

Use of Molecular Assays to Assess Cellular Therapies

**D. Stroncek, P. Jin, M. Sabatino, J. Ren, L.
Castiello, E. Wang, and F.M. Marincola.**

**Cellular Therapies Section
Clinical Center, NIH**

Methods for the Characterization of Cellular Therapies

Tests Required by the FDA

- Safety
- Identity
- Purity
- Potency

Testing useful in manufacturing

- Stability
- Consistency
- Comparability

A Centralized Cell Processing Lab

Assays

Cell Counts

Automated

- Cell dyne 3700
- Saphire
- Multi-sizer

Manual

Viability

- Trypan blue (manual)
- 7AAD (flow cytometry)

Flow Cytometry

Surface markers

Intracellular markers

Sterility

Microbial cultures

Endotoxin

Gram stains

Mycoplasma

Gene and MicroRNA Expression Profiling

Advantages

- Simultaneous expression of thousands of gene and hundred's of micro RNA at one time
- Allows for the global analysis of cells
- Few cells are required 1×10^4 to 1×10^6 cells
- Can be used with cryopreserved cells

Disadvantages

- Transcription doesn't always reflect protein expression
- Platforms are not fully standardized
- Lengthy assay

Need for Comparability Analysis

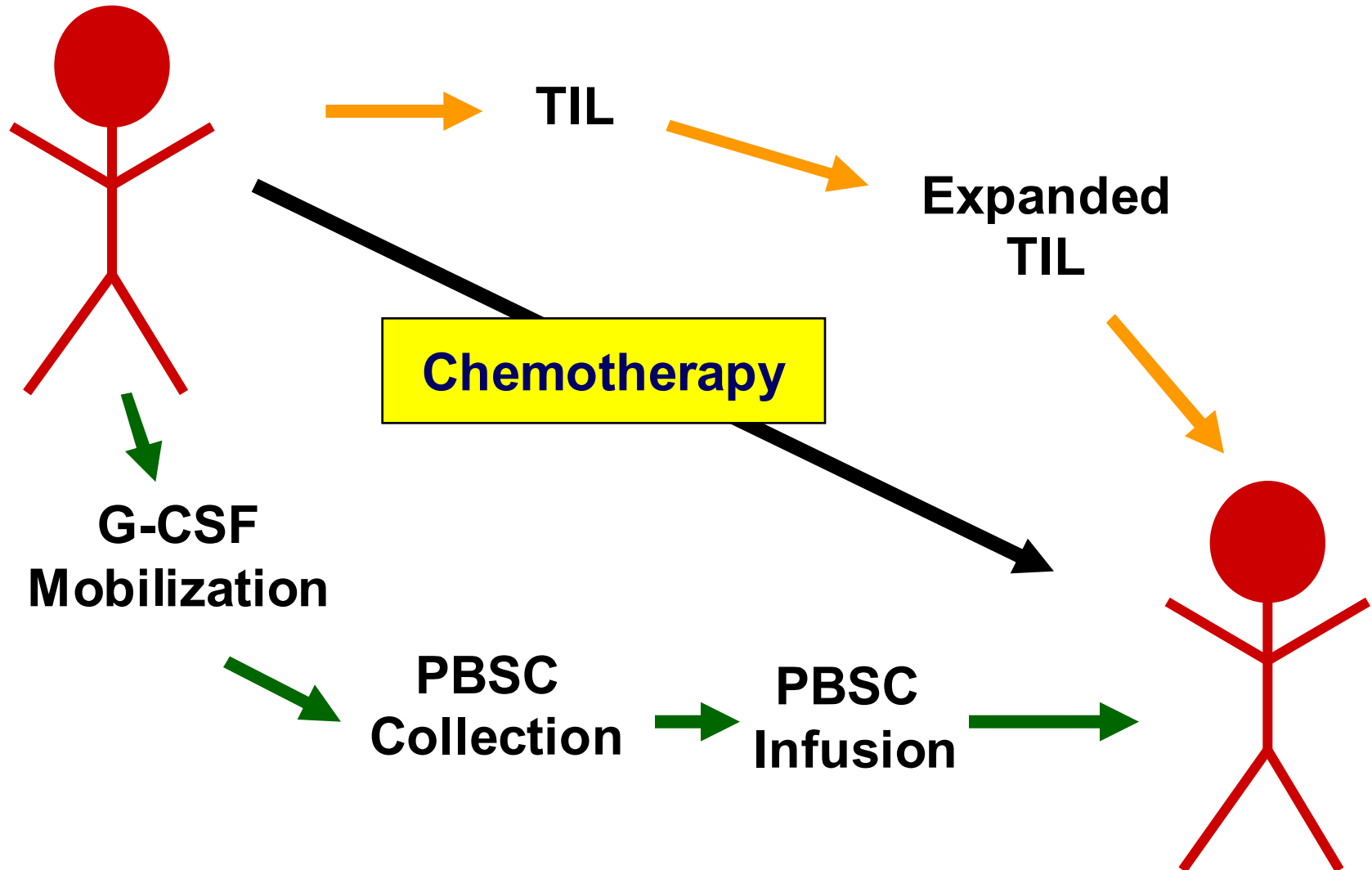
Changes in reagents and supplies

- **Media**
- **Cytokines**
- **Growth factors**
- **Bags**

Changes in starting material

- **Collection procedures**
- **Mobilization agents**
- **Storage duration or conditions**

