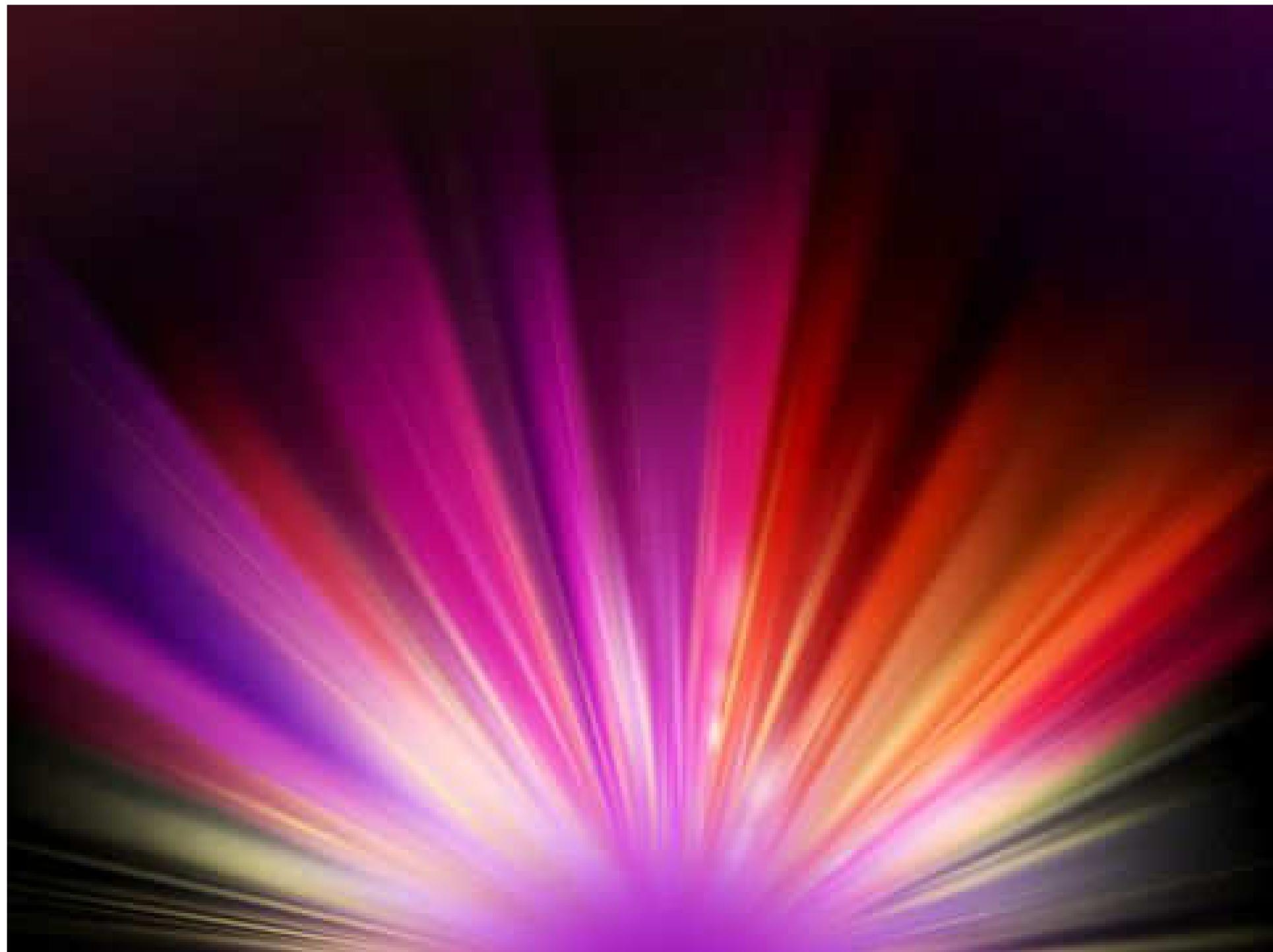


Semmelweis Symposium
2015

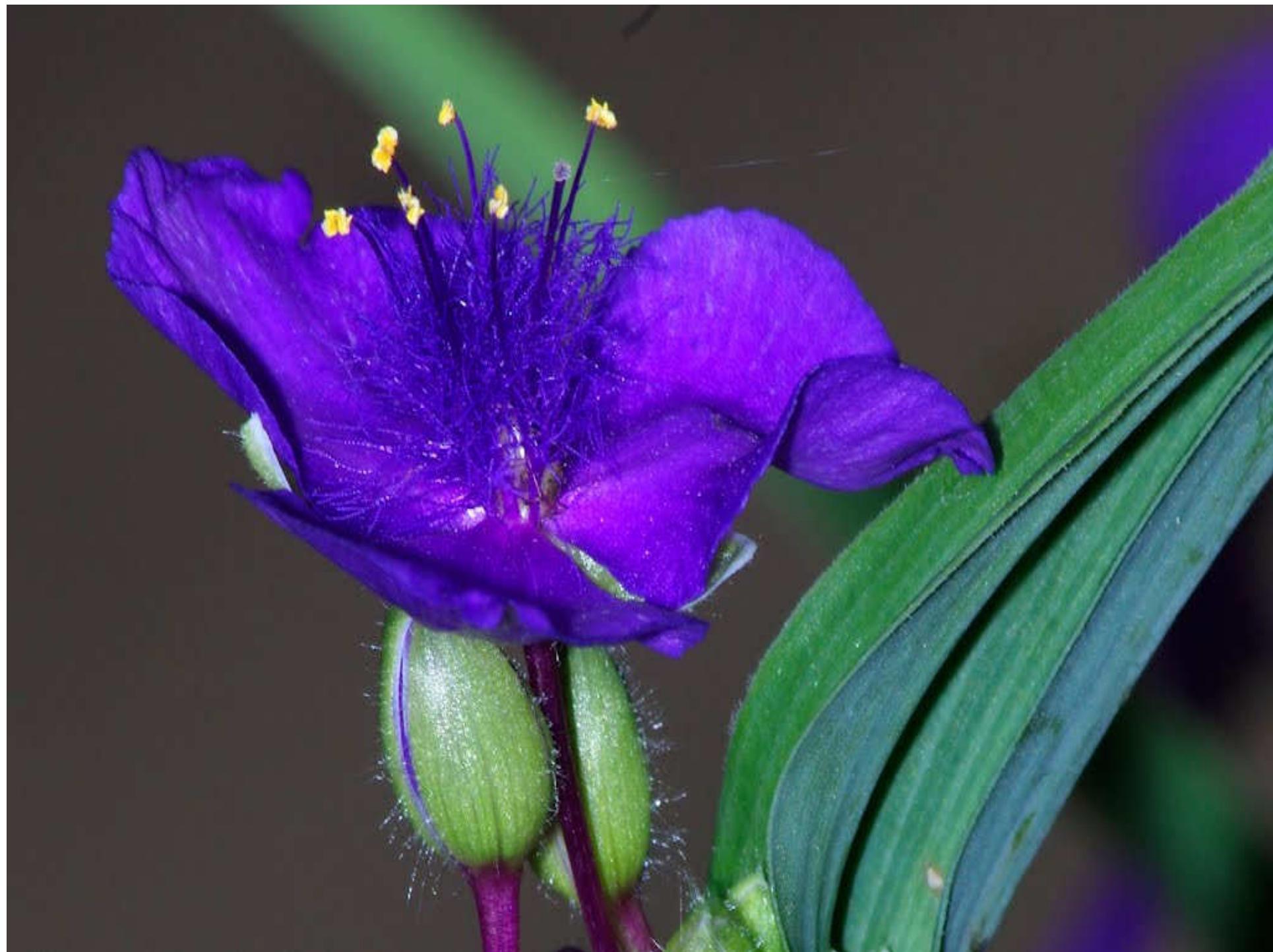
Molecular diagnostics of lymphoma

András Matolcsy

1st. Department of Pathology and Experimental Cancer Research
Semmelweis University
Budapest
Hungary

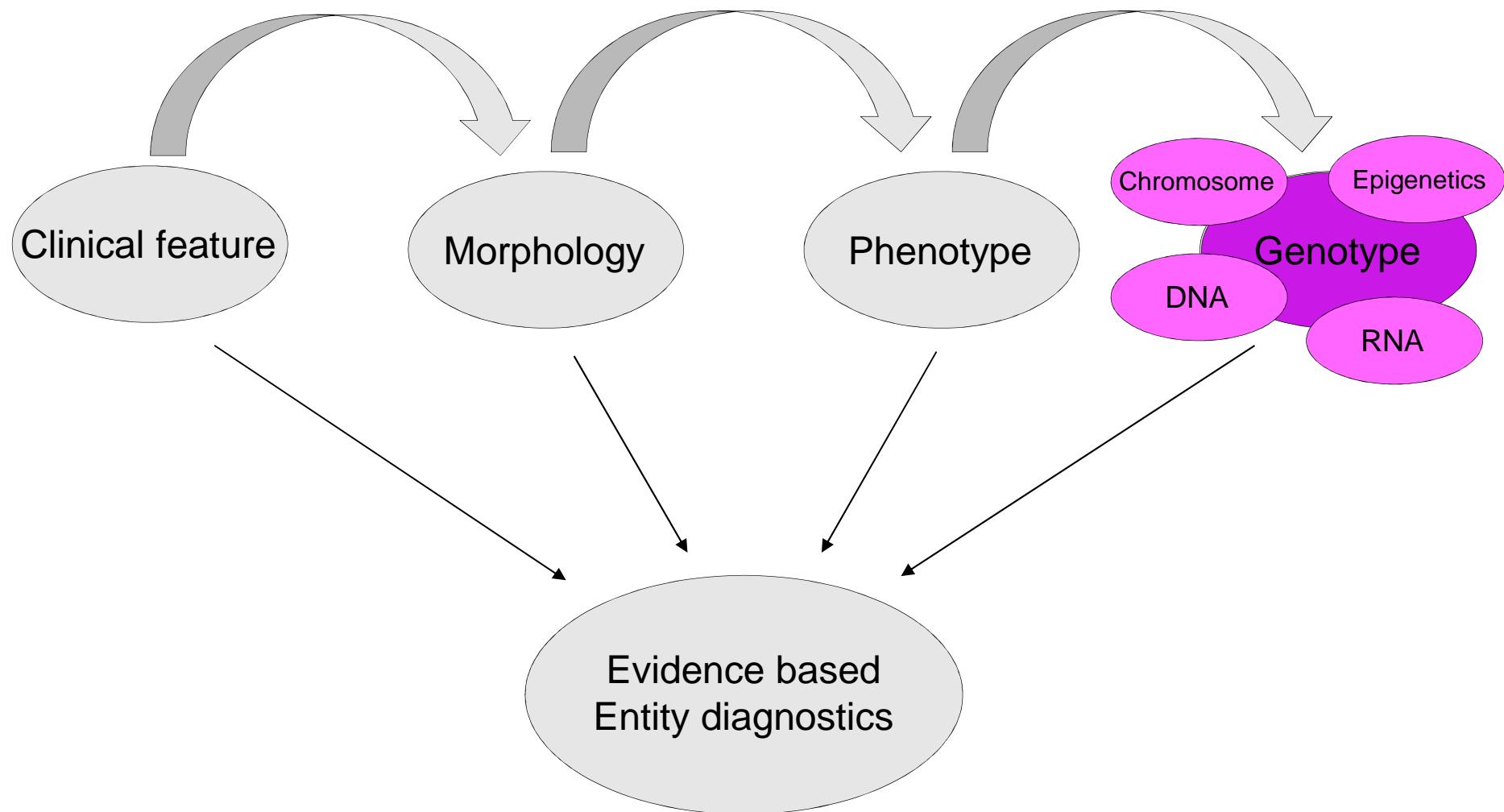








The place of molecular analysis in lymphoma diagnostics





The role of molecular analysis in lymphoma diagnostics

- Determine clonality of the lymphoid cells
- Determine subtype of lymphoma
- Determine prognosis of lymphoma
- Determine effectiveness of treatment (MRD)

