

# Unveiling the Hidden Enemy: The Laboratory's Role in Minimal Residual Disease (MRD) Detection

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1

Trace the evolution of Minimal Residual Disease (MRD) testing from its early concepts to current methodologies.

2

Explain the latest diagnostic techniques for MRD detection including flow cytometry, PCR, and NGS.

3

Assess the impact of advancements in MRD detection technologies on treatment decisions and patient outcomes.

# Objectives

# What is Minimal Residual Disease (MRD)?

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**Small amount of cancer cells that remain after treatment**

Some cells are not responsive to the drug used or develop resistance to the drug therapy used



**Most patients do not have any signs/ symptoms of malignancy**

Traditional methods of detection remission may be negative (peripheral blood smears or serum protein analysis)



**New term**

Measurable Residual Disease

# MRD Testing Interpretation

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## Positive MRD Test

- Residual cancer cells present
- Patient has high risk of relapse

## Negative MRD Test

- No residual cancer cells found
- Below detectable limits of extremely sensitive tests
- Correlates with longer remissions and potentially longer survival rates for many hematological malignancies



# Remission and Relapse

**Complete Remission (CR)** – absence of detectable disease following treatment

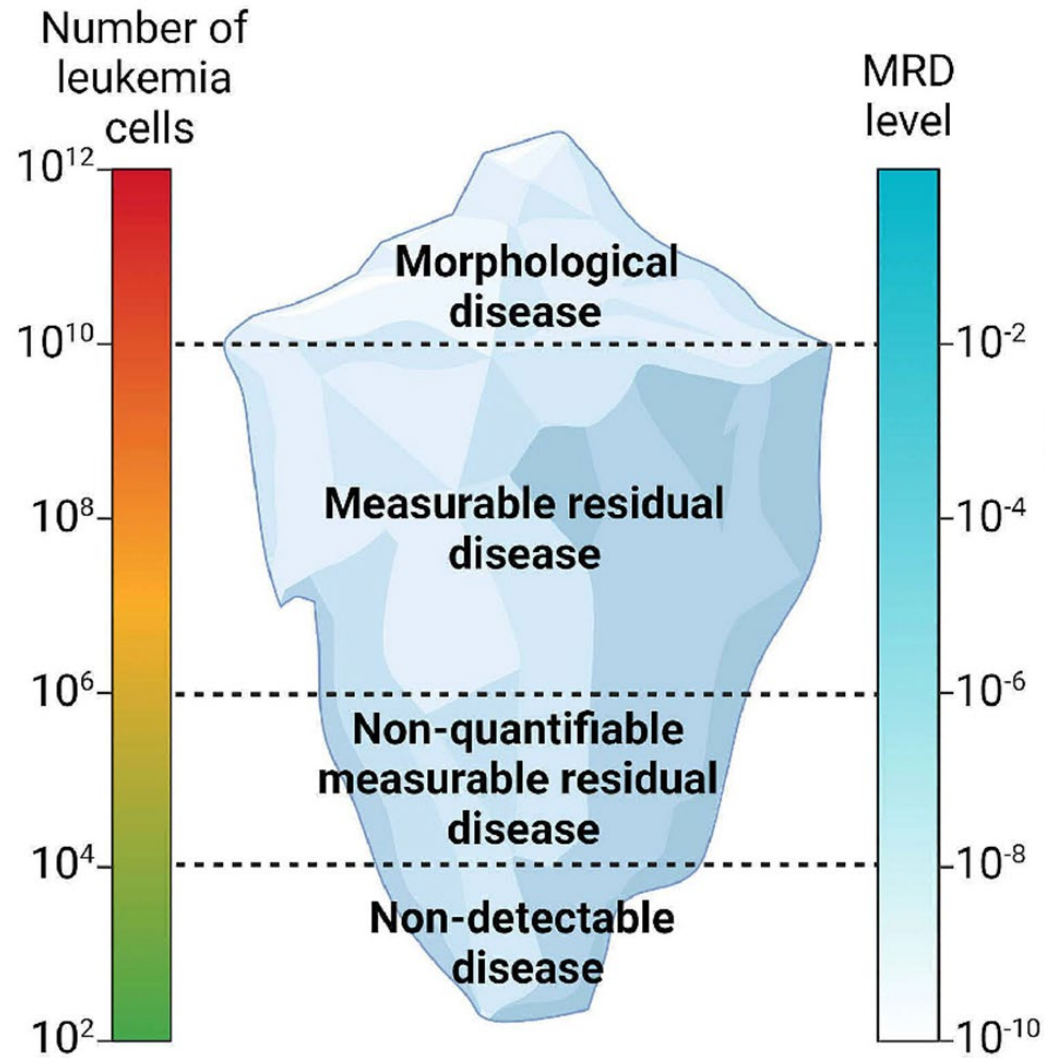
In the context of hematologic malignancy:

- No evidence of cancerous cells in the BM, blood, or other affected tissues based on standard clinical, radiological, and laboratory tests
- Patient's is free of signs/symptoms

Key milestone in cancer treatment

MRD may still be present





# Remission and Relapse

**Molecular Relapse** – residual cancer cells are detected through molecular testing but cannot be detected through routine tests

Methods employed

- Multiparameter flow cytometry (MFC)
- Polymerase chain reaction (PCR)
- Next-generation sequencing (NGS)

Usually occurs before clinical or radiologic evidence of relapse

Predicts higher risk of future relapse and allows for timely intervention

A microscopic view of several cancer cells, appearing as irregular, textured spheres in shades of purple, pink, and blue, set against a dark, grainy background with some light spots.

# Remission and Relapse

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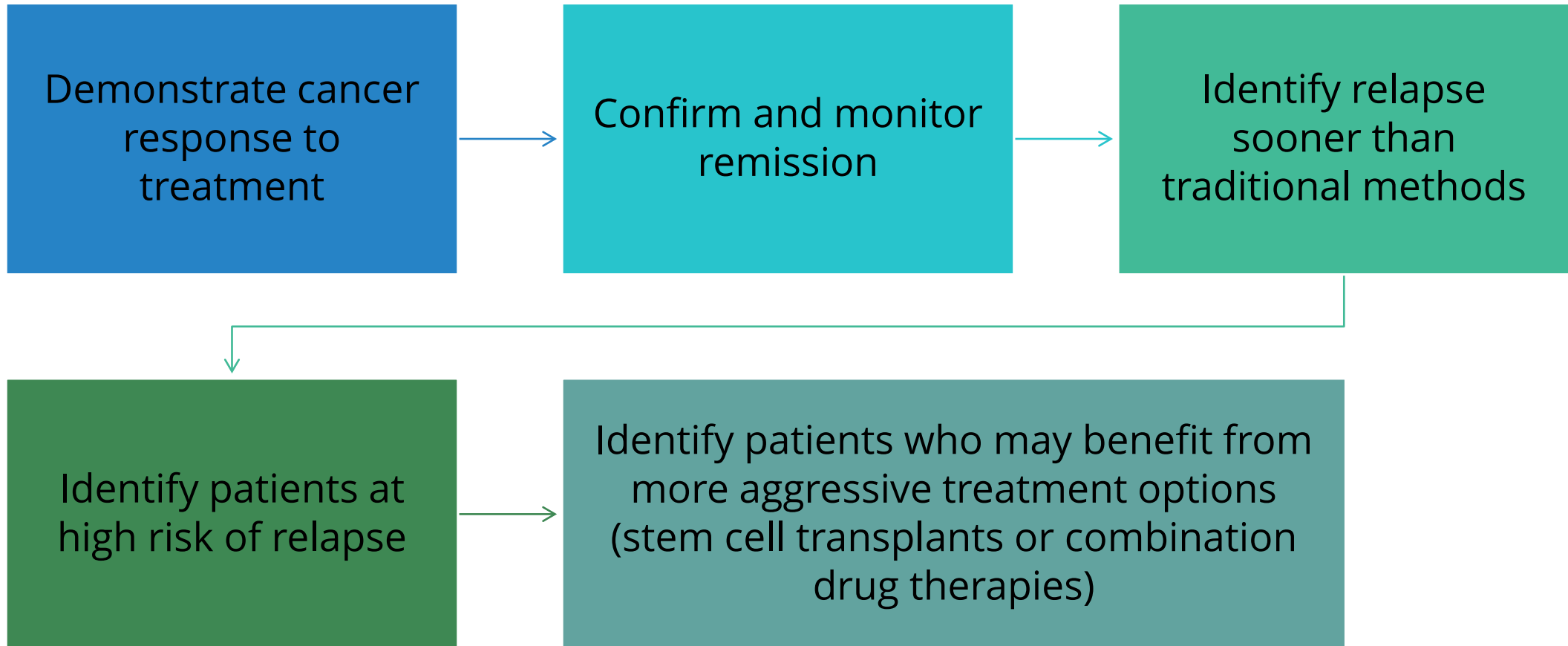
**Clinical Relapse** – re-emergence of disease detected by clinical signs and symptoms

- Abnormal blood counts
- Organ enlargement (liver and spleen)
- Imaging findings
- Symptoms: fever, fatigue, pain

**The goal of MRD testing is to identify and treat MRD before it becomes clinically apparent**

# Purpose of MRD Testing

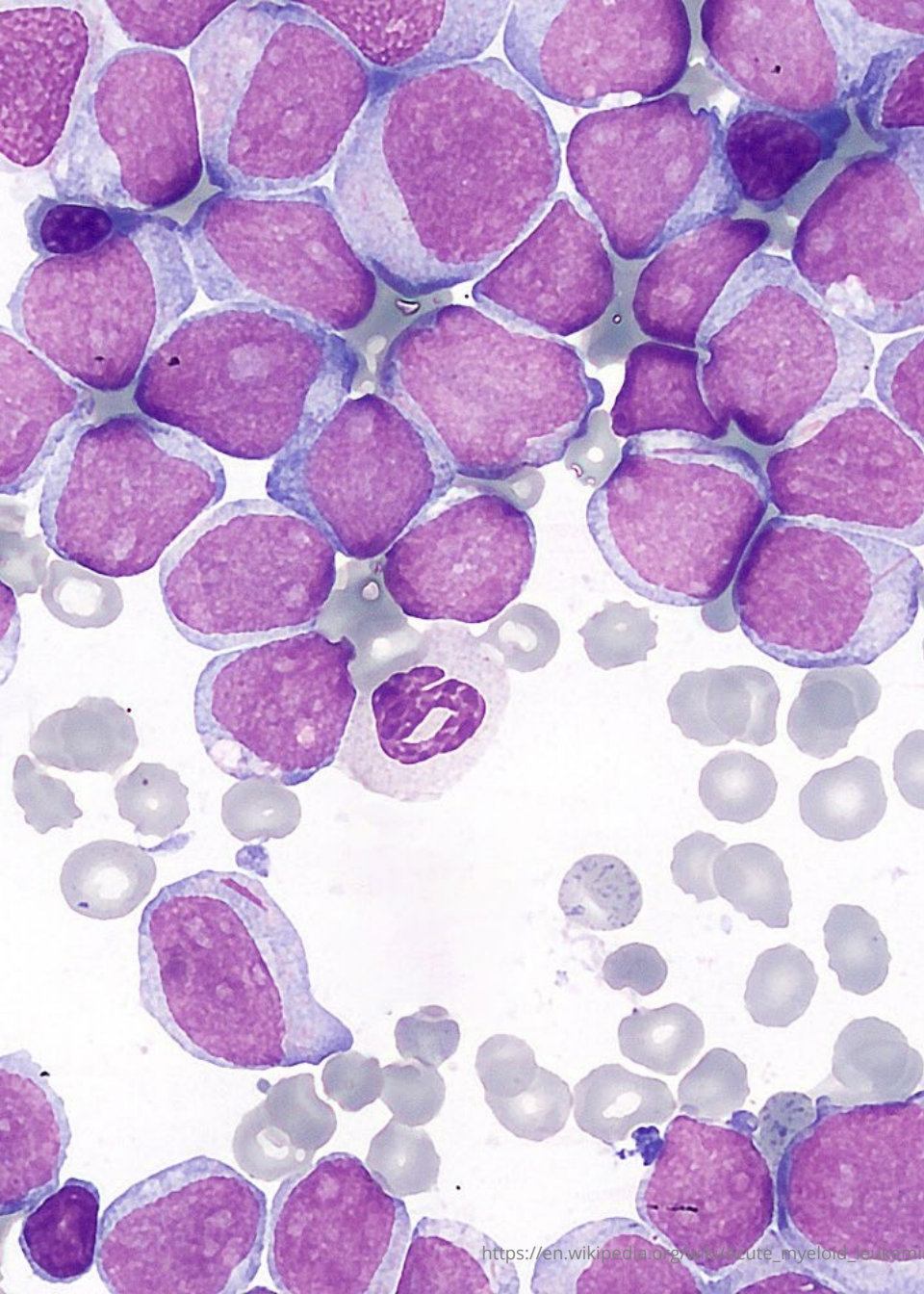
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# Historical Context of MRD Testing



