

Non-clinical Immunogenicity Assessment of Generic Peptide  
Products: Development, Validation, and Sampling

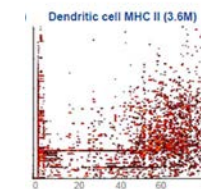
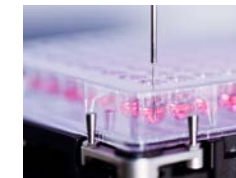
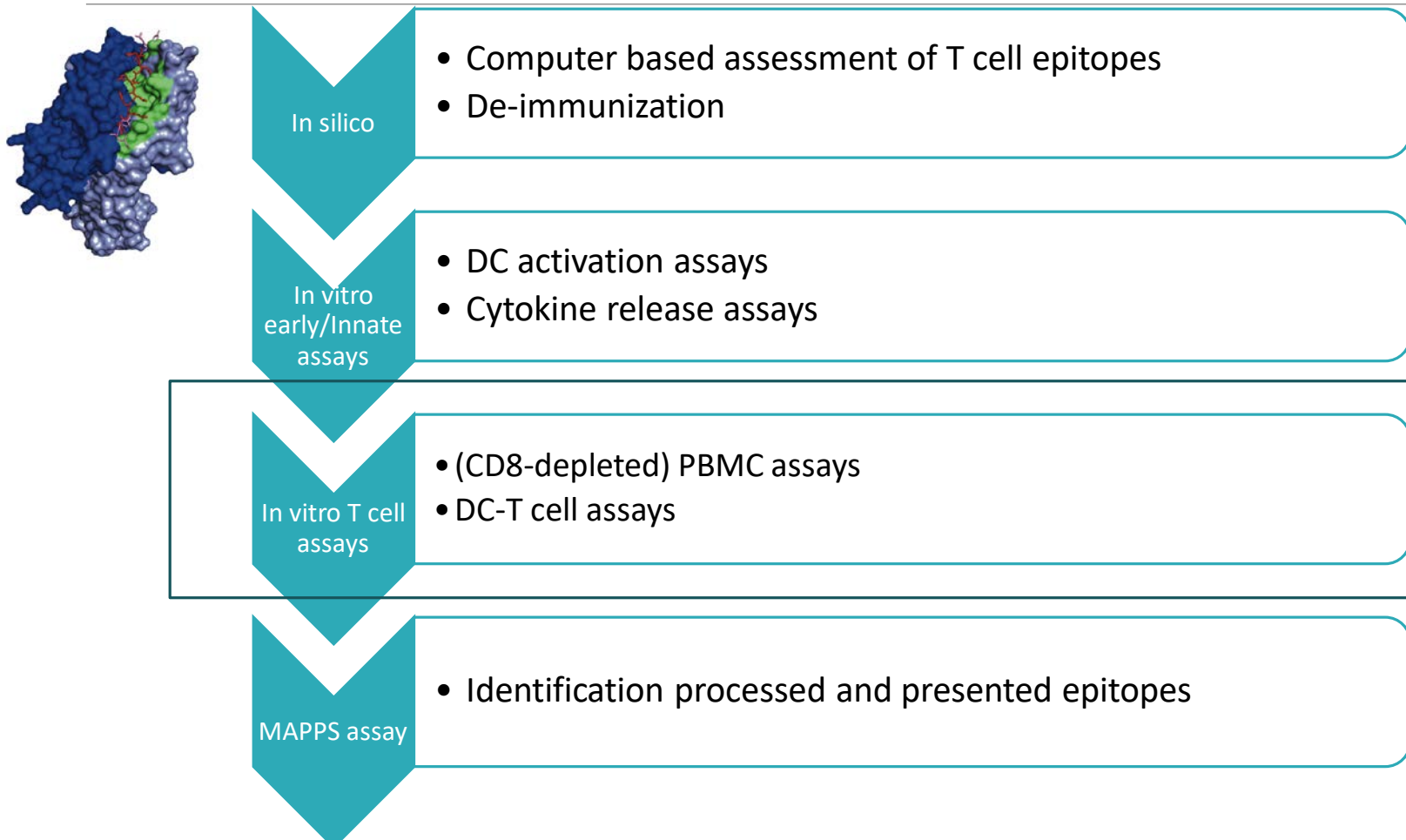
# T cell immunogenicity assays: time for standardization and harmonization

