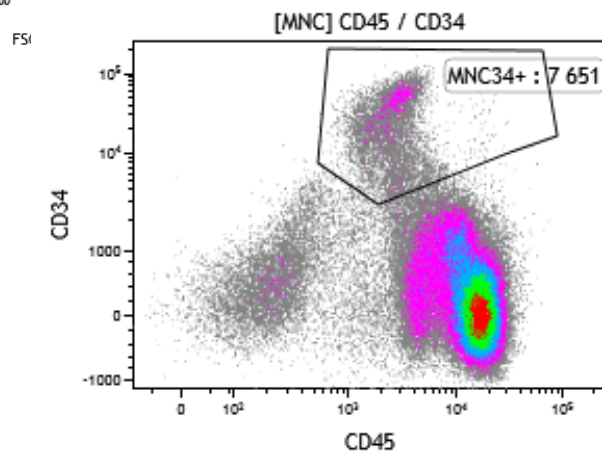
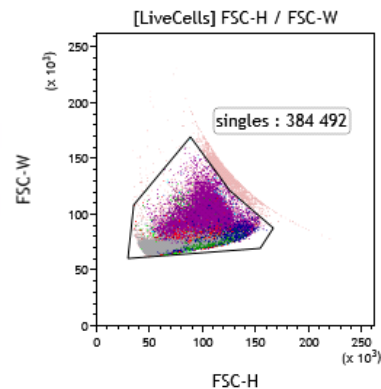
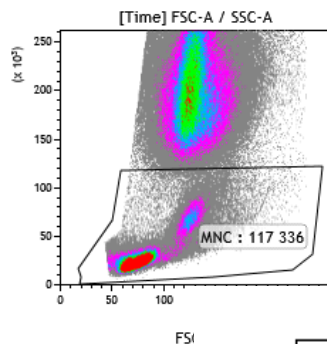
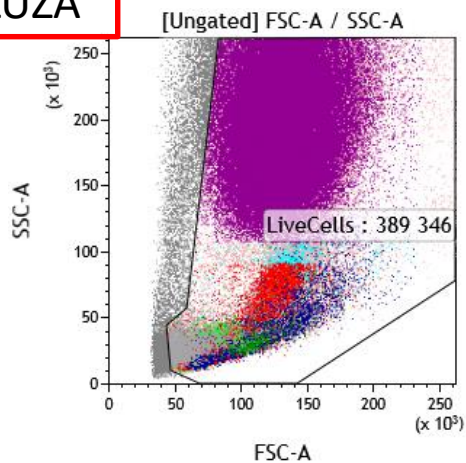
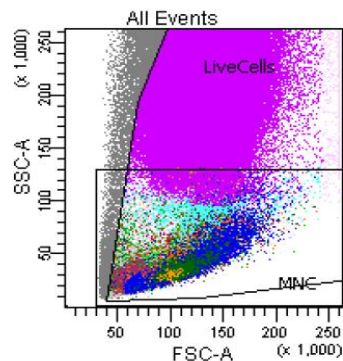
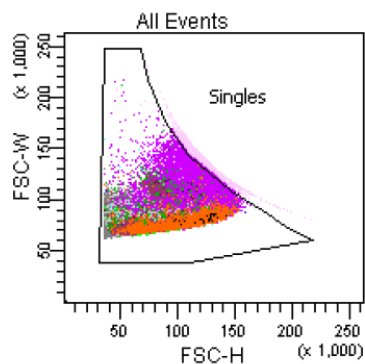
Concordance  
before training