# MRD Detection Before 1970

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Relapse detection occurred through clinical observation of patient symptoms

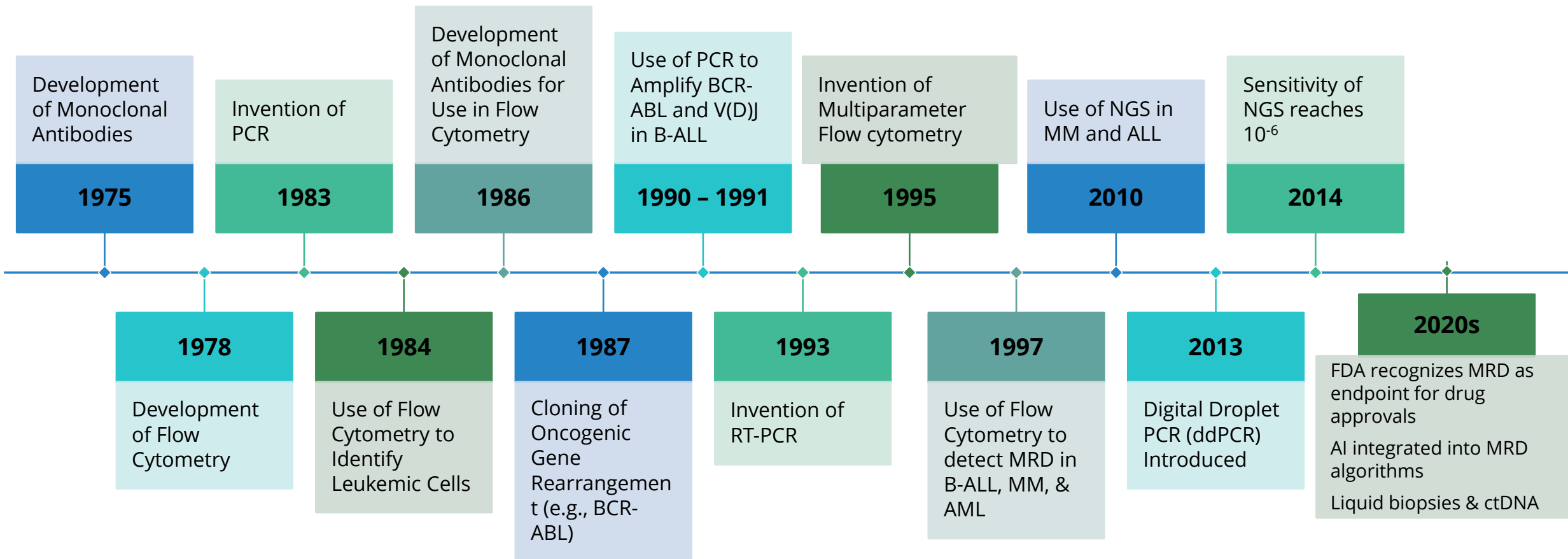
Peripheral Blood Smear

Bone Marrow Biopsy

Radiologic Imaging

Cytogenetics for specific cancers such as Philadelphia chromosome in CML

# Timeline of MRD Detection



# 1970s – 1980s

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## **1975: Development of Monoclonal Antibodies**

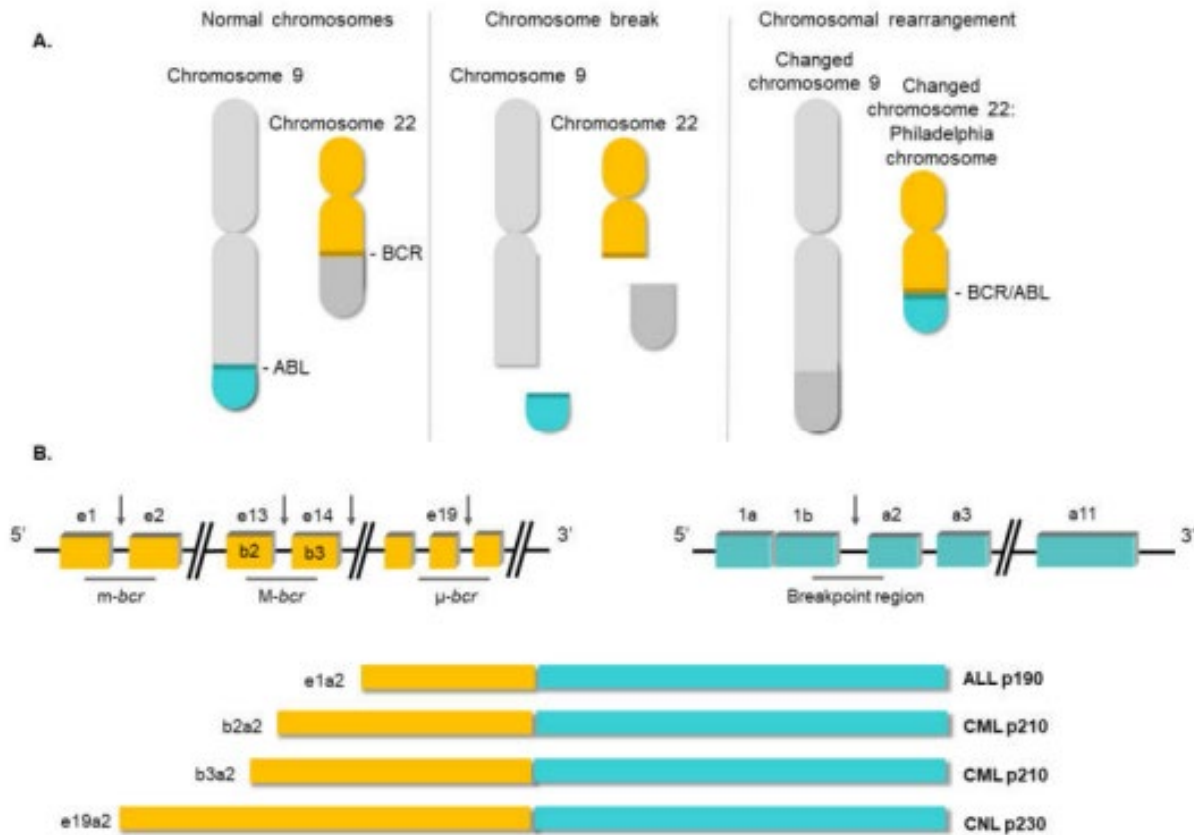
- César Milstein and Georges Köhler developed monoclonal antibodies through hybridoma method
- Bind to specific antigens on leukemic cells
- Used in several laboratory methods
- Often visualized through fluorescence

## **1978: Development of Flow Cytometry**

- Identifies individual cells in a suspension using cell markers



Georges Köhler and César Milstein won Nobel prize in physiology in 1984



# 1980s

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**1983:** Invention of PCR

**1984:** Use of Fluorescent Microscopy to Identify Leukemic Cells

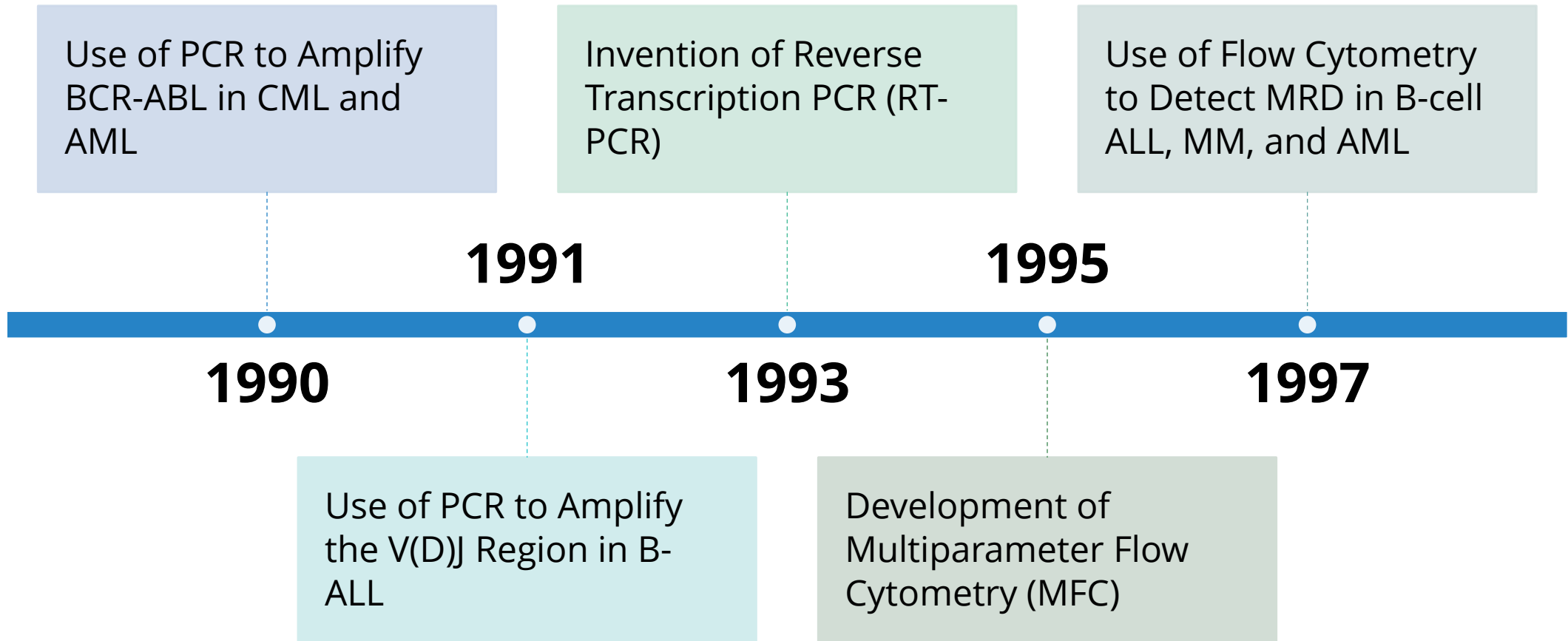
**1986:** Development of Monoclonal Antibodies for Use in Flow Cytometry