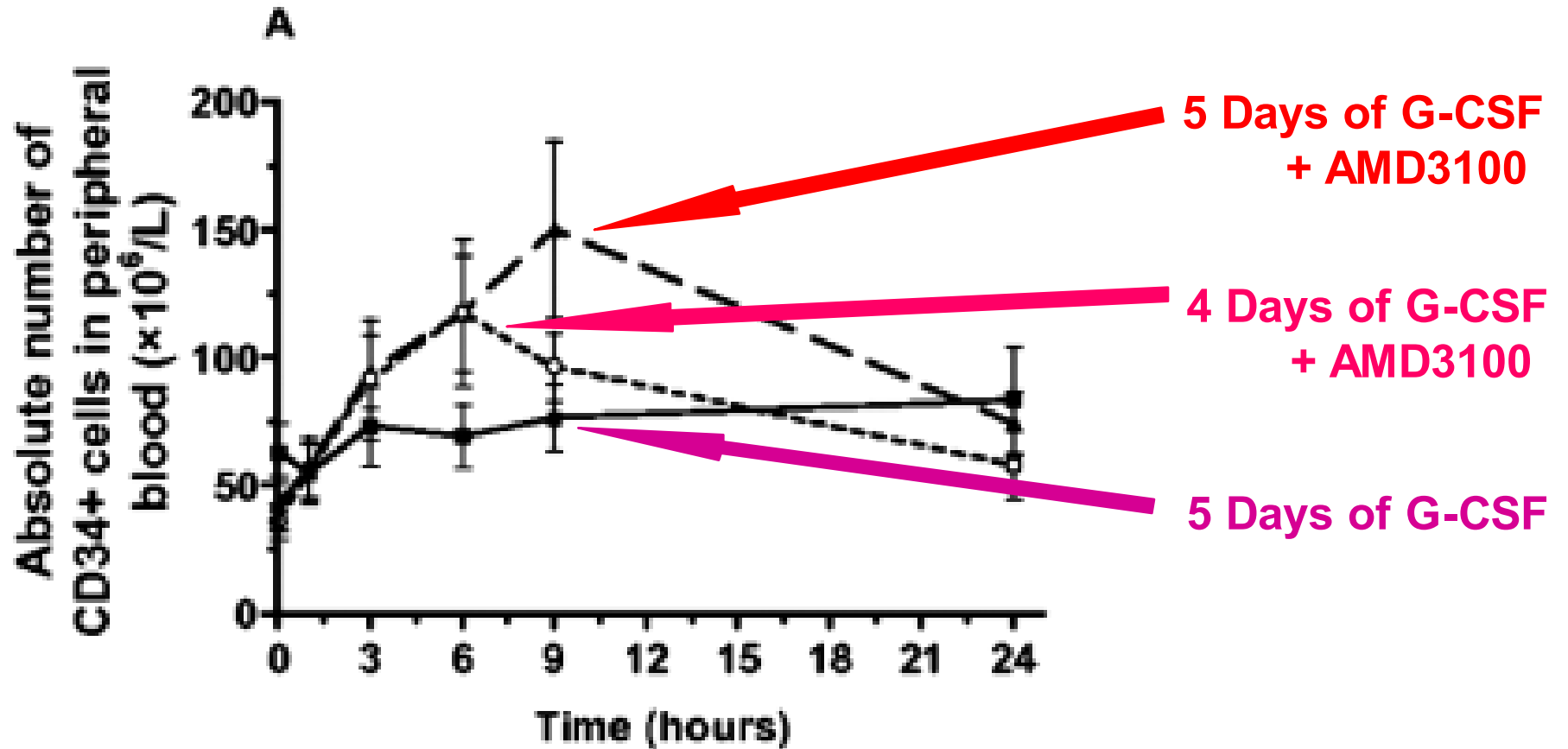
Mobilized Autologous PBSCs in Immune Therapy of Cancer