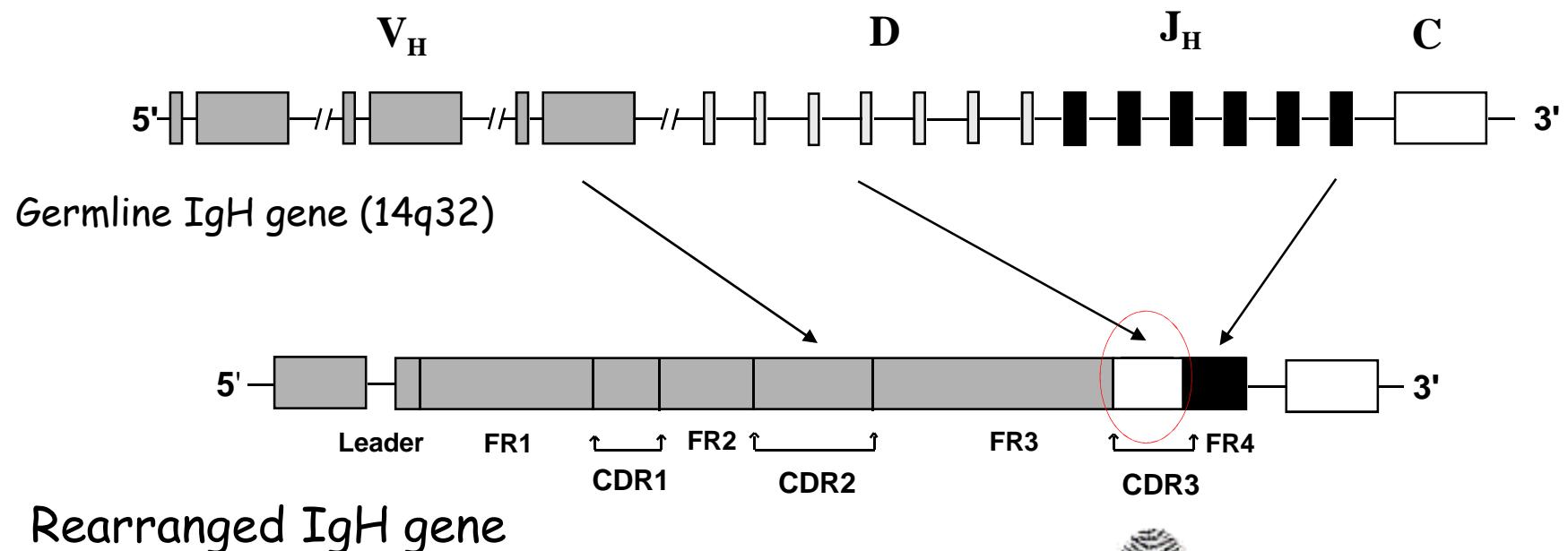


The role of molecular analysis in lymphoma diagnostics

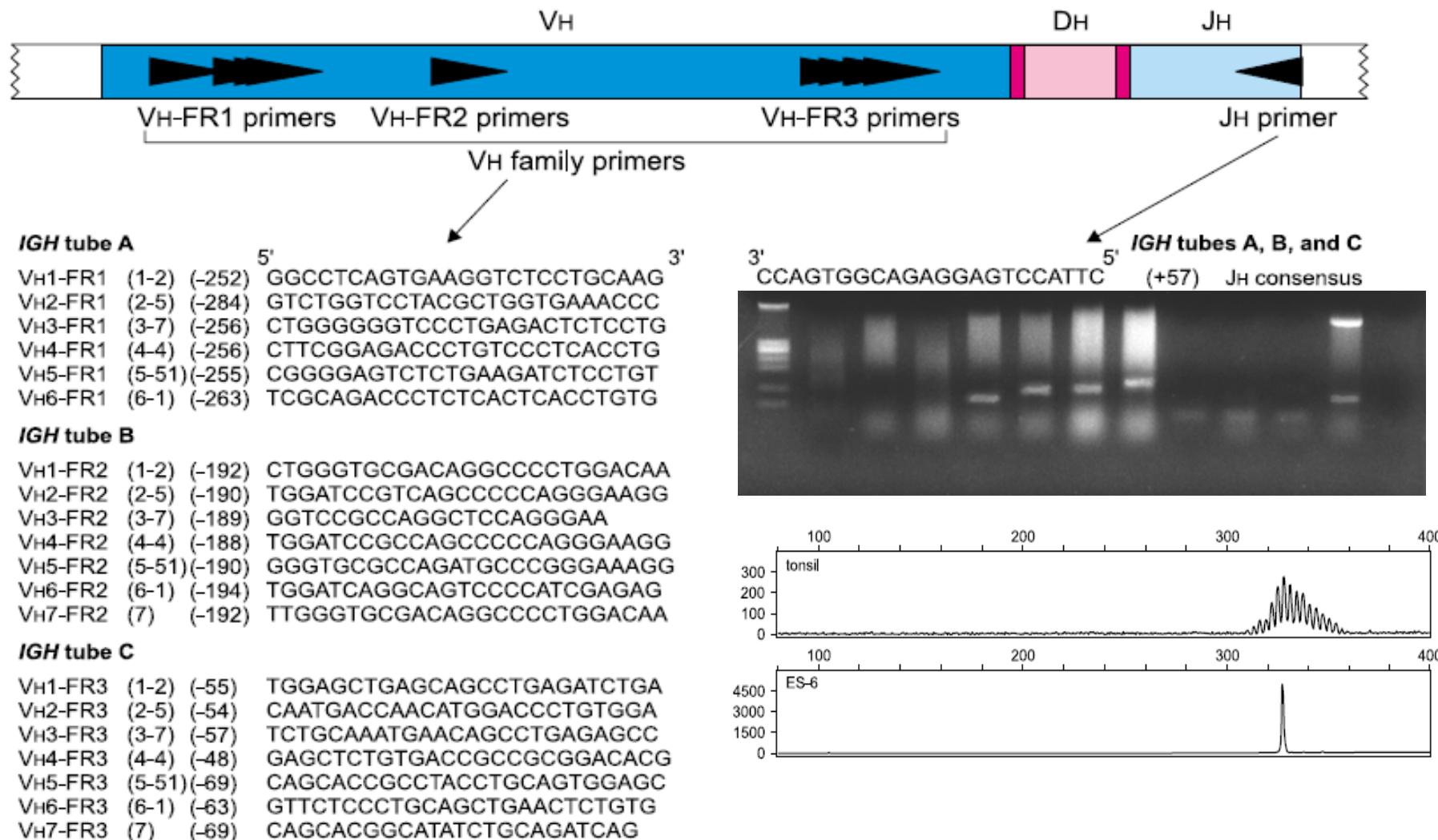
- Determine clonality of the lymphoid cells
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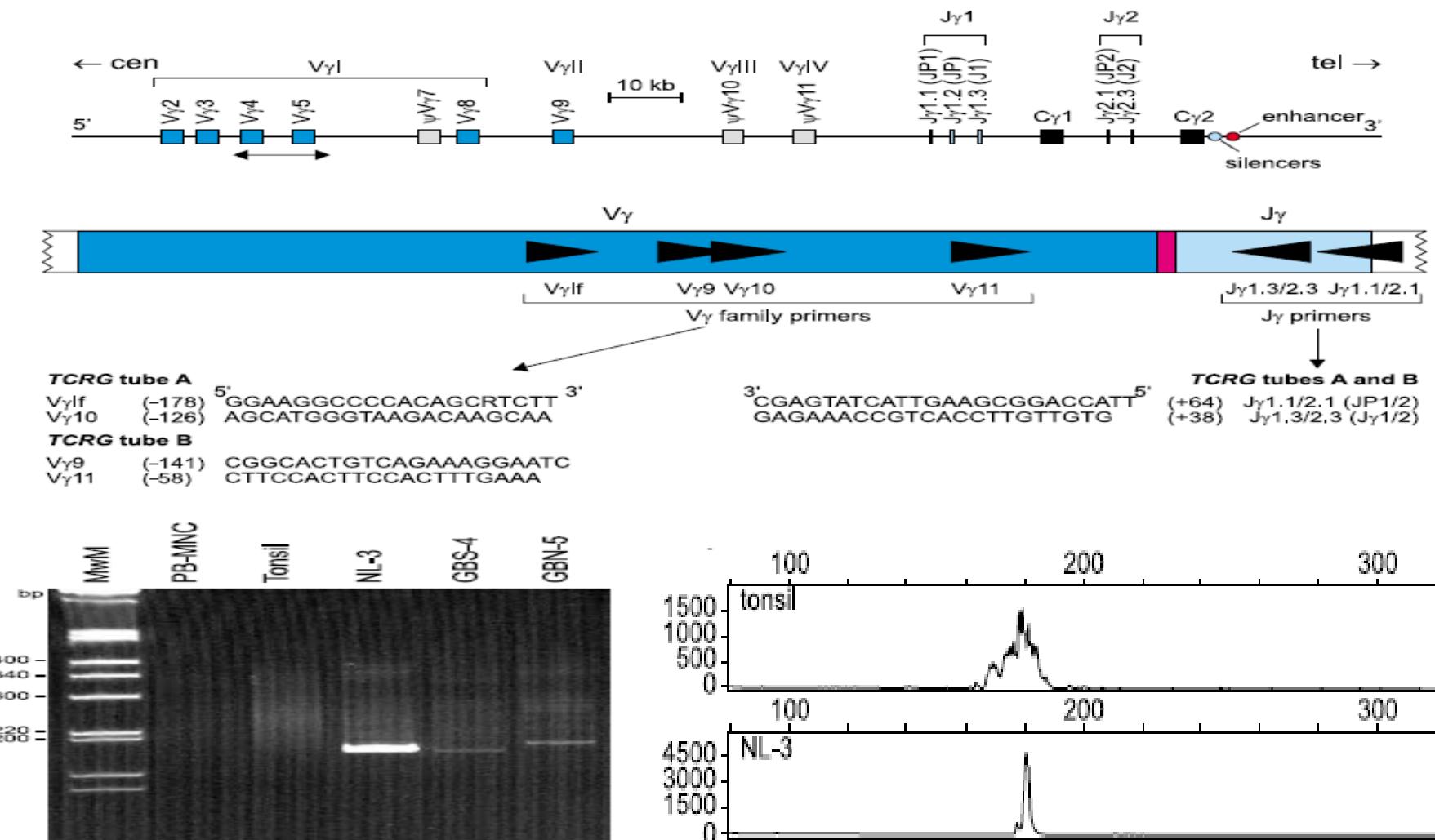
Immunoglobulin heavy chain (IgH) gene rearrangement



PCR analysis of IgH gene rearrangement (BIOMED 2)



PCR analysis of TCR gene rearrangement (BIOMED 2)





Application of clonality analysis in lymphoma diagnostics

Targeted genes for clonality analysis

IgH, Igκ, Igλ
TCRβ, TCRγ TCRδ

Application of Ig/TCR clonality analysis

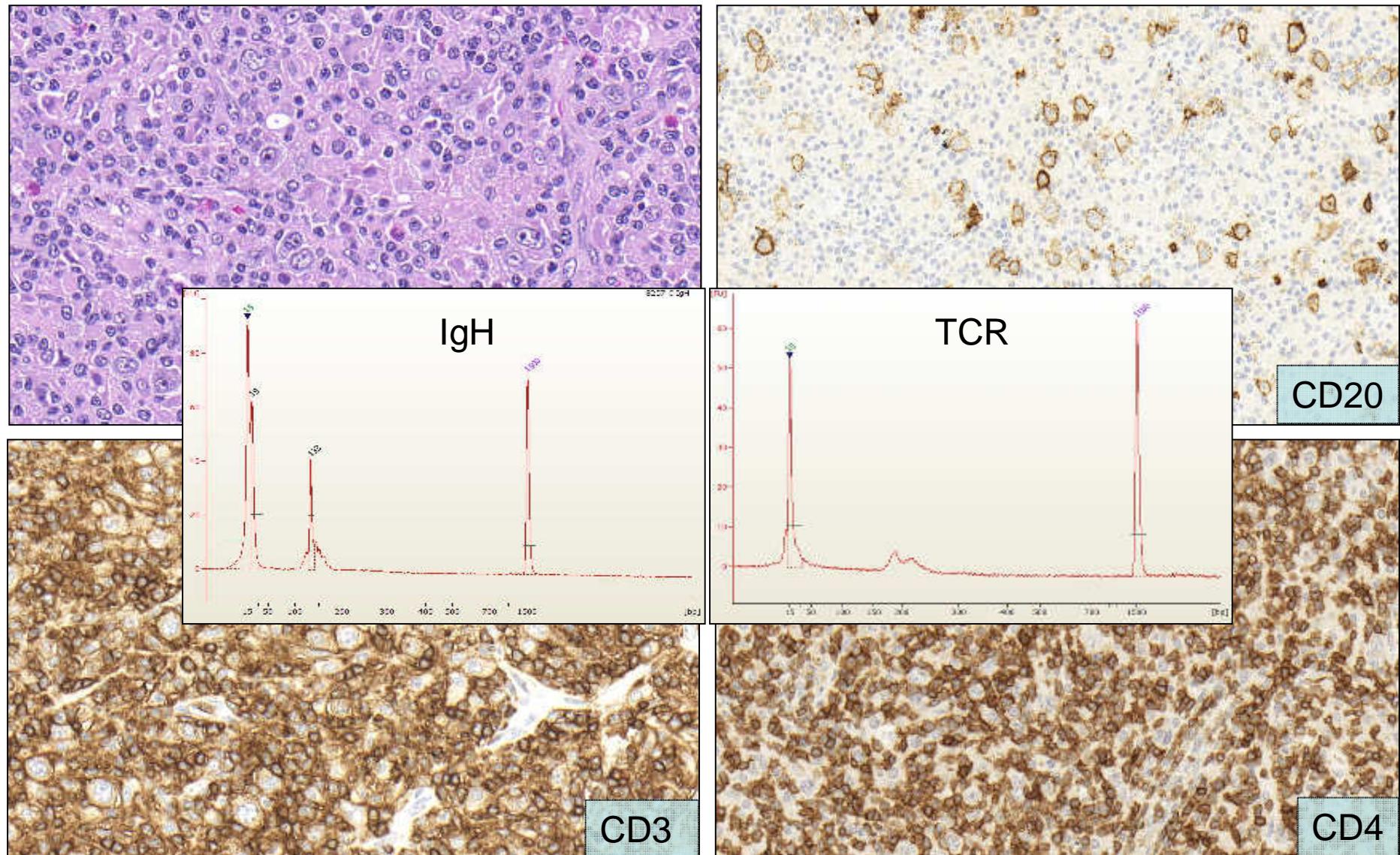
Tumor versus reactive lesions

Lineage (B- versus T-cell) determination (ambiguous lineage)

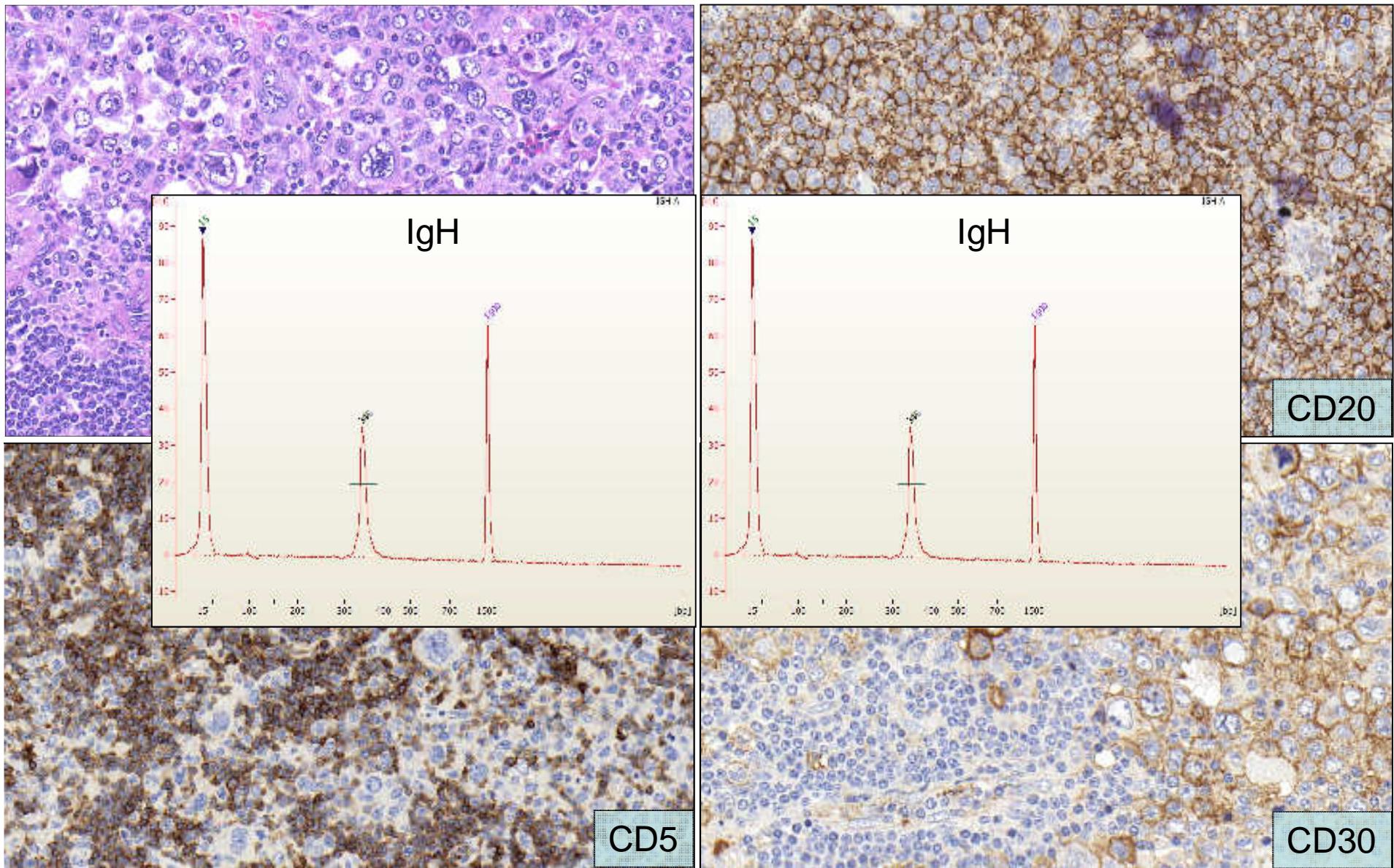
Clonal relationship between different lesions