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SOFIE PATTIJN, CTO – FOUNDER IMMUNXPERTS, A NEXELIS GROUP COMPANY

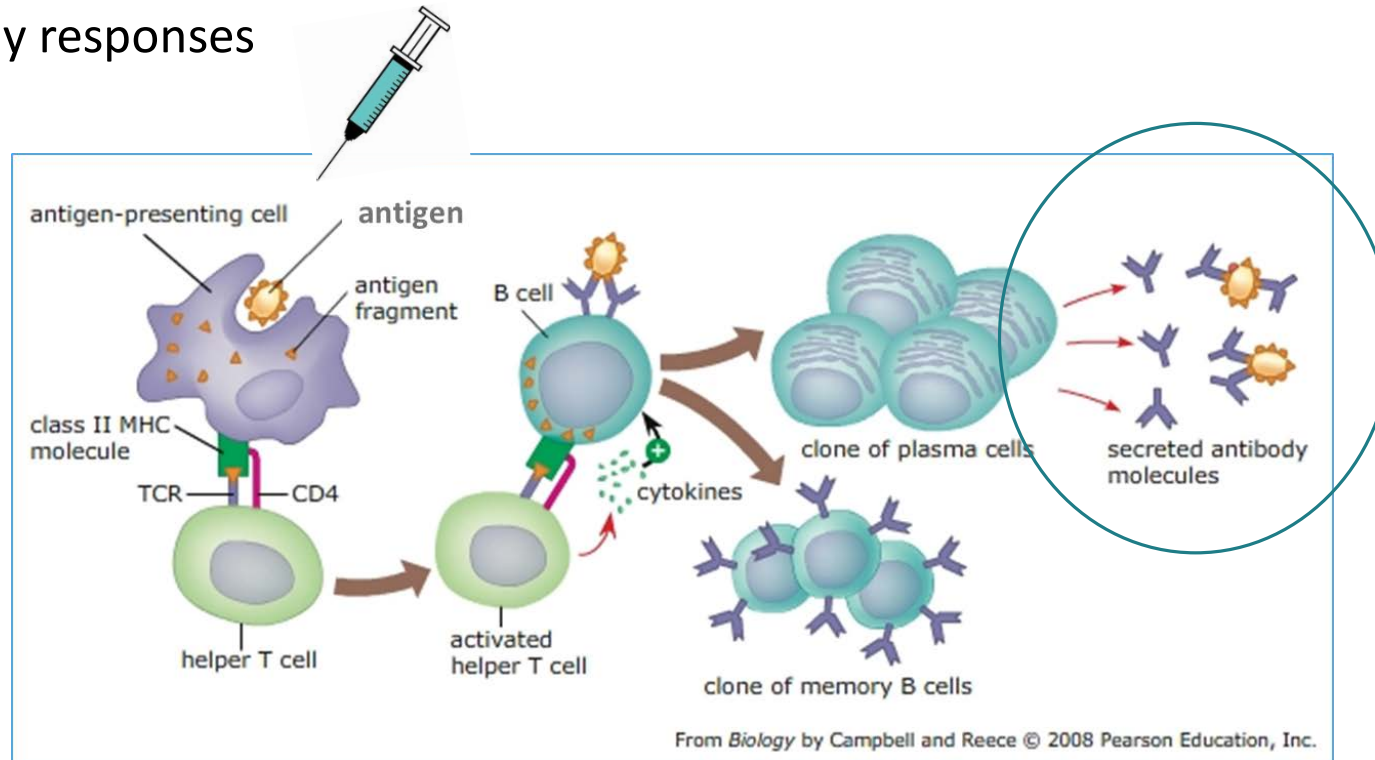
JANUARY 26, 2021

# Tools for early immunogenicity assessment

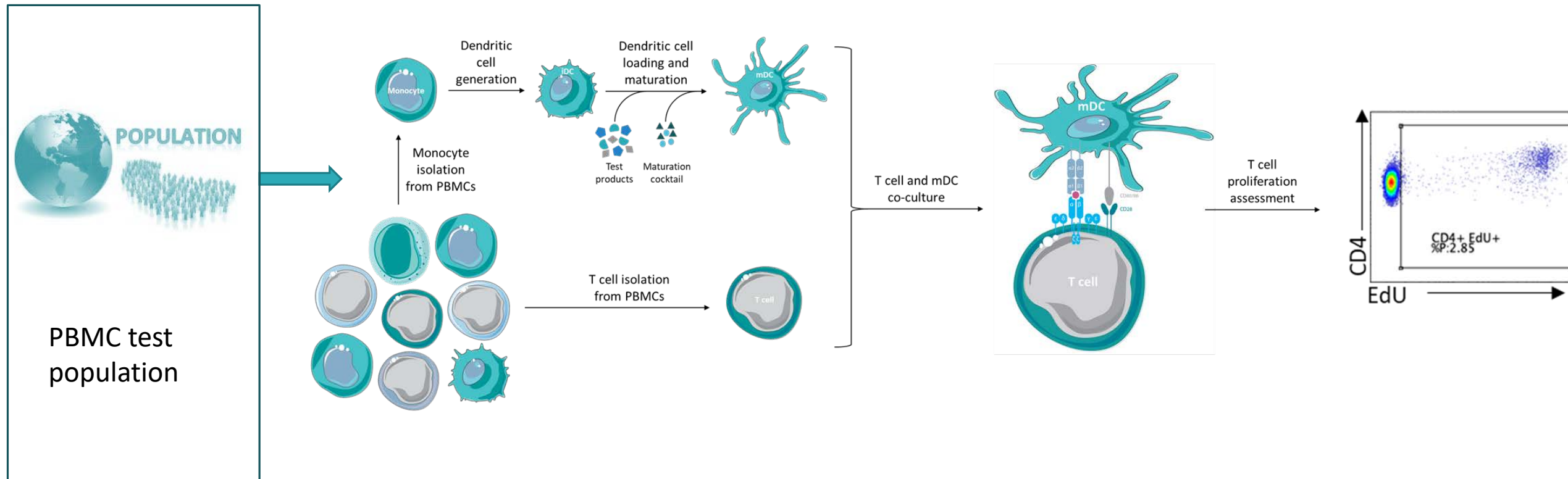


# In vitro tools: T cell proliferation assays

T cell activation/proliferation assays using human PBMC can be used as a **surrogate marker** for antibody responses

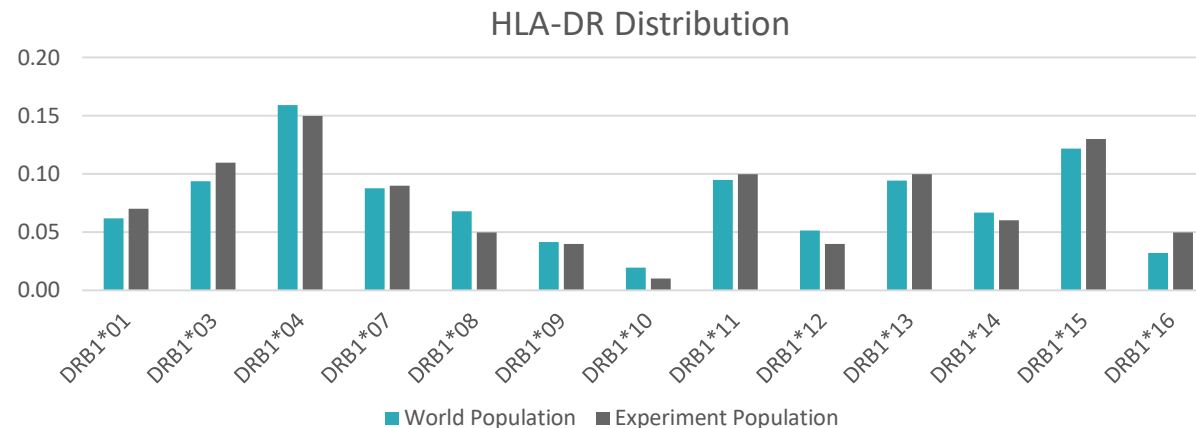


# In vitro tools: DC-T cell assays: procedure



# Test population: Donor selection

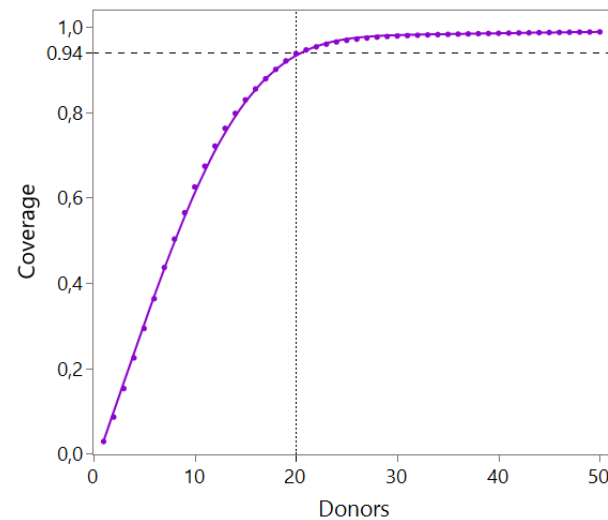
The number of donors should be sufficient to represent the **HLA distribution** of the target population and **allow the detection of a positive and negative response**