**1987:** Cloning of Oncogenic Gene Rearrangements (e.g., BCR-ABL)



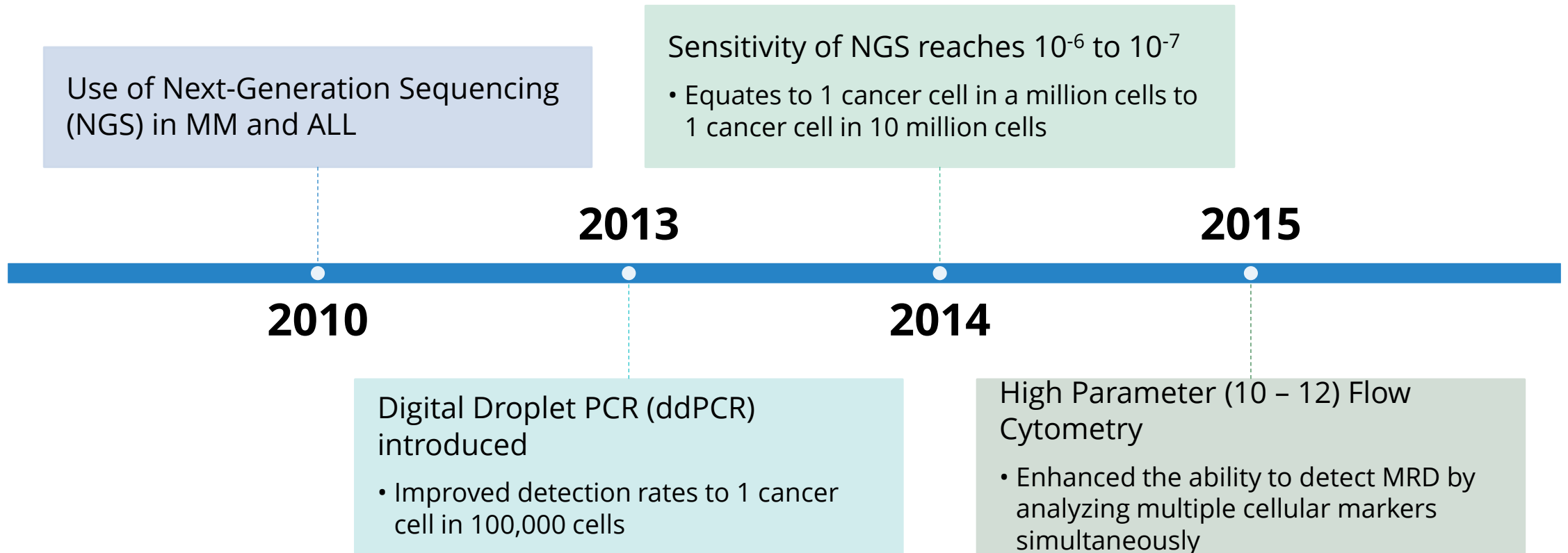
# 1990s

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# 2000s – 2010s

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Use of Next-Generation Sequencing (NGS) in MM and ALL

Sensitivity of NGS reaches  $10^{-6}$  to  $10^{-7}$

- Equates to 1 cancer cell in a million cells to 1 cancer cell in 10 million cells

**2013**

**2010**

Digital Droplet PCR (ddPCR) introduced

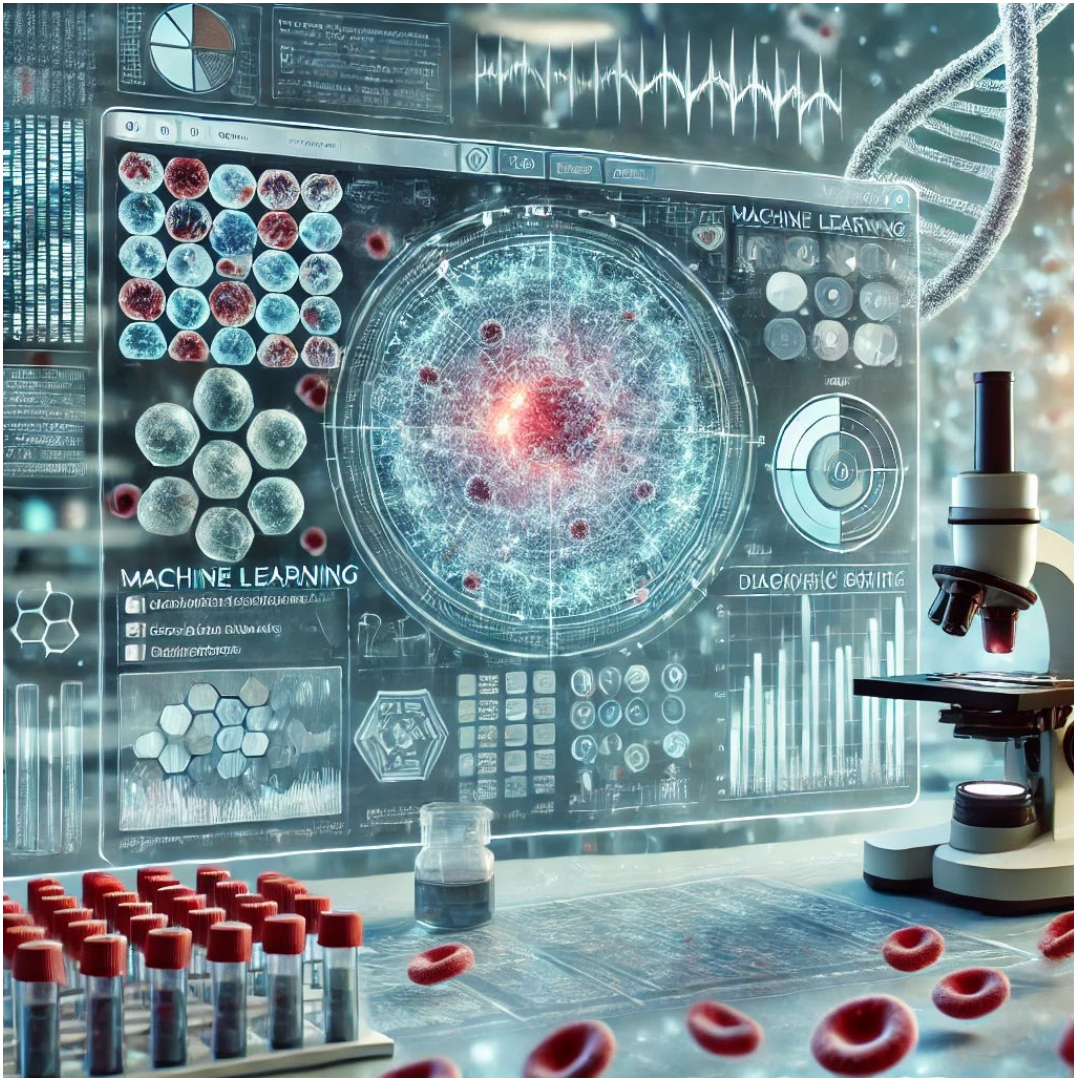
- Improved detection rates to 1 cancer cell in 100,000 cells

**2014**

**2015**

High Parameter (10 – 12) Flow Cytometry

- Enhanced the ability to detect MRD by analyzing multiple cellular markers simultaneously



# 2020s

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**2020:** FDA officially recognized MRD negativity as a surrogate endpoint

- Especially for drug approvals in MM

## **AI Integration**

- Integrated into flow cytometry and NGS analysis
- Algorithms improve consistency and accuracy of MRD identification

## **Liquid Biopsies**

- Analyze circulating tumor DNA (ctDNA)
- Allows for real-time monitoring of disease without need for invasive procedures (BM biopsy)

# Understanding MRD Testing Technologies

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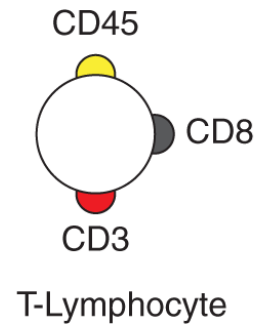
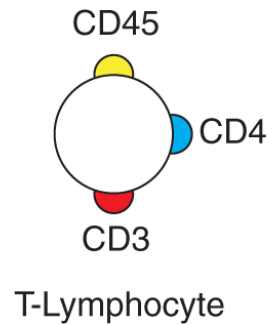
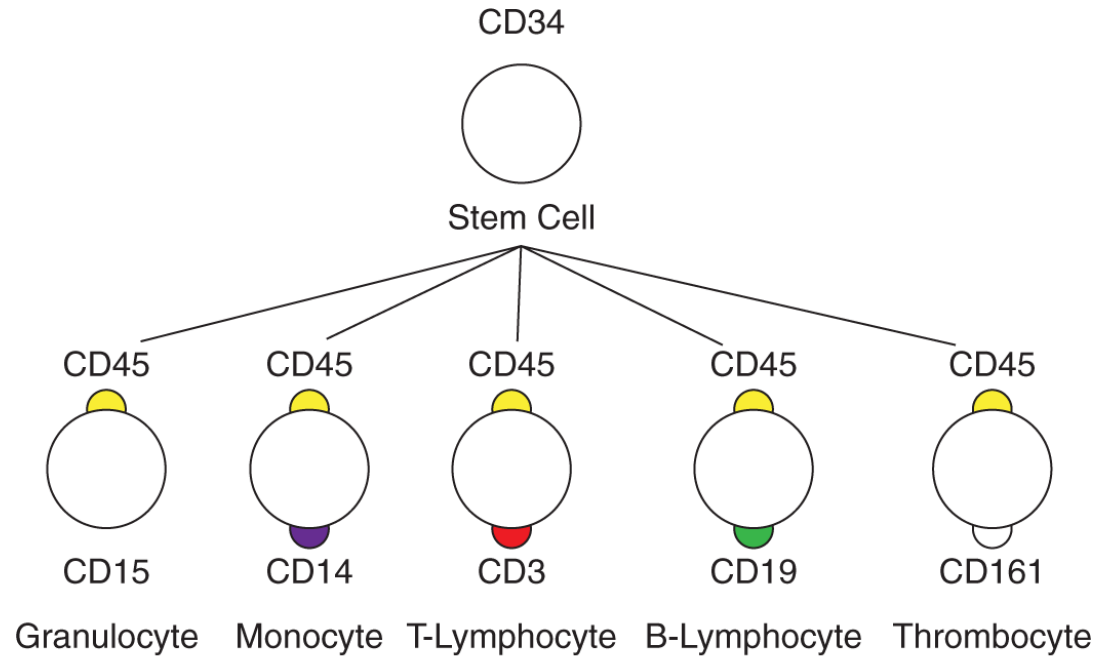
FLOW CYTOMETRY

POLYMERASE CHAIN REACTION (PCR)

NEXT-GENERATION SEQUENCING (NGS)

# Flow Cytometry Overview

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Technique that detects individual cells in a heterogeneous sample by identifying the presence or absence of protein markers on the cell's surface, in the cytoplasm, or on the cell's nucleus

## Applications in Clinical Hematology

- Automated leukocyte differentials
- Counting reticulocytes and platelets
- Immunophenotyping



# Multiparameter Flow Cytometry (MFC) for MRD

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Flow cytometry that can simultaneously recognize several different phenotypic markers

## **Advantages**

- High Sensitivity: Detects MRD down to 0.01% (1 in 10,000 cells)
- Efficient: Results in hours
- Detailed phenotyping available for a large group of malignancies

## **Disadvantages**

- Requires high level of technical expertise in MRD markers and result interpretation
- Lower sensitivity than NGS
- Limited by abnormal markers on malignant cells

# Approaches to Using FC for MRD Detection

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Antigens are expressed differently in cancer cells compared to normal cells

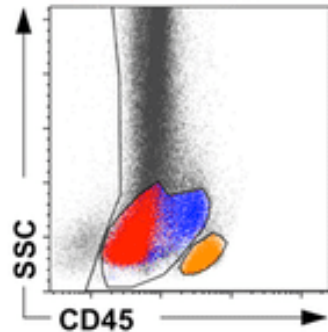
## **Leukemia-associated immunophenotype (LAIP) approach**

- The unique immunophenotype expressed by a patient's cancer cells is identified to create the leukemia's "fingerprint"
- After the patient undergoes treatment, MRD testing will look for cells with that specific immunophenotype

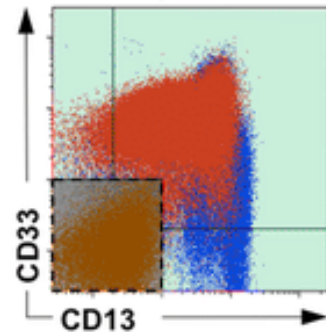
## **Different-from-Normal (DfN) approach**

- A patient's post-treatment cells are compared to a library of normal cells
- Allows detection of residual malignant cells that do not match the initial LAIP due to antigen shifts or clonal evolution
- Interpretation required expertise and well-established normal cell profiles

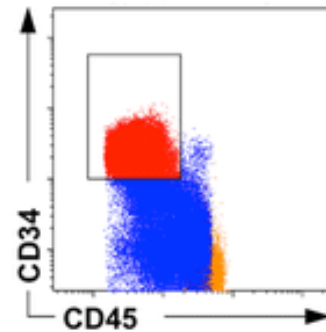
**1** CD45 / SSC  
individual gates:  
blasts/monocytes,  
lymphocytes