Clonal relationship between original and relapsed disease

Clonality analysis of lymphoma with ambiguous lineage



Clonality analysis on progressed CLL





Pitfalls of Ig/TCR clonality analysis

- Fals negativity: Few neoplastic B- or T-cells in sample
Poor DNA quality
Somatic hypermutation in antigen receptors of neoplastic cells

- Fals positivity: Oligoclonality in reactive lesions
Selective amplification of a reactive subclone

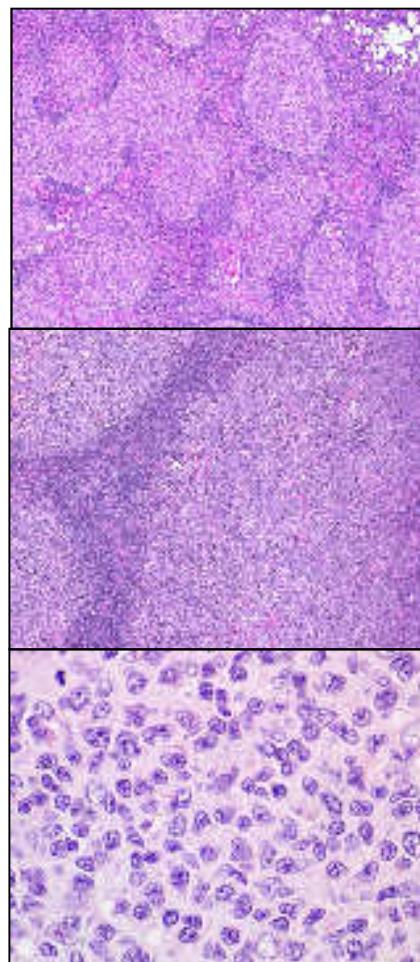


The role of molecular analysis in lymphoma diagnostics

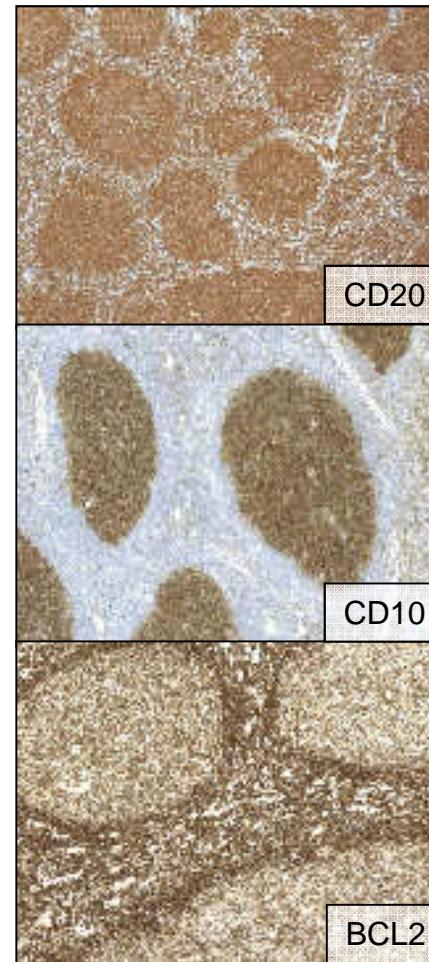
- Determine clonality of the lymphoid cells
- Determine subtype of lymphoma
- Determine prognosis of lymphoma
- Determine effectiveness of treatment (MRD)

WHO classification of lymphomas (entity classification)

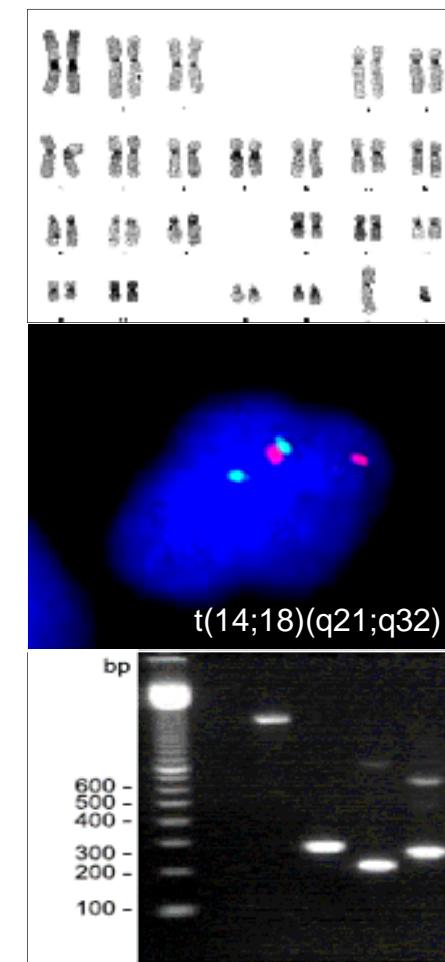
MOPHOLOGY



PHENOTYPE



GENOTYPE



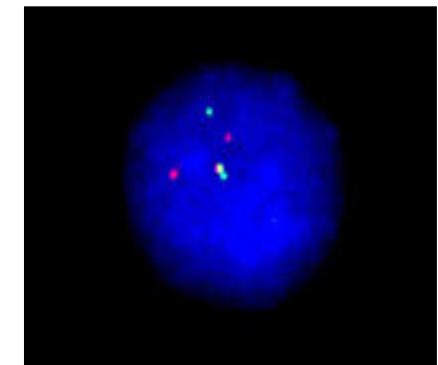
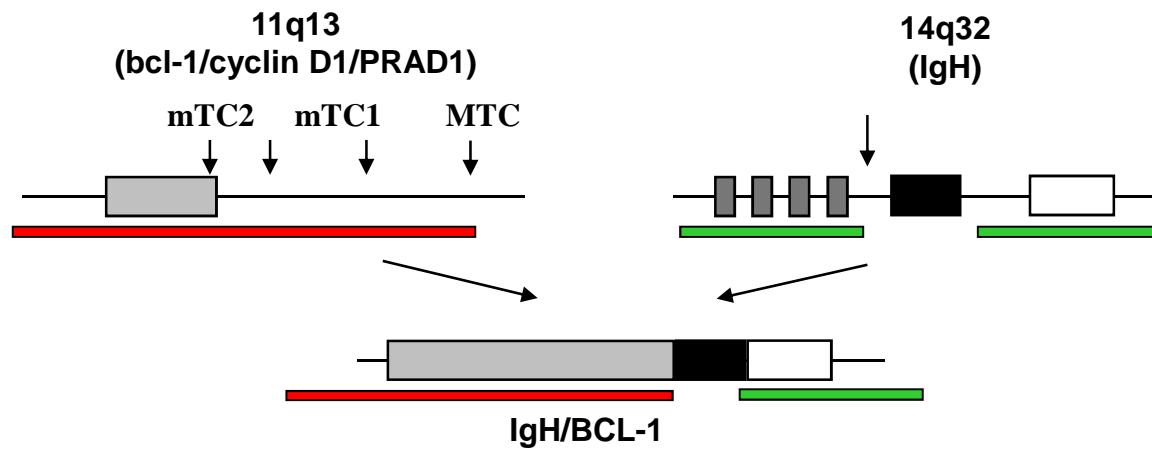
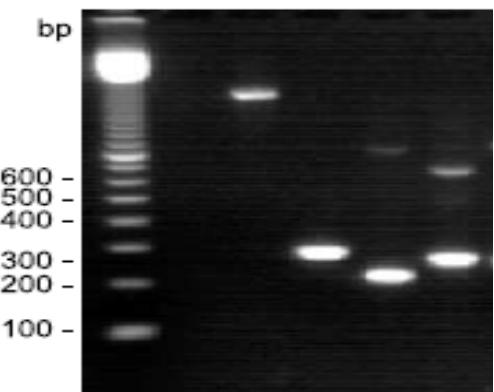
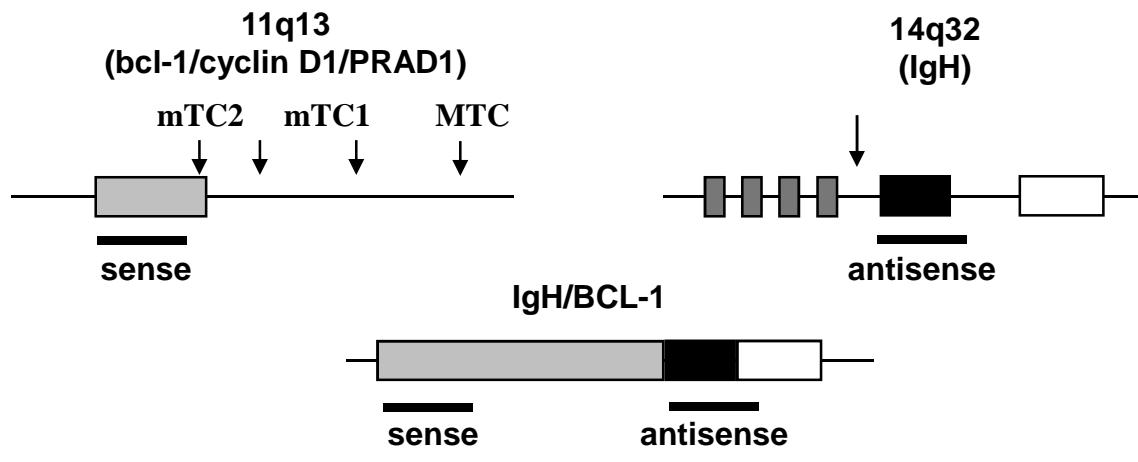


Reccurent genetic lesions in different lymphomas

Lymphoma	Translocation/mutation	Genes involved
Follicular lymphoma	t(14;18)(q21;q32)	<i>IgH/BCL2</i>
Mantle cell lymphoma	t(11;14)q13;q32)	<i>CCND1/IgH</i>
Burkitt lymphoma	t(8;14)(q24;q32)	<i>cMYC/IgH</i>
	t(2;8)(p11;q24)	<i>IgK/cMYC</i>
	t(8;22)(q24;q11)	<i>cMYC/IgL</i>
Anaplastic large cell lymphoma	t(2;5)(p23;q35)	<i>ALK1/NPM</i>
	t(1;2)(q25;p23)	<i>TPM3/ALK1</i>
	inv(2)(p23;q35)	<i>ALK1/ATIC</i>
	t(2;3)(p23;q21)	<i>ALK1/TFG</i>
	t(2;17)(p23;q23)	<i>ALK1/CLTC</i>
	t(2;X)(p23;q11)	<i>ALK1/MSN</i>
	t(2;19)(p23;p13)	<i>ALK1TPM4</i>
Gastric MALT lymphoma	t(11;18)(q21;q22)	<i>API2/MLT1</i>
Diffuse large B-cell lymphoma	t(14;18)(q21;q32)	<i>IgH/BCL2</i>
	t(8;14)(q24;q32)	<i>cMYC/IgH</i>
	3q27	<i>BCL6</i>
Hairy cell leukaemia	7q34	<i>BRAF (V600G)</i>

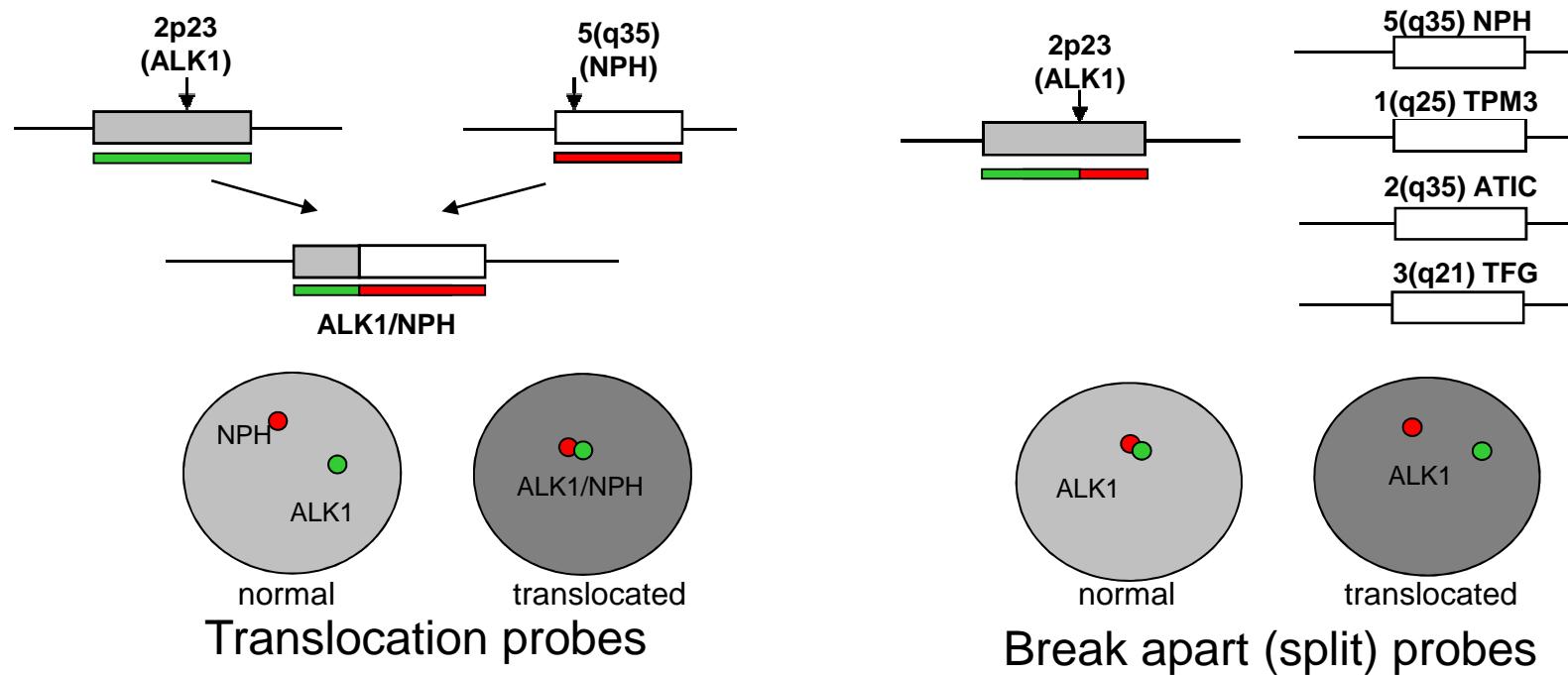


Detection of recurrent genetic lesions in lymphomas



FISH strategies for detection of genetic lesions in lymphomas

Lymphoma	Translocation/mutation	Genes involved
Anaplastic large cell lymphoma	t(2;5)(p23;q35)	<i>ALK1/NPM</i>
	t(1;2)(q25;p23)	<i>TPM3/ALK1</i>
	inv(2)(p23;q35)	<i>ALK1/ATIC</i>
	t(2;3)(p23;q21)	<i>ALK1/TFG</i>
	t(2;17)(p23;q23)	<i>ALK1/CLTC</i>
	t(2;X)(p23;q11)	<i>ALK1/MSN</i>
	t(2;19)(p23;p13)	<i>ALK1/TPM4</i>

