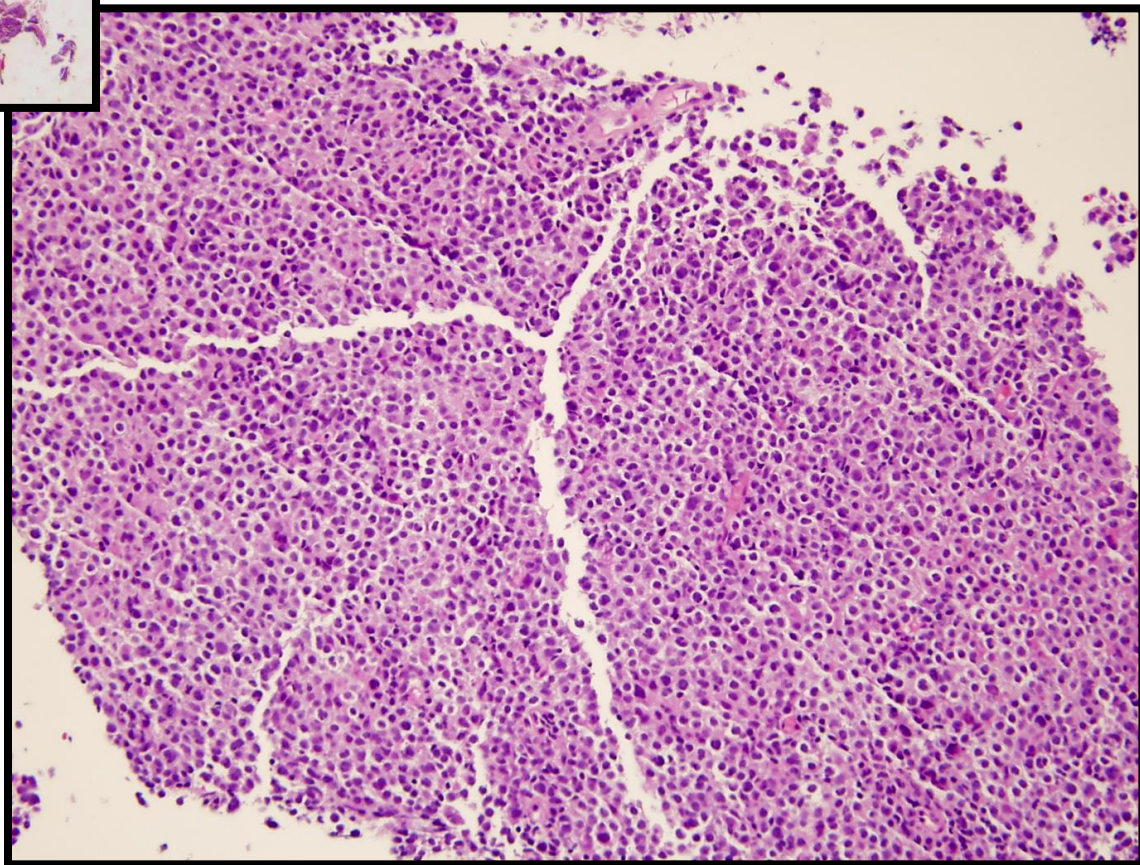
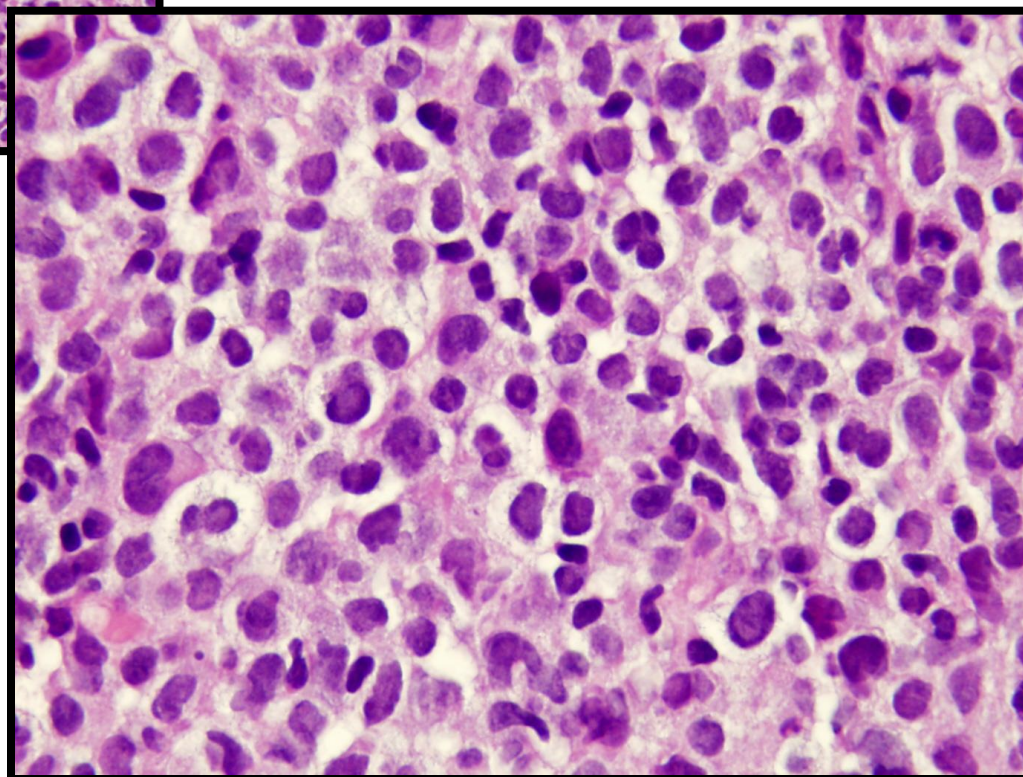