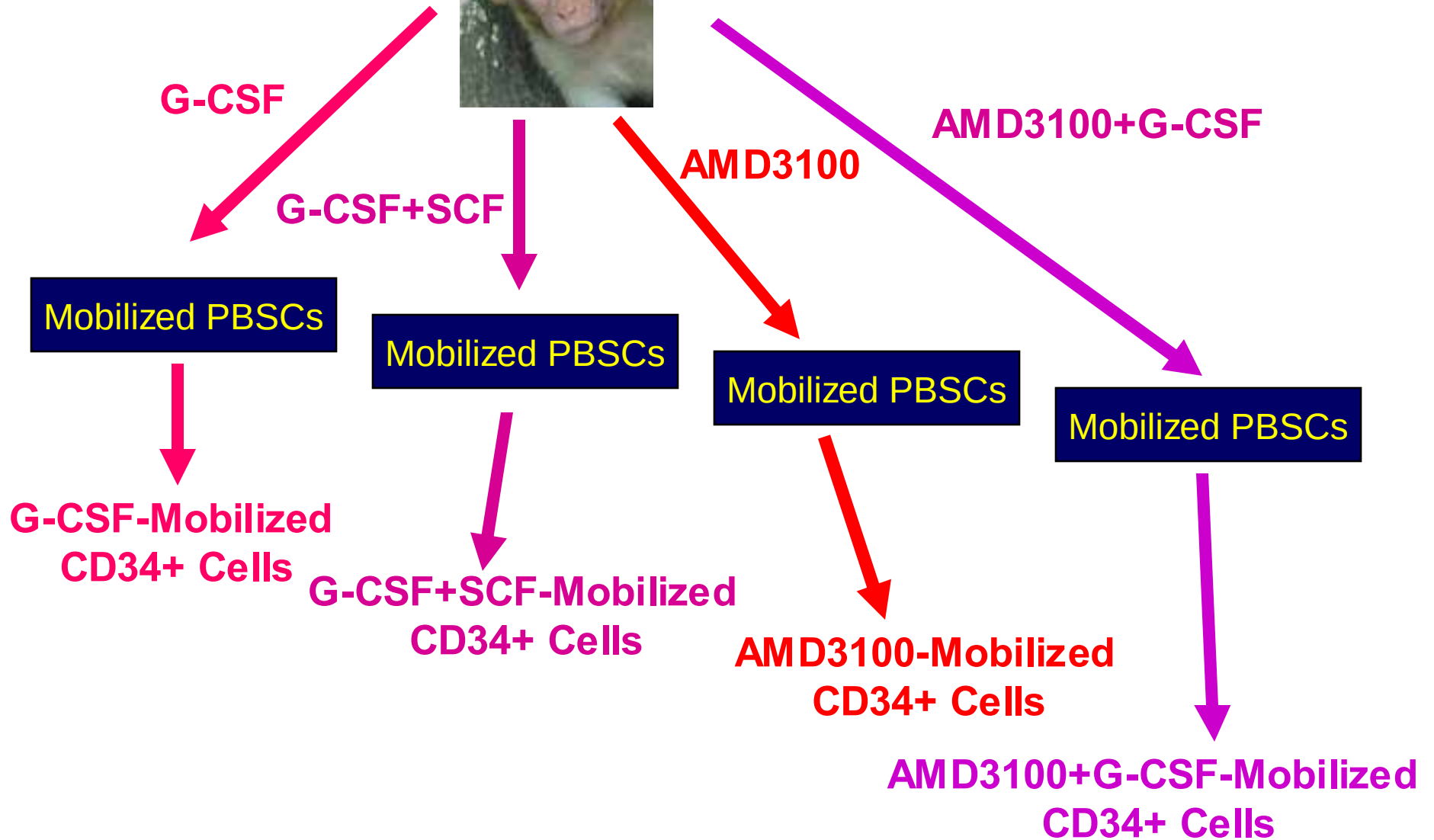


Plerixafor (AMD3100)

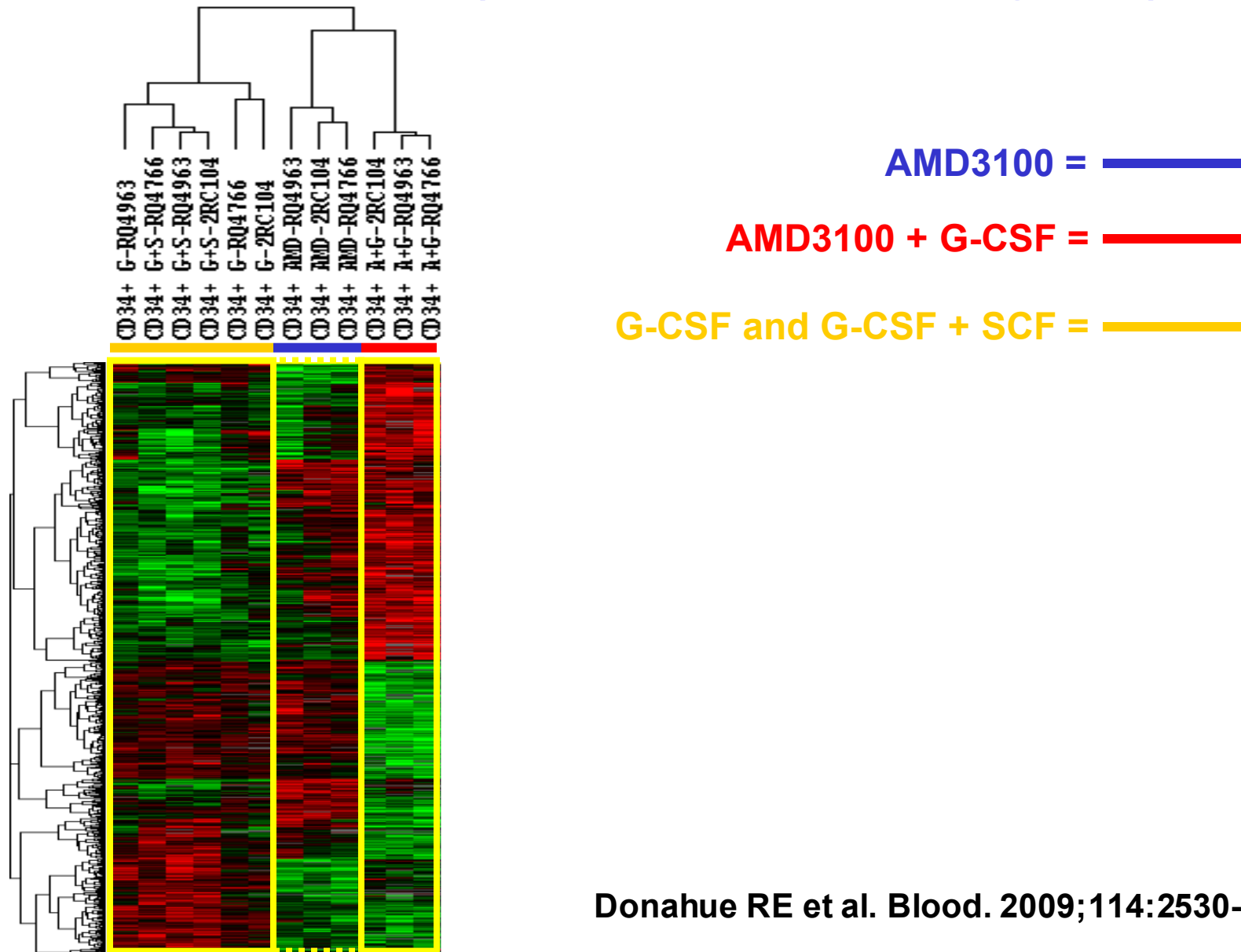
- **Disrupts the binding of CXCR4 and CXCL12**
- **Mobilizes leukocytes and stem cells in 4 to 6 hours**
- **FDA approved for use with G-CSF for stem cell mobilization for autologous transplantation**