# Donor selection: HLA distribution

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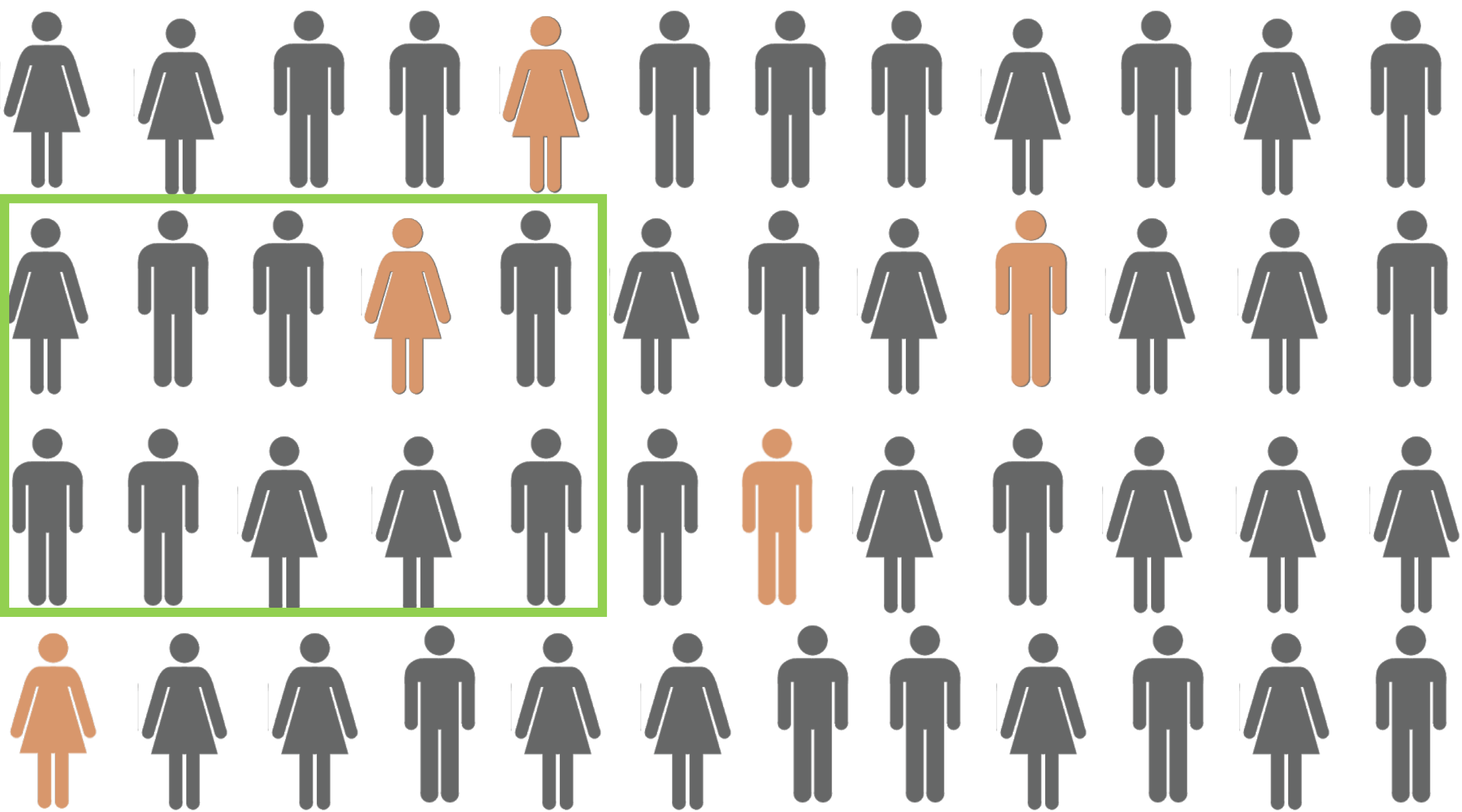
The coverage of world population in function of number of donors.



# Donor selection: rationale number of donors

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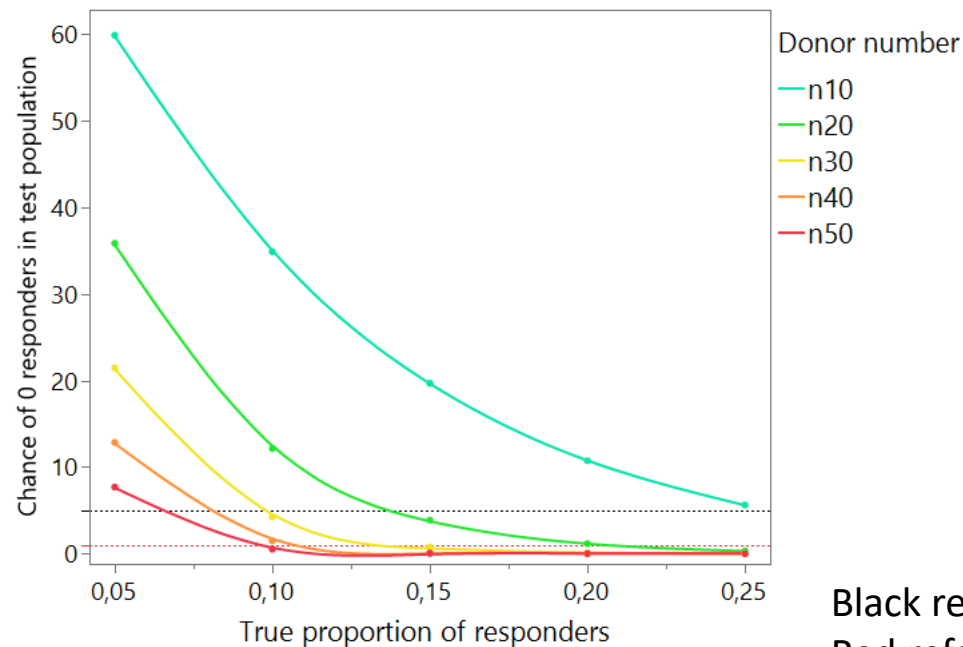
- The number of donors to include in the screening depends on the proportion of positive subjects in the population that would be of concern
- Depending on the proportion of positive subjects in the population, there is a chance that a random/representative sample of the population (test donors) would not include any reactive donors and thus the compound would be classified as negative:
  - ✓ For a fixed number of donors, the chance of including only negative donors increases if the proportion of positive subjects in the population decreases
  - ✓ For a fixed proportion of positive subjects in the population, the chance of not including any positive donor decreases if the number of donors increases
  - ⇒ If it is acceptable to consider a compound negative if less than 25% of subjects in population are negative you need less donors than when you can only consider a compound negative if less than 10% of subjects in the population are negative
- Chance that a random/representative sample of the population (test donors) would not include any reactive donor (cfr. next slide) = basis for the calculation of number of donors





# Donor selection: rationale number of donors

- Graph showing the chance of not including any positive donors (completely missing immunogenicity) in function of sample size (color code) and probability of subjects responding in the population (x-axis) based on the binomial distribution.



With 10 donors, you have 20% chance of not including any reactive donor in test population even if 15% of the population would show a response.

With 40-50 donors you have high sensitivity to detect positive responses if only <10% of population would show a response.