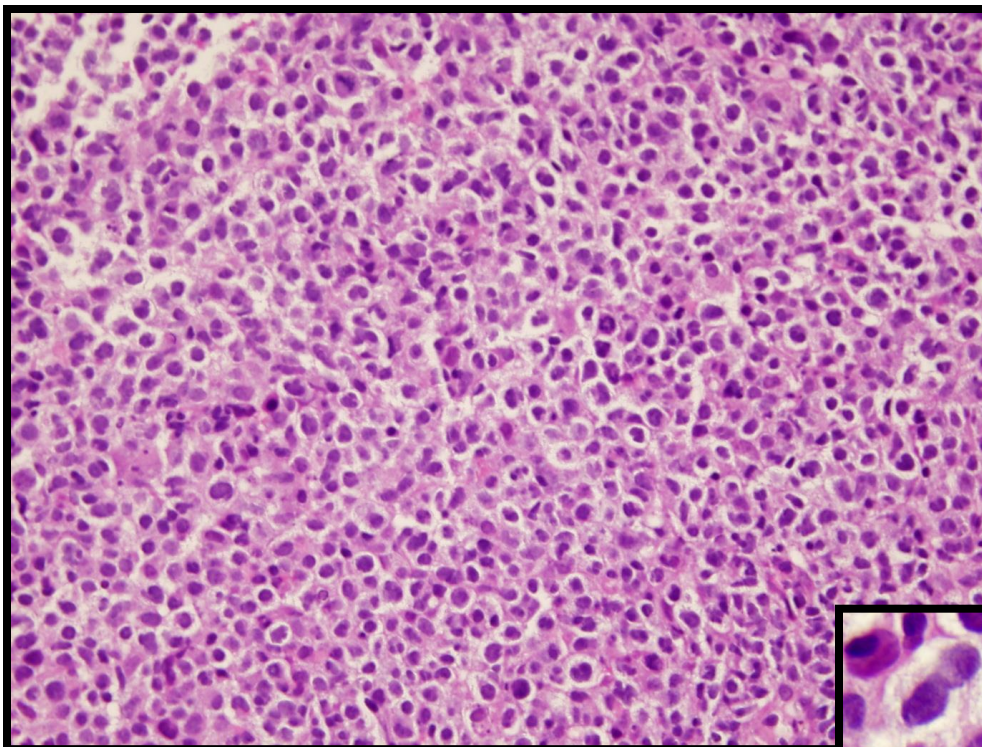
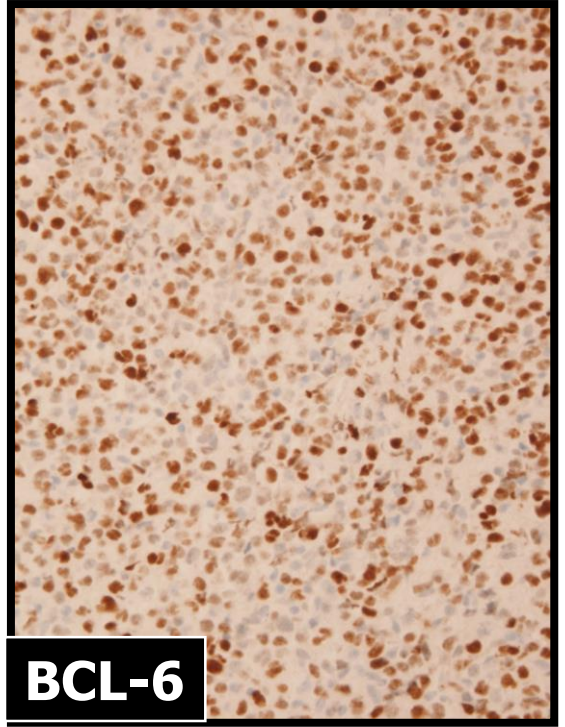
