

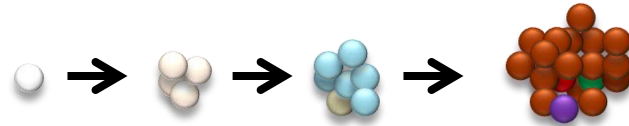
# Introduction to B- and T-cell Clonality Testing & Targets

# What is Clonality?



## Clonality

- A proliferation of cells originating from a single progenitor cell, producing a **pool of identical clonal cells**.



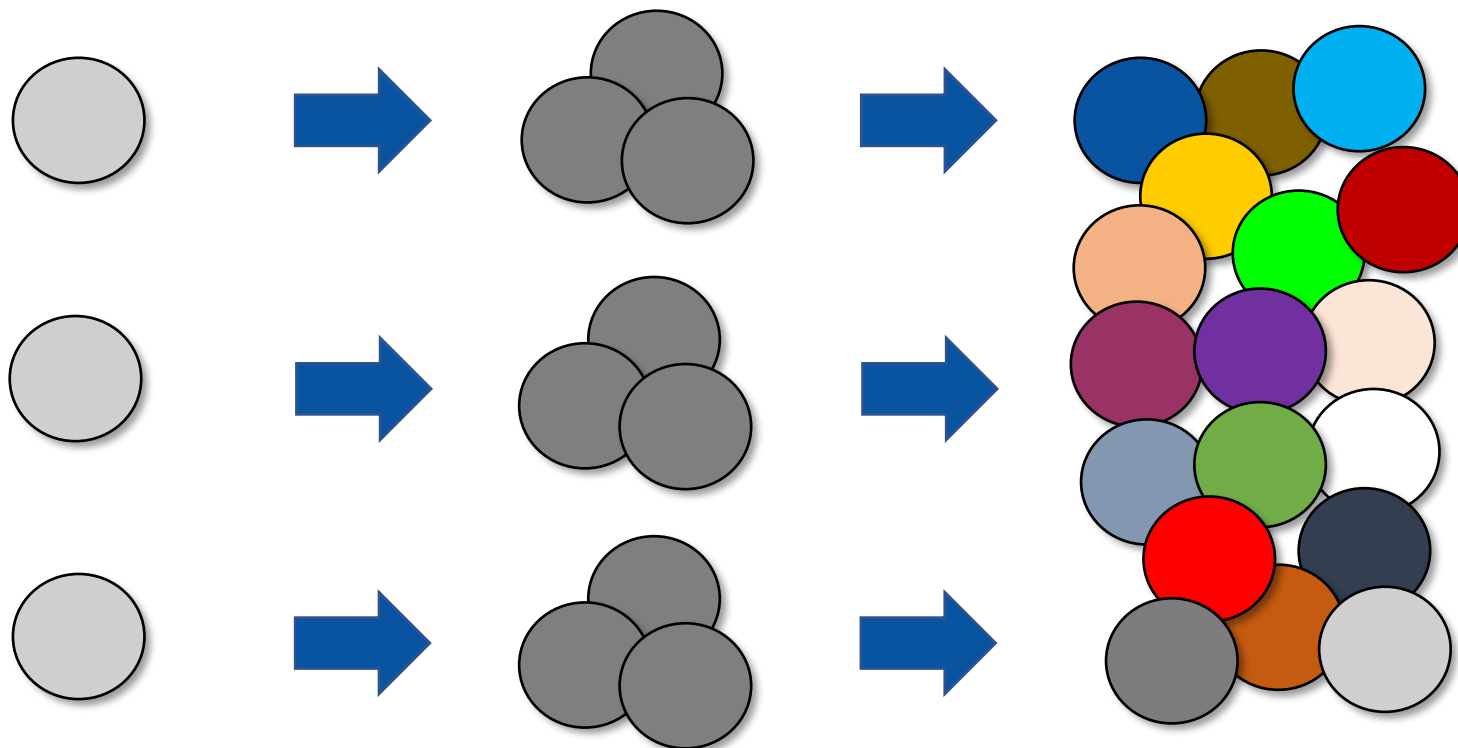
- Clonality testing using molecular techniques is used to **confirm the presence of leukemia or lymphoma**.