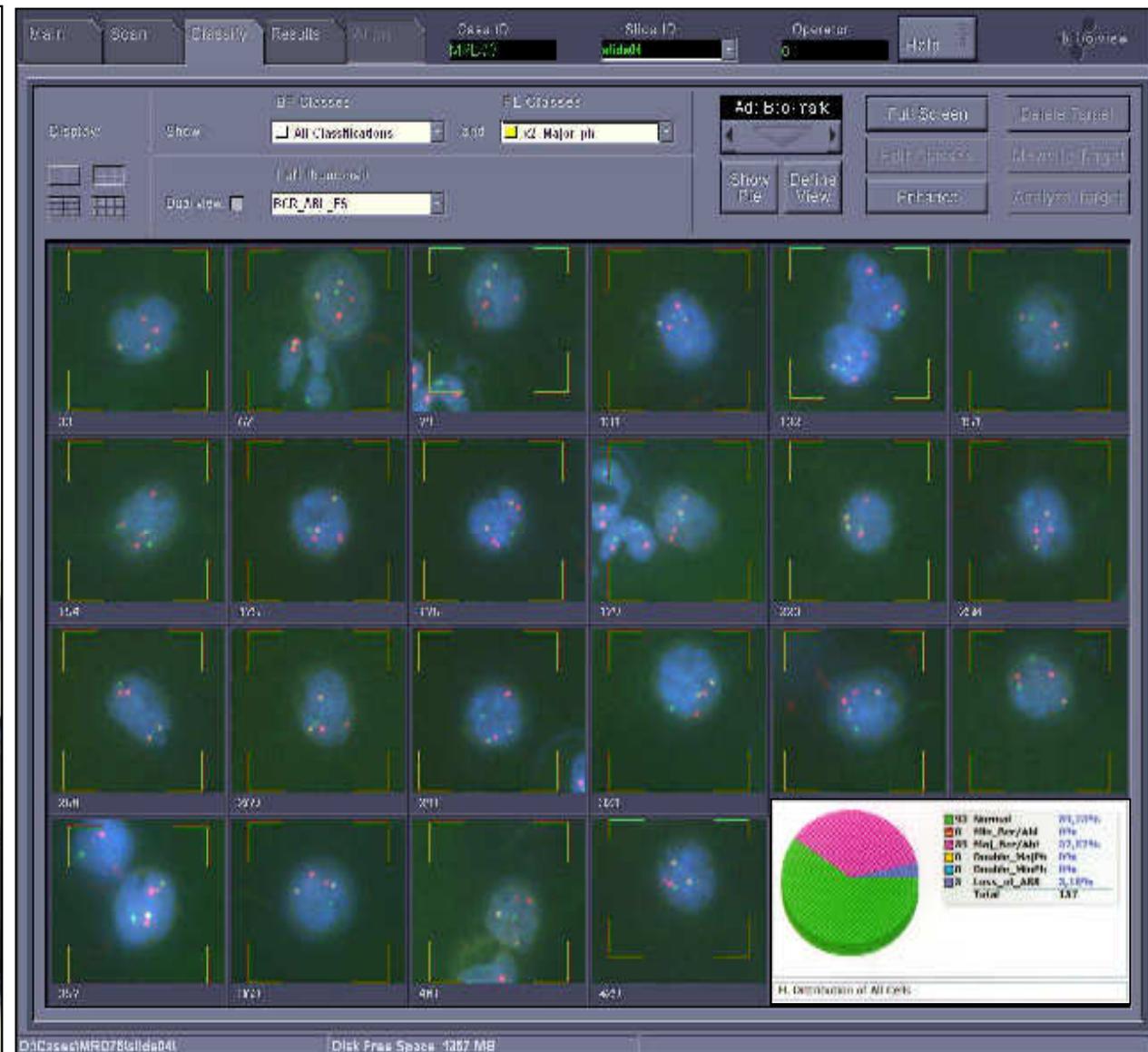




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Automated FISH analysis



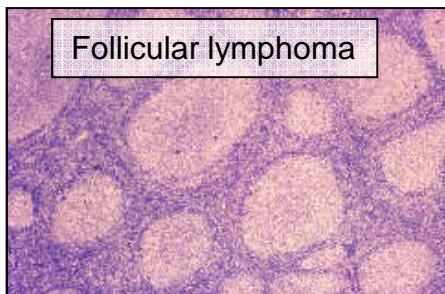


Reccurent genetic lesions with oncoprotein overexpression

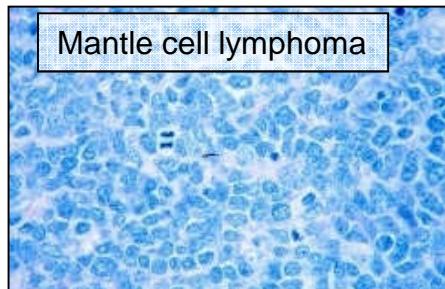
Lymphoma	Translocation/mutation	Genes involved
Follicular lymphoma	t(14;18)(q21;q32)	<i>IgH/BCL2</i>
Mantle cell lymphoma	t(11;14)q13;q32)	<i>CCND1/IgH</i>
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	t(2;8)(p11;q24)	<i>IgK/cMYC</i>
	t(8;22)(q24;q11)	<i>cMYC/IgL</i>
Anaplastic large cell lymphoma	t(2;5)(p23;q35)	<i>ALK1/NPM</i>
	t(1;2)(q25;p23)	<i>TPM3/ALK1</i>
	inv(2)(p23;q35)	<i>ALK1/ATIC</i>
	t(2;3)(p23;q21)	<i>ALK1/TFG</i>
	t(2;17)(p23;q23)	<i>ALK1/CLTC</i>
	t(2;X)(p23;q11)	<i>ALK1/MSN</i>
	t(2;19)(p23;p13)	<i>ALK1TPM4</i>
Gastric MALT lymphoma	t(11;18)(q21;q22)	<i>API2/MLT1</i>
Diffuse large B-cell lymphoma	t(14;18)(q21;q32)	<i>IgH/BCL2</i>
	t(8;14)(q24;q32)	<i>cMYC/IgH</i>
	3q27	<i>BCL6</i>
Hairy cell leukaemia	7q34	<i>BRAF (V600G)</i>



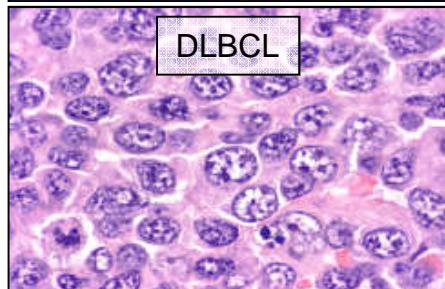
Genetic alterations with oncoprotein overexpression



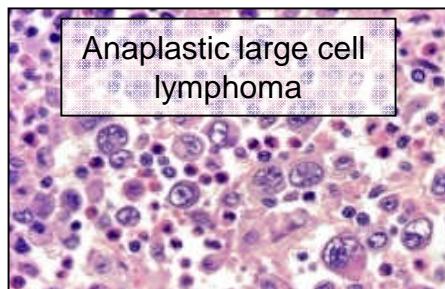
t(14;18)(q21;q32)
IgH/BCL2



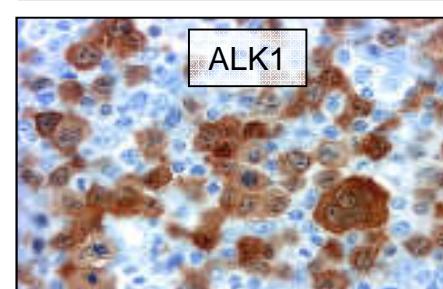
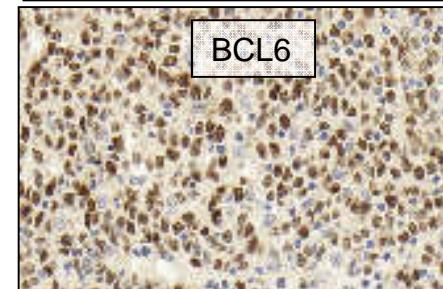
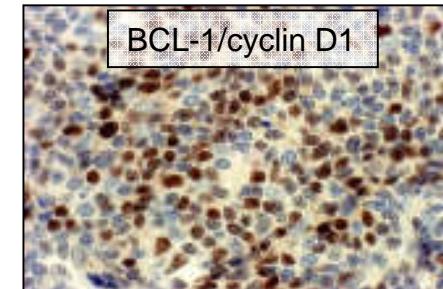
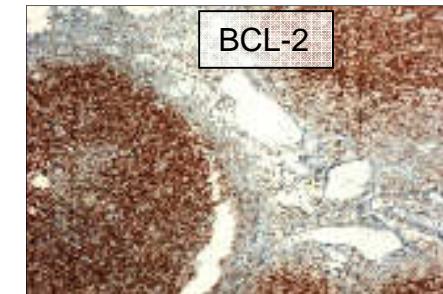
t(11;14)q13;q32
CCND1/IgH



t(3q27; v)
v/BCL6

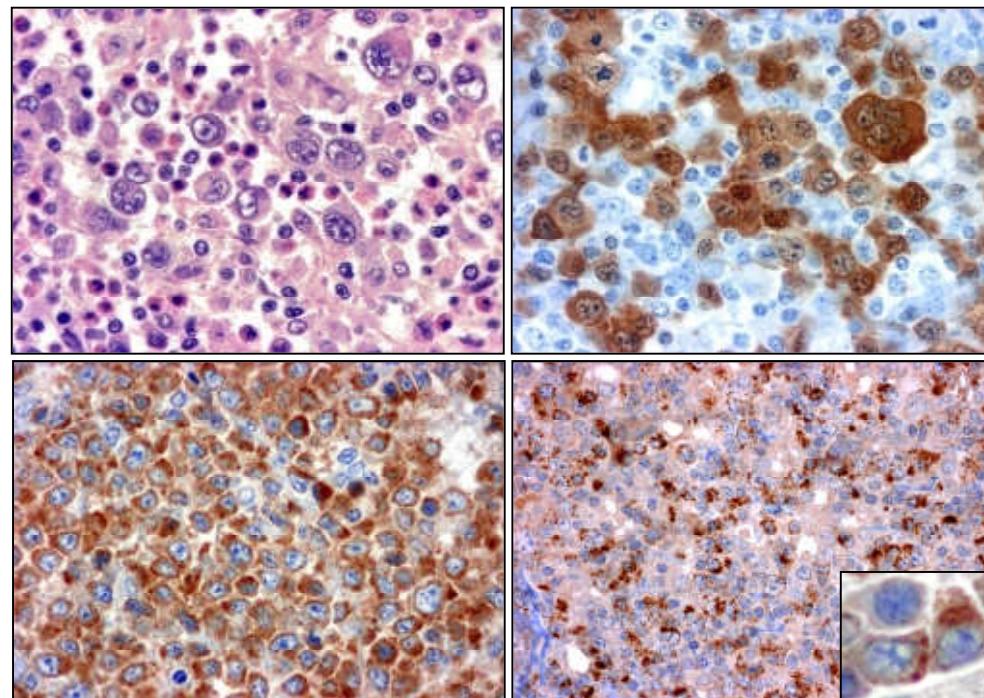


t(2;5)(p23;q35)
ALK1/NPM



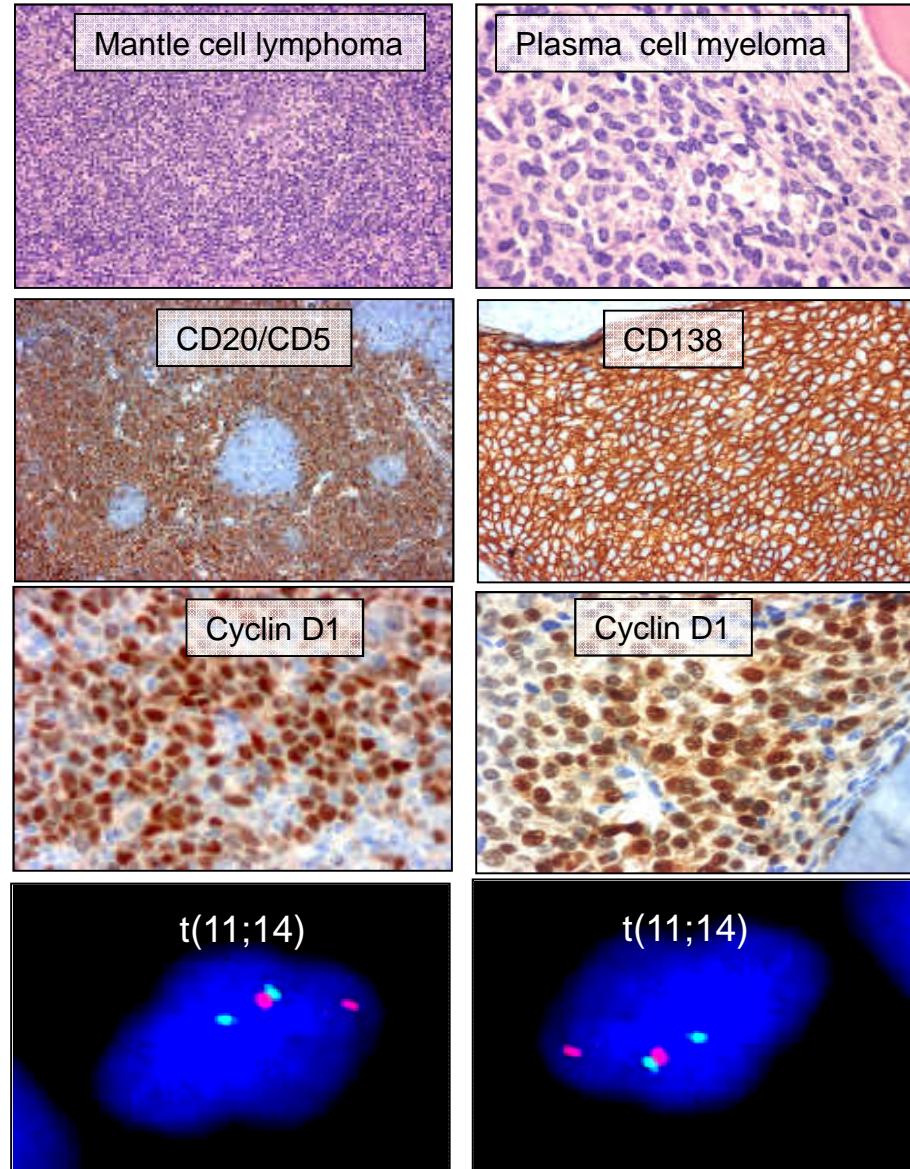
Patterns of ALK1 positivity in anaplastic large cell lymphoma