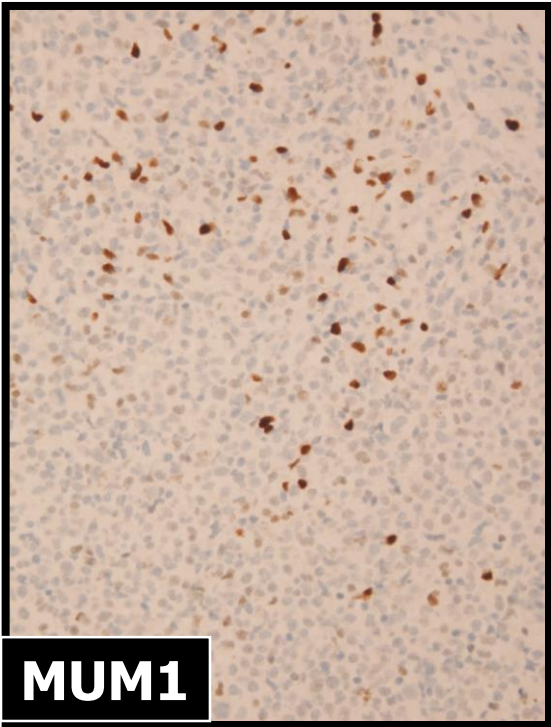
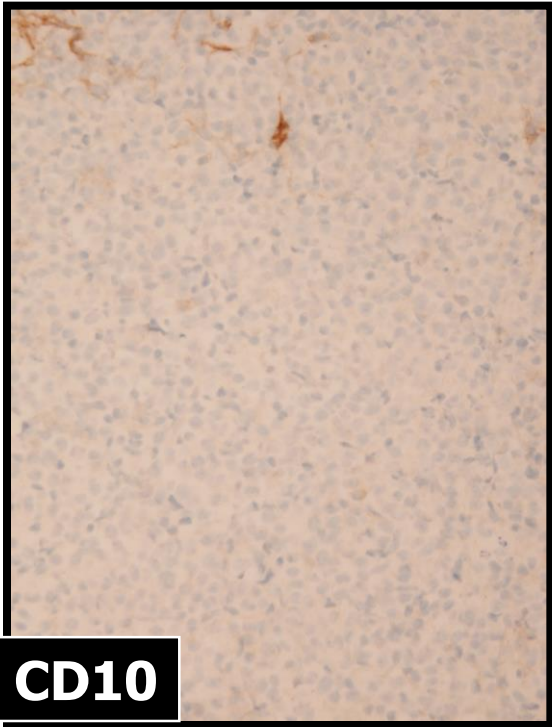
