PRACTICAL APPLICATIONS OF MOLECULAR BIOLOGY IN SURGICAL PATHOLOGY

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Tumor types

Lymphoma

• G.I. tract

I. Lymphoma

- B and T cell monoclonalities
 - Rearrangement of immunoglobin and TCR genes.
- Identification of non-random chromosomal abnormalities detectable by PCR
 - t(14;18) or t(11;14) translocations in FL and MCL respectively.

B and T cell monoclonality

- Genotype does not correspond to phenotype!

Lineage infidelity of Ig and TCR gene rearrangements ("Illegitimate rearrangements"):

- 50-60 % of lymphoblastic B cell malignancies.
- 20-30% of lymphoblastic T cell malignancies.
- -~10% of mature B and T cell malignancies.

Therefore, Ig and TCR gene rearrangements cannot be used as markers for B and T cell lineages, respectively.

B and T cell monoclonality

- Monoclonality is not always equivalent to malignancy!
 - Clinically benign lymphoproliferations may consist of clonal cell populations.
- Although this pitfall is encountered in B cells, it is mainly observed in T cell monoclonality (cf limited combinatorial diversity of TCR-γ and -δ genes)

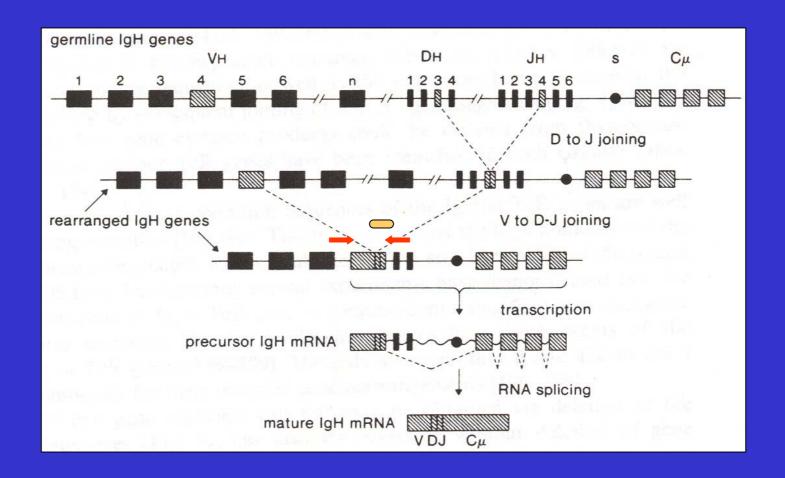
B and T cell monoclonality

Molecular tools

- Southern Blot

- PCR

Schematic diagram of IgH gene rearrangements

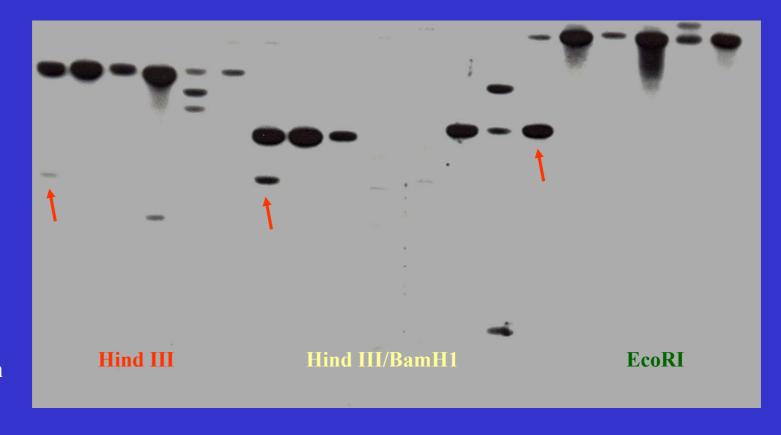


- : indicate the JH probe location for Southern Blot method
- : indicate the primers location for PCR method

B cell monoclonality - Southern Blot Illustration

Patients

1 2 3 4 5 6 1 2 3 4 5 6 6 1 2 3 4 5 6



Restriction **Enzymes**

Southern Blot

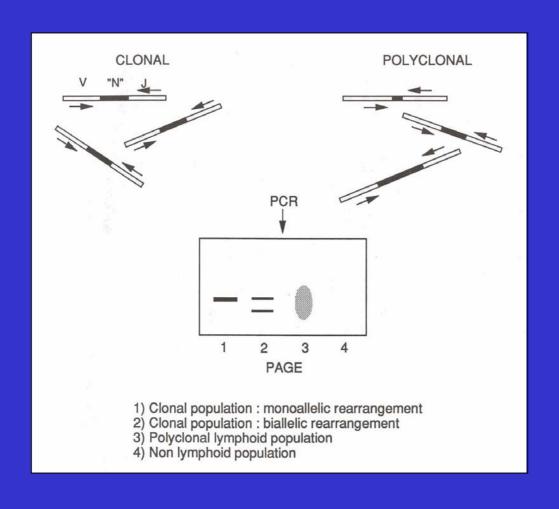
Advantages:

- Very good qualitative sensitivity since ~100% of B and T cell malignancies are detectable by Southern Blot

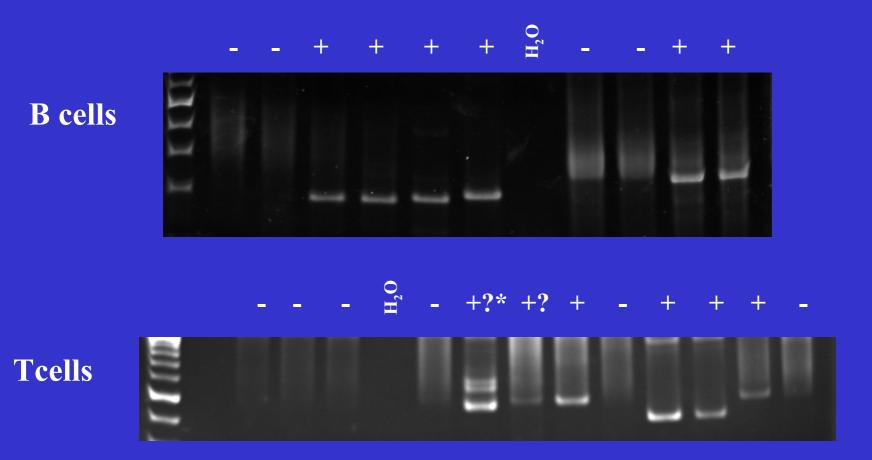
Disadvantages:

- time-consuming
- requires relatively large amounts of pathological material
- low quantitative sensitivity (~ 5 %)

Schematic representation of mono and polyclonal populations detected by PCR.



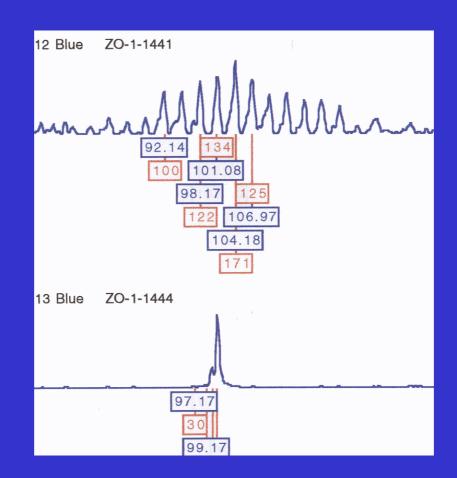
B and T cell monoclonalities - PCR Illustration on paraffin embedded tissue



*? oligoclonality

B cell monoclonality - PCR Illustration (Genescan)

polyclonality



monoclonality

PCR

Advantages: (vs Southern Blot)

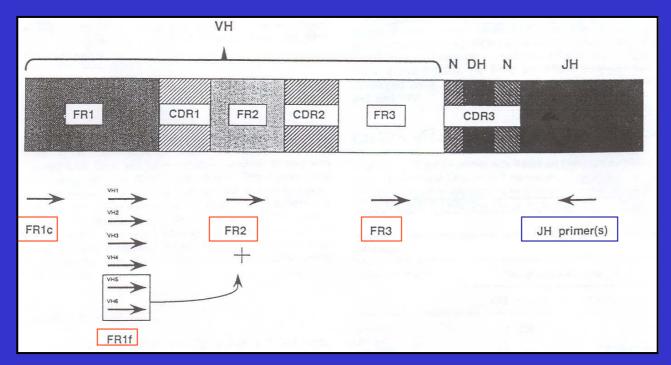
- simple and faster
- requires much less amounts of pathological material
- greater quantitative sensitivity
- can be applied on DNA paraffin-embedded tissue

Disadvantages: (vs Southern Blot)

- lower qualitative sensitivity
- need to use several different PCR strategies in order to increase the overall detection rate.

PCR strategies

• Necessity to use several sets of primers in order to increase the overall detection rate (~90 %) of the PCR method: FR3 -JH FR1c-JH FR1f-JH,...



•This detection rate varies according to the underlying disorders

Detection rates by PCR according to pathological subtypes

- SLL

- MCL

- SNCL

- PCN

- DLCL

- LC (IBL)

- LF

~100 %

~100 %

~80 %

~70 %

~60 %

~50-60 %

~50 %

PCR - Pitfalls

False negative:

- chromosomal translocations into the IgH locus (in FL or DLCL)
- Somatic mutation (in FL and DLCL)
- partial D-J rearrangements (in immature malignancies)
- no VDJ rearrangement produced (in immature malignancies)
- failure of the IgH primers to recognize the VH segment involved

False positive:

- very weak amount of DNA
- reactive lymphoid populations

B and T cell monoclonalities

• Some cases of unequivocal B-cells lymphoma do not generate a clonal signal by PCR despite a demonstrated clonality by Southern Blot.