Black reference line: 5%

Red reference line: 1%

# Donor selection: rationale number of donors

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- Calculations on previous slide are minimal number of donors required
  - It only takes into account sampling variability in selecting donors that are true responders but assumes 100% power of the assay to classify true responders as positive
  - ✓ Power of the assay to classify positive donors as positive depends on the variation specific for the assay and number of replicates per donor

# Test population: PBMC quality

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## Quality of the primary cells:

- Variability and reproducibility of the results highly depends on the initial quality
- Quality = viability and functionality
- Most critical reagent
- Need for a large number of HLA-typed donors in order to represent the wide range of responders (strong-responders versus medium-low responders)



# Test population: preserving PBMC quality

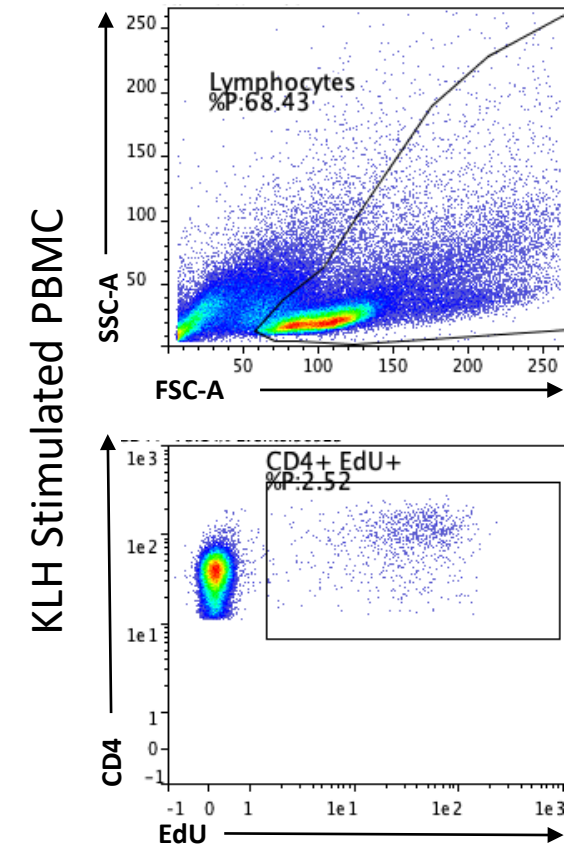
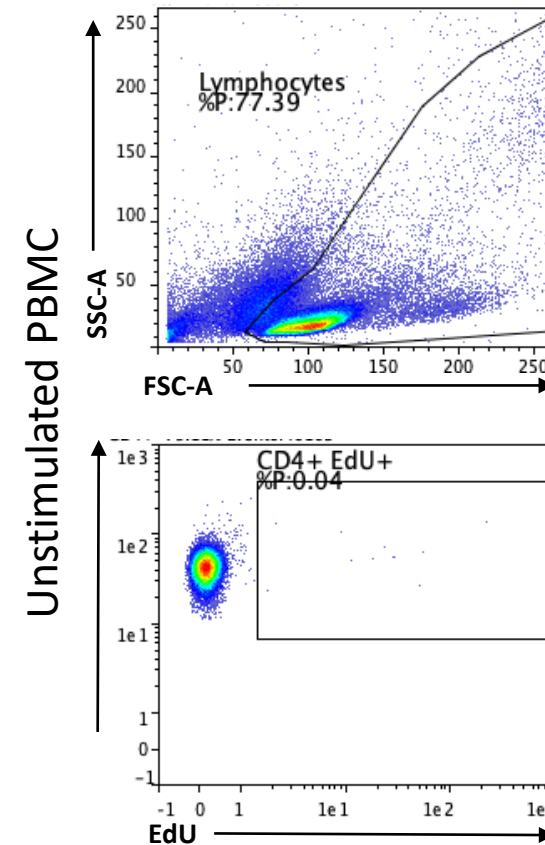
## Preserving Quality of the primary cells:

- Standardized procedures for sampling, shipping, isolation, cryopreservation, thawing, handling, ...
- Start isolation of PBMC within 8 hours after blood draw
- Alternatively, use CPT™ tubes for blood sampling
- Perform an extended quality control on all cell preparations



# Cryopreserved PBMC: functionality assesement

- Assessment of proliferative response towards polyclonal stimulation (anti-CD3 antibody)
- Assessment of proliferative response towards naïve antigen **Keyhole Limpet Hemocyanin (KLH)**



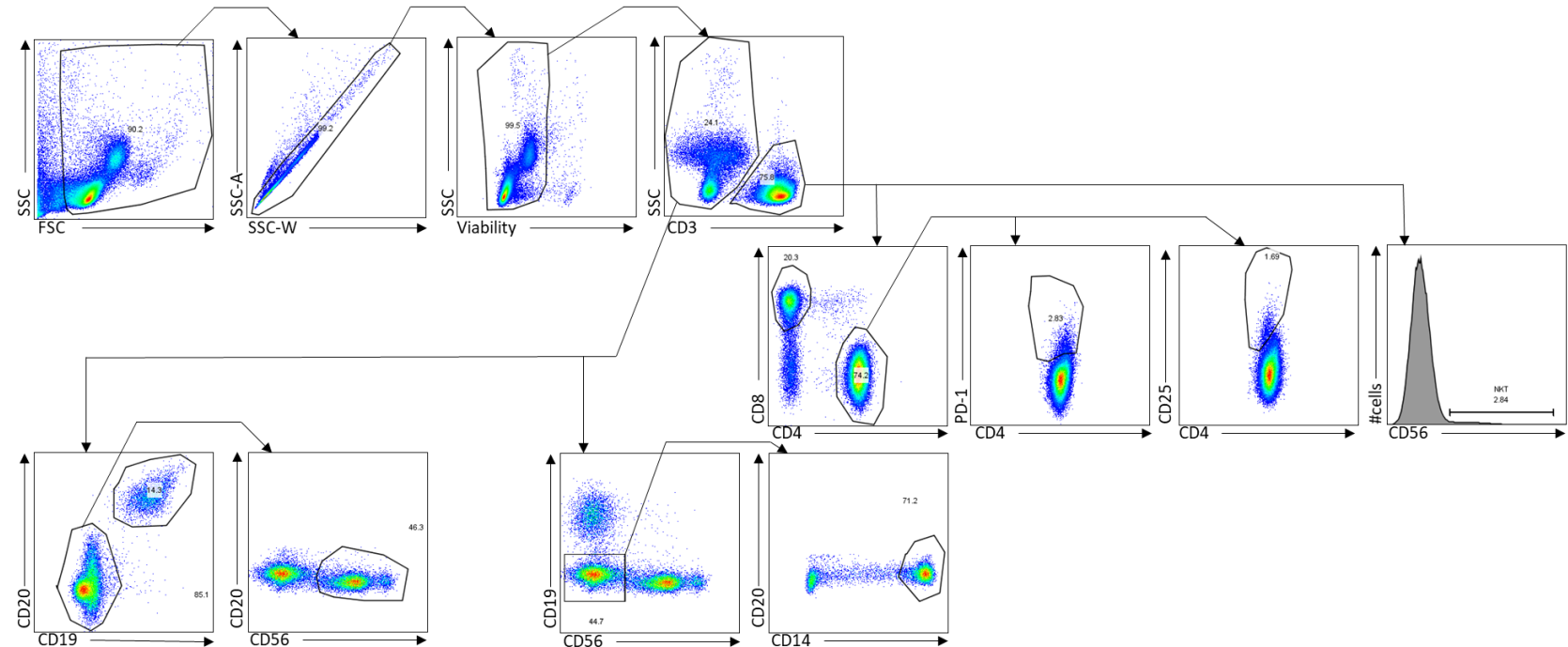
# Cryopreserved PBMC: subpopulation analysis