Translocation/mutation	Genes involved	ALK1 positivity
t(2;5)(p23;q35)	ALK1/NPM	nucleus, cytoplasm
t(1;2)(q25;p23)	TPM3/ALK1	cytoplasm
inv(2)(p23;q35)	ALK1/ATIC	cytoplasm
t(2;3)(p23;q21)	ALK1/TFG	cytoplasm
t(2;17)(p23;q23)	ALK1/CLTC	cytoplasm (granular)
t(2;X)(p23;q11)	ALK1/MSN	membrane
t(2;19)(p23;p13)	ALK1TPM4	cytoplasm



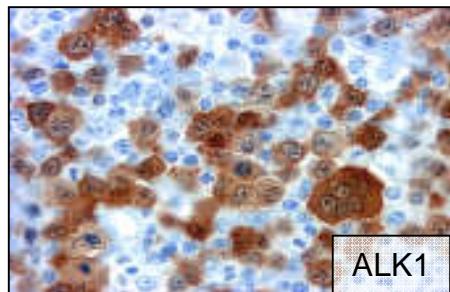


Overlapping oncogene overexpression in different lymphomas

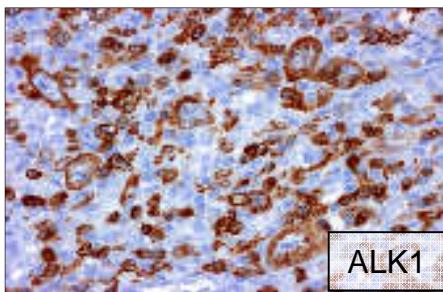




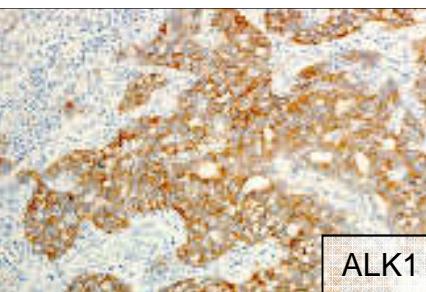
Overlapping oncprotein overexpression in different neoplasms



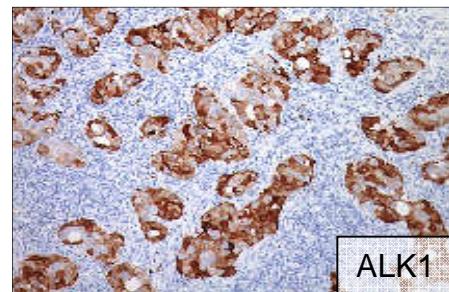
Anaplastic large cell lymphoma



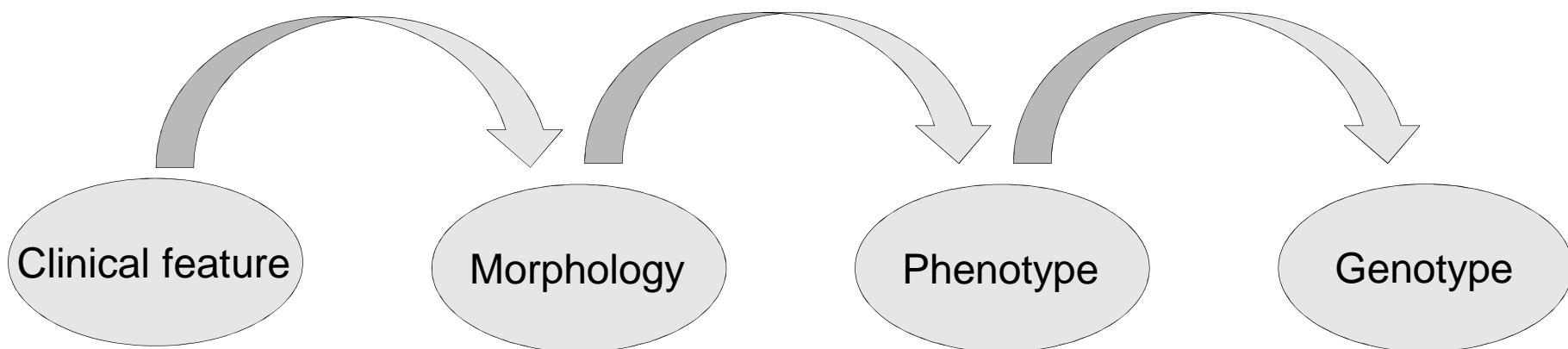
Diffuse large B cell lymphoma



Lung cancer



Gastric cancer





Reccurent genetic lesions in different lymphoma entities

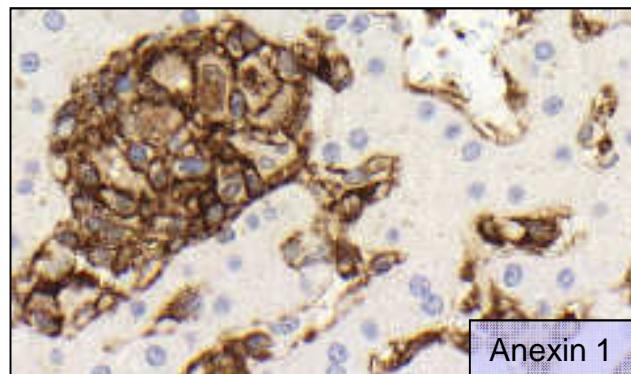
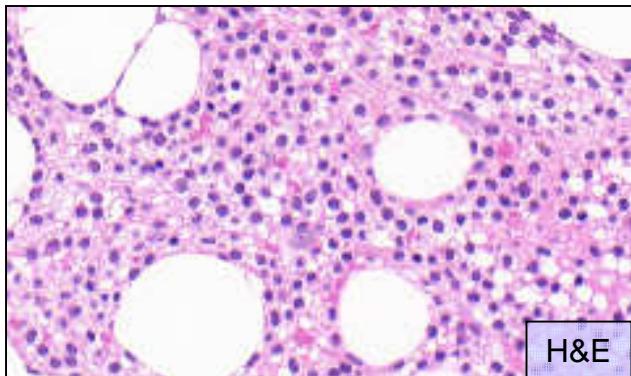
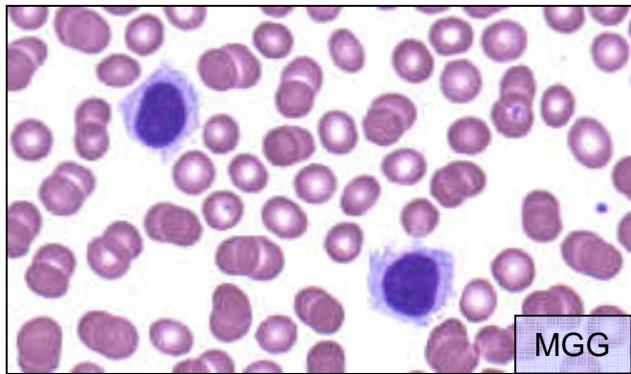
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Mantle cell lymphoma	t(11;14)q13;q32)	<i>CCND1/IgH</i>
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	t(2;8)(p11;q24)	<i>IgK/cMYC</i>
	t(8;22)(q24;q11)	<i>cMYC/IgL</i>
Anaplastic large cell lymphoma	t(2;5)(p23;q35)	<i>ALK1/NPM</i>
	t(1;2)(q25;p23)	<i>TPM3/ALK1</i>
	inv(2)(p23;q35)	<i>ALK1/ATIC</i>
	t(2;3)(p23;q21)	<i>ALK1/TFG</i>
	t(2;17)(p23;q23)	<i>ALK1/CLTC</i>
	t(2;X)(p23;q11)	<i>ALK1/MSN</i>
	t(2;19)(p23;p13)	<i>ALK1TPM4</i>
Gastric MALT lymphoma	t(11;18)(q21;q22)	<i>API2/MLT1</i>
Diffuse large B-cell lymphoma	t(14;18)(q21;q32)	<i>IgH/BCL2</i>
	t(8;14)(q24;q32)	<i>cMYC/IgH</i>
	3q27	<i>BCL6</i>
Hairy cell leukaemia	7q34	<i>BRAF (V600G)</i>



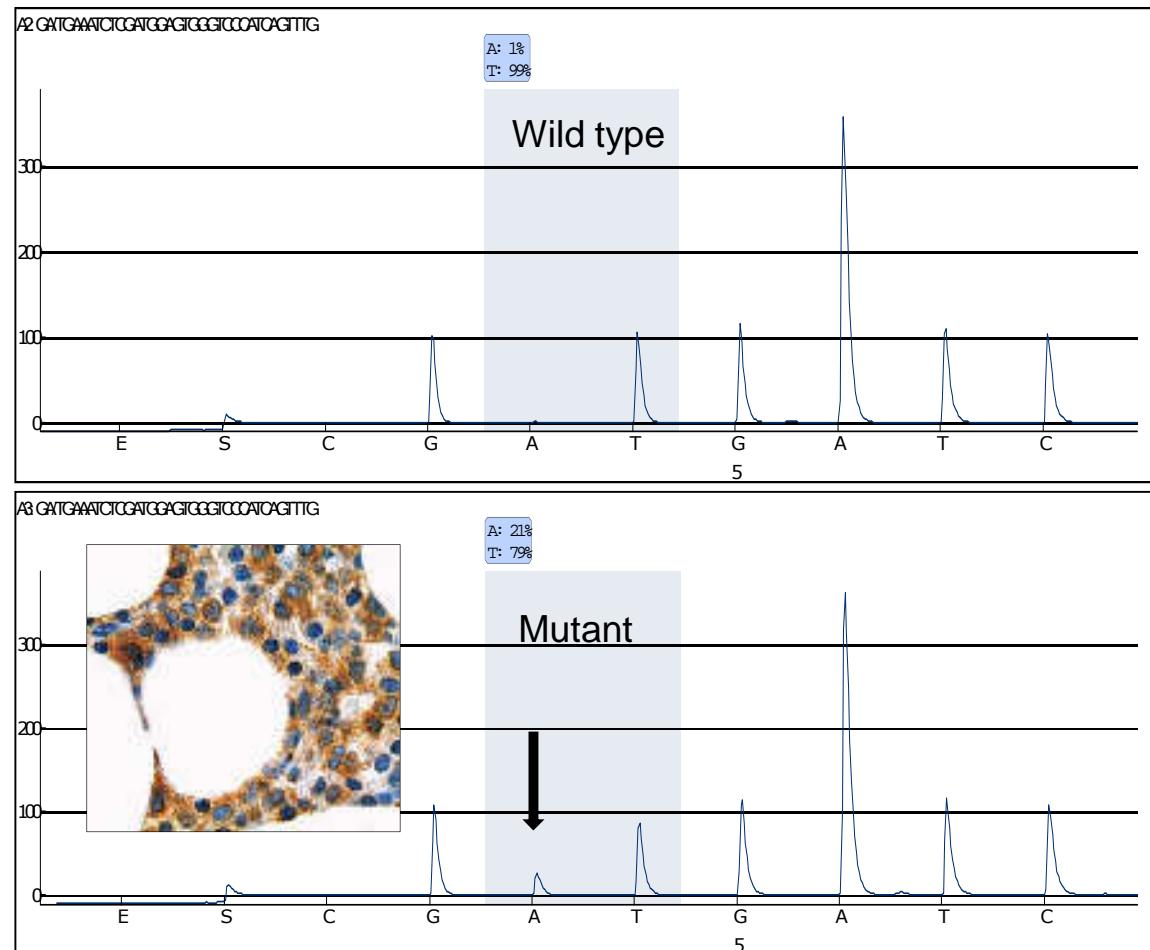
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Molecular diagnostics of hairy cell leukaemia

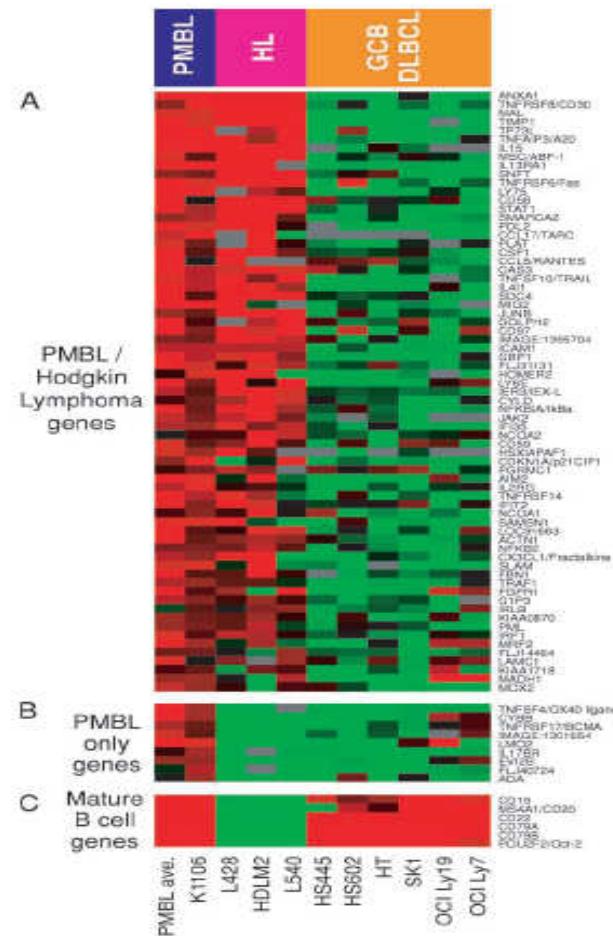
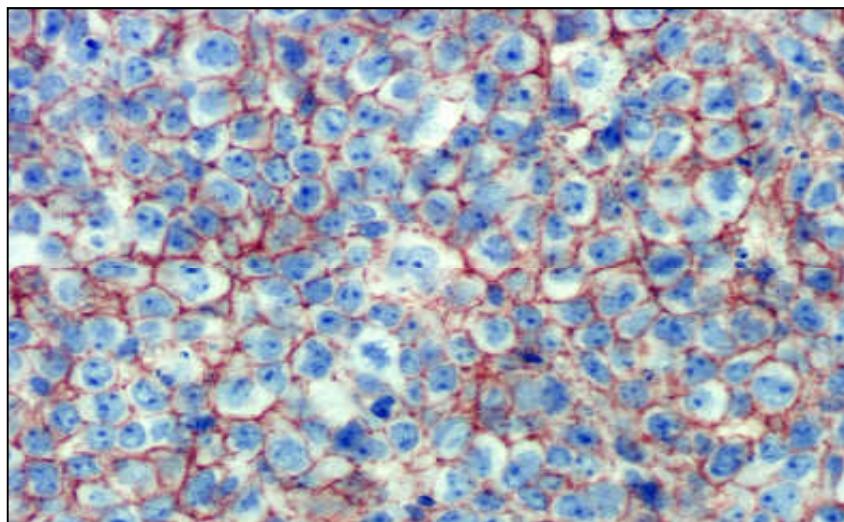
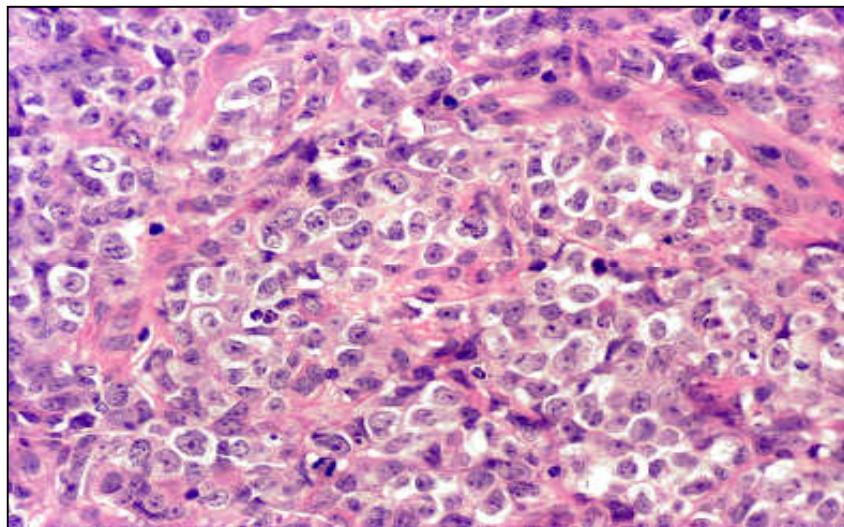