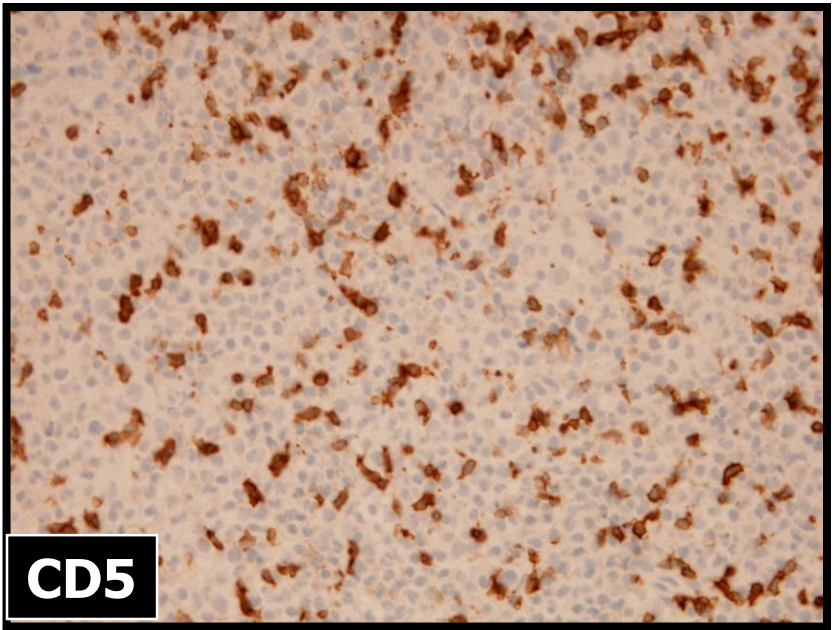
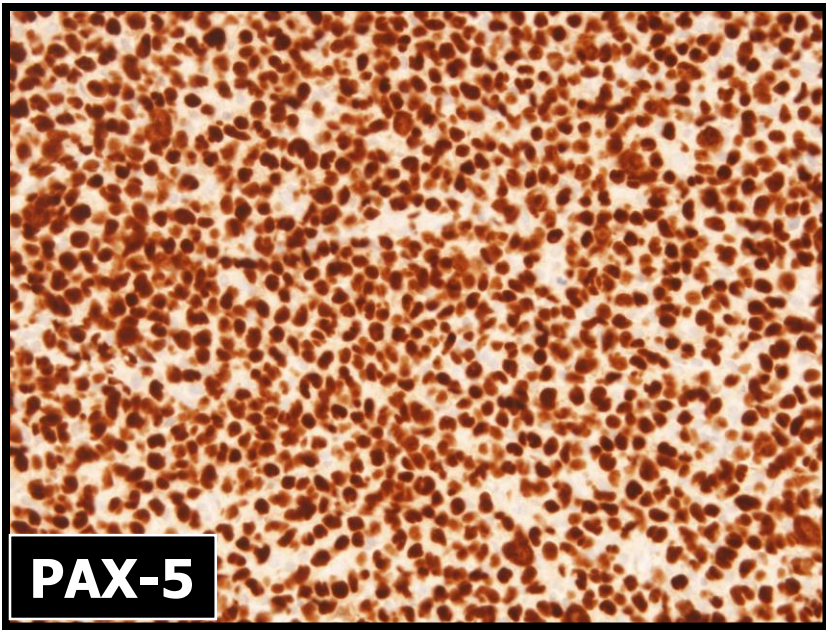