**Highly indicative of B- or T-cell malignancy**

# What is Clonality?



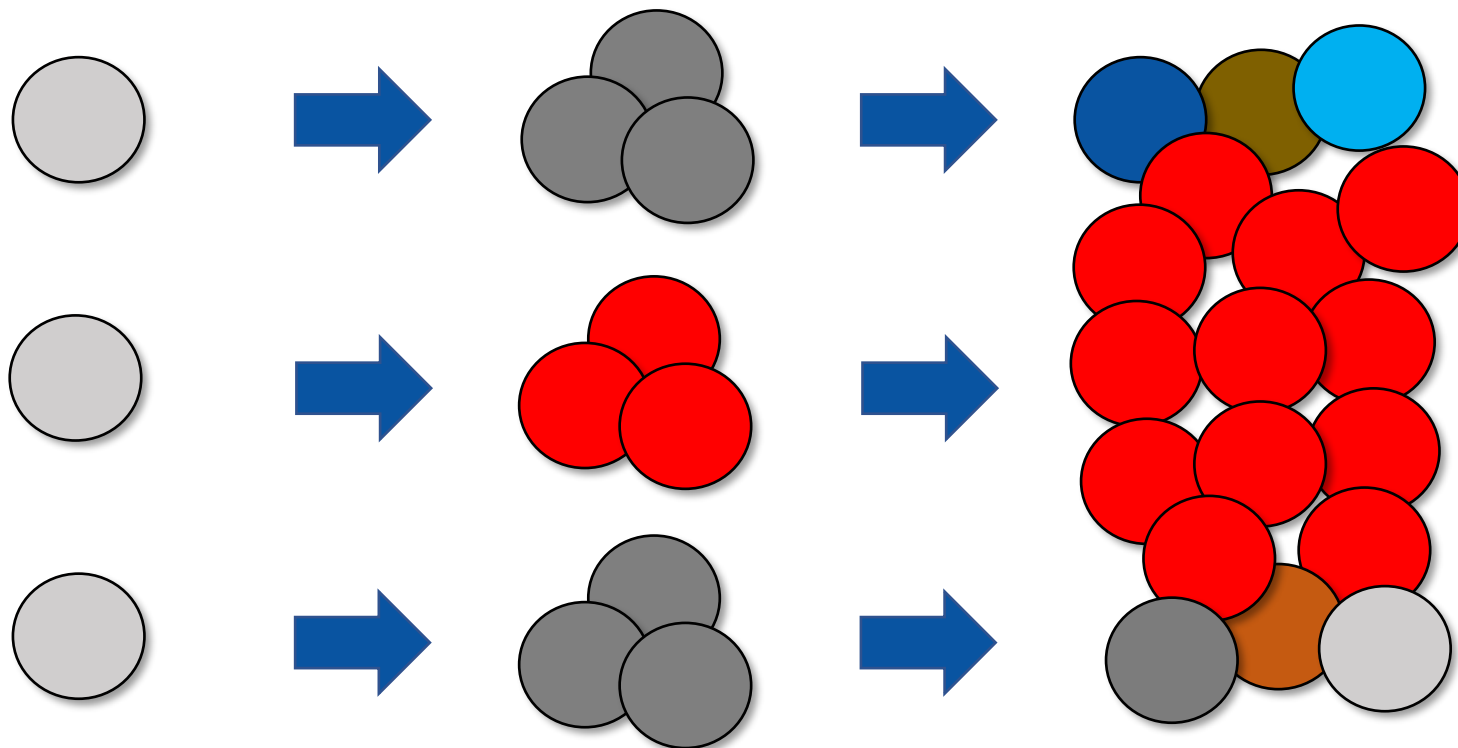
## Polyclonal Progression



# What is Clonality?



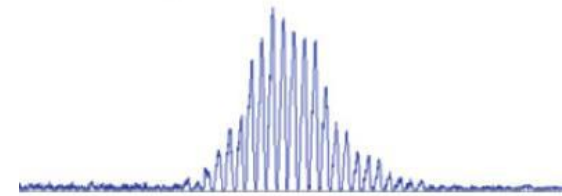
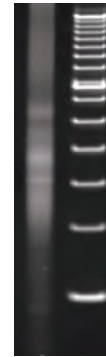
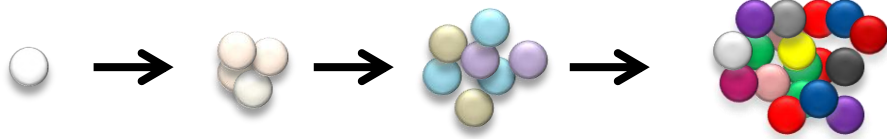
## Clonal Progression



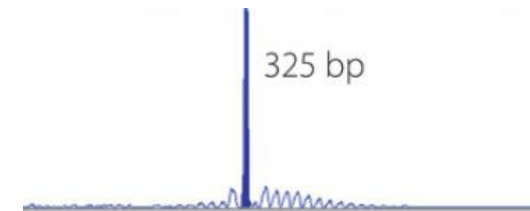
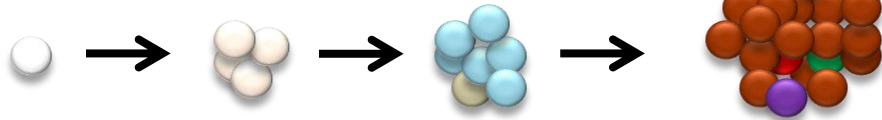
# What is Clonality?



- Polyclonal Progression



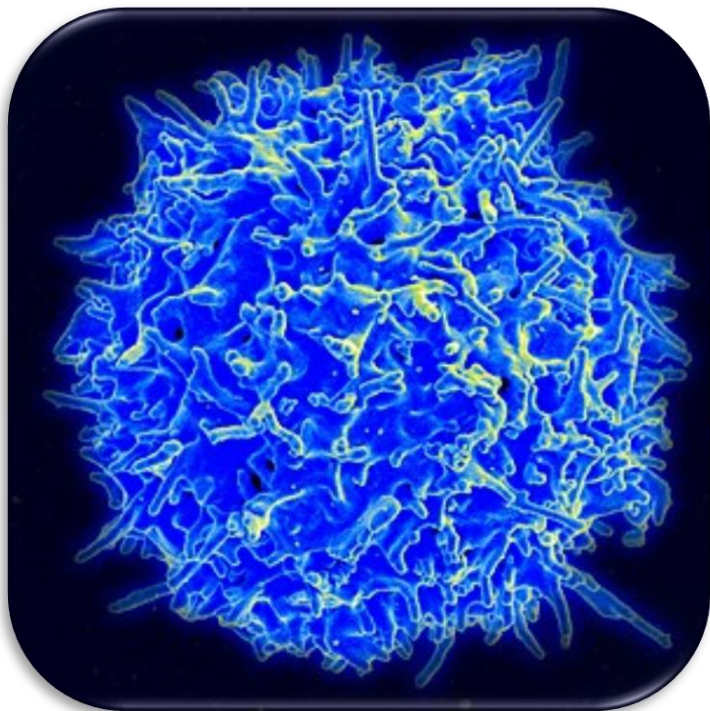
- Clonal Progression



Highly Indicative of B- or T-Cell Malignancy



# Why test for B- and T-cell clonality?



Leukemias and lymphomas can be challenging to diagnose by

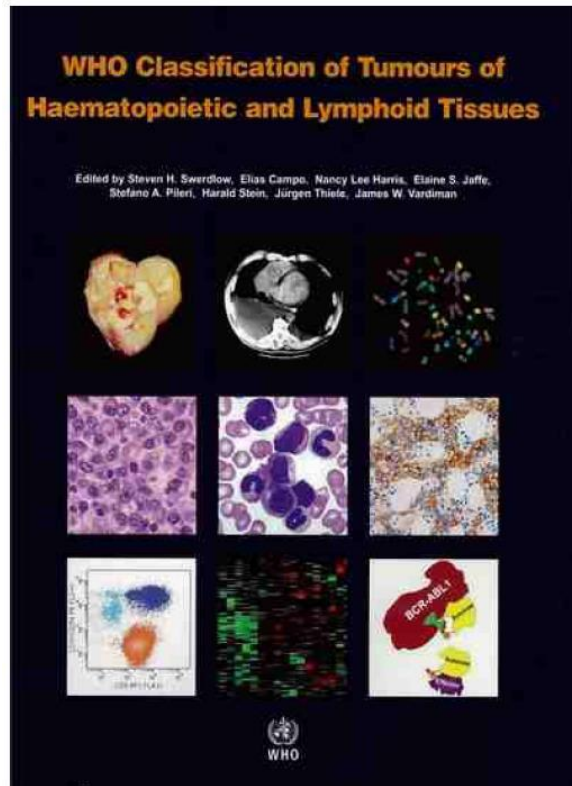
- morphology
- immunohistochemistry
- flow cytometry

**5-15%** of above cases result in inconclusive diagnoses

Diagnosis of lymphoid malignancies is greatly supported and facilitated by **clonality testing**

**Adopted in routine diagnostics for further MRD testing**

# Why test for B- and T-cell clonality?



We are facing a growing number of hematopoietic tumors with specific or characteristic molecular changes.