## Classic Surface marker staining for:

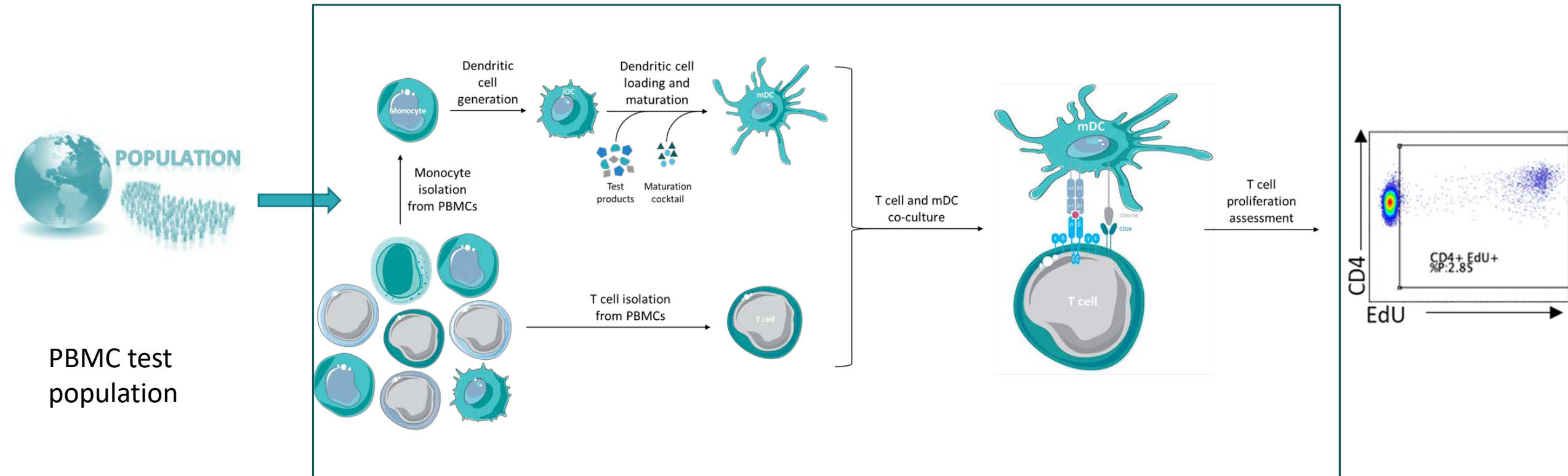
- CD14: Monocytes
- CD3: T cells
- CD4: Helper T cells
- CD8: Cytotoxic T cells

## Extended

- CD14: Monocytes
- CD3: T cells
- CD4: Helper T cells
  - PD-1+
  - CD25+
- CD8: Cytotoxic T cells
- CD56: NK and NKT
- CD19/20: B cells

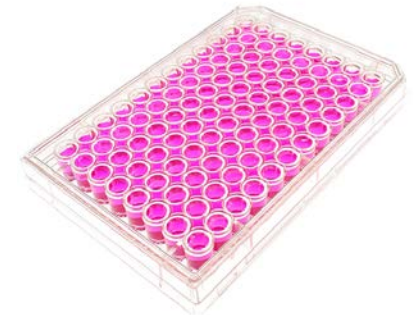


# In vitro tools: DC-T cell assays: procedure



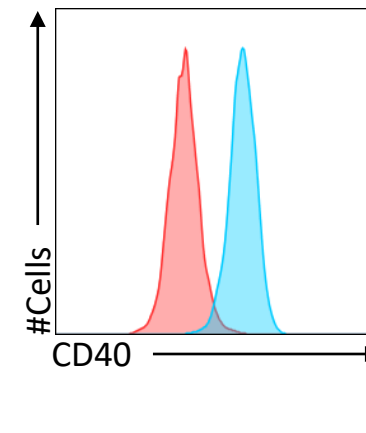
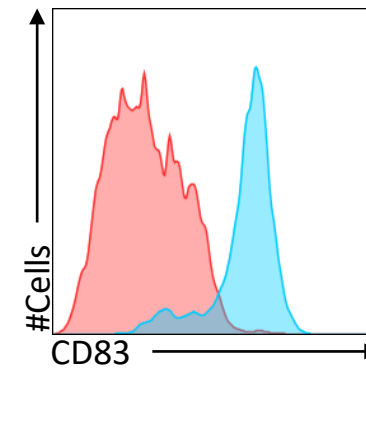
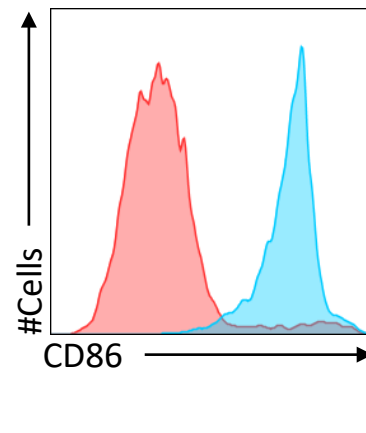
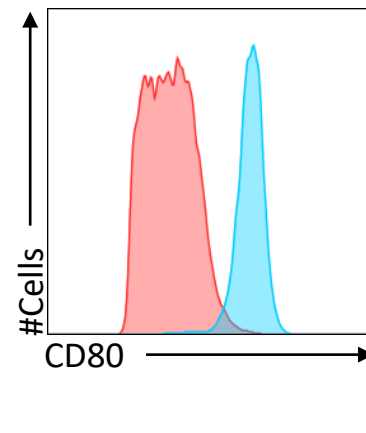
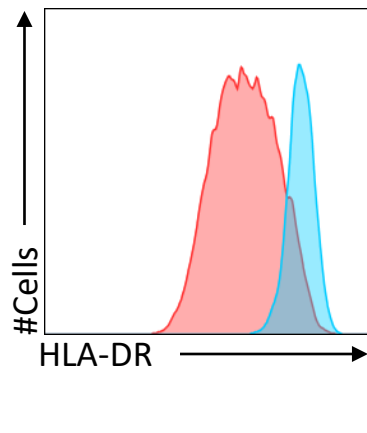
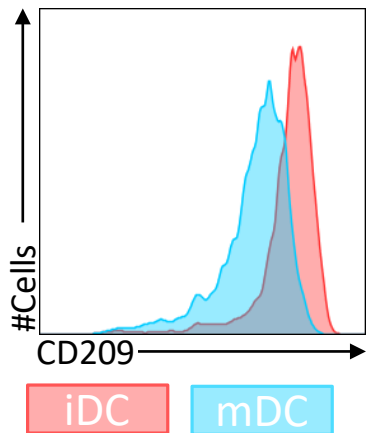
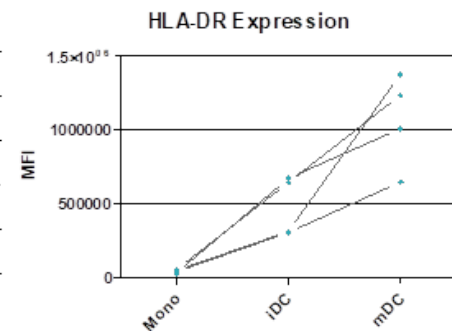
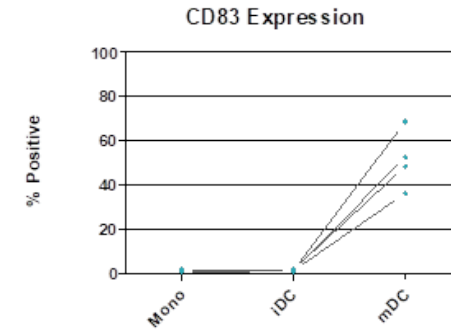
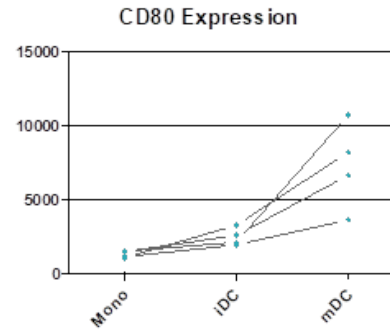
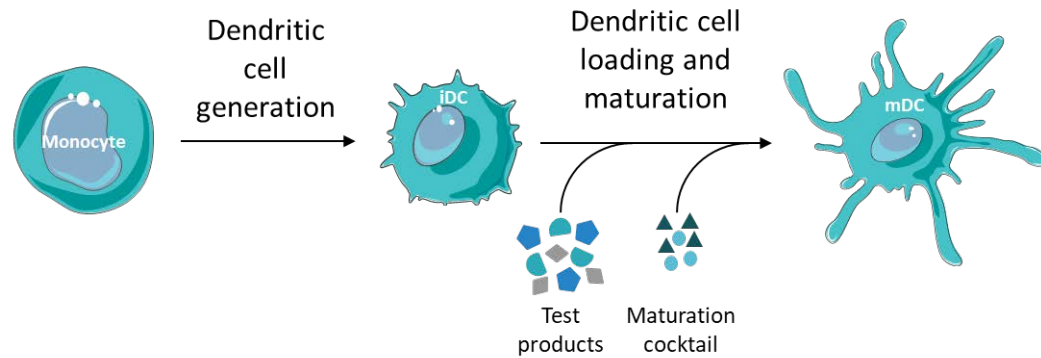
# Assay optimization and standardization