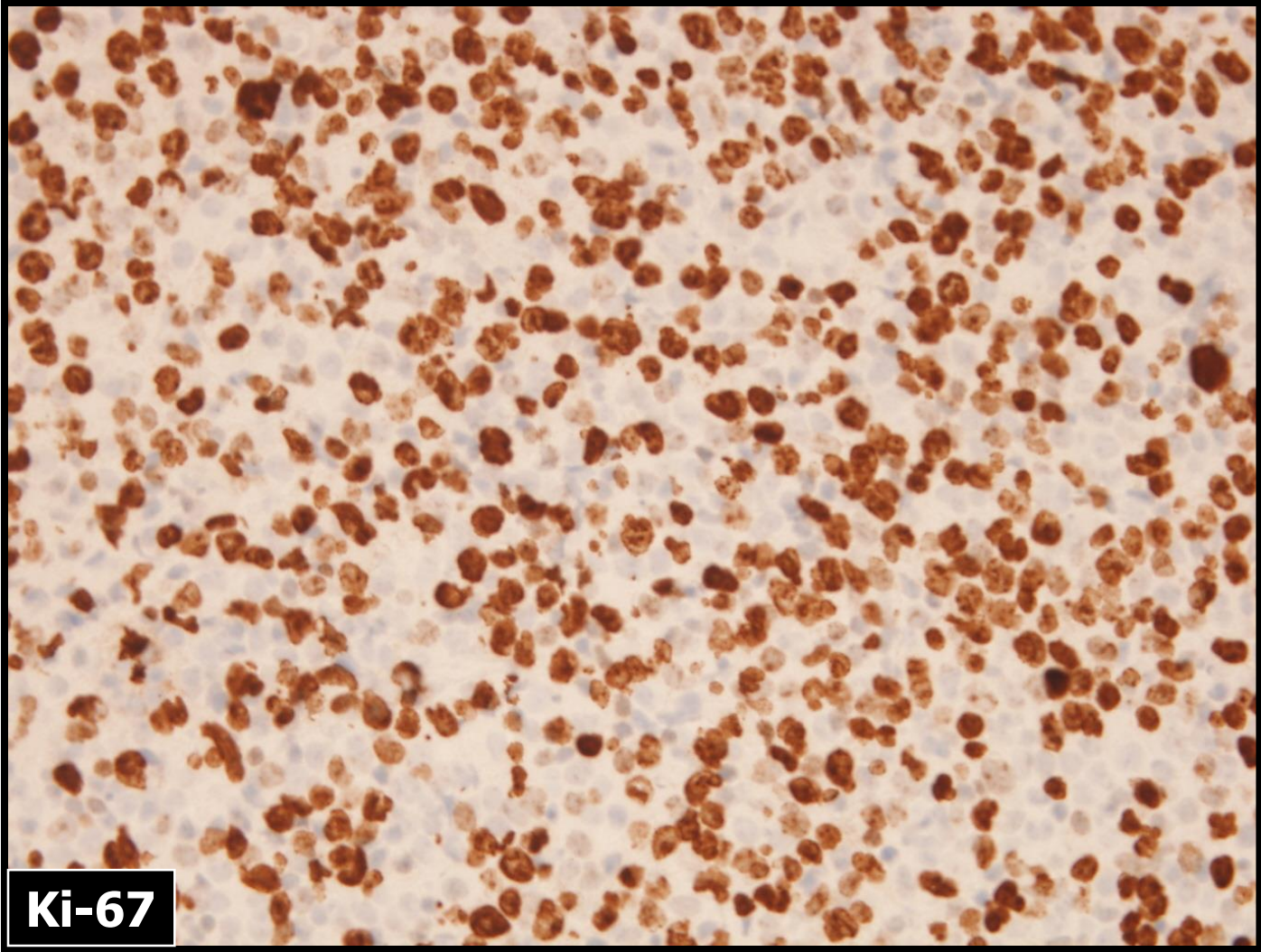
## **CASE 4**

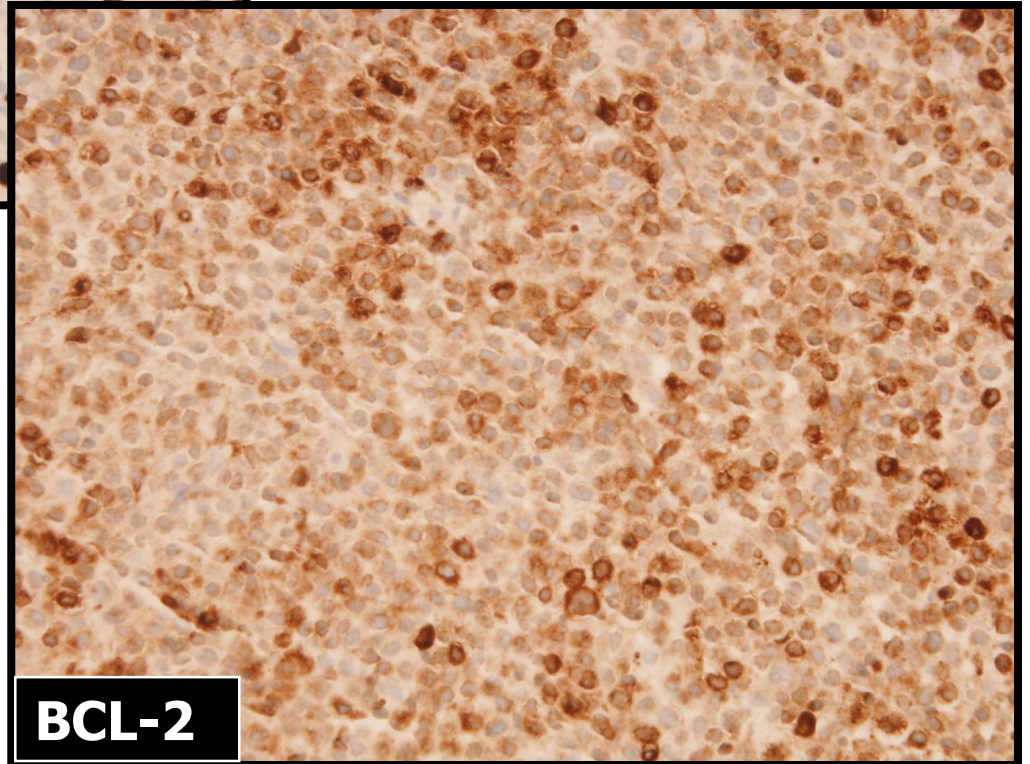
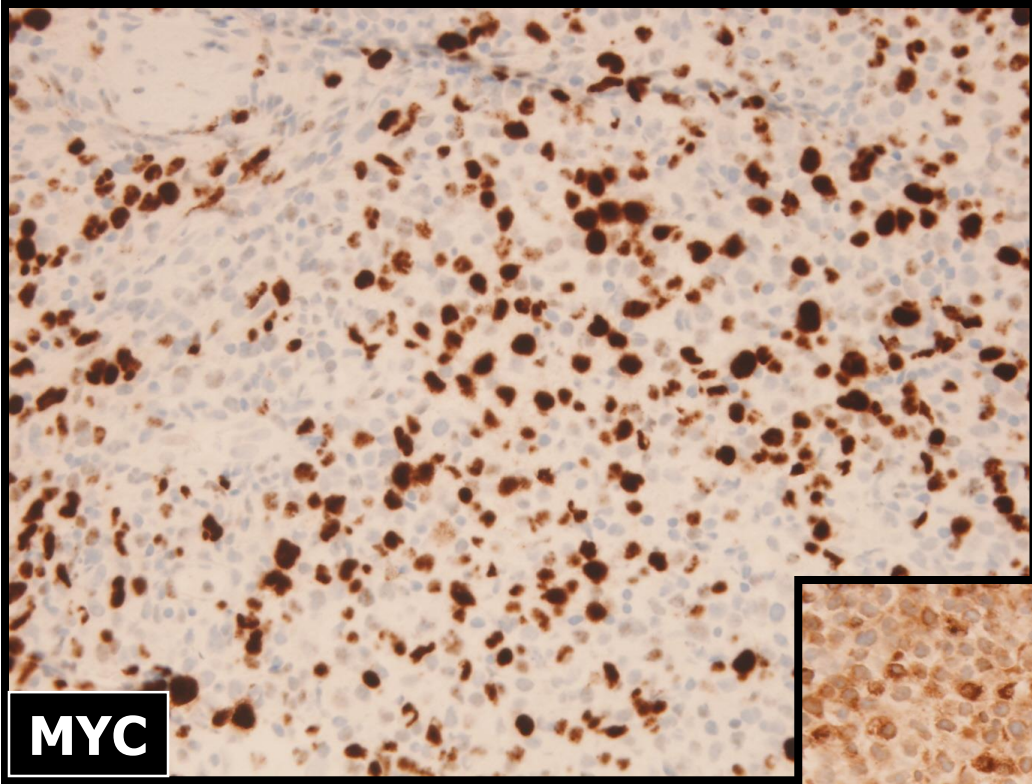
**A 70-year-old woman with a history of diabetes, hypertension and kidney failure presented with acute onset of fatigue and dizziness. Physical examination showed B-type symptoms. Laboratory evaluation showed hypercalcemia. PET/CT showed lymphadenopathy, splenomegaly and bone lesions. This is a right axillary lymph node needle biopsy.**