Diagnosis of lymphoid malignancies is greatly supported and facilitated by **clonality testing**

# Why test for B- and T-cell clonality?



Monoclonality is a dominant characteristic of cancer

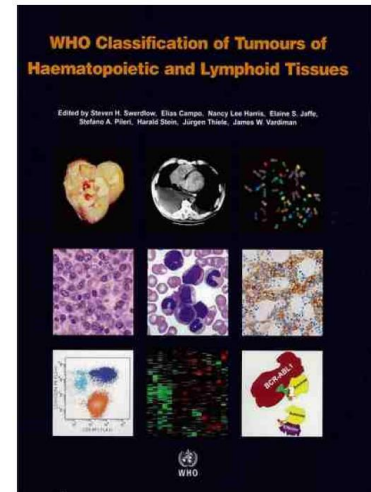
Allows for discrimination between :

- Reactive lesions (**polyclonal**) - generally considered **benign**
- Hematologic malignancies (**clonal**) - generally considered **neoplastic**

Aids in diagnosis:

- of Minimal tumor infiltration
- on limited diagnostic tissue when the architecture is not evaluable
- of neoplastic proliferations without specific cytological, histological or immunohistochemical criteria

Helps to identify tumor-specific markers for post-treatment monitoring

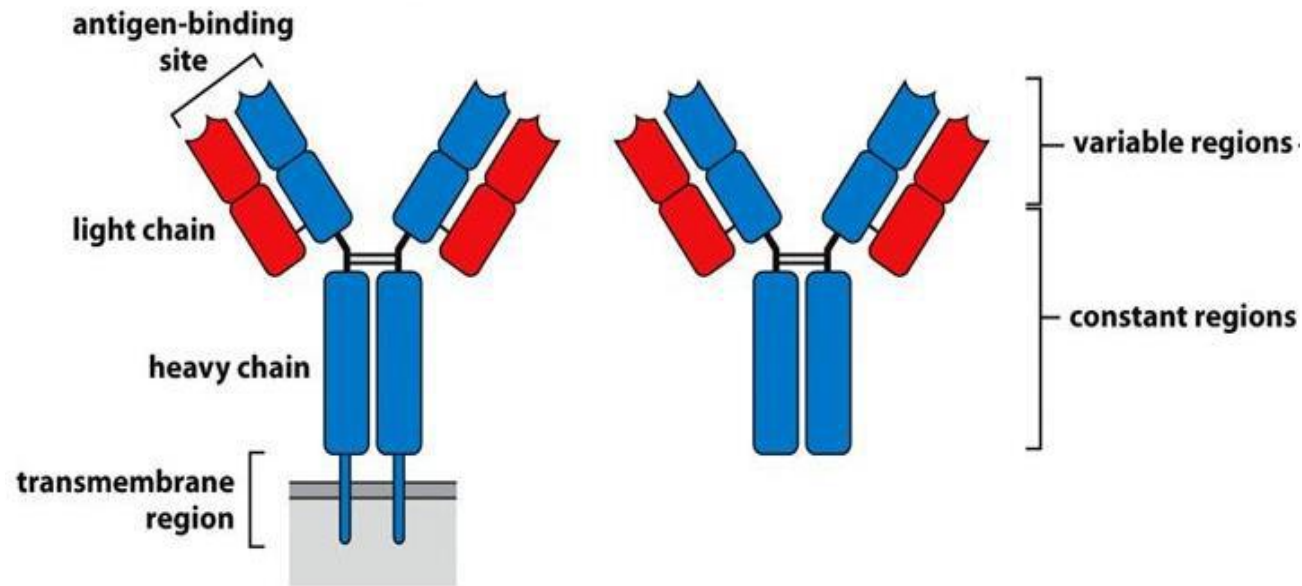




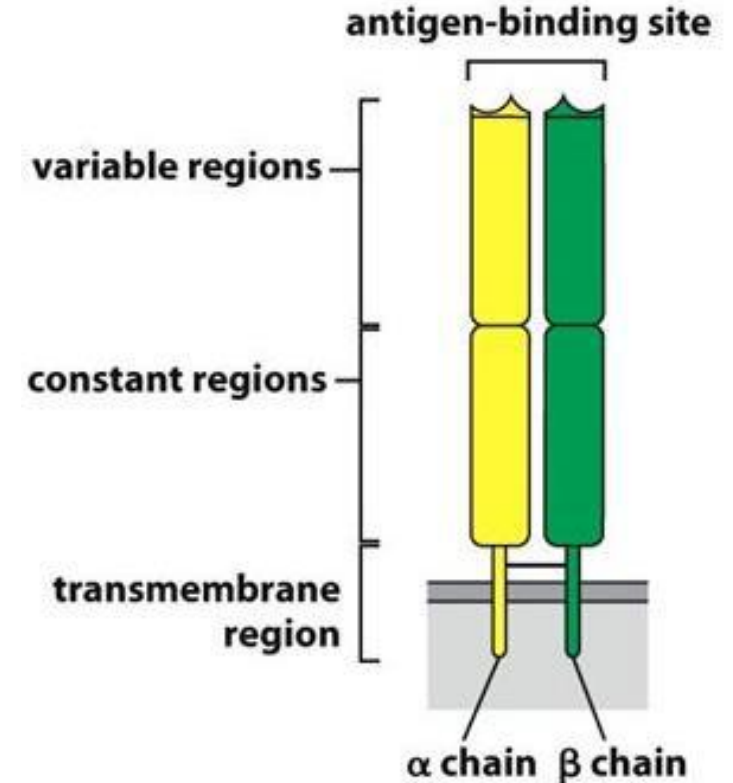