Concordance  
after training

DIVA



KALUZA

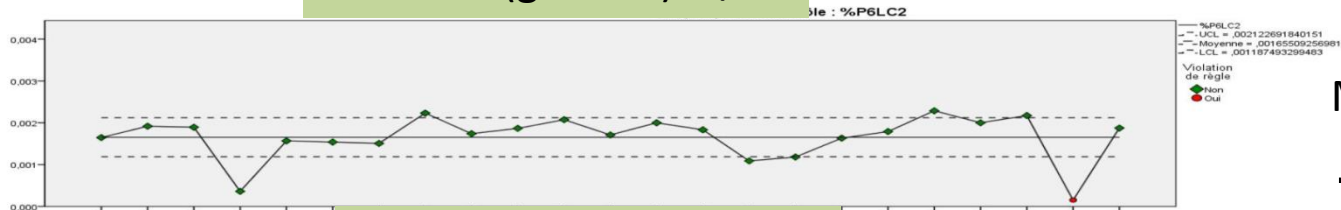


CQA 23 centers (2 sets of data not integrated)  
DIVA vs KALUZA  
LMD nBM



# CQA LMD Files shared among 23 centers

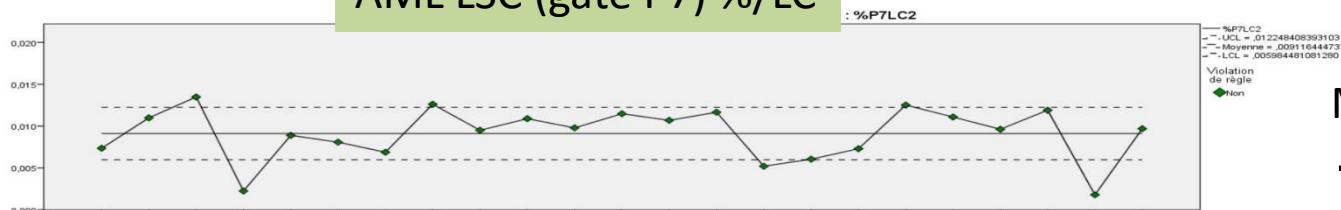
AML LSC (gate P6) %/LC



Mean :  $1,6 \cdot 10^{-3}$

--- +/- 1 SD

AML LSC (gate P7) %/LC

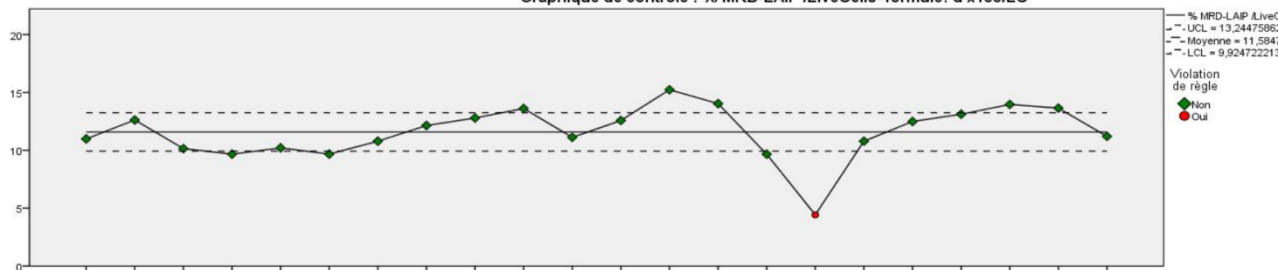


Mean :  $9,1 \cdot 10^{-3}$

--- +/- 1 SD

AML LAIP %/LC

Graphique de contrôle : % MRD-LAIP /LiveCells- formule: d x100/LC

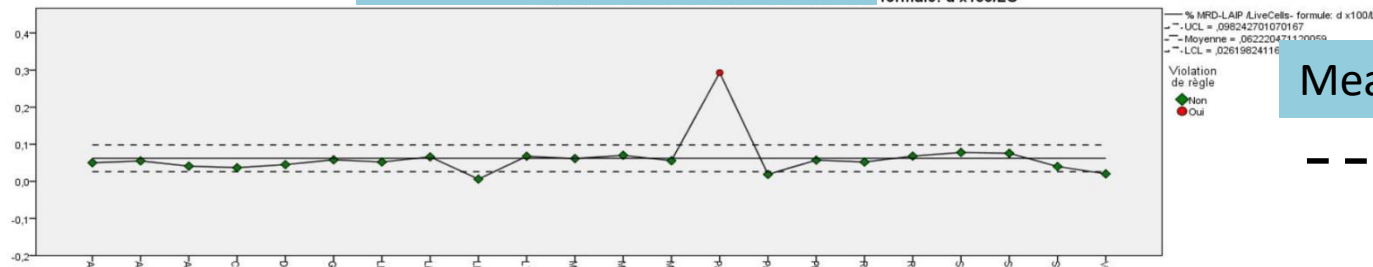


Mean 11,9%

--- +/- 1 SD

LOD AML LAIP in NBM

formule: d x100/LC