**2** CD33 / CD13  
fixed gates: CD13<sup>+</sup>  
and/or CD33<sup>+</sup> blasts/  
monocytes



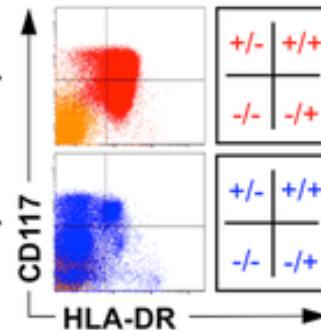
**3** CD45 / CD34  
fixed gates: CD34<sup>+</sup> or  
CD34<sup>-</sup> myeloid blasts/  
monocytes



CD34<sup>+</sup>

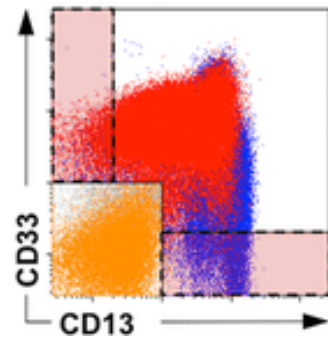
CD34<sup>-</sup>

**4** HLA-DR / CD117  
fixed gates: HLA-DR<sup>+/-</sup>  
and CD117<sup>+/-</sup> myeloid  
blasts/monocytes

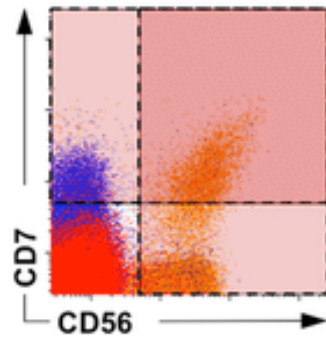


8 different myeloid  
blast populations  
by CD34, CD117  
and HLA-DR

**5** CD33 / CD13 and CD7 / CD56  
fixed gates: CD33<sup>-</sup> or  
CD13<sup>-</sup> myeloid blasts/  
monocytes



fixed gates: CD7<sup>+</sup> or  
CD56<sup>+</sup> myeloid  
blasts/monocytes



32 different  
subpopulations  
by CD13, CD33,  
CD7 and CD56

**6** Applying reference values  
*upper limit of the one-sided  
97.5% reference range*

Bone marrow donor	30
ALL in molecular CR	19
PCNSL	9
Hip surgery	32

**7** Results

4 categories of MRD  
• CD13 deficiency  
• CD33 deficiency  
• cross-lineage CD7  
• cross-lineage CD56

# Clinical Relevance of LAIP-DfN in MRD Detection

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Sensitivity  
and  
Specificity

Personalized  
Medicine

Broad  
Applications

# Flow Cytometry Process

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## 1. Sample Preparation

- Acceptable specimens: Whole blood, bone marrow, CSF, and lymph node and tissue biopsies
- Cell count performed to optimize cellular ratio for analysis
- Varies by type of specimen
  - BM aspirates passed through cell strainer to remove clumps and debris
  - CSF may have to be centrifuged more than once to concentrate cells
  - RBC lysing agents may be added to reduce interference from non-leukocyte cells

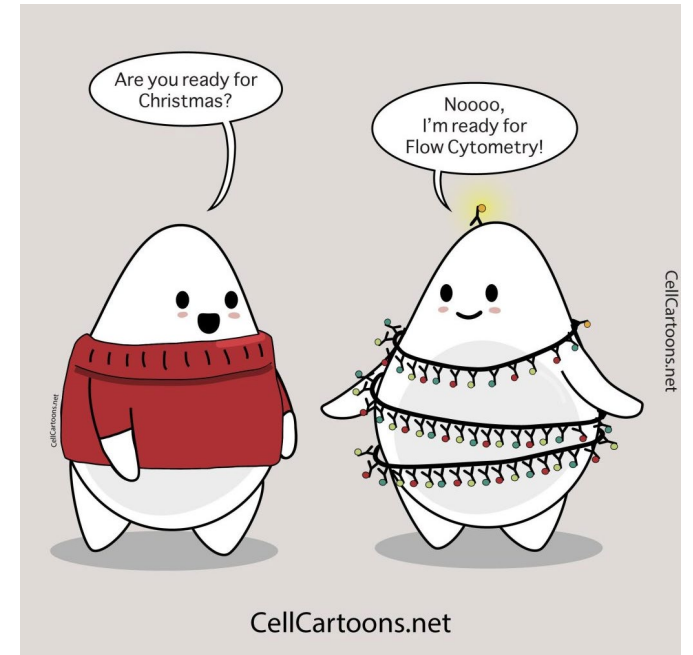


# Flow Cytometry Process

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## 2. Staining with Fluorescent Antibodies

- Cells are incubated with a panel of fluorescently labeled antibodies
- Antibodies can target antigens on the cell's surface, cytoplasm, or nucleus
- The sample is incubated for approximately 20 to 30 minutes in the dark at 4°C to avoid photobleaching
- After staining, cells are washed to remove unbound antibody and resuspended in buffer for analysis



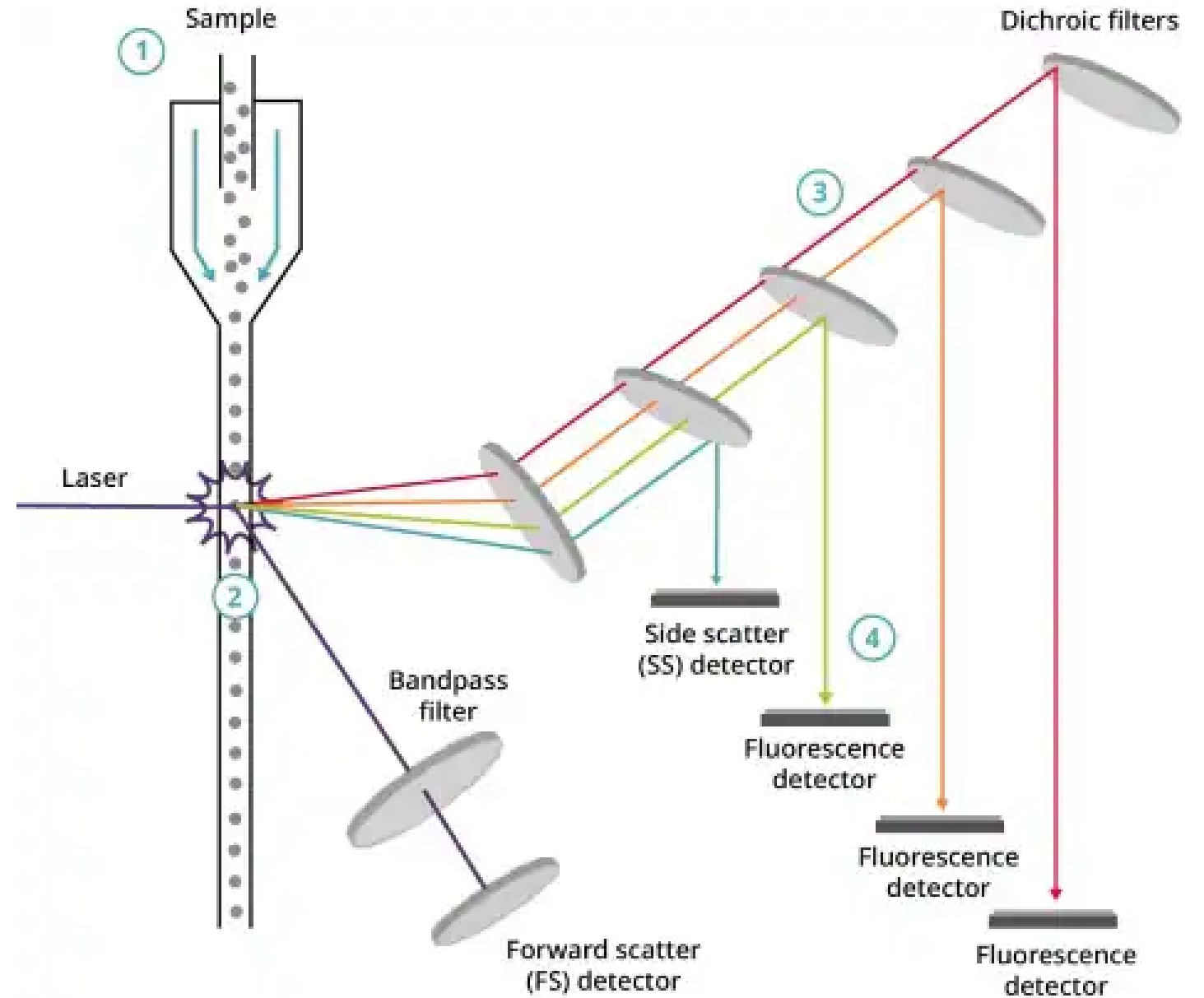
**Table 1 Surface or intracellular antigens commonly assessed by flow cytometry for lineage classification and immunophenotyping of hematolymphoid malignant neoplasms**

<b>Lineage</b>	<b>Antigens</b>
Stem cells	CD34, CD38, CD45
B cells	CD5, CD10, <b>CD19, CD20, CD22</b> , CD23, CD25, CD34, CD38, CD43, CD45, <b>CD79a</b> , CD103, CD200, FMC2, <b>cIgM, Kappa, Lambda</b> , LEF1, TdT
Plasma cells	CD19, CD20, CD38, CD45, CD56, CD117, CD138, <b>cKappa, cLambda</b>
T cells/NK cells	CD1a, <b>CD2, CD3</b> , CD4, <b>CD5</b> , CD7, <b>CD8</b> , CD10, CD16, CD25, CD26, CD30, CD34, CD45, CD56, <b><math>\alpha\beta</math>-TCR, <math>\gamma\delta</math>-TCR</b> , TdT, <b>TRBC1</b>
Myelomonocytic cells	CD4, CD7, CD10, CD11b, CD11c, <b>CD13, CD14</b> , CD15, CD16, <b>CD33</b> , CD34, CD36, CD38, CD45, CD56, <b>CD64, CD65</b> , CD71, CD117, CD123, <b>cMPO, cLyso</b> , HLA-DR
Erythroblasts	CD34, CD36, CD38, CD45, CD71, CD117, <b>CD235a</b>
Megakaryoblasts	CD33, CD34, CD38, <b>CD41, CD42</b> , CD45, <b>CD61</b> , CD117, HLA-DR

The antigens in bold are lineage-specific markers. cIgM, cytoplasmic IgM; cKappa, cytoplasmic kappa; cLambda, cytoplasmic lambda; cLyso, cytoplasmic lysozyme; cMPO, cytoplasmic peroxidase; TCR, T-cell receptor; TdT, terminal deoxynucleotidyl transferase; TRBC1, TCR  $\beta$  chain constant 1.