# Clonality Targets



## B-Cell Receptors (BCR)/Immunoglobulins (Ig)



## T-Cell Receptor (TCR)



**Each Lymphocyte has a Unique Antigen Receptor**

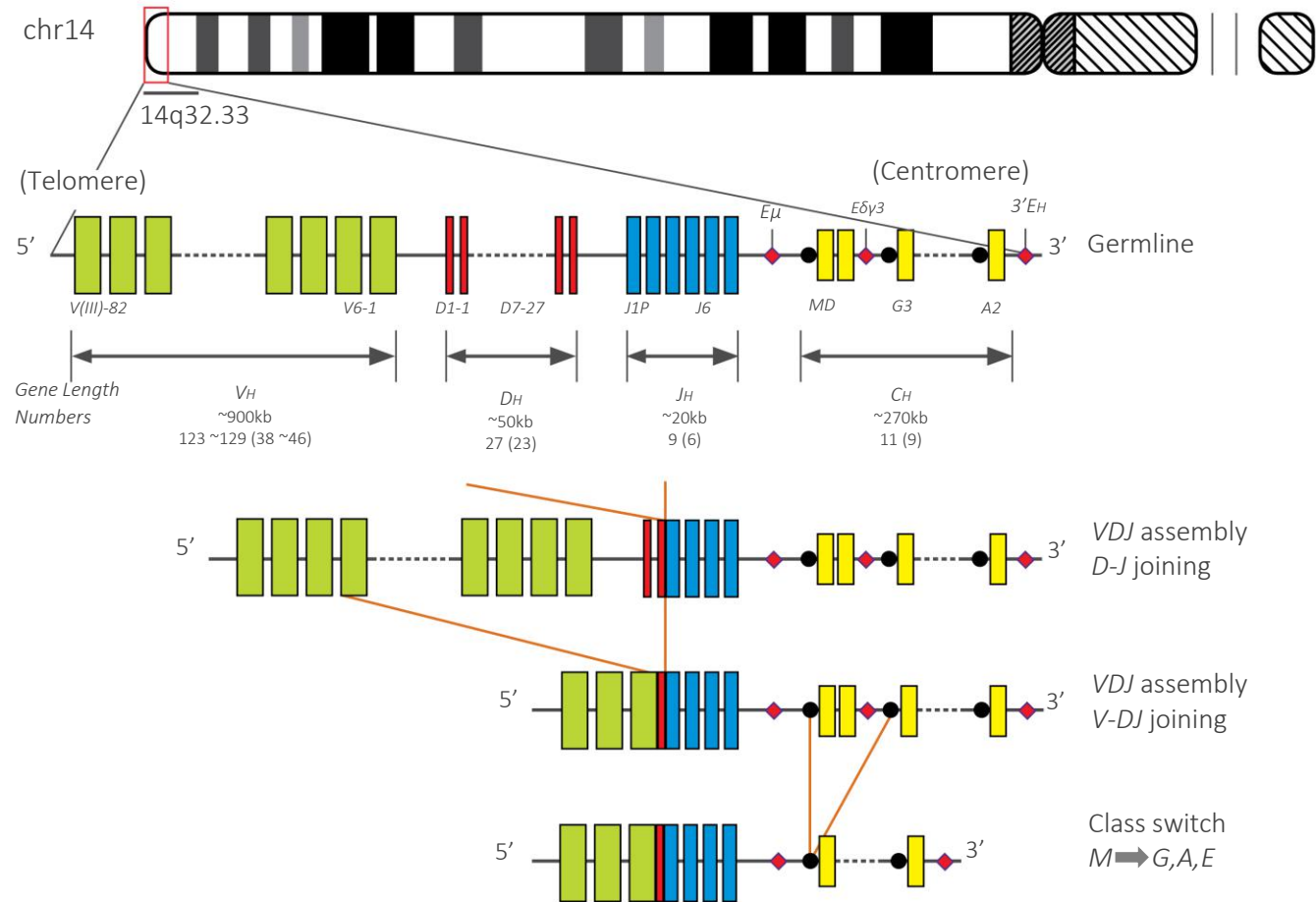
Magnitude  $\sim 10^{12}$  (i.e., 1,000,000,000,000) different Ig or TCR molecules



## Antigen Receptor Molecules (AgRs) are the Molecular Targets

- **B- and T- Cell Receptors**
  - *IGH, IGK, IGL* (B-Cell Receptors)
  - *TRG, TRB, TRD* (T-Cell Receptors)
- **Each lymphocyte has a unique AgR** (single specificity)

# Human *IGH* Gene Locus

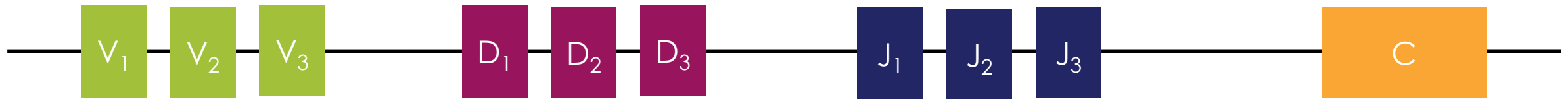


Dyer, M. et al., *Blood* 115:1490-1499 (2010).

# Generation of Diversity



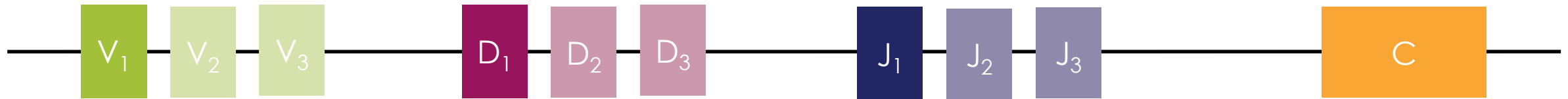
## Rearranged V-D-J gene :



# Generation of Diversity



## Rearranged V-D-J gene :

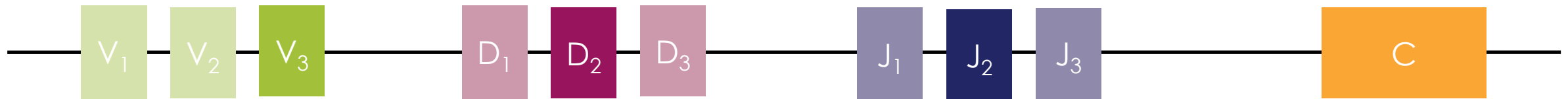




# Generation of Diversity



## Rearranged V-D-J gene :



B-cell "X"



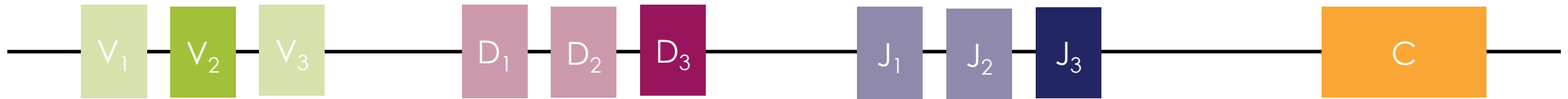
B-cell "Y"