- Optimized procedures for generation of dendritic cells, use of premium grade cytokines
- Quality control of dendritic cells at each stage (monocytes, iDC and mDC)
- Optimized procedures for T cell isolation and in vitro culture (format, cell number, culture media, number of replicates ...)
- Selection of positive and negative controls and benchmark controls





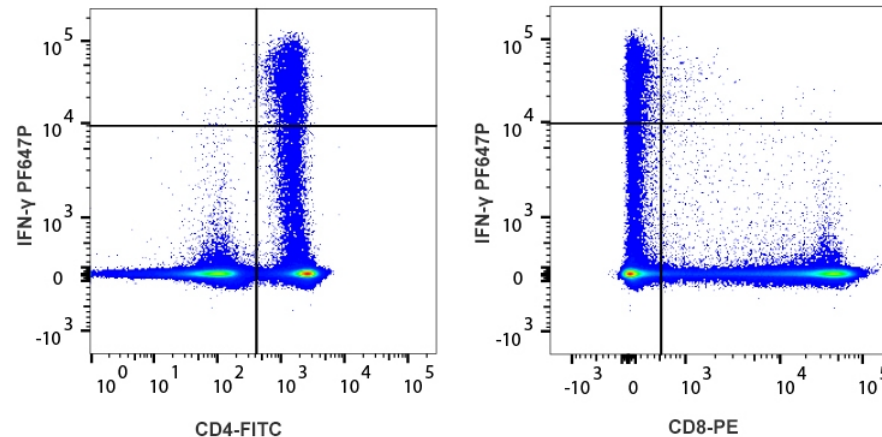
# Quality control DC at each step of the DC-T assay



# Selection of positive and negative controls

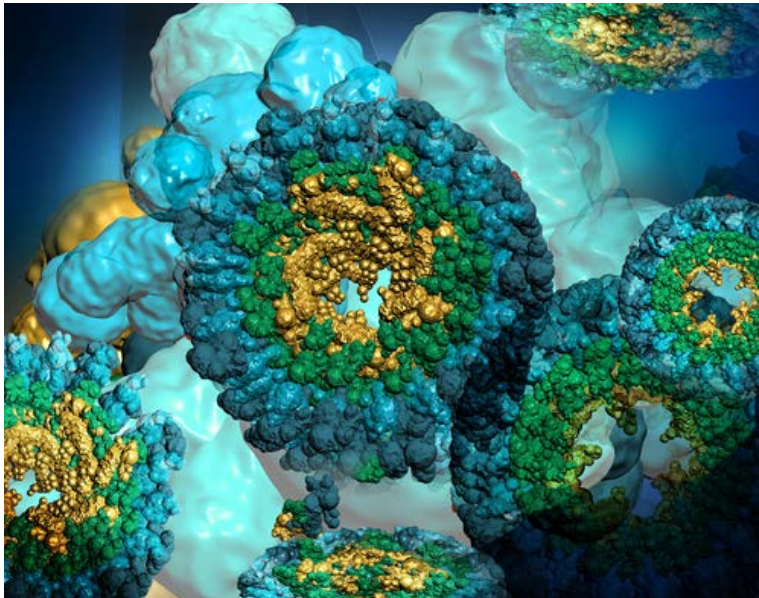
- Positive control(s): peptide mix of recall peptides: **CEFTA** Peptide Mix Mabtech and DMSO control or a strong immunogenic antigen like CMV or KLH

## Stimulation with CEFTA in Flow cytometry



# Selection of **positive** and **negative** controls

- Positive control(s): peptide mix of recall peptides: CEFTA Peptide Mix Mabtech and DMSO control or a strong immunogenic antigen like CMV or **KLH**



- The outcome with different brands/types of KLH can be varying a lot
- (Some) toxicity observed with certain lots/brands
- Always perform a lot specific titration
- Stability KLH, storage temperature
- Endotoxin levels