BRAF G600V

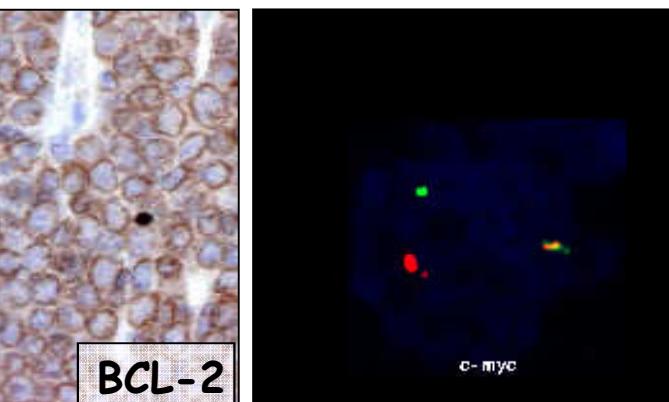
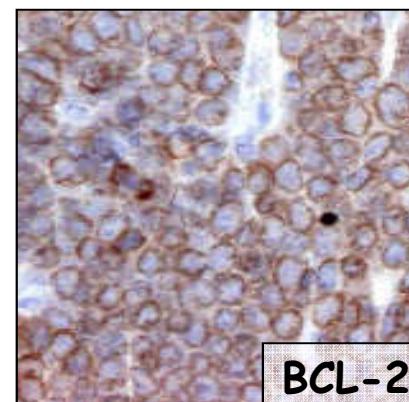
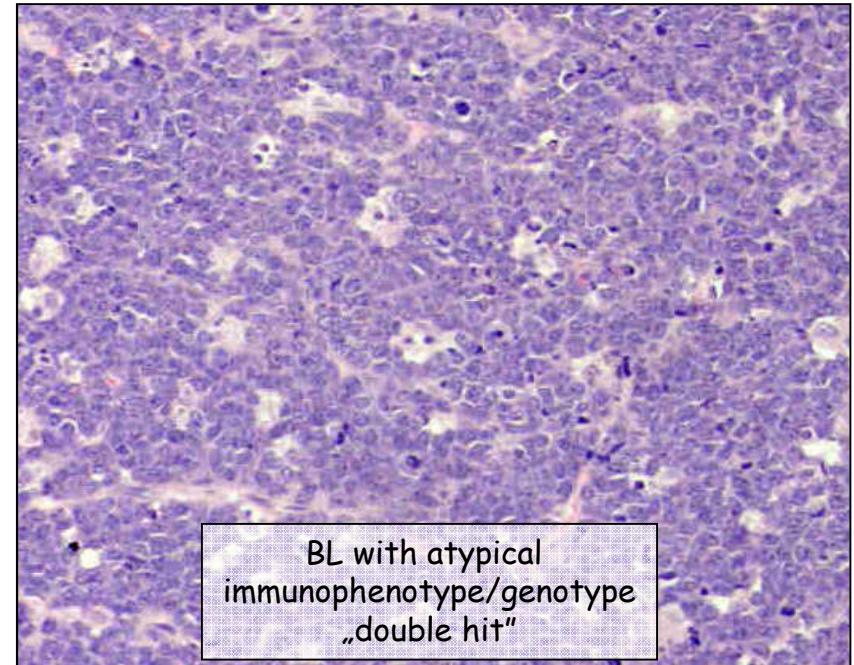
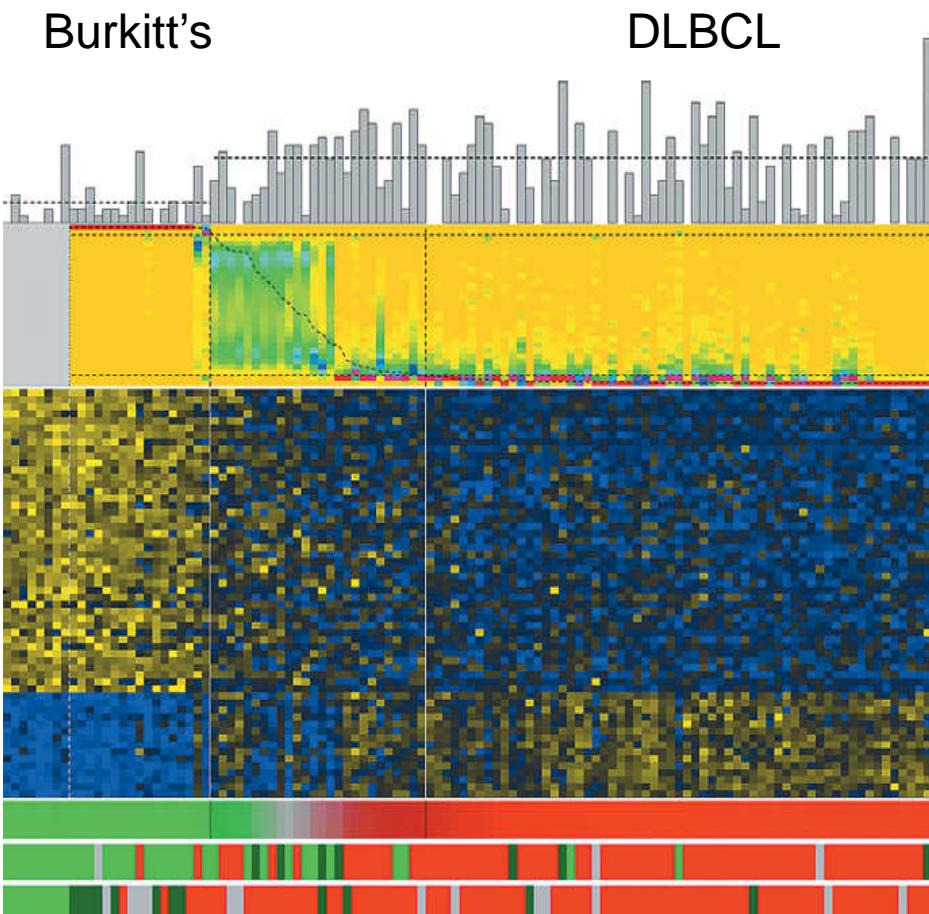




Expression profile analysis of primary mediastinal large B-cell lymphomas



Expression profile analysis of Burkitt's lymphoma



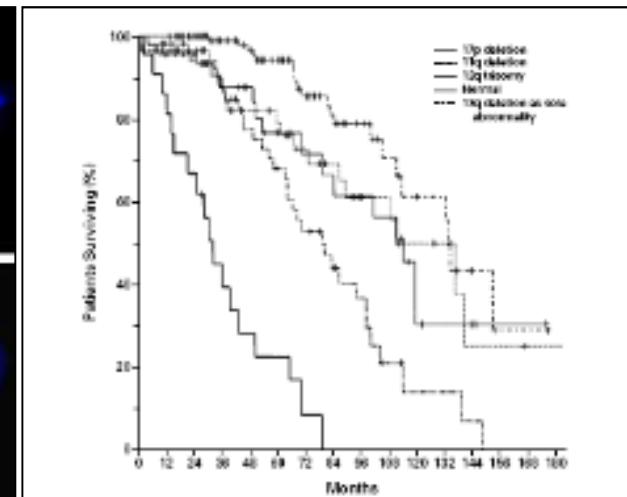
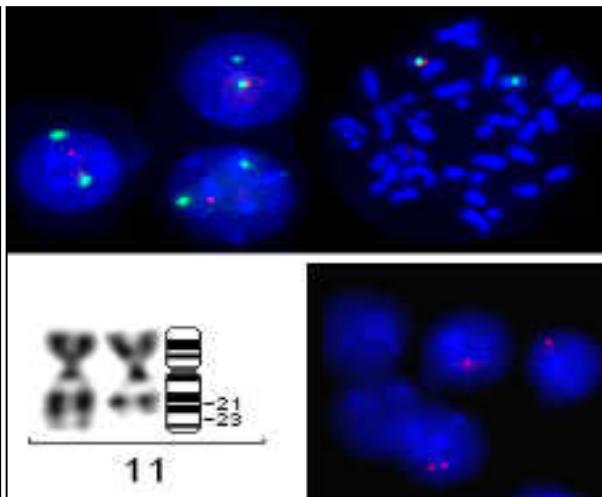
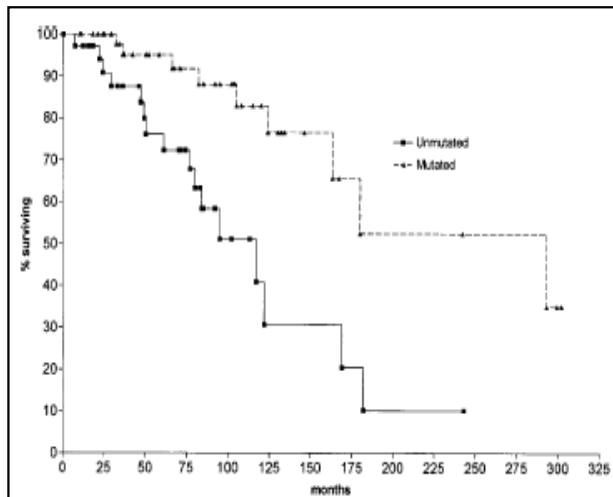


The role of molecular analysis in lymphoma diagnostics

- Determine clonality of the lymphoid cells
- Determine subtype of lymphoma
- **Determine prognosis of lymphoma**
- Determine effectiveness of treatment (MRD)

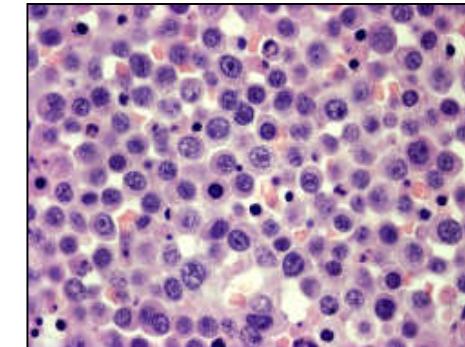
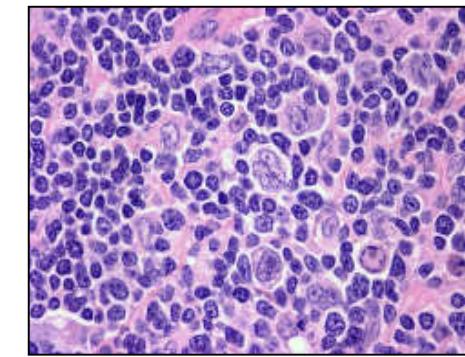
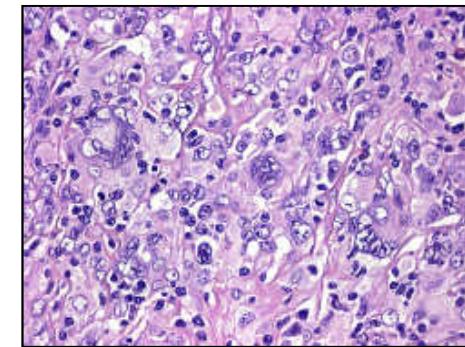
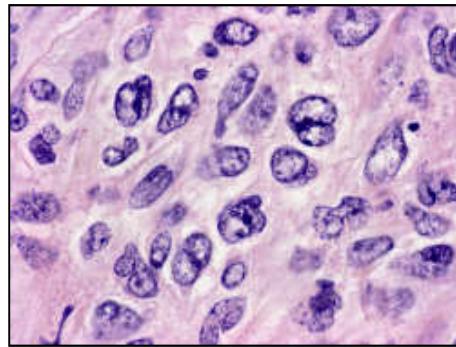
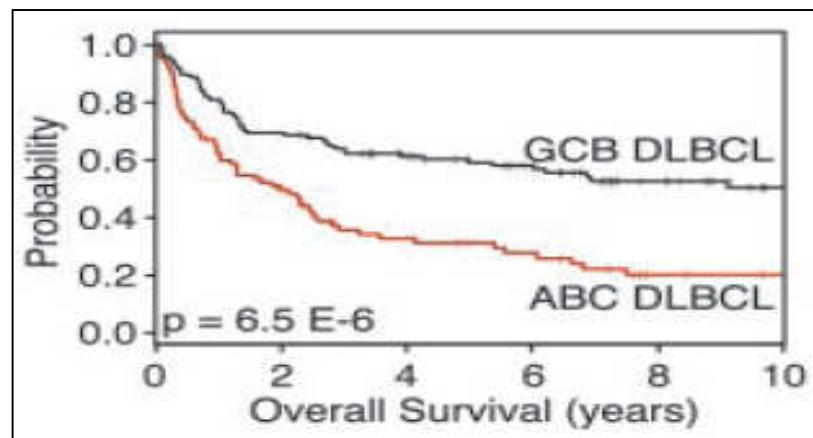
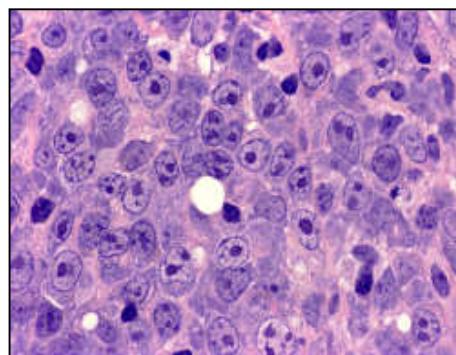
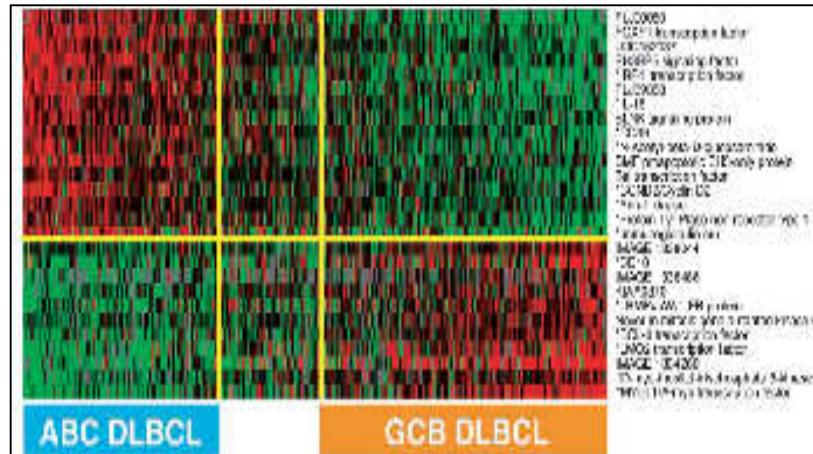
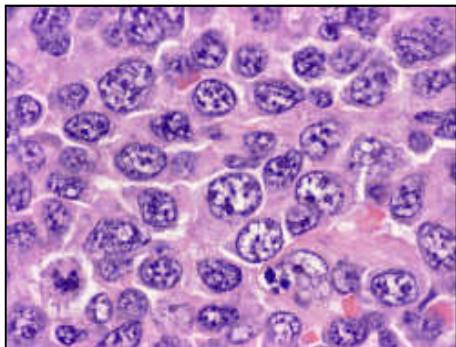
Prognostic groups of chronic lymphocytic leukaemia/lymphocytic lymphoma