# Generation of Diversity



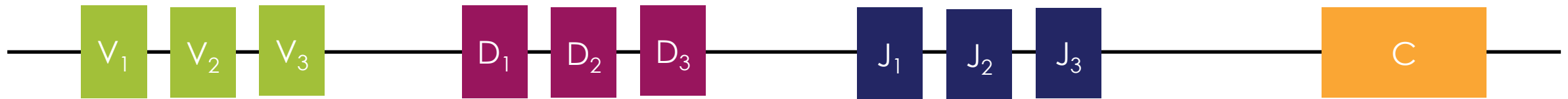
## Rearranged V-D-J gene :



# Generation of Diversity



Rearranged V-D-J gene :



B-cell "X"



B-cell "Y"



B-cell "Z"



Unique sequences

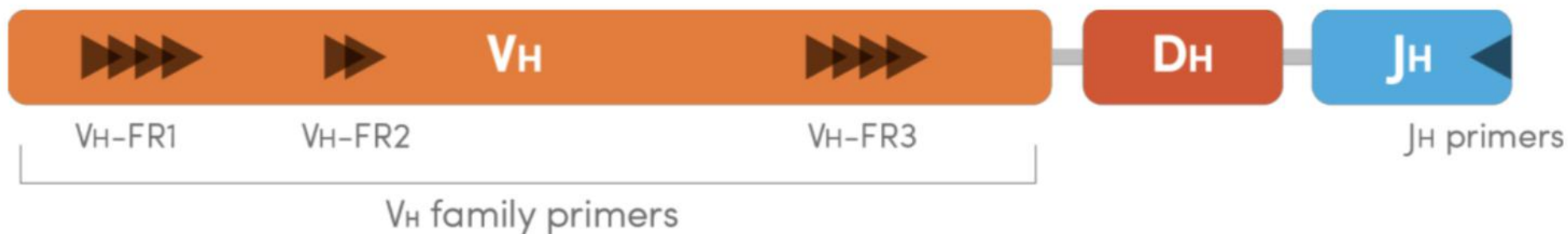


Unique fingerprints

# B- and T-cell Clonality Simplified



*IGH* Locus (14q32.33)



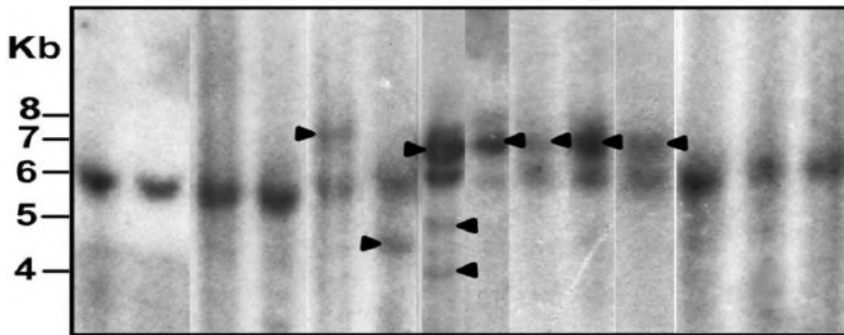
*TRG* Locus (7p14)





## Clonality Testing was originally performed by Southern Blots

- Labor intensive
- Restricted repertoire
- High DNA quality and quantity required
- Time consuming
- Moderate limits of detection
- Subjective interpretation



**Southern Blot**

**Fragment Analysis / CE**

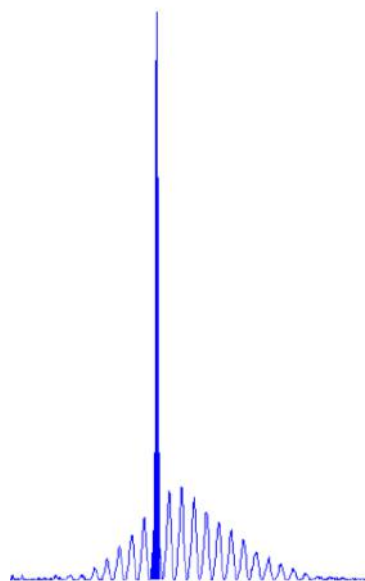
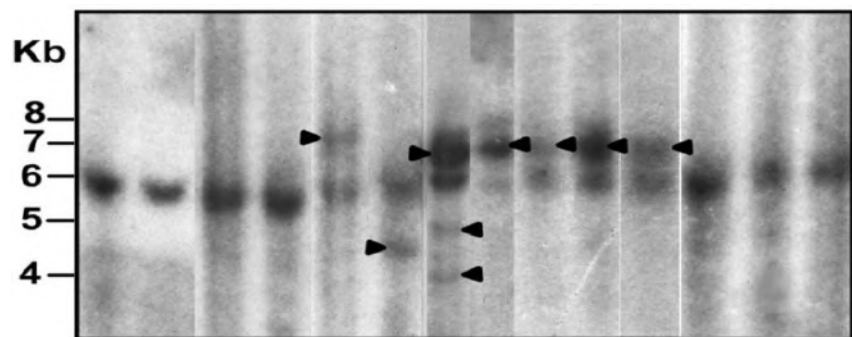


# Evolution of Clonality Testing



PCR-based clonality assays are now widely accepted as the gold standard, with improved :

- Sensitivity
- Testing time
- Coverage
- Limit of detection
- Sample type suitability