# Selection of benchmark/reference peptide controls

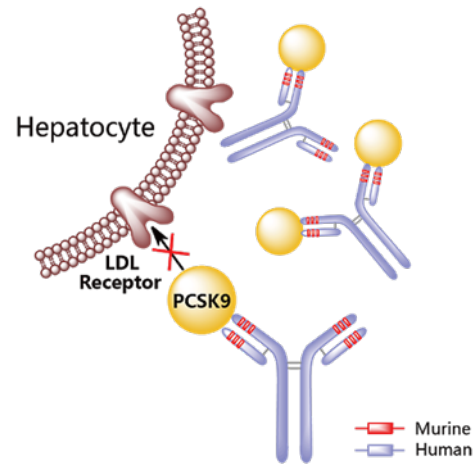
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- Benchmark/reference peptides : PADRE (universal T cell epitope sequence AKFVAAWTLKAAA) or internal positive and negative peptide mix
- Internal **reference positive peptide** mix:
  - binding of a set of 250,000 random natural non-human 15mer peptides was evaluated for 43 HLA-II alleles with NetMHCIIpan-4.0 and **strong binders** were identified. Out of these, 5 peptides with the highest degree of promiscuity were selected.
- Internal **reference negative peptide** mix:
  - binding of a set of 250,000 random natural non-human 15mer peptides was evaluated for 43 HLA-II alleles with NetMHCIIpan-4.0 and **non binders** were identified.

*In silico work performed by Prof. Morten Nielsen – DTU Denmark*

# Selection of benchmark/reference antigen/antibody controls

- 2 reference monoclonals, Bococizumab and ATR-107

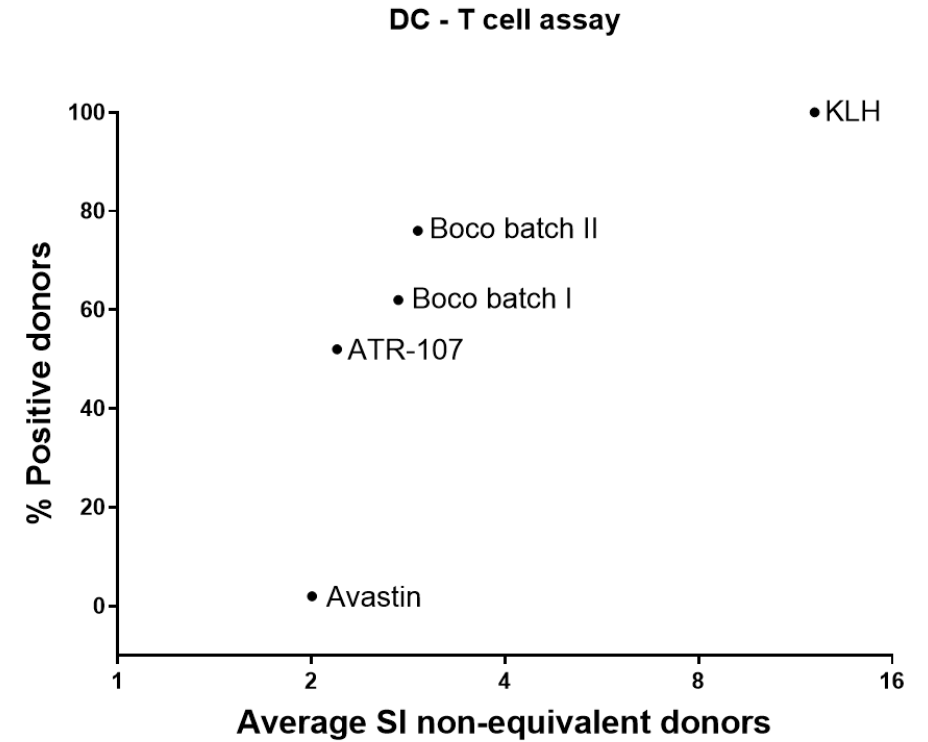
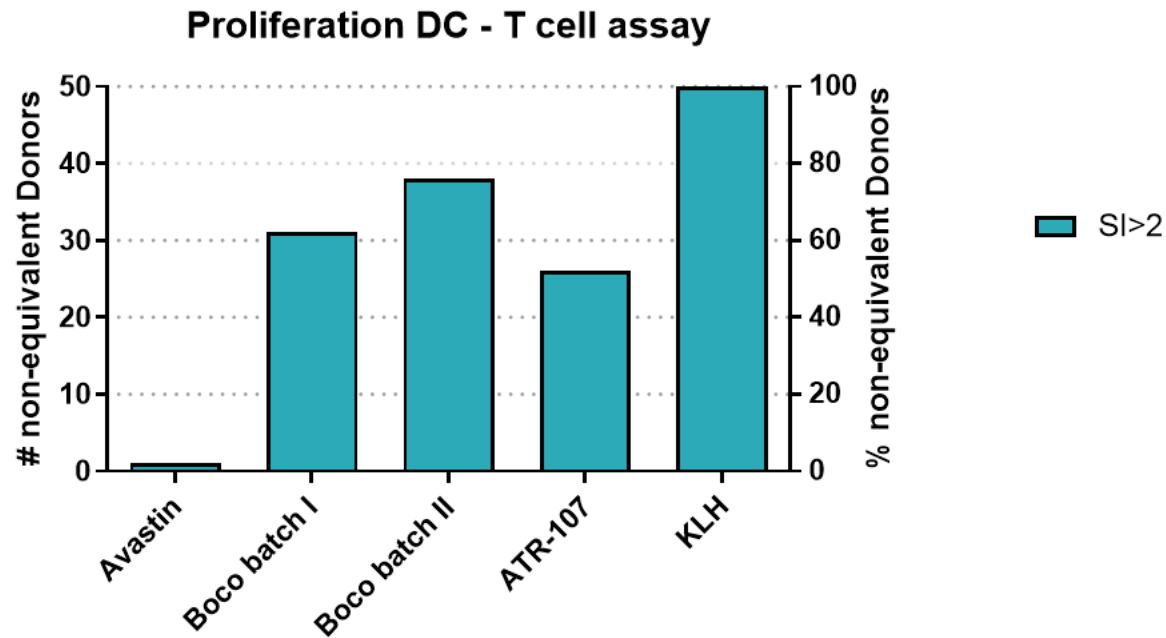


**Bococizumab**

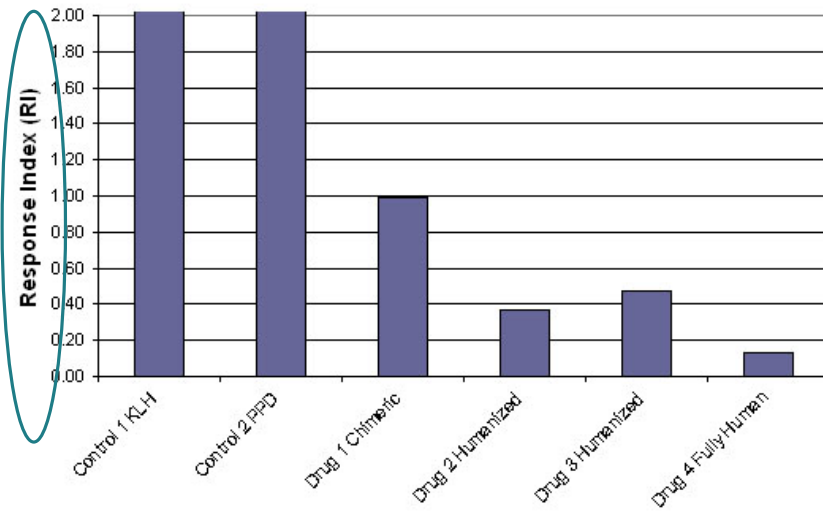
**ATR107**

Anti-IL21 Receptor Monoclonal Antibody (ATR-107): Safety, Pharmacokinetics, and Pharmacodynamic Evaluation in Healthy Volunteers: A Phase I, First-in-Human Study  
DOI: 10.1002/jcph.158