G-CSF Plus AMD3100 Mobilization

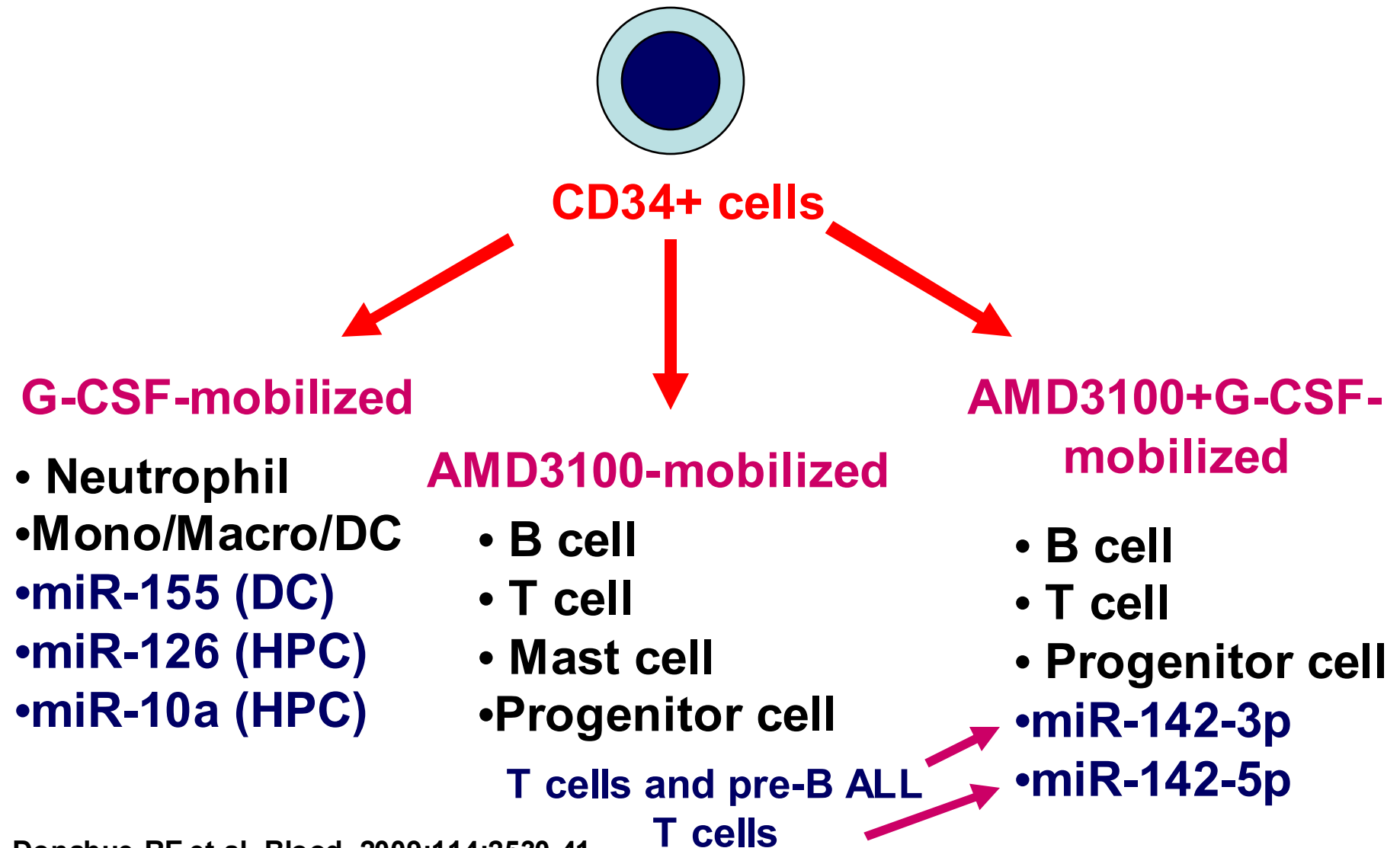




Gene Expression Profiling of Differentially Expressed CD34+ Cell Genes (F-test, $p \leq 0.005$, 1,097 genes)