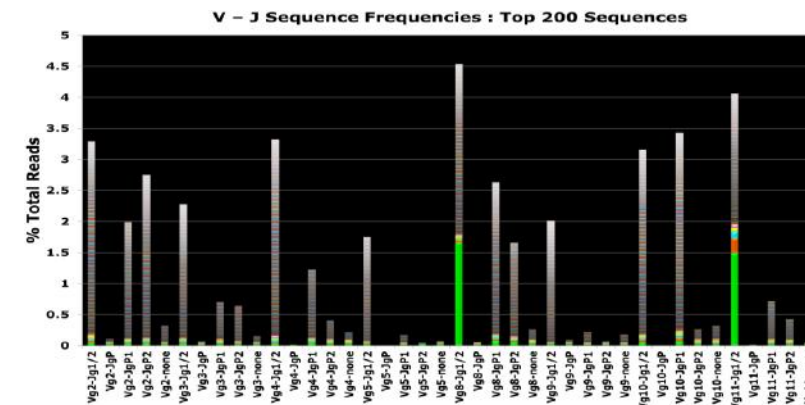
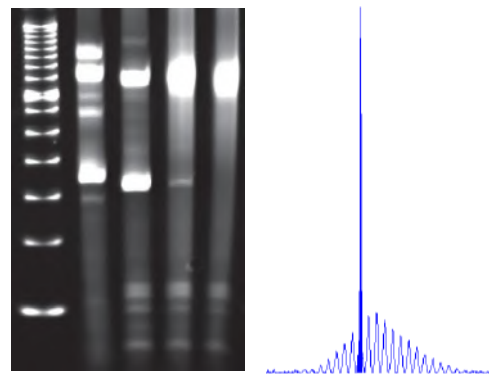
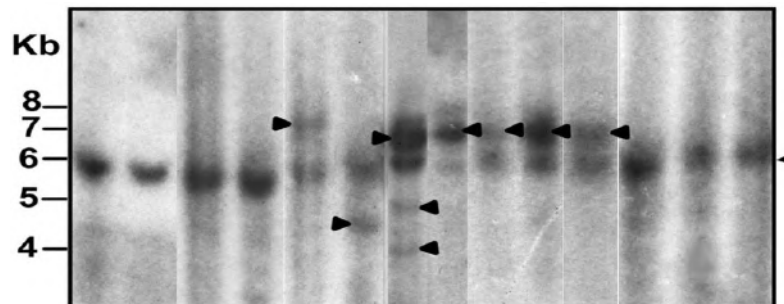


**Southern Blot**

**Fragment Analysis / CE**

**NGS**

# Evolution of Clonality Testing



- Labor intensive
- Time consuming
- Restricted repertoire
- Requires high DNA quantity & quality

- Increased sensitivity
- Reduced testing time
- Better coverage
- Lower limits of detection
- More sample types

- DNA sequence of clones
- Highest sensitivity
- Ability to track clones

**Southern Blot**

**Fragment Analysis / CE**

**NGS**

# Hematologic Malignancy Research & Testing Guide



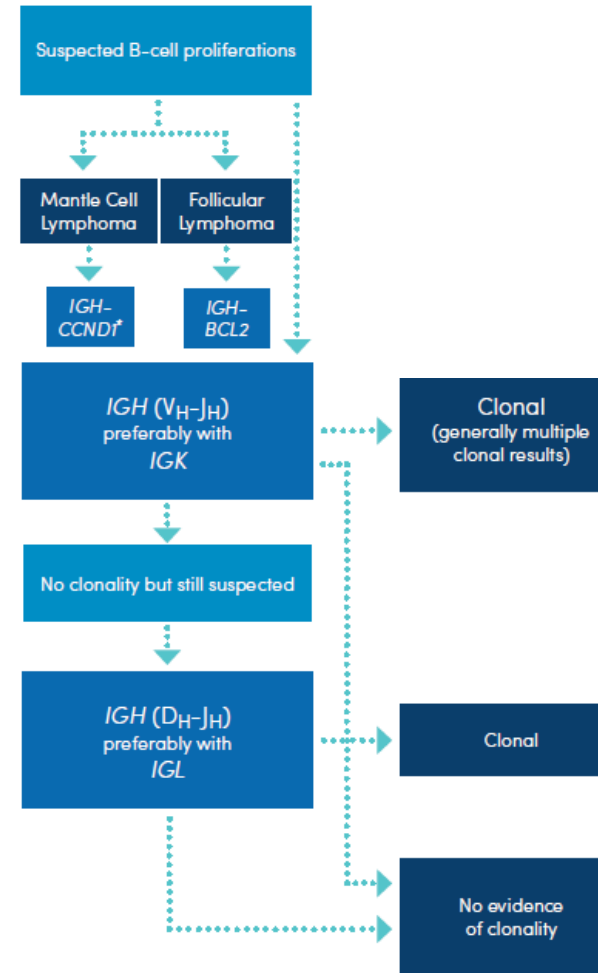
Disease	Gene Rearrangement									Translocation				Mutations	
	IGH (V <sub>H</sub> -J <sub>H</sub> )	IGH (D <sub>H</sub> -J <sub>H</sub> )	IGK	IGL	IGHV SHM	TRB	TRD	TRG	IGH-BCL1 (CCND1)	IGH-BCL2	BCR-ABL1*	PML-RARα*	FLT3	NPM1	
Lymphoid/ Lymphoma	Marginal Zone Lymphoma (MZL), extranodal <sup>12,13,27</sup>	88%	58%	84%	29%		23%	10%	16%						
	Marginal Zone Lymphoma (MZL), nodal <sup>13</sup>	100%	30%	80%	30%		10%	20%	10%						
	Mantle Cell Lymphoma (MCL) <sup>2,6,7,12,13,27,37</sup>	100%	11%	100%	44%	*	9%	4%	11%	75%					
	Follicular Lymphoma (FL) <sup>3,7,12,13,27,28</sup>	84%	19%	84%	21%		6%	5%	2%		90%				
	Diffuse Large B-cell Lymphoma (DLBCL) <sup>3,12,13,27</sup>	80%	30%	80%	28%		21%	14%	15%		30%				
	Multiple Myeloma (MM) and other Plasma Cell Neoplasms (PCN) <sup>2,9,10,20,25</sup>	84%	60%	57%	97%					20%					
	Chronic Lymphocytic Lymphoma (CLL) <sup>11,12,13,15,23,27,35</sup>	100%	43%	100%	30%	*	25%	12%	18%						
	B-cell Acute Lymphoblastic Leukemia (B-ALL) <sup>4,12,14,18,21,22,27,29,30,31,32,33,34</sup>	96%	57%	95%	20%		81%	86%	75%			30%			
	Suspect B-cell Proliferations <sup>12,26,27,33</sup>	93%	93%	90%	40%		20%		20%						
	Peripheral T-cell Lymphoma (PTCL) <sup>12,13,14,24</sup>	35%	4%		2%		98%		94%						
	T-cell Acute Lymphoblastic Leukemia (T-ALL) <sup>12,14,21,22,29,31</sup>	24%	25%	4%			92%	68%	95%						
	Angioimmunoblastic T-cell Lymphoma (AITL) <sup>12,13,14</sup>	19%	11%	30%	5%		99%	35%	92%						
	Adult T-cell Leukemia/Lymphoma <sup>39</sup>						97%		96%						
	Anaplastic Large-Cell Lymphoma (ALCL) <sup>12,13,14</sup>						74%	12%	74%						
	T-cell Prolymphocytic Leukemia (T-PLL) <sup>12,13,14</sup>	3%	3%	3%	3%		100%	6%	94%						
	T-cell Large Granular Lymphocytic Leukemia (T-LGL Leukemia) <sup>12,13,14</sup>			4%	4%		97%	29%	96%						
Suspect T-Cell Proliferations <sup>12,26,40</sup>	10%		10%			90%	11%	90%							
Myeloid	Acute Myeloid Leukemia (AML) <sup>6,16</sup>												33%	64%	
	Acute Promyelocytic Leukemia (APL) <sup>1,5,16,17</sup>											90%			
	Chronic Myeloid Leukemia (CML) <sup>7,10,19,21,38</sup>										87%				
	Myeloproliferative Neoplasms (MPNs) <sup>38</sup>										10%				