		CDR1	CDR2	
VH251 Case 1	EVQLVQSGAEVKPGESLKISCKGSGYSFT	SYWIG WVRQMPGKGLEWMG	IIYPGDSDTRYSPSFQG	QVTISADKSISTAYLQWSSLKASDTAMYYCAR
DP-42 Case 2	EVQLVETGGGLIQPAGSLRLSCAASGFTVS	SNYMS WVRQAPGKGLEWVS	VIY-SGGSTYYADSVKG	RFTISRDNSKNLTYLQMNSLRAEDTAVYYCAR
DP-14 Case 3	QVQLVQSGAEVKPGASVKVSCKASGYTFT	SNGIS WVRQAPGQGLEWMG	WISAYNGNTNYAQKLQG	RVTMTTDSTSTAYMELRSLSRSDDTAVYYCAR
VH4.21 Case 4	QVQLQQWGAGLLKPSETLSLTCAVYGGFS	GYYWS WIRQPPGKGLEWIG	EII-HSGSTNYNPSLKS	RVTISVDTSKNQFSLKLSSVTAADTAVYYCAR
VH26 Case 5	EVQLLESGGGLVQPGGSLRLSCAASGFTFS	SYAMS WVRQAPGKGLEWVS	AISGSGGSTYYADSVKG	RFTISRDNSKNLTYLQMNSLRAEDTAVYYCAR





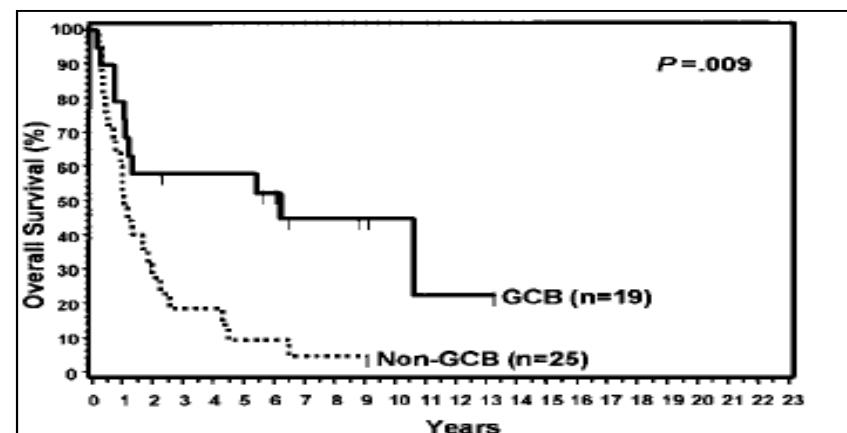
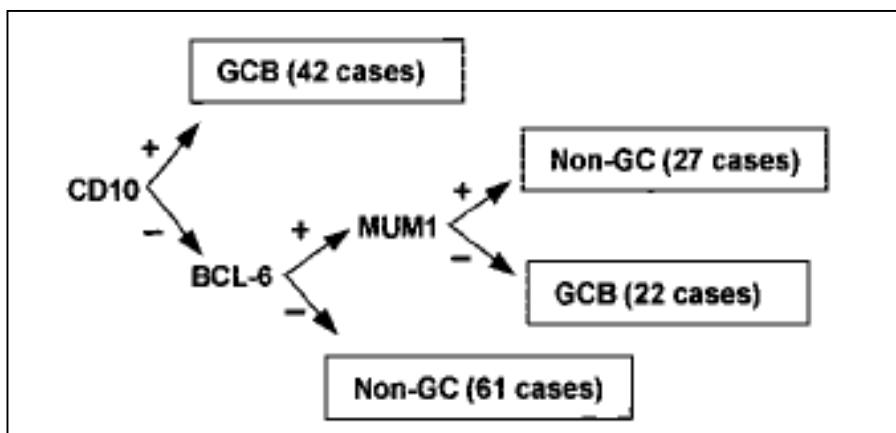
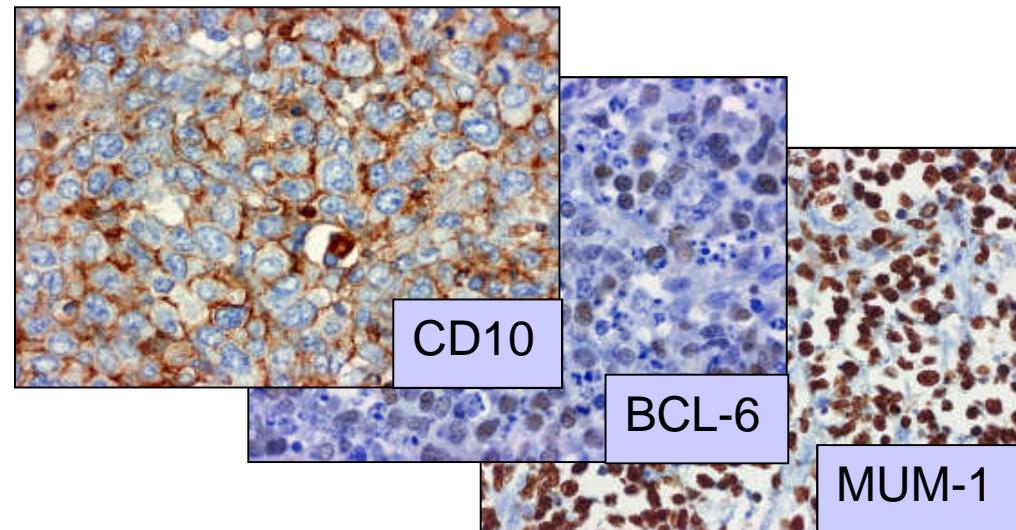
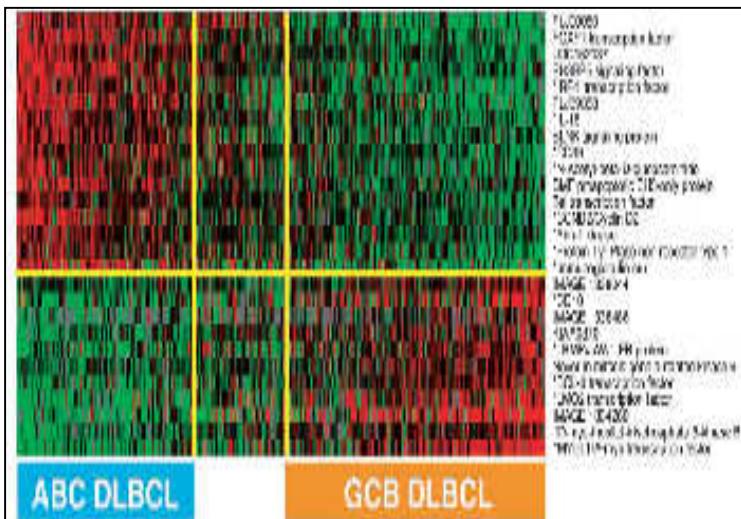
Prognostic groups of DLBCL based on expressional profile analysis



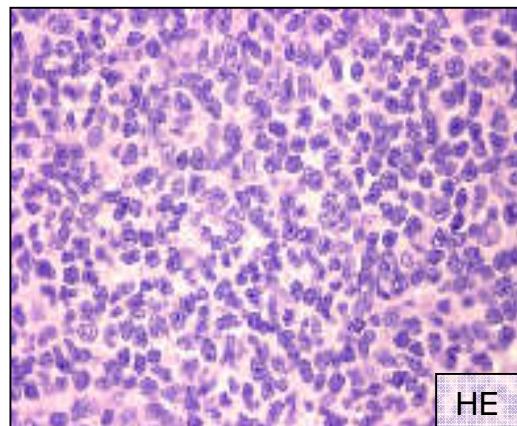
PNAS 2003, 100:9991



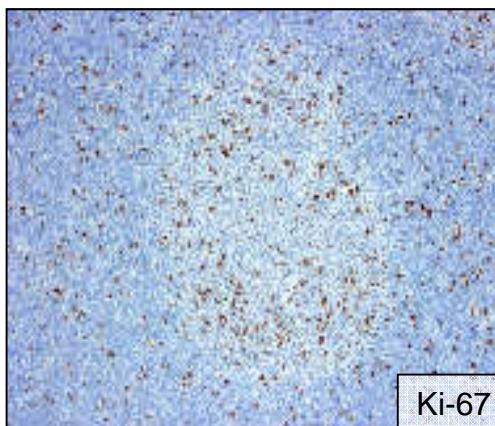
Diagnostic application of the results of expressional profile analysis in DLBCL



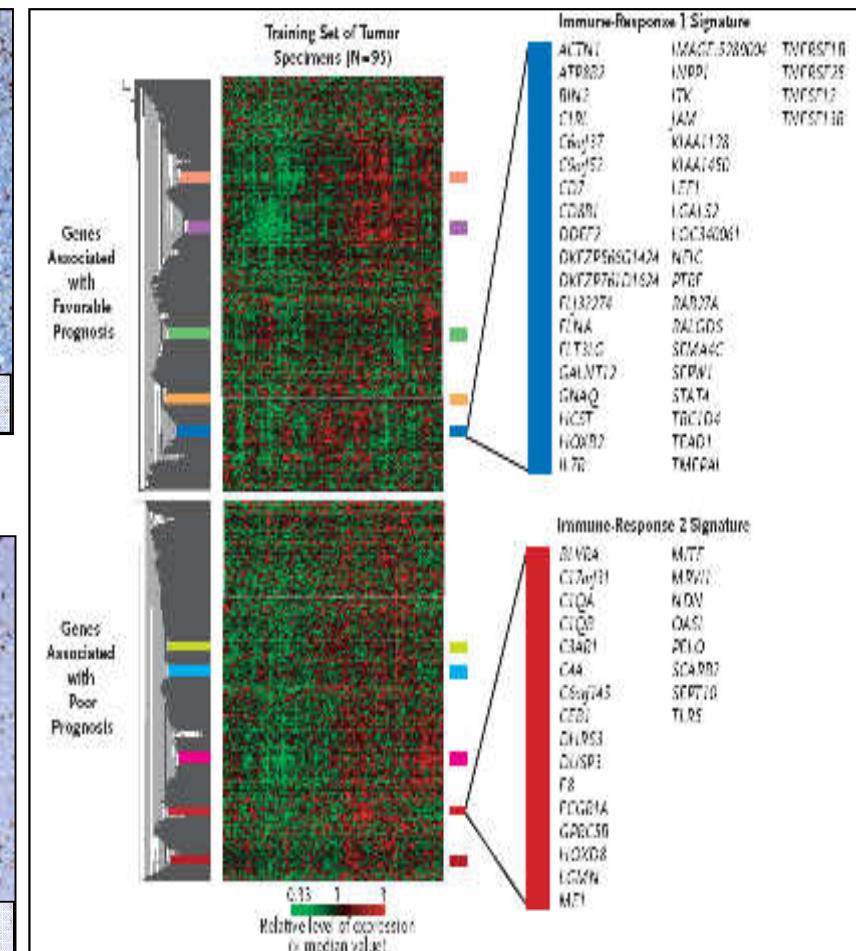
Prognostic factors of follicular lymphoma based on expressional profile analysis