• Any result must be interpreted in view of other findings and clinical informations

Recurrent molecular abnormalities in lymphoma

 $t(14;18) / Bcl2 - J_H$

in follicular lymphoma

t(11;14) / Bcl1 - J_H

in Mantle Zone lymphoma

 $t(3;14) / Bcl6 - J_H$

in Diffuse Large Cell lymphoma

 $t(8;14) / cMyc - J_H$

in Burkitt lymphoma

t(2,5) / ALK-NPM

in Anaplastic Large Cell Lymphoma

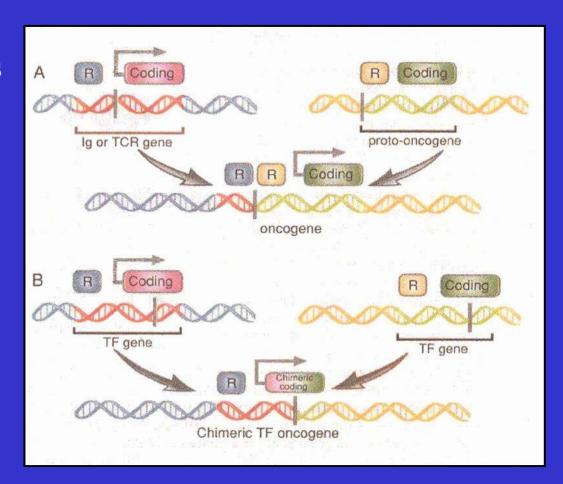
Two distinct types of chromosomal translocations at molecular level

A. Quantitative changes

 $Bcl2-J_H$, $Bcl1-J_H$,...

B. Qualitative changes

ALK-NPM,...

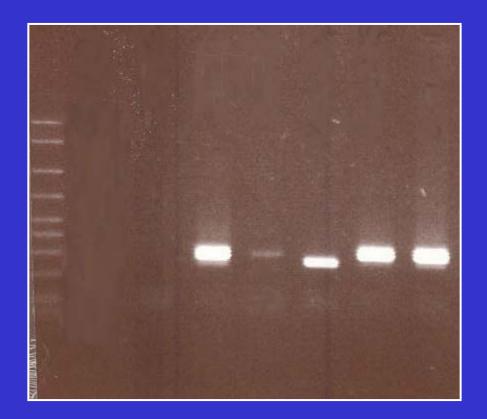


Bcl2 in Follicular lymphoma

- t(14;18) detectable by cytogenetic in ~ 90 % of cases
- this translocation gives rise to an overexpression of the antiapoptotic Bcl2 protein
- four different breakpoints on Bcl2 gene

mbr	in $\sim 45 \%$ of cases		
mer	in $\sim 7 \%$ of cases		
3'UTR	in $\sim 10 \%$ of cases		
icr	in $\sim 10\%$ of cases		

PCR Bcl2-JH in follicular lymphoma Illustration



mbr breakpoint

Bcl2 in Follicular lymphoma

Method

Detection rate

FISH

Cytogenetic

S.B.

PCR Bcl2 / mbr-JH

PCR Bcl2 / mcr-JH

>95%

 $\sim 80-90\%$

65-80%

40-50%

~10%

Bcl2 in Follicular lymphoma

False positivity in normal patients ($\sim 23\%$) and in benign follicular hyperplasia where a very low percentage of positive cells (10^{-3} - 10^{-4}) are detectable by nested PCR \longrightarrow need to use a less sensitive method (standard PCR) to avoid false positive cases

Bcl2 in Diffuse Large Cell Lymphoma

Bcl2-JH rearrangement is detectable in ~35 % of DLCL and seems to be associated with a better prognosis

Bcl1 in Mantle Cell lymphoma

- t(11;14) detectable in ~ 100 % of MCL observed also in MM (~ 20 %) in SLVL (~ 20 %)

- this translocation gives rise to an overexpression of the Bcl1 gene encoding the cyclin D1 protein(positive cell cycle regulatory protein)
- the majority of the Bcl1 breakpoints are clustered in the MTC region

Bcl1 in Mantle Cell lymphoma

70 /	1	
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Detection rate

FISH

Cytogenetic

S.B.

PCR Bcl1/MTC - $J_{\rm H}$

Northern Blot

RT-PCR (cyclinD1 overexpression)

 $\geq 95 \%$

50-80 %

60-70 %

40 %

100 %

100 %

BCL1 Overexpression in MCL

Positive Control



Competitive RT-PCR:
Comparison of the different expression profiles of the three cyclins D1, D2 and D3

This method avoids any false positive results

Cyclin D1

Cyclin D2 →

Cyclin D3 —