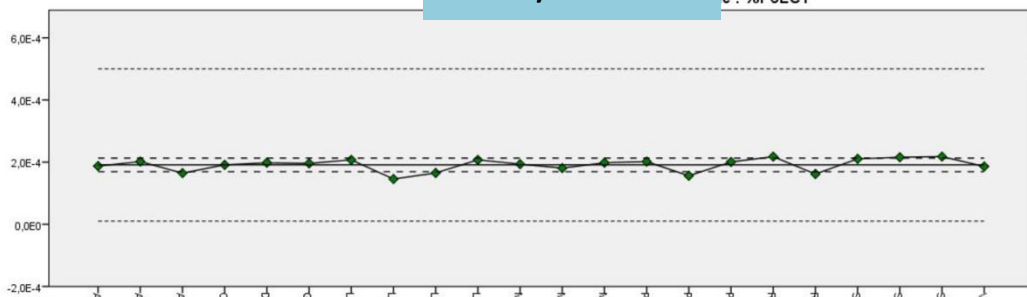


Mean :  $6,2 \cdot 10^{-4}$

--- +/- 1 SD

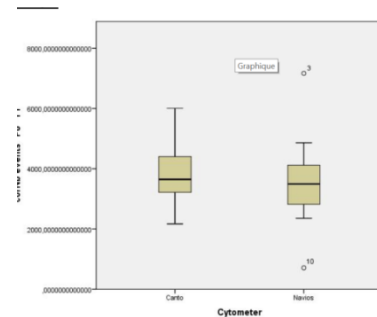
# CQE nBM sample shared among 22 centers

% P6/LC T1



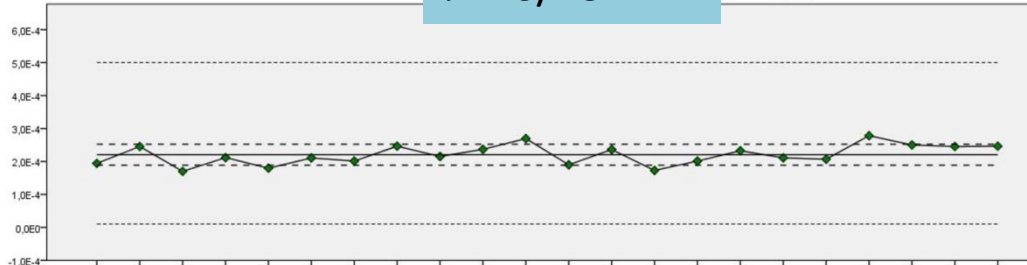
%P6LC1  
 - UCL = .000213116726474  
 - Spéc. U = 0.00005  
 - Moyenne = .000191158758942  
 - Spéc. L = 0.00001  
 - LCL = .000169200791411

Mean :  $1,92 \cdot 10^{-4}$   
 - - - +/- 1 SD



Canto Navios

% P6/LC T2

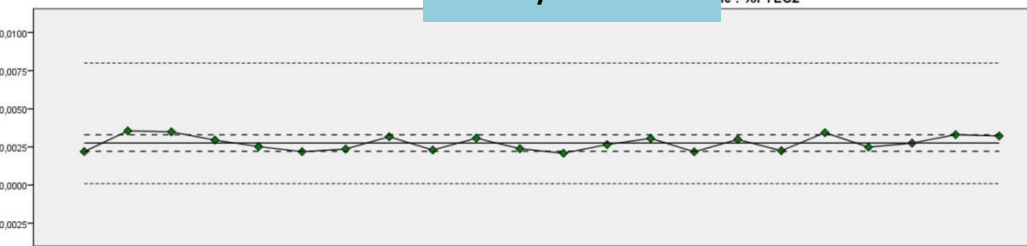


%P6LC2  
 - UCL = .000252313438686  
 - Spéc. U = 0.00005  
 - Moyenne = .00022058264  
 - Spéc. L = 0.00001  
 - LCL = .000198851854227