Note: The percentage of samples within a given disease category were detected using each gene target. Percentages indicate the highest referenced value.

# Clonality Testing Guide



- A test algorithm for suspect B-cell lymphoproliferations
- Developed in concert with the EuroClonality/BIOMED-2 group for PCR-based clonality assessment of suspected B-cell lymphoproliferative disorders



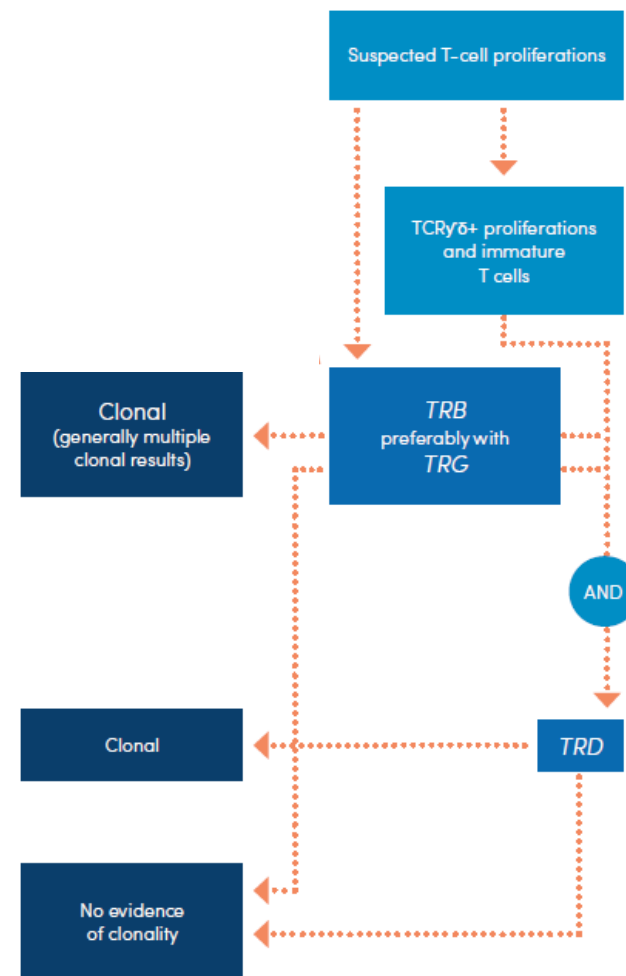
J.J.M. van Krieken *et al.*, *Leukemia* 2007 21: 201-206.

A.W. Langerak *et al.*, *Leukemia* 2012 26: 2159-71.

# Clonality Testing Guide



- A test algorithm for suspect T-cell lymphoproliferations
- Developed in concert with the EuroClonality/BIOMED-2 group for PCR-based clonality assessment of suspected T-cell lymphoproliferative disorders



J.J.M. van Krieken *et al.*, *Leukemia* 2007 21: 201-206.

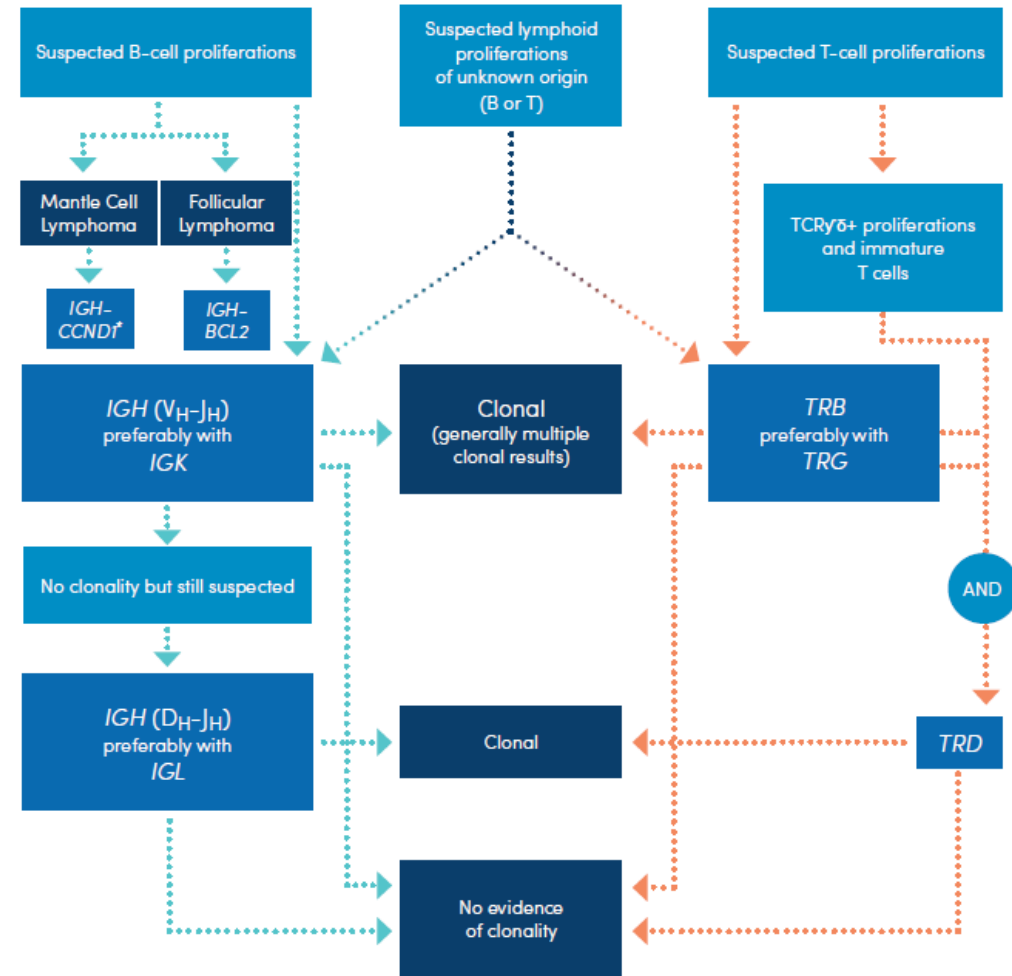
A.W. Langerak *et al.*, *Leukemia* 2012 26: 2159-71.