# **DIAGNOSIS (CASE 4)**

**Diffuse large B-cell lymphoma**

**The neoplasm also has:**

**Germinal center B-cell immunophenotype**

**Double expressor immunophenotype**

**Ki-67 ~70%**

***MYC* not rearranged**

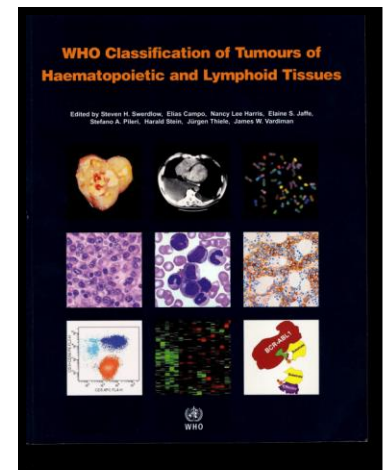
***BCL6* rearranged**

***BCL2* not rearranged**

# Diffuse Large B-cell Lymphoma

## Definition

**DLBCL is a neoplasm with a diffuse growth pattern composed of medium or large B lymphoid cells with nuclear size equal to or exceeding normal macrophage nuclei, or more than twice the size of normal lymphocyte nuclei**





# **WHO Classification of Diffuse Large B-cell Lymphoma (2017)**

## **Diffuse large B-cell lymphoma, NOS**

**GCB versus ABC/non-GCB**

**CD5**

## **Other lymphomas of large B-cells**

**T-cell/histiocyte-rich large B-cell lymphoma**

**Primary DLBCL of the central nervous system**

**Primary cutaneous DLBCL, leg-type**

**Primary mediastinal (thymic) large B-cell lymphoma**

**Intravascular large B-cell lymphoma**

**DLBCL associated with chronic inflammation**

**Lymphomatoid granulomatosis**

**EBV+ diffuse large B-cell lymphoma**

**ALK+ large B-cell lymphoma**

**Plasmablastic lymphoma**

**HHV8+ lymphoproliferative disorders**

**Primary effusion lymphoma**

## **Borderline cases**

**High-grade B-cell lymphoma (NOS versus double hit)**

**B-cell lymphoma, unclassifiable, intermediate between DLBCL & CHL**

# Diffuse Large B-cell Lymphoma

## R-CHOP is Standard Frontline Therapy

**R**ituximab  
**C**yclophosphamide  
**H**ydroxydaunorubicin/Adriamycin  
**O**ncovin/vincristine  
**P**rednisone

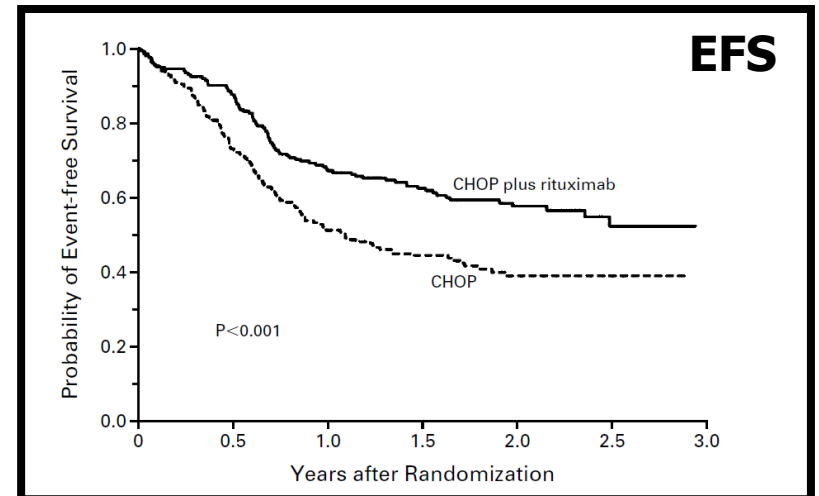
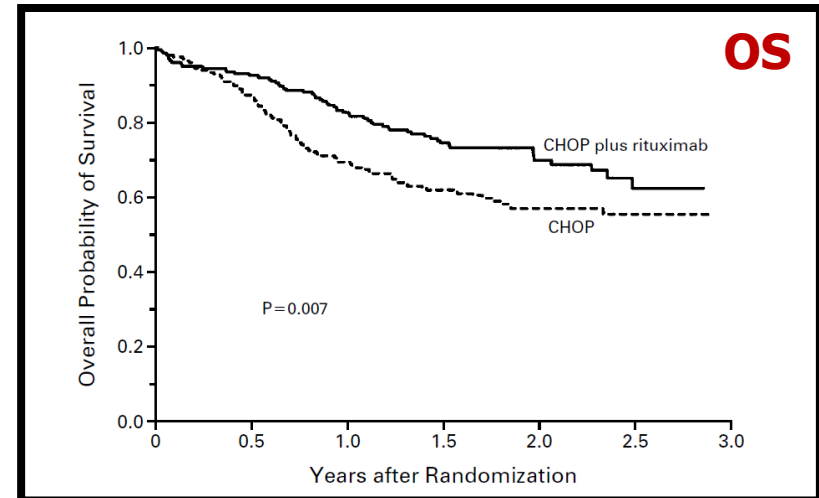


**Bertrand Coiffier, MD**

CHOP CHEMOTHERAPY PLUS RITUXIMAB COMPARED WITH CHOP ALONE  
IN ELDERLY PATIENTS WITH DIFFUSE LARGE-B-CELL LYMPHOMA

BERTRAND COIFFIER, M.D., ERIC LEPAGE, M.D., PH.D., JOSETTE BRIÈRE, M.D., RAOUL HERBRECHT, M.D., HERVÉ TILLY, M.D.,  
REDA BOUABDALLAH, M.D., PIERRE MOREL, M.D., ERIC VAN DEN NESTE, M.D., GILLES SALLES, M.D., PH.D.,  
PHILIPPE GAULARD, M.D., FELIX REYES, M.D., AND CHRISTIAN GISSELBRECHT, M.D.