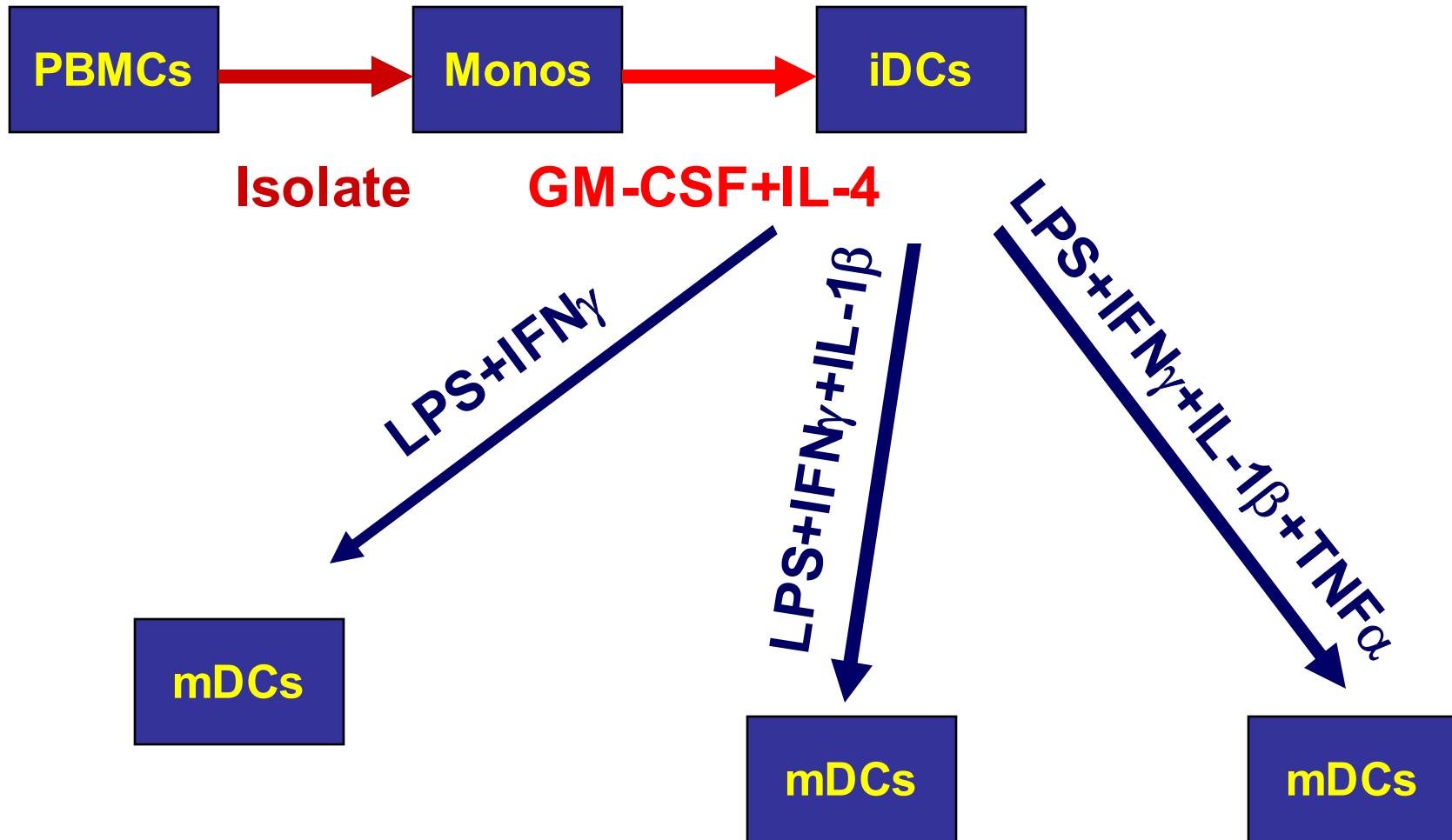


Comparison of Different Types of CD34+ Cells: Summary of Genes Expressed by each type of CD34+ cells