Follicular lymphoma, grade 1



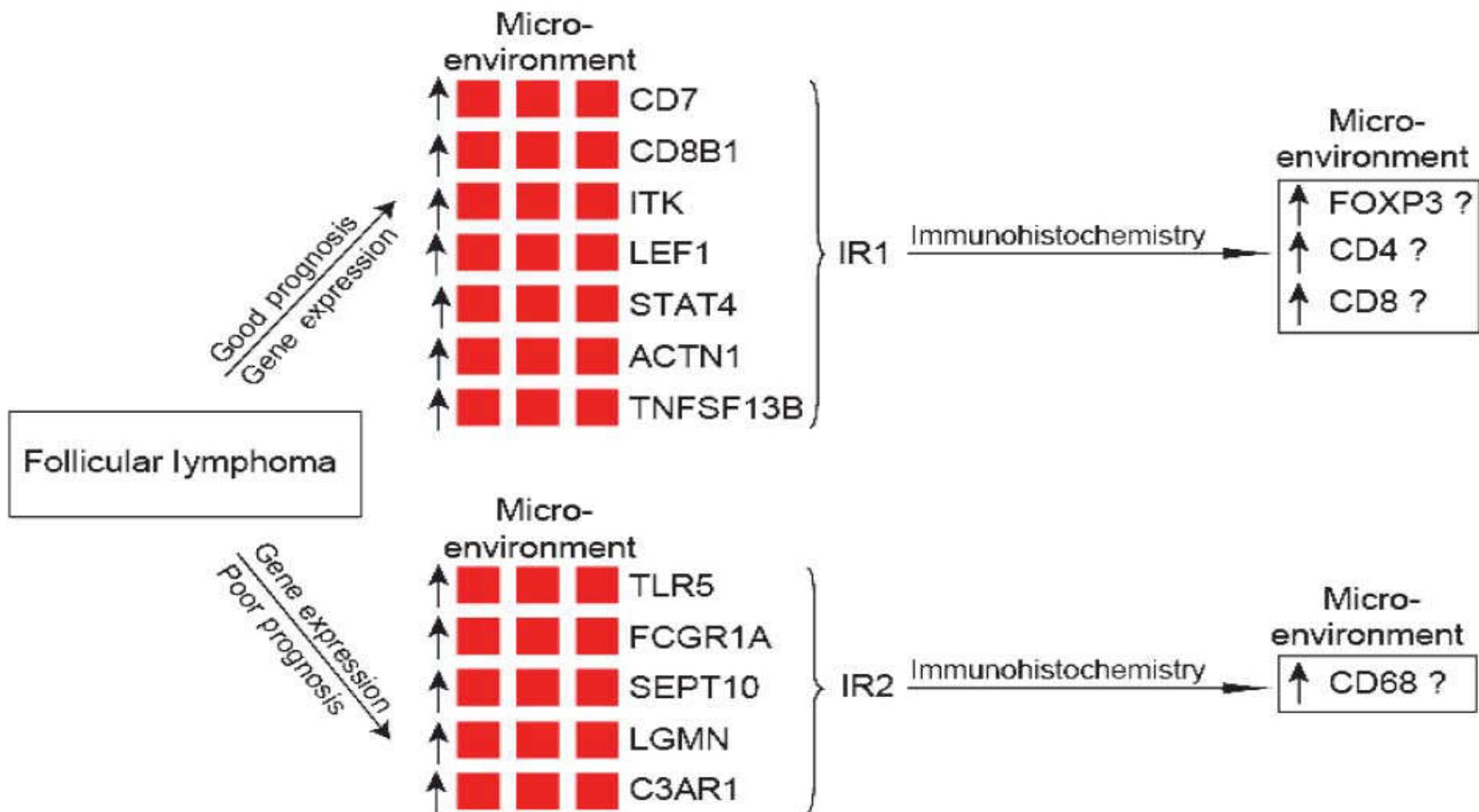
Follicular lymphoma, grade 3



N. Engl. J. Med. 2004, 351:2159



Prognostic factors of follicular lymphoma based on expressional profile analysis



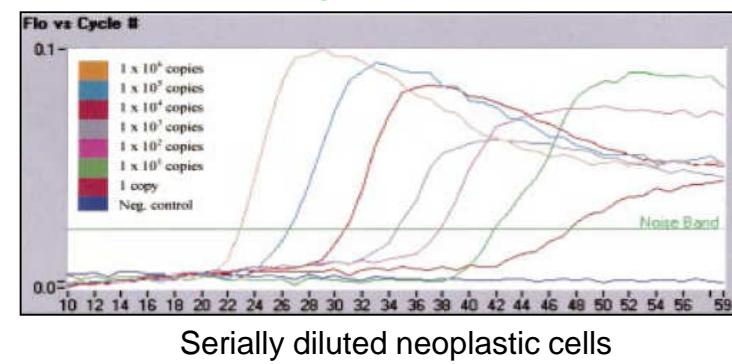
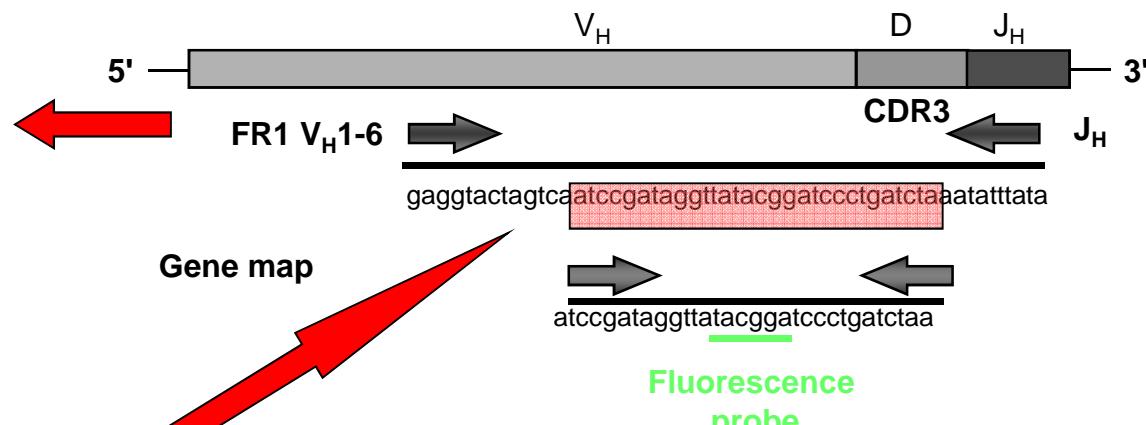
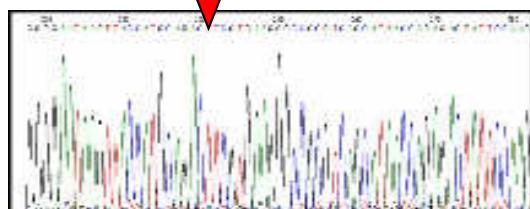
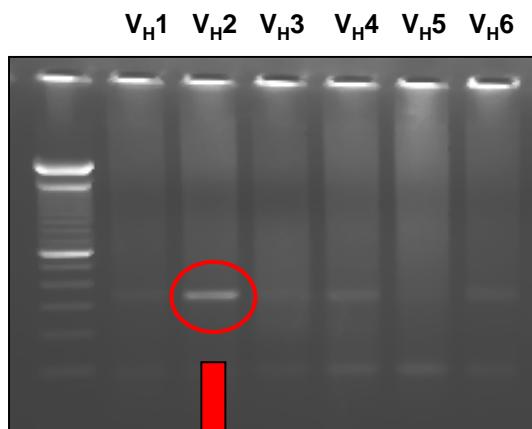
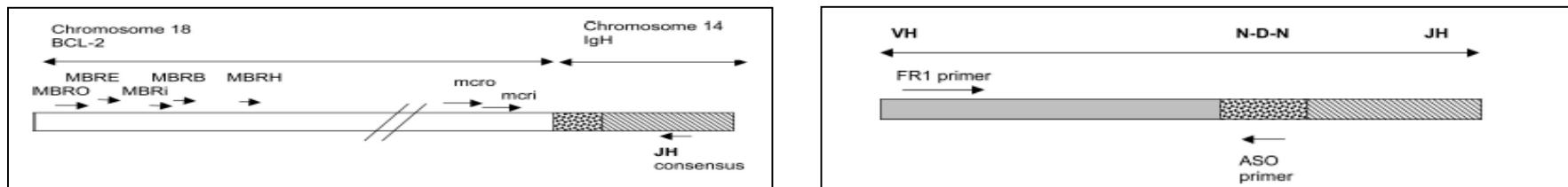


The role of molecular analysis in lymphoma diagnostics

- Determine clonality of the lymphoid cells
- Determine subtype of lymphoma
- Determine prognosis of lymphoma
- **Determine effectiveness of treatment (MRD)**



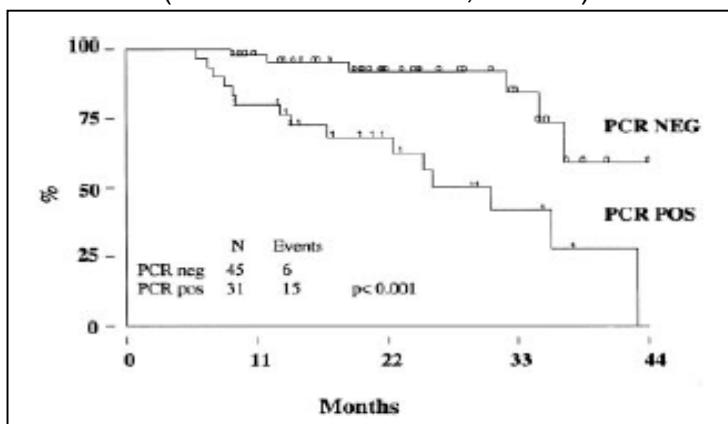
Real-time quantitative allele-specific oligonucleotide PCR (RQ-ASO PCR) for assessing MRD in lymphoma



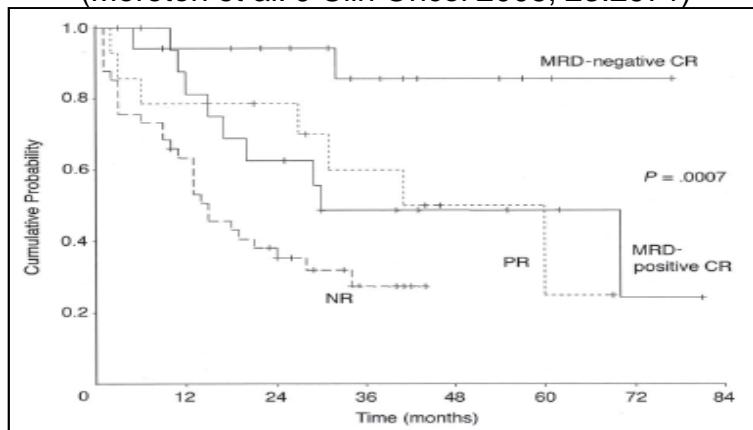


MRD detection following immunochemotherapy of different lymphomas

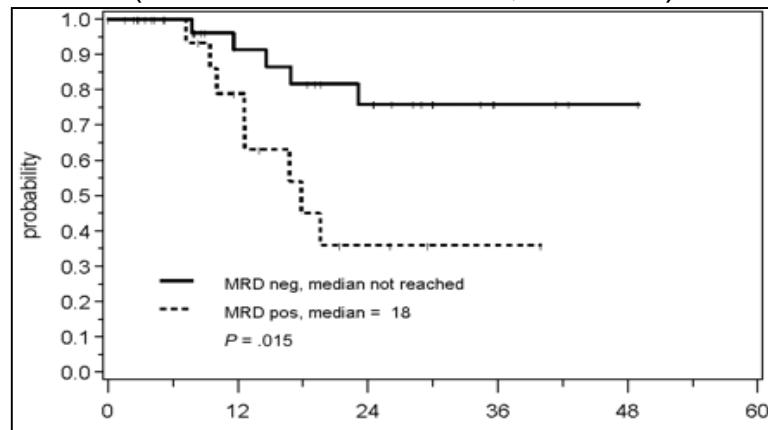
CHOP and Rituximab therapy in FL
(Rambaldi et al. 2002, 99:856)



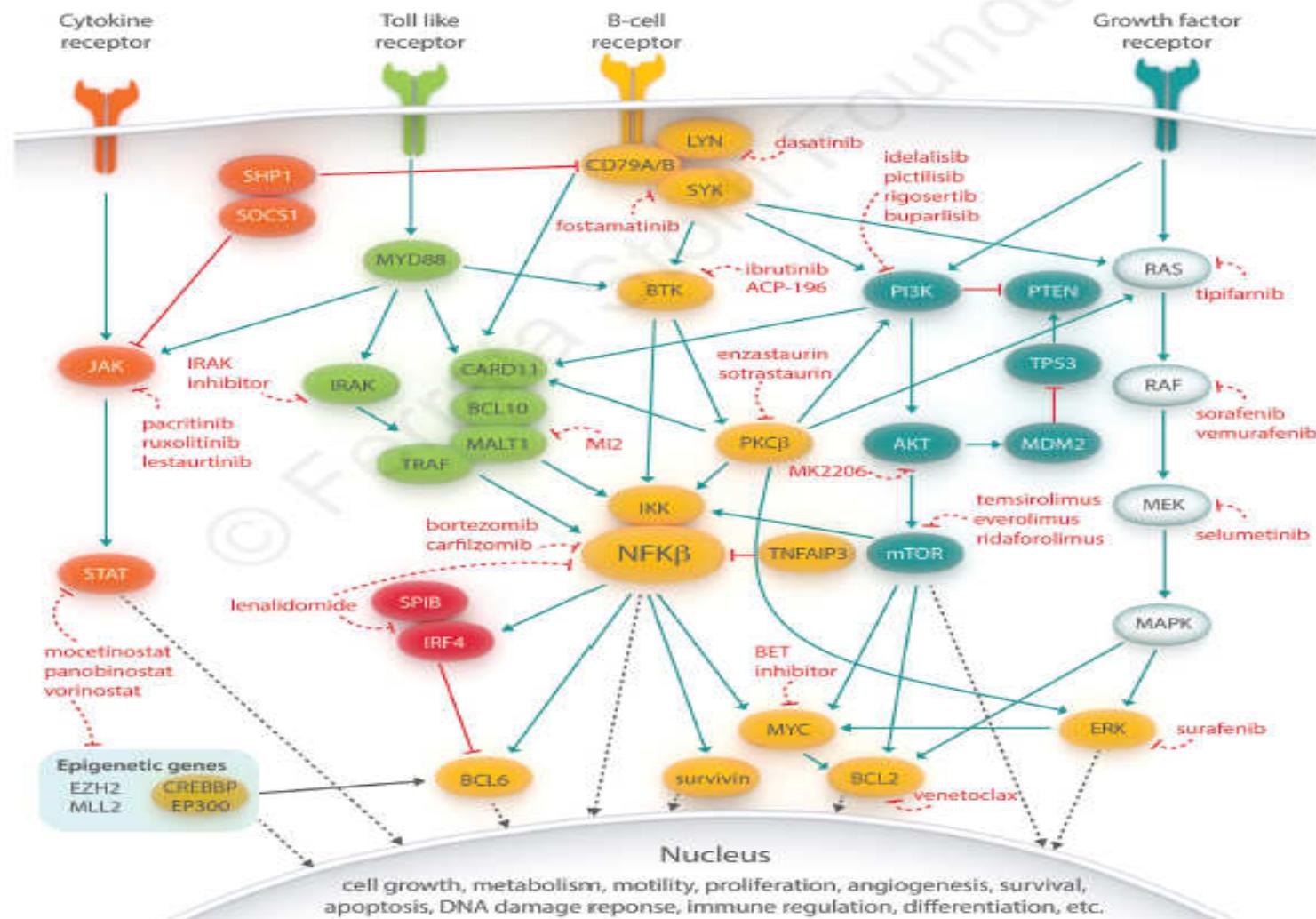
Alemtuzumab therapy in CLL
(Moreton et al. J Clin Oncol 2005, 23:2971)



Immunotherapy in MCL
(Pott et al. Blood. 2010, 22:3215)



The future of lymphoma diagnostics



The effect of molecular analysis in lymphoma diagnostics