Mean :  $2,2 \cdot 10^{-4}$   
 - - - +/- 1 SD

- staining validation  
 - stability at 24h  
 - homogeneity of the results

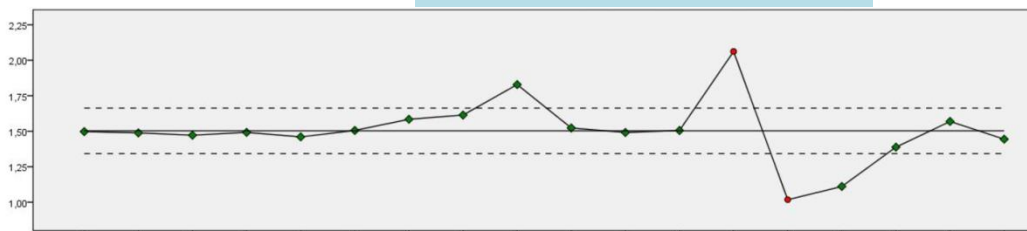
% P7/LC T2



%P7LC2  
 - UCL = .003294683034556  
 - Spéc. U = 0.008  
 - Moyenne = .002755408169147  
 - Spéc. L = 0.0001  
 - LCL = .002216133303738

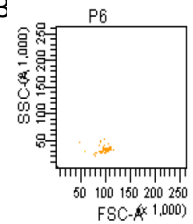
Mean :  $2,75 \cdot 10^{-3}$   
 - - - +/- 1 SD

Ratio FSC of P6/HTG

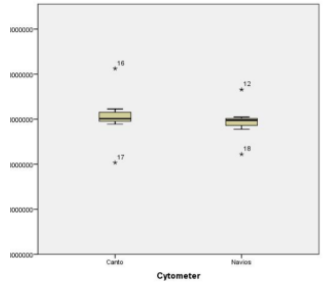
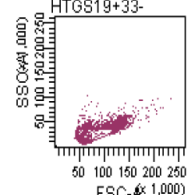


Ratio FSC P6/HTG19+33-  
 - UCL = 1.65317389453730  
 - Moyenne = 1.502861930096268  
 - LCL = 1.342586551736805

Mean : 1,50  
 Up Limit: 1,66  
 Low Limit: 1,34



Canto Navios



# Ready to start

Minimal information should be given in **Clinical FlowReport**:

- Quality of BM (dilution)
- Strategy of identification of MRD: LAIP/DFN/LSC
- Description of the LAIP used
- LOD based on nBM
- Thresholds  $< 10^{-3}$  for LAIP or  $< 10^{-4}$  for LSC
- Interpretation of the result:
  - MRD+
  - MRD- with LOD value
  - MRD detectable but nonquantifiable

# Take home message

- This methodological validation protocol is a mandatory step to consider the use of MRD flow in AML clinical trials;
- 3 MRD approach are complementary: LAIP/DFN/LSC
- Prospectively Lyophilised Ab approach
- Choosing a multicentric approach could be challenging, but our first results are promising and showed the feasibility of this concept when:
  - (i) a straight harmonisation of the instruments sensitivity and samples preparation are established, and
  - (ii) training and systematic education among the analytical operators are regularly performed.

# Thanks!

Flow Cytometric Labs

AFC (French Association of Cytometry)

**Adriana Plesa & Christophe Roumier**

**Florent Dumezy, Francois Vergez, Valerie Bardet, Orianne Wagner Ballon, Stephanie Mathis, Anna Raimbault, Victoria Ragueneau, Agnes Charpentier, Veronique Harrivel, Elsa Bera, Veronique Salaun, Edouard Cornet, Julien Guy, Veronique Saada, Isabelle Arnoux, Magali Le Garf Tavernier, Remi Letestu, Jean Feuillard, Estelle Guerin, Christine Arnoulet, Anne Catherine Lhoumeau, Lydia Campos, Carmen Aanei, Tiphane Picot, Marie Christine Jacob, Tatiana Raskovalova, Camille Lours, Delphine Manzoni, Richard Veyrat Masson, Franck Geneviève, Nicolas Chapuis, Mikael Roussel, Hélène Lapillonne**

Clinical coordinators: **Hervé Dombret, Cristian Recher**

Biological coordinator: **Claude Preudhomme**

ALFA coordinator: **Karine Celli-Lebras**



Collaborative InterGroup for Acute Leukemia