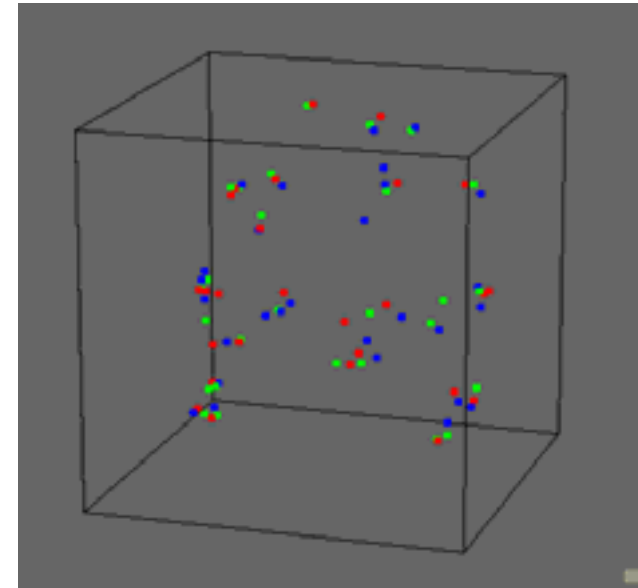
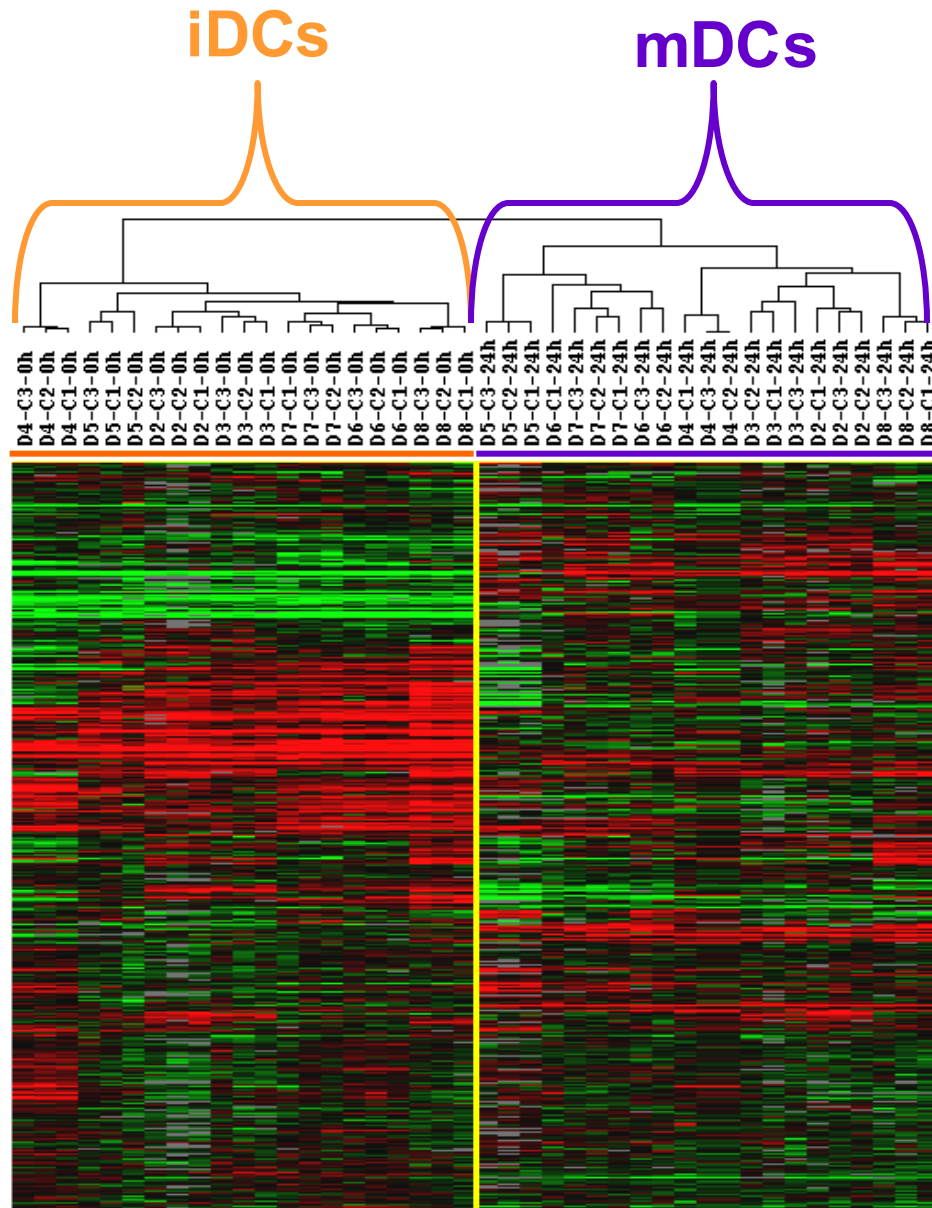
Dendritic Cells

Comparability Testing and DCs: Comparison of DC Maturation Cocktails



Assessed iDCs and mDCs by gene expression profiling

*Multidimensional scaling,
using centered correlation*



The ANOVA model: $\log \text{ expression} \sim \text{Donor} + \text{Treatment} + \text{Time}$

Table 1: Number of significant genes for testing fixed effects.

Effect	P-value < 0.001	FDR < 0.1
Donor	9590	16103
Treatment	13	0
Time course	9576	15822

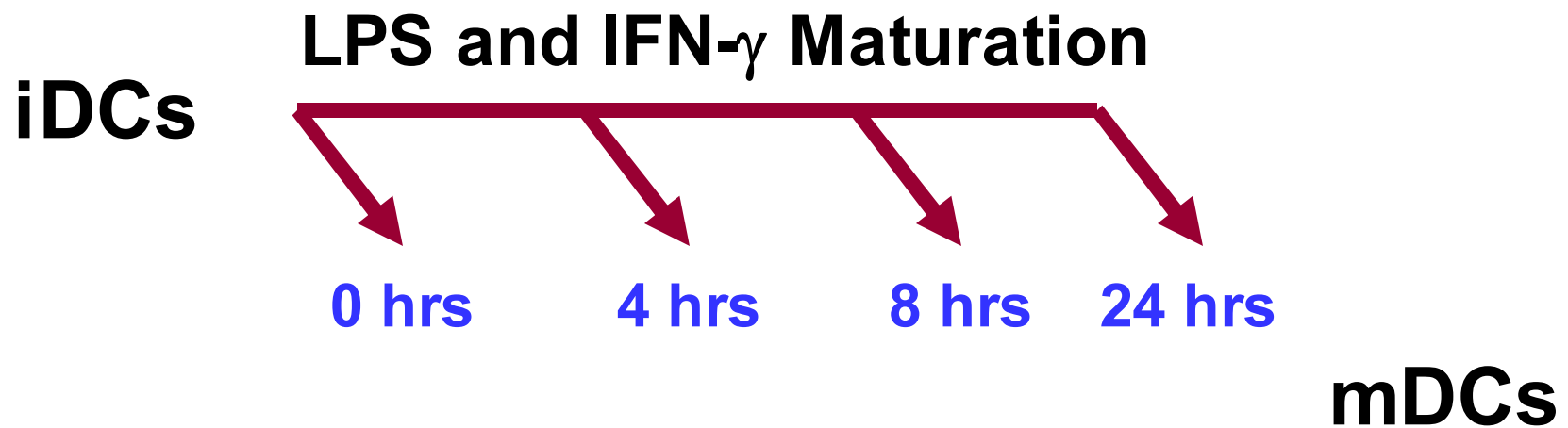
Table 2: Number of significant genes for testing the model.