# Flow Cytometry Process

## 4. RUNNING THE FLOW CYTOMETER



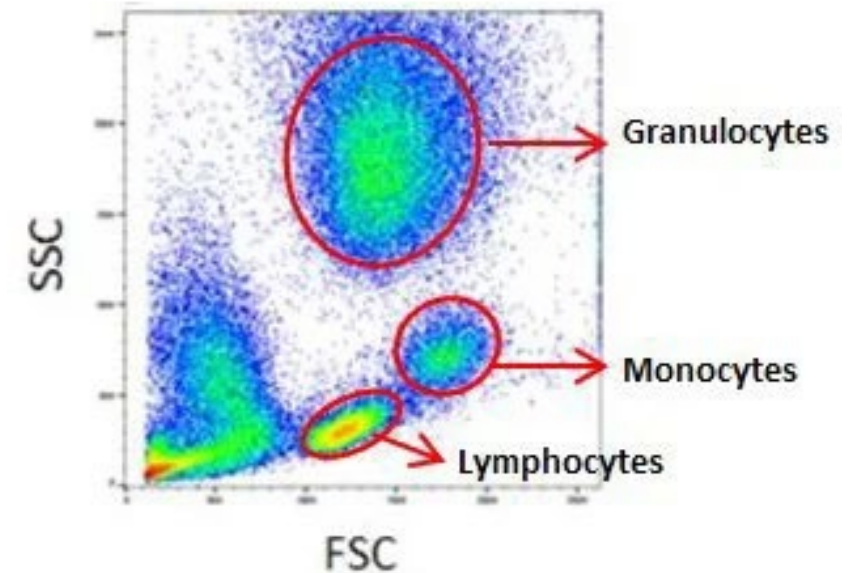
# Flow Cytometry Process

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## 5. Data Analysis

### Cell Gating

- Sequential Gating
  - Unwanted cells are eliminated based on scatter properties to isolate the cell population of interest
  - This is adjusted to exclude dead cells, debris, and doublets
- Population-Specific Gating
  - Based on antigen expression patterns (LAIP and DfN approaches)
- Boolean Gating
  - AND, OR, NOT logic



# Flow Cytometry Process

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## 5. Data Analysis

### Techniques

- Algorithms are used to cluster cells based on their marker expression profiles to help find small clusters of abnormal cells that may not be evident through manual gating alone
- Histograms and dot plots are used to graphically visualize normal from abnormal cell populations
- Specialized software and machine learning applications are emerging to assist with complex or rare hematological malignancies

Score	B-Lymphoid	T-Lymphoid	Myeloid
2	CD79a	CD3	MPO
	cCD22	TC Ralpha/beta	
	clgM		
1	CD19	CD2	CD117
	CD20	CD5	CD13
	CD10	CD8	CD33
		CD10	CD65
0.5	TdT	TdT	CD14
	CD34	CD7	CD15

**Abbreviation:** TCR, T-cell receptor.

	Precursor B-ALL	Common ALL	Pre-B-ALL	Mature-B-ALL
HLA-DR	Positive			
cCD22				
CD79a				
CD19				
TdT	Positive			Negative
CD10	Negative	Positive		Negative
clgM	Negative		Positive	Negative
slg	Negative			Positive

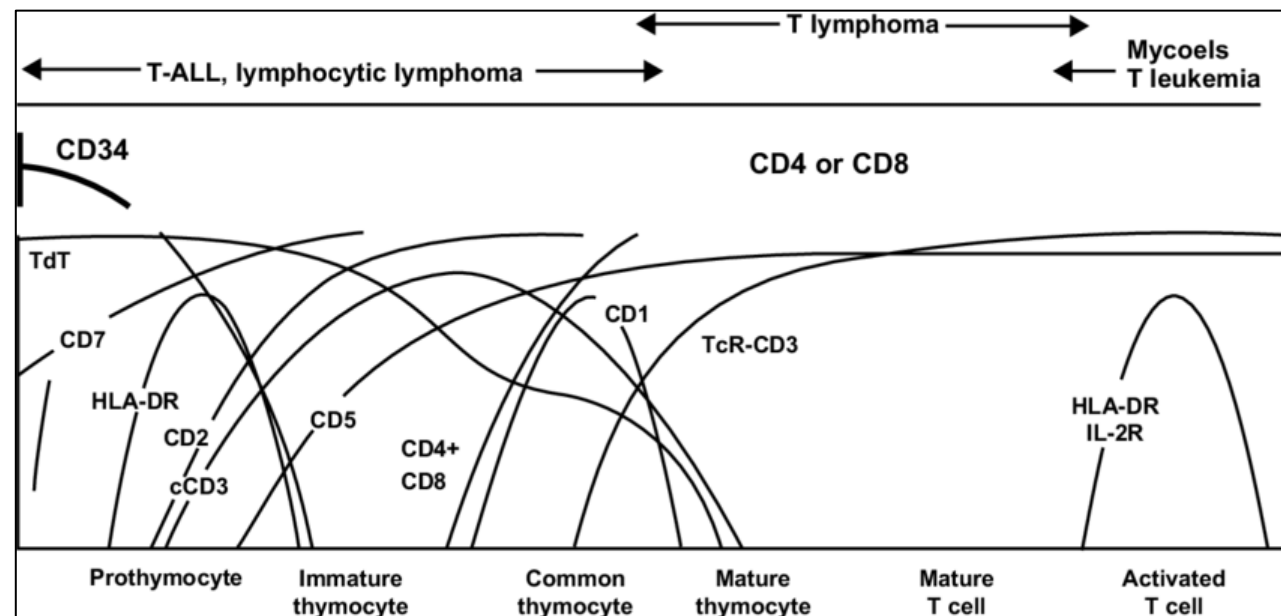
**Note:** Reproduced from NewBU.<sup>65</sup>

**Abbreviations:** ALL, acute lymphoblastic leukemia; B-ALL, B-acute lymphoblastic leukemia.

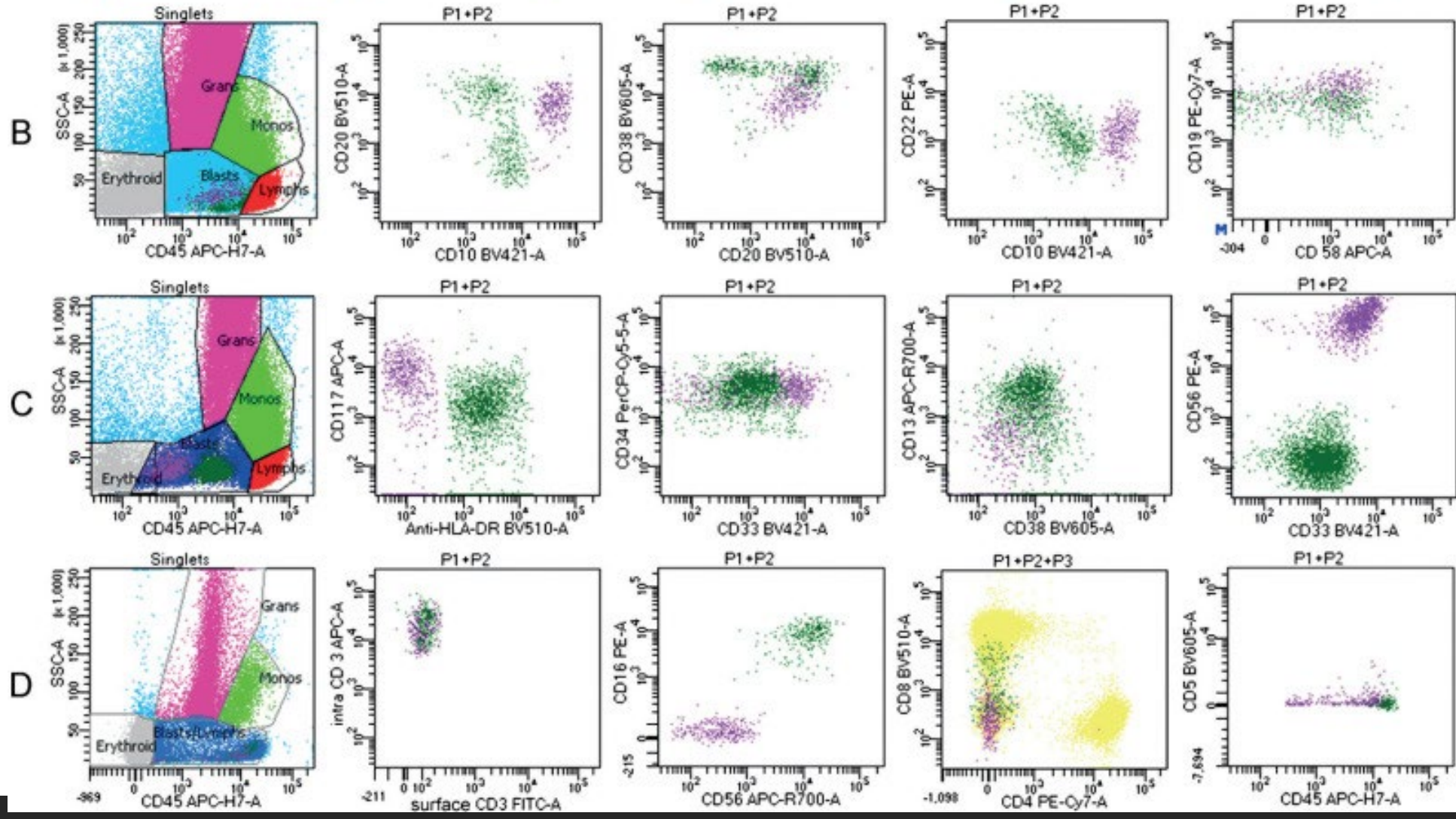
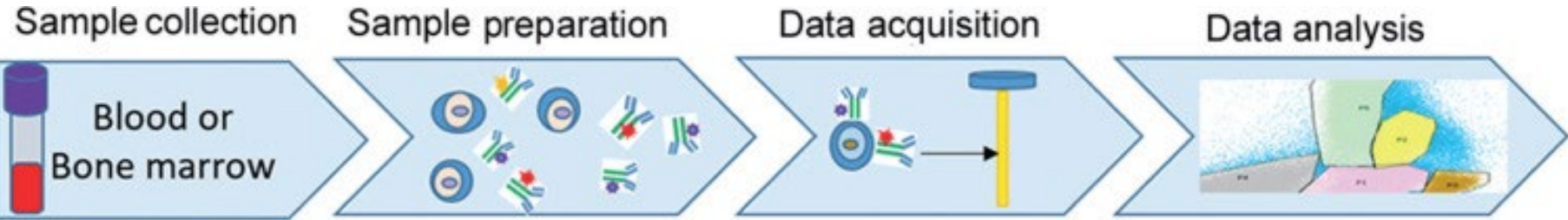
	Pro-T-ALL	Pre-T-ALL	Cortical-T-ALL	Mature-T-ALL
TdT	Positive			Negative
cCD3	Positive			
CD7	Positive			
CD2	Negative	Positive		
CD5	Negative	Positive		
CD4	Negative		Positive for CD4 and CD8	Positive for CD4 or CD8
CD8	Negative			
CD1a	Negative		Positive	Negative
sCD3	Negative			Positive

**Note:** Reproduced from NewBU.<sup>65</sup>

**Abbreviation:** T-ALL, T-acute lymphoblastic leukemia.







Pacific Blue™	OC515™	FITC	PE	PerCP-Cyanine5.5	PE-Cyanine7	APC	APC-C750
CD20+CD4	CD45	CD8 + SmIgLambda	CD56 + SmIgKappa	CD5	CD19 + TCRγδ	SmCD3	CD38



**Reagent composition:**

**Lyophilized tubes:**

- 12 antibodies against **surface antigens** (5 vials, 5 tests per vial)

**Compensation vials** (4 tubes in lyophilized format):

- PerCP-Cyanine 5.5-CD5
- PE-Cyanine7-CD19+TCRγδ
- APC-C750-CD38
- OC515-CD45

“The LST kit is an 8-color antibody combination designed by the EuroFlow™ consortium aiming to detect populations of mature B-, T- and NK-cells lineage for research purposes.”