# Clonality Testing Guide



- A test algorithm for suspect B- and T-cell lymphoproliferations
- Developed in concert with the EuroClonality/BIOMED-2 group for PCR-based clonality assessment of suspected B- and T-cell lymphoproliferative disorders



J.J.M. van Krieken *et al.*, *Leukemia* 2007 21: 201-206.

A.W. Langerak *et al.*, *Leukemia* 2012 26: 2159-71.

# Multiplexing Targets



## Why combine testing of *IGH* V-J & *IGK*?

The majority of mature B-cell malignancies can be identified by targeting three *IGH* ( $V_H$ - $J_H$ ) frameworks\*

	<i>IGH</i> FR1 ( $V_H$ - $J_H$ )	<i>IGH</i> FR2 ( $V_H$ - $J_H$ )	<i>IGH</i> FR3 ( $V_H$ - $J_H$ )	<i>IGH</i> (FR 1, 2 & 3)
MCL (n=54)	100%	98%	96%	100%
B-CLL/SLL (n=56)	95%	91%	93%	100%
FL (n=109)	73%	76%	52%	84%
MZL (n=41)	73%	85%	68%	87%
DLBCL (n=109)	68%	61%	50%	79%
Total (n=369)	79%	78%	66%	88%

Abbreviations: B-CLL, B-cell chronic lymphocytic leukemia; DLBCL, diffuse large B-cell lymphoma; MZL, marginal zone B-cell lymphoma.

\*PA Evans et al. Leukemia. 2007 21:207-214

# Multiplexing Targets



## Why combine testing of *IGH* V-J & *IGK*?

The majority of mature B-cell malignancies can be identified by targeting three *IGH* ( $V_H$ - $J_H$ ) frameworks and two *IGK* ( $V_k$ - $J_k$  and Kde) rearrangements\*

	<i>IGH</i> FR1 ( $V_H$ - $J_H$ )	<i>IGH</i> FR2 ( $V_H$ - $J_H$ )	<i>IGH</i> FR3 ( $V_H$ - $J_H$ )	<i>IGH</i> (FR 1, 2 & 3)	<i>IGK</i> ( $V_k$ - $J_k$ & Kde)	Total (FR1, 2, 3 & <i>IGK</i> )
MCL (n=54)	100%	98%	96%	100%	100%	100%
B-CLL/SLL (n=56)	95%	91%	93%	100%	100%	100%
FL (n=109)	73%	76%	52%	84%	84%	100%
MZL (n=41)	73%	85%	68%	87%	83%	97%
DLBCL (n=109)	68%	61%	50%	79%	80%	96%
Total (n=369)	79%	78%	66%	88%	88%	98%

Abbreviations: B-CLL, B-cell chronic lymphocytic leukemia; DLBCL, diffuse large B-cell lymphoma; FL, follicular lymphoma; MCL, mantle cell lymphoma; MZL, marginal zone B-cell lymphoma.

\*PA Evans et al. Leukemia. 2007 21:207-214

Clonality can be Identified in 98% of all B-Cell Malignancies

## Testing Complementary Gene Targets in Parallel Improves Confidence!

# Multiplexing Targets



## Why combine testing of TRB & TRG?

	TRB	TRG	TRB+TRG
T-PLL(%)	100	94	100
T-LGL(%)	96	96	100
PTCL-U(%)	98	94	100
AILT(%)	89	92	95
ALCL(%)	74	74	79*
Total(%)	91	89	94 (99)*

\*Approximately 20–25% of ALCL are known to have no TCR gene rearrangements and are defined as null ALCL; J.J.M. van Krieken et al. *Leukemia*. 2007 21:201-206.

Clonality can be Identified in 94% of all T-Cell Malignancies

## Testing Complementary Gene Targets in Parallel Improves Confidence!

# Why Multiplexing Targets?



Easy to combine:

- IGH and IGK
- TRB and TRG

Advantages of Combining Targets:

- Highest sensitivity
- Helps confirm diagnosis in difficult cases
- Improves Reliability

## B-Cell Targets

	IGH (FR1, 2 & 3)	IGK (Vk - Jk & Kde)	IGH+IGK
MCL(%)	100	100	100
B-CLL/SLL(%)	100	100	100
FL(%)	84	84	100
MZL(%)	87	83	97
DLBCL(%)	79	80	96
Total(%)	88	88	98

## T-Cell Targets

	TRB	TRG	TRB+TRG
T-PLL(%)	100	94	100
T-LGL(%)	96	96	100
PTCL-U(%)	98	94	100
AITL(%)	89	92	95
ALCL(%)	74	74	79*
Total(%)	91	89	94 (99)*

**Testing Complementary Gene Targets in Parallel Improves Confidence!**



**Clonality is defined as a proliferation of cells originating from a single progenitor cell, producing a pool of identical clonal cells. True or False?**

- **TRUE**

**Why test for clonality?**

- Monoclonality is a dominant feature of cancer.
- Clonality testing facilitates the diagnosis of leukemias and lymphomas.
- Clonality testing allows for discrimination between reactive lesions and hematologic malignancies.
- **All of the above**



The molecule targets for clonality testing are B- and T-Cell Receptors.  
True or False?

- **TRUE**

Which detection method is **NOT** available in Invivoscribe assay kits?

- Gel Electrophoresis
- **Flow Cytometry**
- Capillary Electrophoresis (ABI)
- Next-Generation Sequencing (NGS)





## True or False? Multiplexing targets is important because:

- It increases the detection rate of clonal rearrangements
- It increases the test sensitivity
- It improves confidence in results
  
- **TRUE**

# Take Home Message



- **PCR-based Clonality Testing** of B- and T- Cell Gene Rearrangements is the worldwide Gold Standard.
- **Clonality** testing should be performed **at the minimum**, in all cases where the pathological results contradict the clinical findings.
- **Combining targets** results in excellent sensitivity.
- **NGS** allows for unprecedented **detection levels** and information.