	P-value < 0.001	FDR < 0.1
Model	12505	18559

C1 = LPS+IFN- γ ●; C2 = LPS+IFN- γ +IL-1 β ●; C3 = LPS+IFN- γ +IL-1 β +TNF- α ●

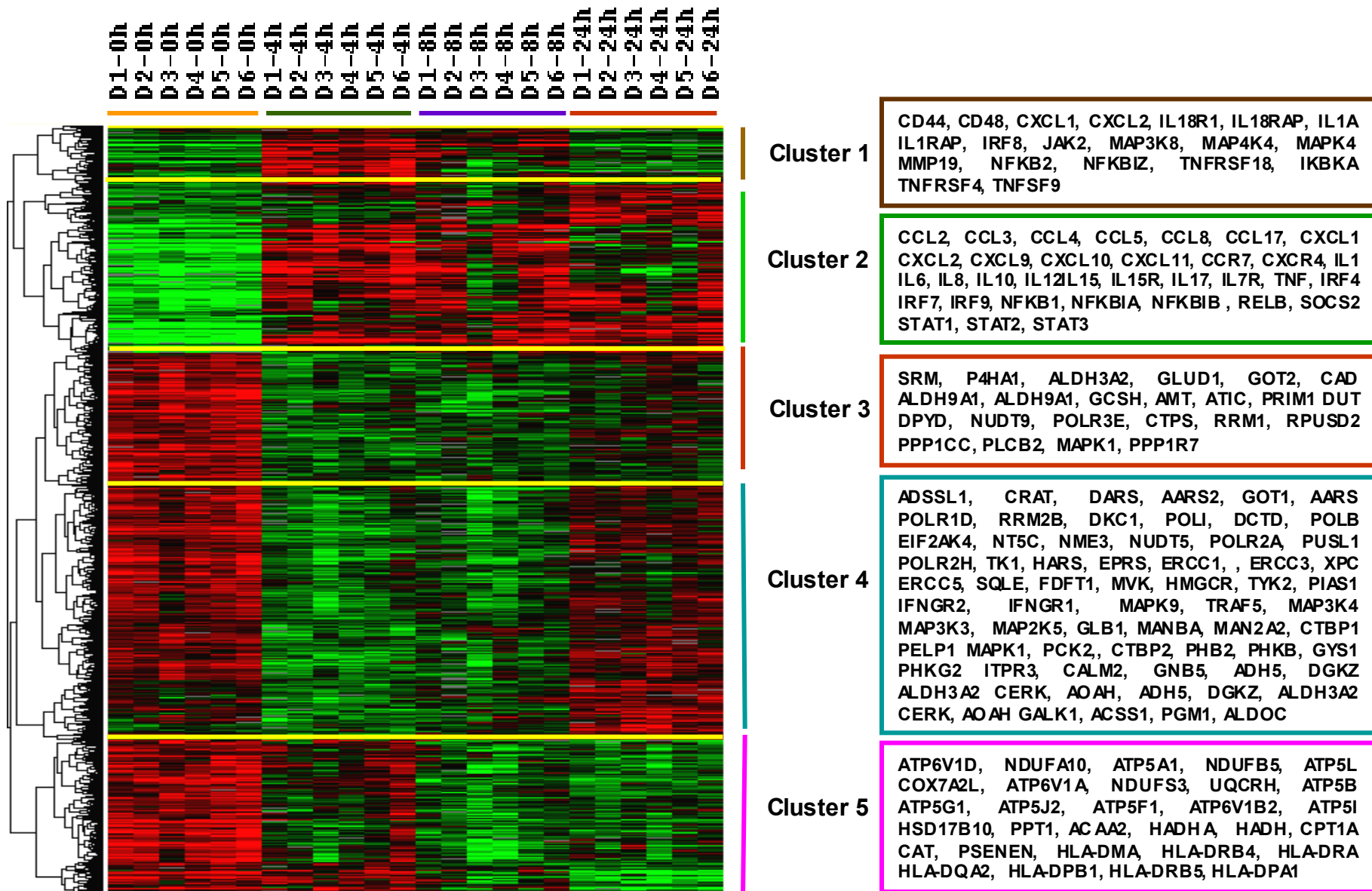
Han TH, Jin P, Ren J, Slezak S, Marincola FM, Stroncek DF. J Immunother. 2009;32:399-407

mDC Biomarker Discovery: Kinetics of the molecular changes associated with DC maturation



Assessed Global Gene and MicroRNA Expression

Kinetics of LPS and IFN- γ Mediated DC Maturation: Transcriptome analysis of DCs from 6 healthy subjects