# Standardized Protocols

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## EuroFlow External Quality Assessment Program

- Developed by Children’s Oncology Group (COG)
- Different protocols created for specific malignancies
  - EuroFlow Lymphoid Screening Tube
  - Multiple Myeloma MRD tube
  - B Cell Precursor-ALL MRD tube
  - Primary Immunodeficiency Orientation tube

# PCR Overview

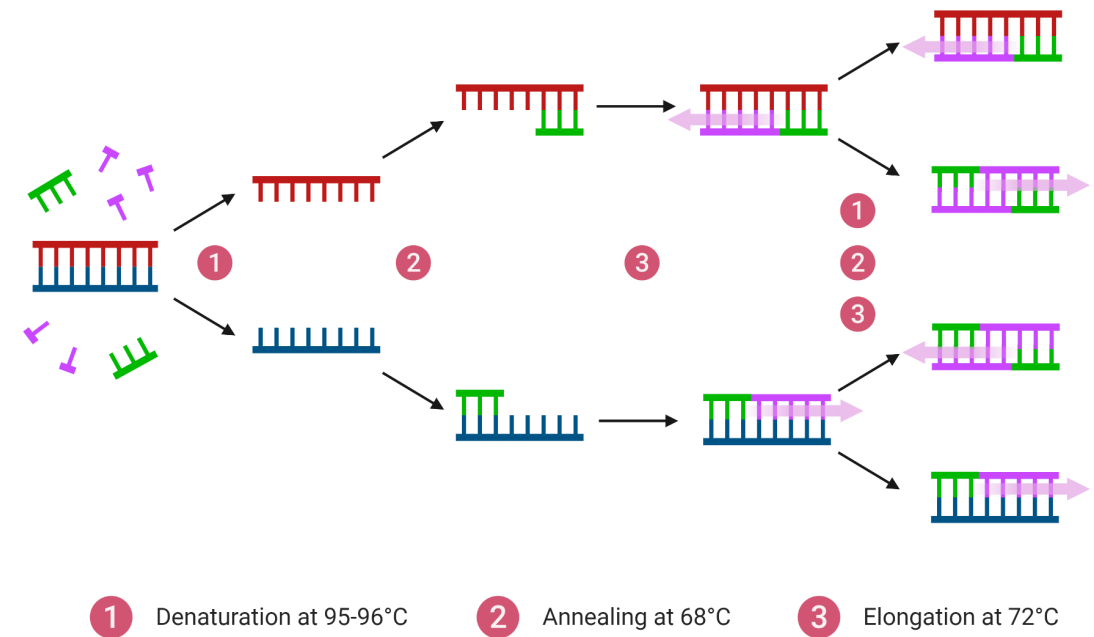
Technique used to make copies of a piece of DNA

Able to target gene rearrangements, fusion genes, and mutated genes

## Two PCR Techniques Used in MRD Testing

- Real-time quantitative PCR (RT-qPCR)
- Droplet Digital PCR (ddPCR)

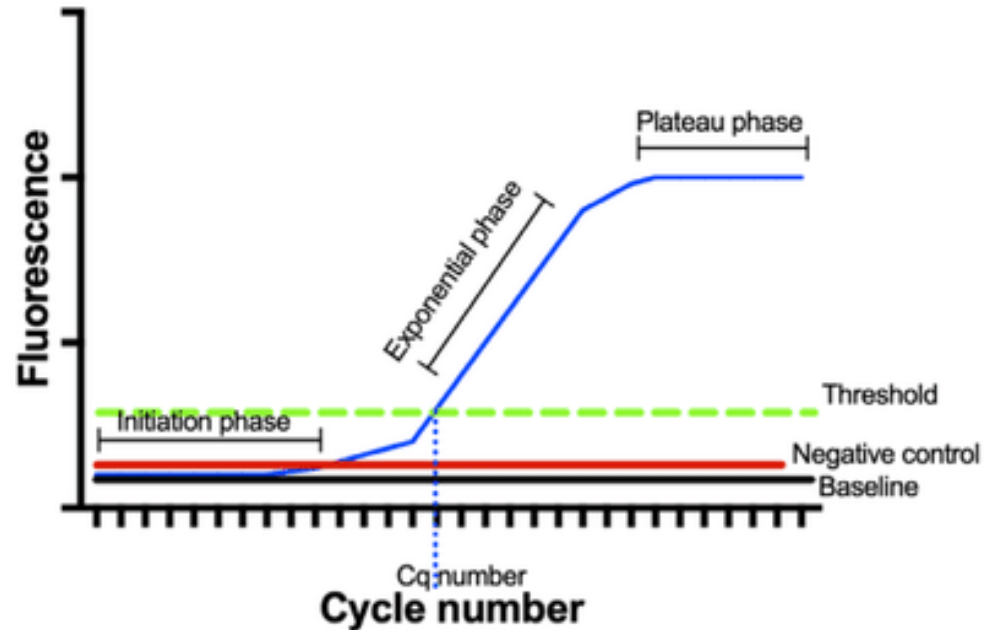
### Polymerase chain reaction - PCR



# Real-time quantitative PCR (RT-qPCR)

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Amplification curve



Variation of PCR where fluorescent probes quantify amplified product at the end of each cycle

Used to detect antigen-receptor gene rearrangements in lymphoid malignancies

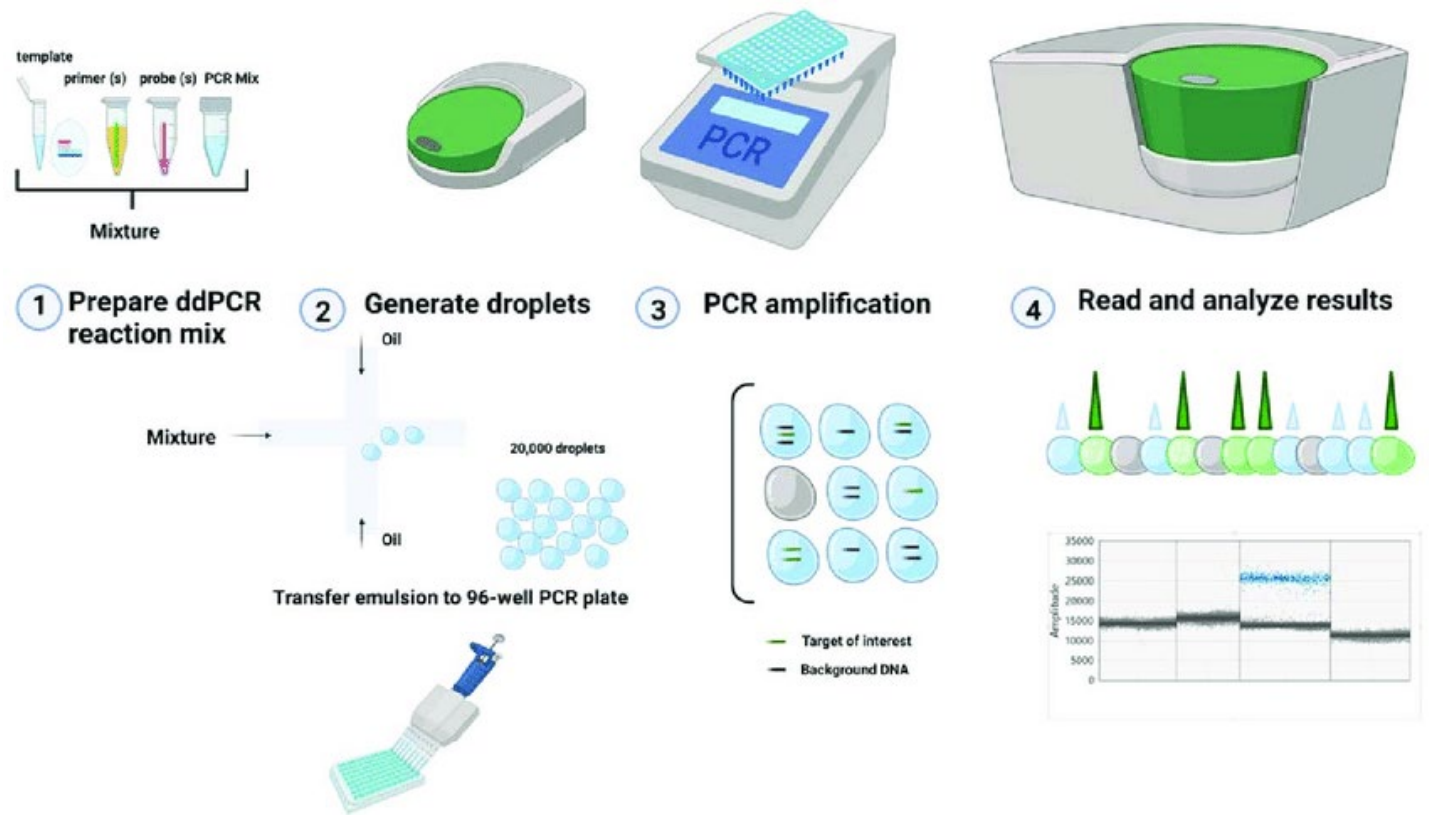
Used to detect fusion genes in AML and ALL such as BCR:ABL1 and ETV6:RUNX1

# Droplet Digital PCR (ddPCR)

Each PCR sample is partitioned into thousands of microscopic droplets before amplification

Each droplet is an individual PCR reaction

Fluorescence detects droplets containing the target sequence





# Droplet Digital PCR (ddPCR)

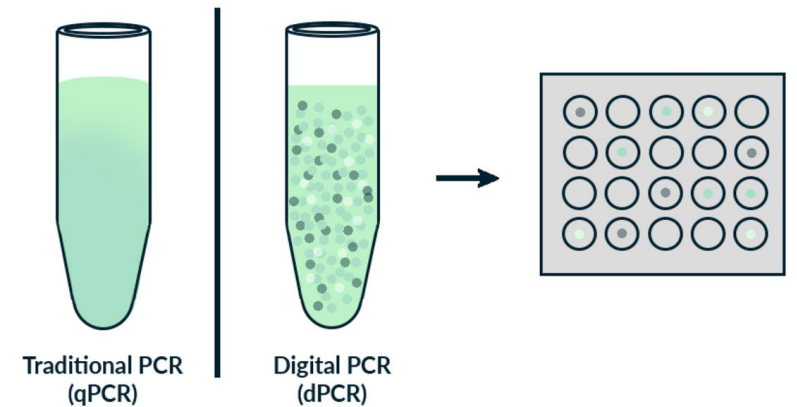
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## Advantages

- More accurate than RT-qPCR
- Higher amplification efficiency
- Less affected by PCR inhibitors
- Multiplexing capable
- No calibration curve needed – absolute quantification
- Able to quantify low concentration targets

## Disadvantages

- Expensive
- Not as efficient for large scale screening of numerous samples/targets
- Sample preparation and data analysis are complex
- Needs standardization





# Next-Generation Sequencing (NGS) Overview

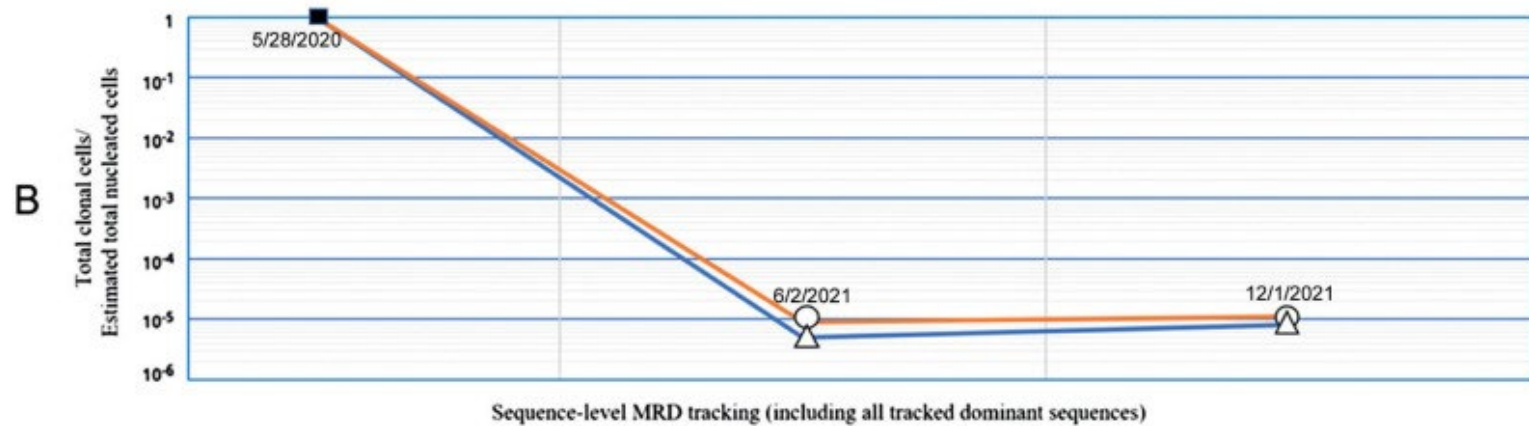
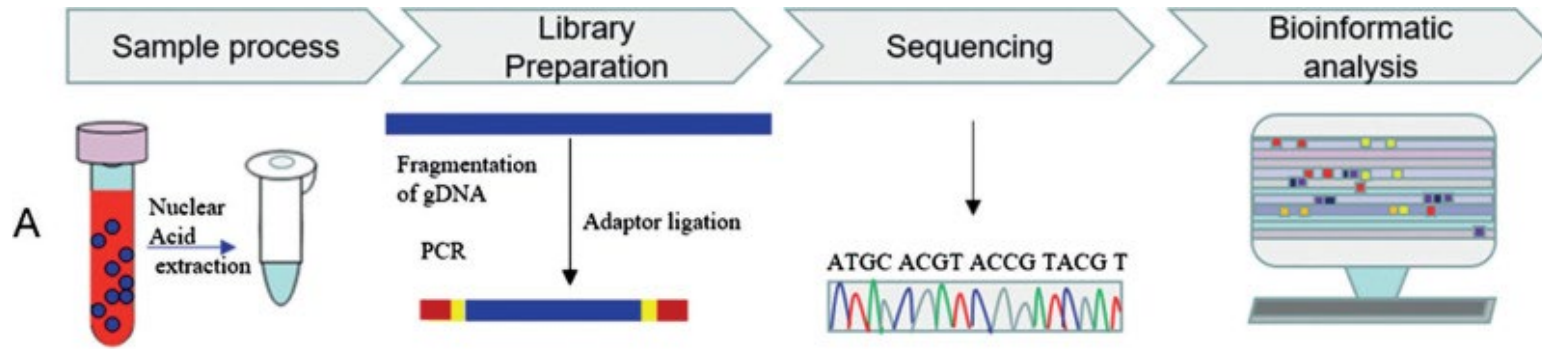
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Technique that read and analyze millions of DNA pieces at once