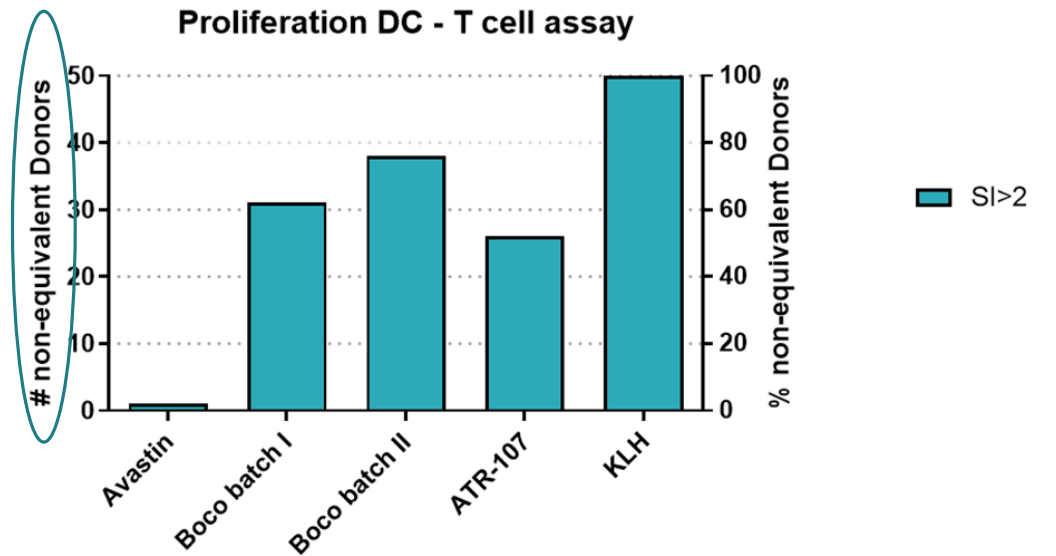
# Stability/performance bench mark controls



# Definition of positive responses

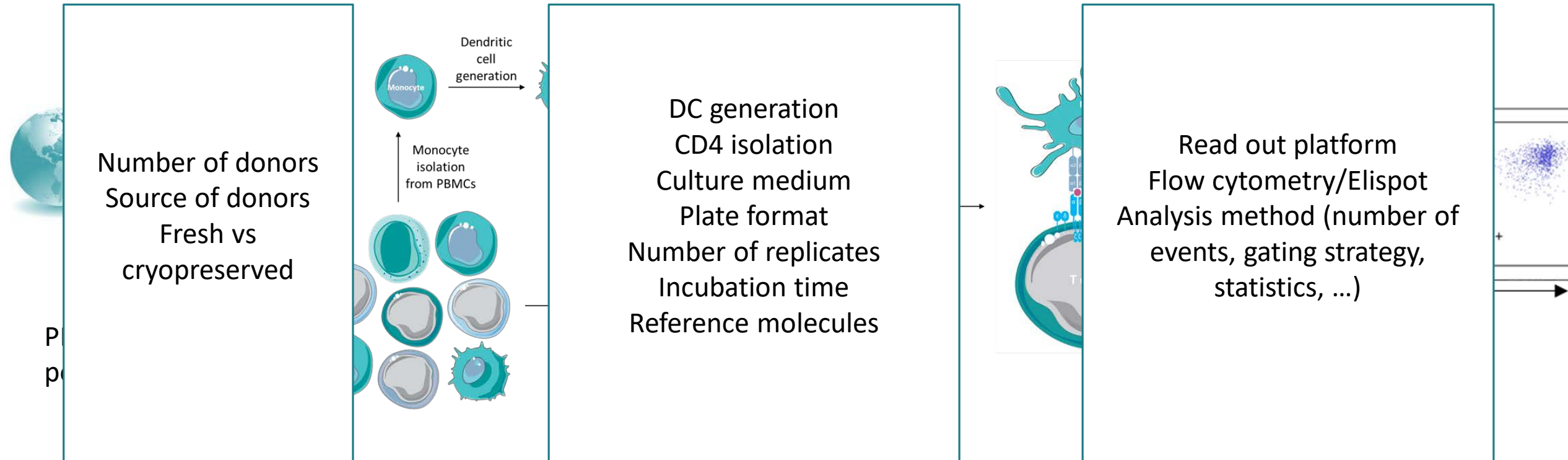


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# Harmonisation/standardisation DC-T cell assays at every level





# Harmonisation/standardization initiatives

## Cancer Vaccine Consortium - History

2002	2003	2004	2005	2006	2007
◆ CVC Founded			◆ 1st Elispot Panel	◆ 2nd Elispot Panel	◆ Immuno-Assessment Satellite Symposium
◆ Colloquium and Annual Meeting					
		◆ Potency Testing Workshop	◆ Clinical Trials Working Group		◆ FDA Workshop
				◆ Commercialization Strategy Meeting	◆ 3rd Elispot, 1st ICS, CFSE, and Multimer
				◆ Phase 3 Clinical Trials Workshop	



2015

## Guidelines for the automated evaluation of Elispot assays

Authors:

Sylvia Janetzki <sup>6</sup>, Leah Price <sup>7</sup>, Helene Schroeder <sup>1</sup>, Cedrik M Britten <sup>3,4</sup> ... Axel Hoos <sup>5</sup>

[show more details](#)

# Harmonisation/standardization initiatives






## Cytokine: X

Volume 2, Issue 4, December 2020, 100042



Development of the first reference antibody panel for qualification and validation of cytokine release assay platforms – Report of an international collaborative study

Sandrine Vessillier <sup>a</sup>  , Madeline Fort <sup>b</sup>, Lynn O'Donnell <sup>c</sup>, Heather Hinton <sup>d</sup>, Kimberly Nadwodny <sup>e</sup>, Joseph Piccotti <sup>f</sup>, Peter Rigsby <sup>a</sup>, Karin Staflin <sup>g</sup>, Richard Stebbings <sup>h</sup>, Divya Mekala <sup>i</sup>, Aaron Willingham <sup>j</sup>, Babette Wolf <sup>k</sup>, participants of the study <sup>a, b, c, d, e, f, g, h, i, j, k</sup>

Show more 



# Thank you

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