Criteria for Biomarker Selection for Gene Expression Analysis

Important in DC function

- Potency, comparability, consistency

Increased at least 10-fold in mDCs

- All tests

Increased variably during maturation

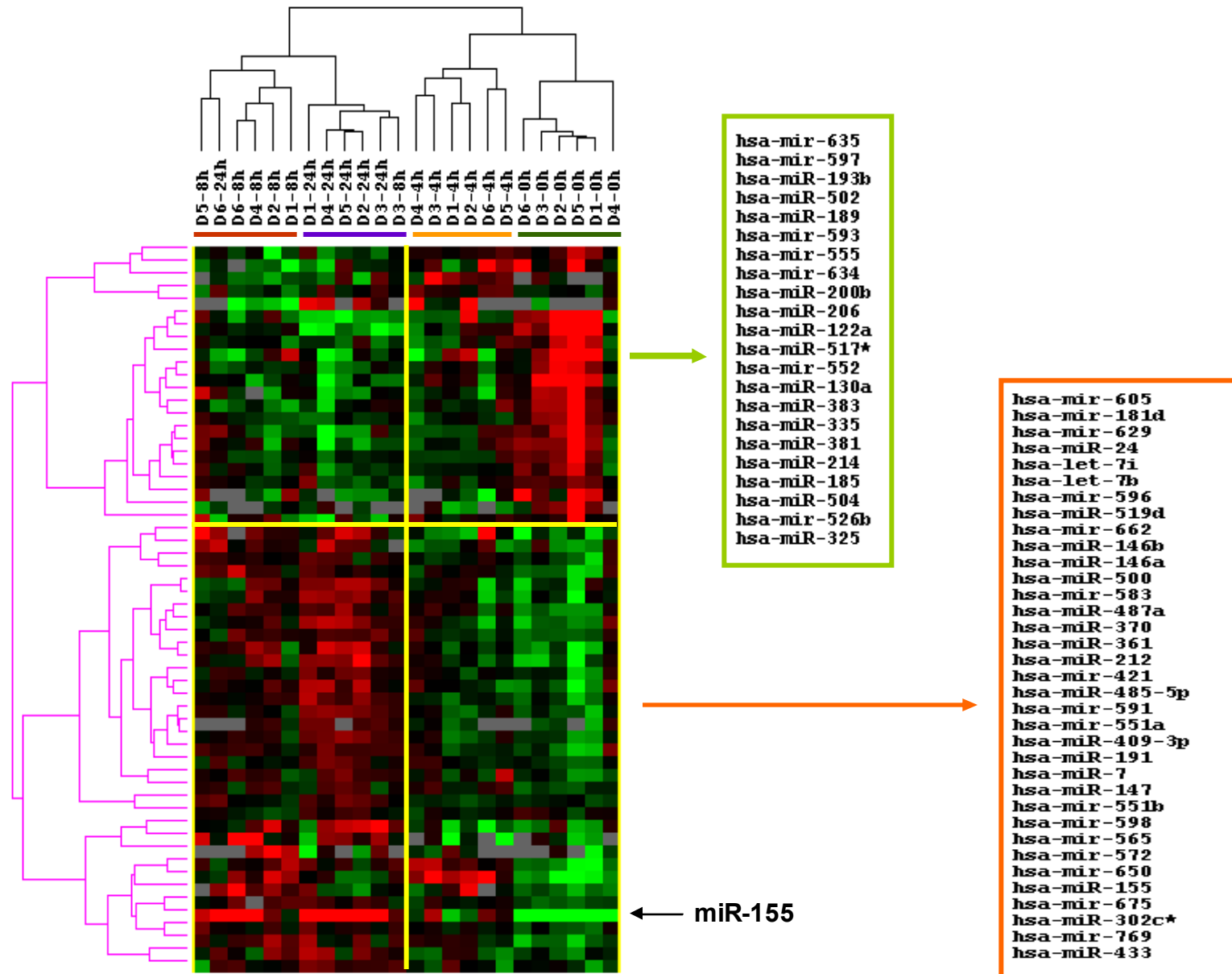
- Consistency, comparability

Potential mDC Biomarkers:

Genes up-regulated during DC maturation

Fold increase in gene expression for each maturation time*							
Gene	4 hours	8 hours	24 hours	Gene	4 hours	8 hours	24 hours
Up-regulated to a similar degree throughout maturation				Up-regulated most early in DC maturation			
CCL5	108	148	94.1	IL6	13.1	11.3	5.15
CXCL10	28.5	31.8	21.2	IL8	78.0	70.3	12.7
CCR7	10.5	11.5	18.2	IL7R	29.1	29.9	10.5
IL15	7.12	5.54	8.13	CCL4	92.3	53.4	6.91
IFI27	6.99	7.62	10.2	TNFAIP6	30.0	18.7	10.7
IFI44L	14.8	16.7	20.5	IFIT3	36.2	22.5	10.5
IFIH1	16.9	9.42	11.3	OASL	68.1	44.1	30.7
IFIT1	29.8	27.0	21.7	GBP1	66.2	35.3	30.2
MX1	18.4	15.6	14.1	HES4	229	115	37.4
ISG15	50.6	58.3	41.8	Up-regulated most late in DC maturation			
ISG20	94.1	87.9	62.9	CCL8	11.3	31.8	31.2
IRF7	9.77	9.38	12.0	EBI3	17.6	21.8	34.6
GBP4	36.3	21.2	20.2	IFITM1	13.2	22.5	48.6
DUSP5	21.7	15.8	22.6	MT1B	10.2	10.5	20.6
NFKBIA	11.7	13.3	10.4	MT1E	NS	1.78	46.1
ATF3	10.2	5.38	11.4	MT1G	NS	2.77	42.3
TNFSF10	19.4	14.8	13.8	MT1H	22.7	20.5	62.6
TNFRSF9	8.13	6.39	10.2	GADD45A	NS	11.6	50.7

MicroRNA changes during DC maturation with LPS and IFN- γ



DC Potency Testing

- **Assay validation**
- **All clinical DCs produced at the Clinical Center**

Luciano Castiello, Marianna Sabatino, and Ping Jin

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