It works by breaking DNA into smaller fragments, reading each piece, and then putting them back together to understand the full sequence

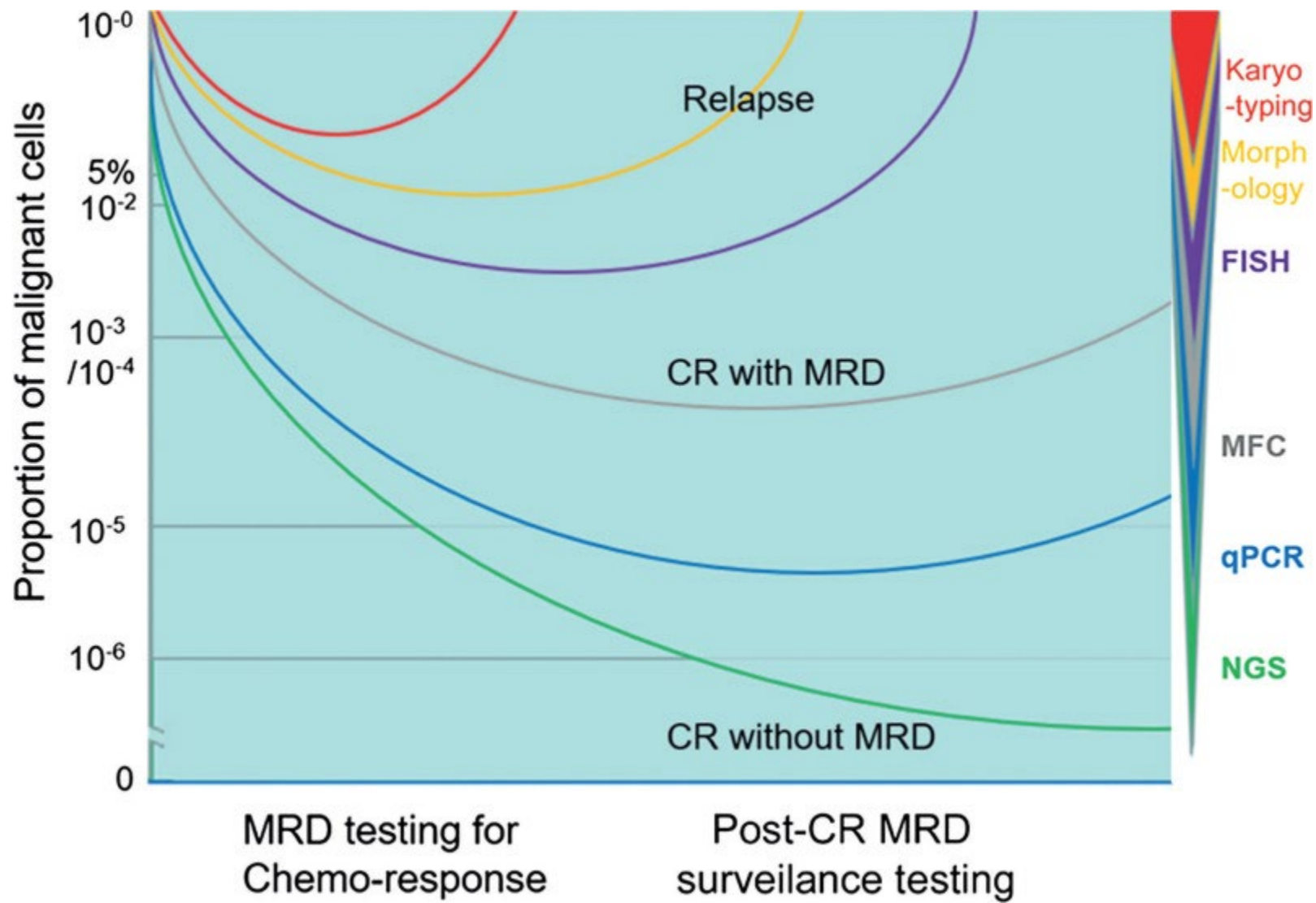
## **Role in MRD Testing**

- Detects very small amounts of cancer cells that remain after treatment
- Identifies genetic mutations and rare MRD markers often missed in other methods
- Monitor responses to treatment
- Treatment plans can be tailored to patient's specific genetic profile



# NGS

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# Comparative Analysis of MRD Detection Technologies

Method	Sensitivity	Advantages	Disadvantages
<b>Multiparameter Flow Cytometry using LAIP &amp; DfN</b>	$10^{-3}$ to $10^{-5}$	<ul style="list-style-type: none"> <li>Fast (within hours)</li> <li>High applicability</li> <li>Relatively inexpensive</li> <li>Personalized</li> <li>Detects phenotypic shifts</li> </ul>	<ul style="list-style-type: none"> <li>Requires fresh sample/viable cells</li> <li>High level of expertise needed</li> <li>Limited standardization</li> </ul>
<b>RT-qPCR for gene fusions</b>	$10^{-4}$ to $10^{-5}$	<ul style="list-style-type: none"> <li>Sensitive</li> <li>Relatively simple</li> <li>Standardized</li> </ul>	<ul style="list-style-type: none"> <li>Limited applicability</li> <li>Risk of cross contamination</li> <li>Can't detect clonal evolution</li> </ul>
<b>Digital PCR</b>	$10^{-3}$ to $10^{-5}$	<ul style="list-style-type: none"> <li>Sensitive</li> <li>Absolute quantification</li> <li>No standard curve needed</li> <li>Not affected by PCR inhibitors</li> </ul>	<ul style="list-style-type: none"> <li>Lacks standardization</li> <li>Can't detect clonal evolution</li> <li>May need patient specific primers</li> </ul>
<b>Next-Generation Sequencing</b>	$10^{-6}$	<ul style="list-style-type: none"> <li>Highly sensitive</li> <li>No need for patient specific primers</li> <li>Wide applicability</li> <li>Can sometimes track clonal evolution</li> </ul>	<ul style="list-style-type: none"> <li>Requires specimen pretreatment</li> <li>No standardization</li> <li>Requires high expertise</li> <li>Expensive</li> </ul>

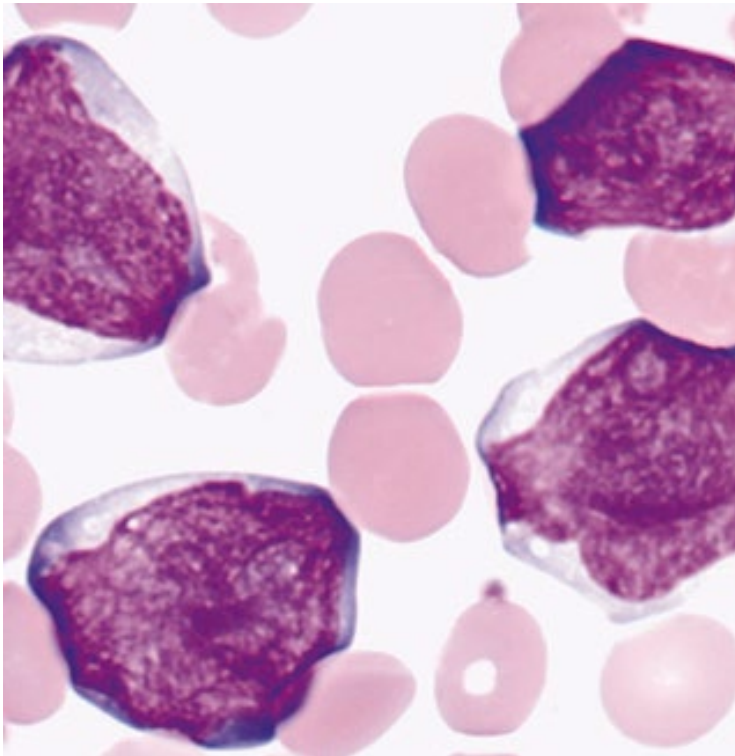


# Clinical Applications of MRD Testing

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# Acute Lymphoblastic Leukemia (ALL)

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Blood cancer with aggressive growth of immature lymphoblasts

Primarily affects children between ages 2 to 5

- Accounts for 25% of pediatric cancers and up to 75% of childhood leukemias

Rare in adults, but risk of ALL increases with age

Morphology

- Lymphoblasts appear in two forms: small with scant blue cytoplasm and indistinct nucleoli, or larger with prominent nucleoli and irregular nuclear membranes, which can resemble AML blasts



# ALL WHO

## Classification

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### B-cell ALL

- Several subtypes with known gene translocations
- **ETV6:RUNX1 fusion**
  - Common in children
  - Excellent prognosis
  - Formed from t(12;21)(p13;q22)
- **BCR:ABL1 fusion (Philadelphia chromosome)**
  - More common in adults
  - Poor prognosis; improved with the development of tyrosine kinase inhibitors
  - t(9;22)(q34;q11)

### T-cell ALL

- Classified into early T-precursor ALL and T-ALL not otherwise specified (NOS)

# MRD Testing in Acute Lymphoblastic Leukemia (ALL)

Detected using MFC, PCR, and NGS

Regular MRD checks during induction and maintenance chemotherapy (1.5 to 2.5 years) is standard

- First evaluated 2 to 4 weeks after induction therapy
- MRD negativity after first test indicates good response and less invasive treatments needed
- Recommended to test every 3 months to catch relapses early
- MFC & NGS most frequently used in B-ALL
- RT-PCR added for Ph+ ALL

# MRD Strong Predictor of ALL Relapse

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Large scale studies of over 13,000 ALL patients found that reaching MRD negative status improved chances of event-free survival regardless of the method used for MRD testing

MRD-negative patients had a 75% lower risk of relapse or death compared to those with detectable levels of MRD



Allogeneic hematopoietic cell transplantation is indicated for adults with relapsed or refractory ALL or patients with MRD positivity after induction therapy



Studies have demonstrated that patients with pre-transplant MRD levels of  $>10^{-4}$  have 7.7 times higher risk of relapse after transplantation



These patients need additional chemotherapy to reduce cancer loads before Allo HCT

# MRD Therapeutic Implications for ALL

# MRD Role in Monitoring New Therapies

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Blinatumomab: CD19/CD3 bispecific T-cell engager (BiTE) antibody

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Given to patients with refractory or relapsed Precursor B-ALL with poor prognosis

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Targets CD-19 positive B-cells to be destroyed by T-cells

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43% of patients with relapsed/refractory B-precursor ALL achieved complete remission after 2 cycles of treatment

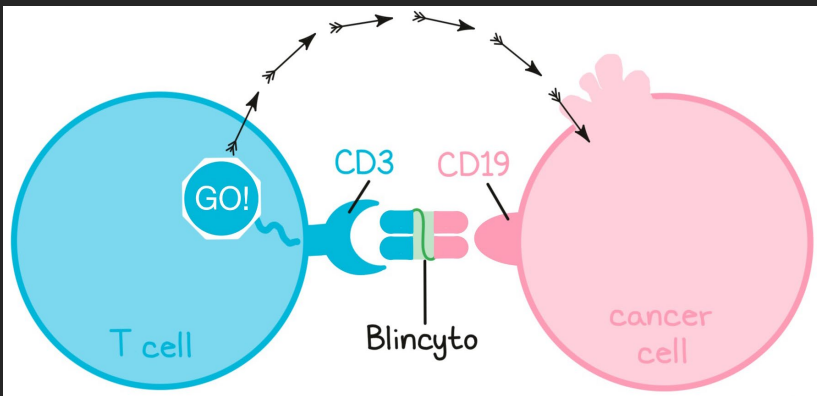
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This increased the median overall survival from 4.5 months with chemo alone to 13 months for patients who took BiTE

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MRD negativity showed strong correlation to long-term survival and improved T-cell expansion

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# Multiple Myeloma (MM)

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**C**alcium  
**R**enal complications  
**A**nemia  
**B**one pain



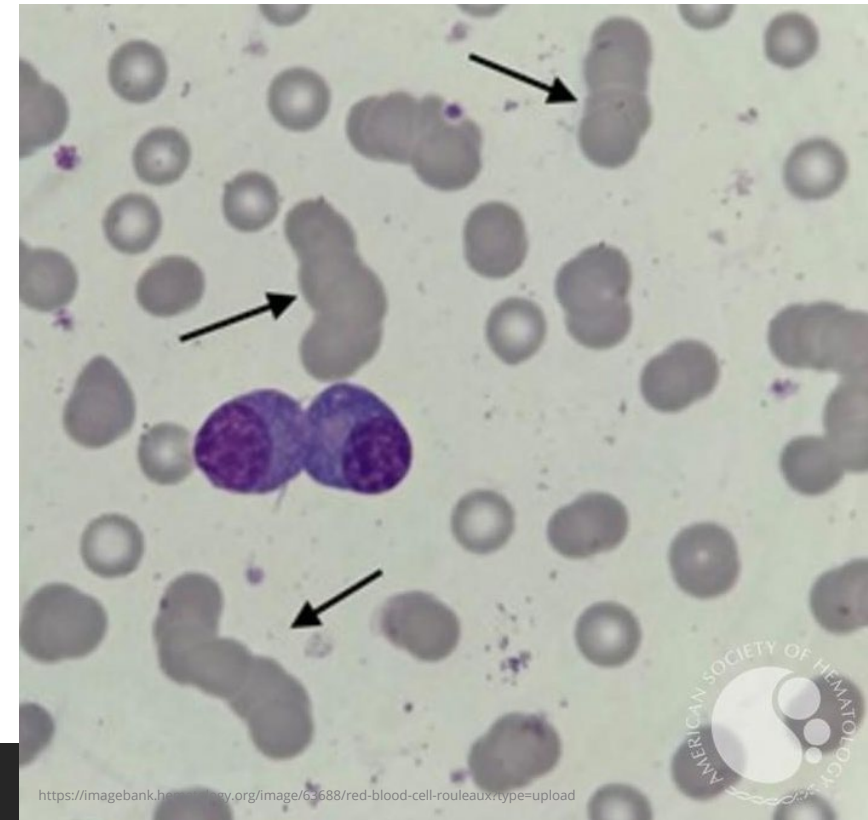
Accumulation of malignant plasma cells in the bone marrow that create lytic bone lesions

Primarily affects middle-aged to elderly

Symptoms: hypercalcemia, bone pain, anemia, kidney problems and frequent infections

## Laboratory Tests

- CBC (anemia) & Chem panel (high calcium)
- SPE – monoclonal gammopathy (usually IgG)
- Bence Jones protein in urine
- Hyperviscosity to plasma





# MRD Testing in Multiple Myeloma (MM)

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**FDA approved MRD testing as clinical endpoint for the accelerated approval of new drugs therapies in MM**



**Replaces overall response rates; which typically took 10 years of research to see real lasting benefit for patients**



**While the average survival has increased from 3.5 years to 10 years, 60% of MM patients will survive 5 years after diagnosis**



**MRD testing has the potential to speed up drug discovery and improve patient outcomes with efficient, sensitive, and personalized detection**

# Future Questions in MRD Detection

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Who should be evaluated?

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What is the optimal time for MRD evaluation?

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What are the precise thresholds for MRD testing that correlate with clinically significant outcomes?

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Does the clinical benefit outweigh testing costs?

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Which malignancies benefit the most from MRD testing?

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What role does MRD testing have in liquid biopsy testing for circulating tumor DNA and circulating tumor cells?

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Are MRD positive cells truly dormant or potentially capable of reactivating disease?

# Summary

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Minimal residual disease testing plays a crucial role in detecting small numbers of cancer cells that remain after treatment, which may not be visible through routine clinical tests

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Techniques, such as multiparameter flow cytometry, PCR, and next-generation sequencing, allow for early relapse detection

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MRD testing guides treatment decision, predicts relapse, and enables timely interventions

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In certain hematological malignancies, such as ALL and MM, MRD testing is a marker for treatment response and survival outcomes

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The FDA now recognizes MRD negativity as a key endpoint for drug approvals

# References

1. Advancements in machine learning (ML): Transforming the future of blood cancer detection and outcome prediction. (2024). Healthbook TIMES Oncology Hematology, (20). <https://doi.org/10.36000/hbt.oh.2024.20.146>
2. Alkan, S. S. (2004). Monoclonal antibodies: The story of a discovery that revolutionized science and medicine. *Nature Reviews Immunology*, 4(2), 153–156. <https://doi.org/10.1038/nri1265>
3. Harmening, D. (2024). *Clinical Hematology and Fundamentals of Hemostasis*. F. A. Davis Company. <https://bookshelf.vitalsource.com/books/9781719653138>
4. Keohane, E. M., Butina, M., Mirza, K. M., & Walenga, J. M. ([Insert Year of Publication]). *Rodak's Hematology* (7th ed.). Elsevier Health Sciences (US). <https://bookshelf.vitalsource.com/books/9780323937634>
5. Lambrescu, I., Popa, A., Manole, E., Ceafalan, L. C., & Gaina, G. (2022). Application of droplet digital PCR technology in muscular dystrophies research. *International Journal of Molecular Sciences*, 23(9), 4802. <https://doi.org/10.3390/ijms23094802>
6. Li, W. (2022, October 16). Measurable residual disease testing in acute leukemia: Technology and clinical significance. *Leukemia* [Internet]. <https://www.ncbi.nlm.nih.gov/books/NBK586210/>
7. Measurable residual disease (MRD). Measurable Residual Disease (MRD) | Leukemia and Lymphoma Society. (n.d.). <https://www.lls.org/booklet/measurable-residual-disease-mrd>

# References

8. Ngai, L. L., Hanekamp, D., Kelder, A., Scholten, W., Carbaat-Ham, J., Fayed, M. M. H. E., Snel, A. N., Bachas, C., Tettero, J. M., Mocking, T. R., Van Der Velden, V. H. J., Slomp, J., Hobo, W., Breems, D., Fischer, T., Gjertsen, B. T., Griškevičius, L., Juliusson, G., Maertens, J., ... Cloos, J. (2023). The LAIP-based-dfn approach is superior in terms of useful MRD results as compared to the LAIP approach after Cycle II in acute myeloid leukemia. *Blood*, 142(Supplement 1), 1572–1572. <https://doi.org/10.1182/blood-2023-181709>
9. Olaniyi. (2011). Flow cytometric immunophenotyping of hematological malignancies: The way forward in nigeria. *Pathology and Laboratory Medicine International*, 17. <https://doi.org/10.2147/plmi.s20709>
10. Medina-Herrera, A., Sarasquete, M. E., Jiménez, C., Puig, N., & García-Sanz, R. (2023). Minimal residual disease in multiple myeloma: Past, present, and future. *Cancers*, 15(14), 3687. <https://doi.org/10.3390/cancers15143687>
11. Riva, G., Nasillo, V., Ottomano, A. M., Bergonzini, G., Paolini, A., Forghieri, F., Lusenti, B., Barozzi, P., Lagreca, I., Fiorcari, S., Martinelli, S., Maffei, R., Marasca, R., Potenza, L., Comoli, P., Manfredini, R., Tagliafico, E., Trenti, T., & Luppi, M. (2021). Multiparametric flow cytometry for MRD monitoring in hematologic malignancies: Clinical applications and new challenges. *Cancers*, 13(18), 4582. <https://doi.org/10.3390/cancers13184582>
12. Roberts K. G. (2018). Genetics and prognosis of ALL in children vs adults. *Hematology. American Society of Hematology. Education Program*, 2018(1), 137–145. <https://doi.org/10.1182/asheducation-2018.1.137>
13. Röhnert, M. A., Kramer, M., Schadt, J., Ensel, P., Thiede, C., Krause, S. W., Bücklein, V., Hoffmann, J., Jaramillo, S., Schlenk, R. F., Röllig, C., Bornhäuser, M., McCarthy, N., Freeman, S., Oelschlägel, U., & von Bonin, M. (2022). Reproducible measurable residual disease detection by multiparametric flow cytometry in acute myeloid leukemia. *Leukemia*, 36(9), 2208–2217. <https://doi.org/10.1038/s41375-022-01647-5>
14. Salama, M. E., Otteson, G. E., Camp, J. J., Seheult, J. N., Jevremovic, D., Holmes, D. R., 3rd, Olteanu, H., & Shi, M. (2022). Artificial Intelligence Enhances Diagnostic Flow Cytometry Workflow in the Detection of Minimal Residual Disease of Chronic Lymphocytic Leukemia. *Cancers*, 14(10), 2537. <https://doi.org/10.3390/cancers14102537>

# References

15. Saygin, C., Cannova, J., Stock, W., & Muffly, L. (2022). Measurable residual disease in acute lymphoblastic leukemia: Methods and clinical context in adult patients. *Haematologica*, 107(12), 2783–2793. <https://doi.org/10.3324/haematol.2022.280638>
16. Thursday, A. 8. (2024, August 8). A new endpoint for accelerated approvals in multiple myeloma. Memorial Sloan Kettering Cancer Center. <https://www.mskcc.org/clinical-updates/new-endpoint-for-accelerated-approvals-in-multiple-myeloma>
17. Turgeon, M. L. (2020). *Clinical Hematology: Theory & Procedures, Enhanced Edition* (6th ed.). Jones & Bartlett Learning. <https://bookshelf.vitalsource.com/books/9781284348996>
18. Vuelta, E., García-Tuñón, I., Hernández-Carabias, P., Méndez, L., & Sánchez-Martín, M. (2021). Future approaches for treating chronic myeloid leukemia: CRISPR therapy. *Biology*, 10(2), 118. <https://doi.org/10.3390/biology10020118>
19. Wijnands, C., Noori, S., Donk, N. W. C. J. van de, VanDuijn, M. M., & Jacobs, J. F. M. (2023). Advances in minimal residual disease monitoring in multiple myeloma. *Critical Reviews in Clinical Laboratory Sciences*, 60(7), 518–534. <https://doi.org/10.1080/10408363.2023.2209652>
20. Yu, Z., Xie, L., Zhang, J., Lin, H., & Niu, T. (2023, July 18). *The evolution of minimal residual disease: Key insights based on a bibliometric visualization analysis from 2002 to 2022*. *Frontiers in oncology*. <https://pmc.ncbi.nlm.nih.gov/articles/PMC10391156/>
21. Zugmaier, G., Gökbuget, N., Klinger, M., Viardot, A., Stelljes, M., Neumann, S., Horst, H.-A., Marks, R., Faul, C., Diedrich, H., Reichle, A., Brüggemann, M., Holland, C., Schmidt, M., Einsele, H., Bargou, R. C., & Topp, M. S. (2015). Long-term survival and T-cell kinetics in relapsed/refractory all patients who achieved MRD response after Blinatumomab treatment. *Blood*, 126(24), 2578–2584. <https://doi.org/10.1182/blood-2015-06-649111>



Questions?

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