**N Engl J Med 346: 235, 2002**



# **Diffuse Large B-cell Lymphoma NOS**

## **Clinical Findings**

<b>Median age</b>	<b>64 y (wide range)</b>
<b>Male</b>	<b>55%</b>
<b>Stage I-II</b>	<b>54%</b>
<b>III-IV</b>	<b>46%</b>
<b>B symptoms</b>	<b>33%</b>
<b>BM involved</b>	<b>16%</b>
<b>IPI 0-1</b>	<b>35%</b>
<b>2-3</b>	<b>46%</b>
<b>4-5</b>	<b>19%</b>

# Diffuse Large B-cell Lymphoma

## International Prognostic Index

<b>A</b> ge	$\leq 60$ vs. $>60$ years
<b>P</b> erformance status	0-1 vs. 2-4
<b>L</b> DH	Normal vs elevated
<b>E</b> xtranodal sites	$\leq 1$ vs $>1$ site
<b>S</b> tage	I-II vs III-IV

# An enhanced International Prognostic Index (NCCN-IPI) for patients with diffuse large B-cell lymphoma treated in the rituximab era

Zheng Zhou,<sup>1</sup> Laurie H. Sehn,<sup>2</sup> Alfred W. Rademaker,<sup>1</sup> Leo I. Gordon,<sup>1</sup> Ann S. LaCasce,<sup>3</sup> Allison Crosby-Thompson,<sup>3</sup> Ann Vanderplas,<sup>4</sup> Andrew D. Zelenetz,<sup>5</sup> Gregory A. Abel,<sup>3</sup> Maria A. Rodriguez,<sup>6</sup> Auayporn Nademanee,<sup>7</sup> Mark S. Kaminski,<sup>8</sup> Myron S. Czuczman,<sup>9</sup> Michael Millenson,<sup>10</sup> Joyce Niland,<sup>4</sup> Randy D. Gascoyne,<sup>2</sup> Joseph M. Connors,<sup>2</sup> Jonathan W. Friedberg,<sup>11</sup> and Jane N. Winter<sup>1</sup>

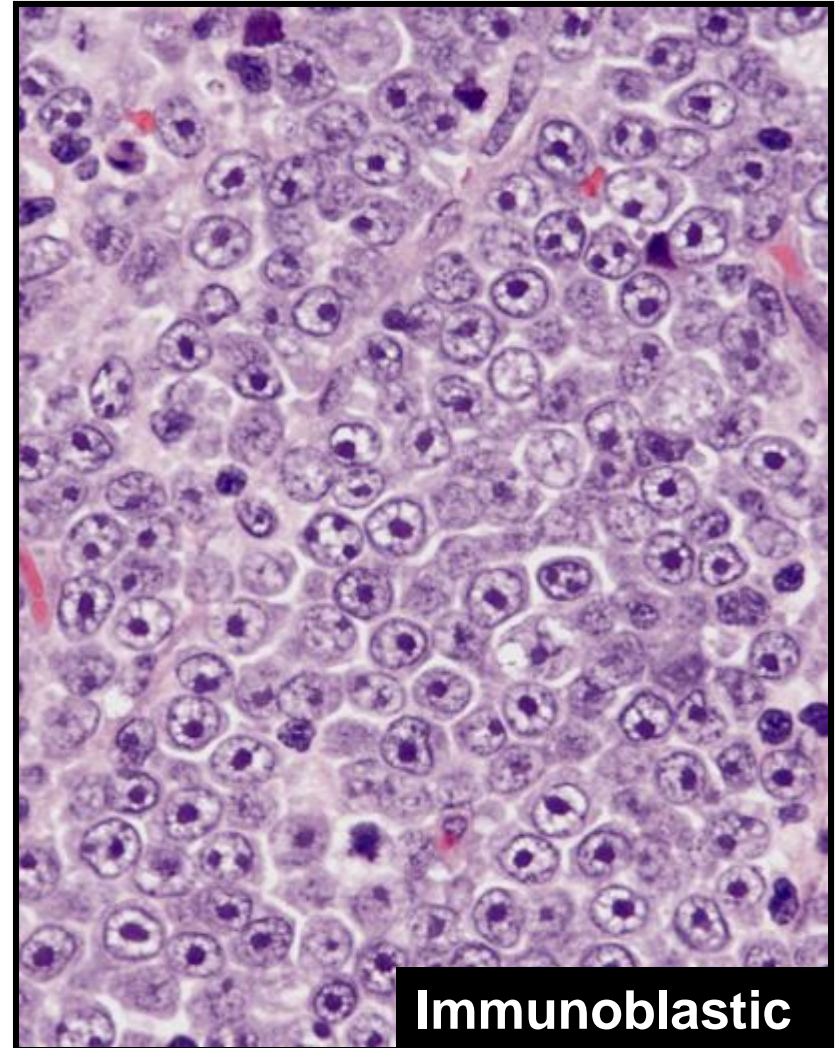
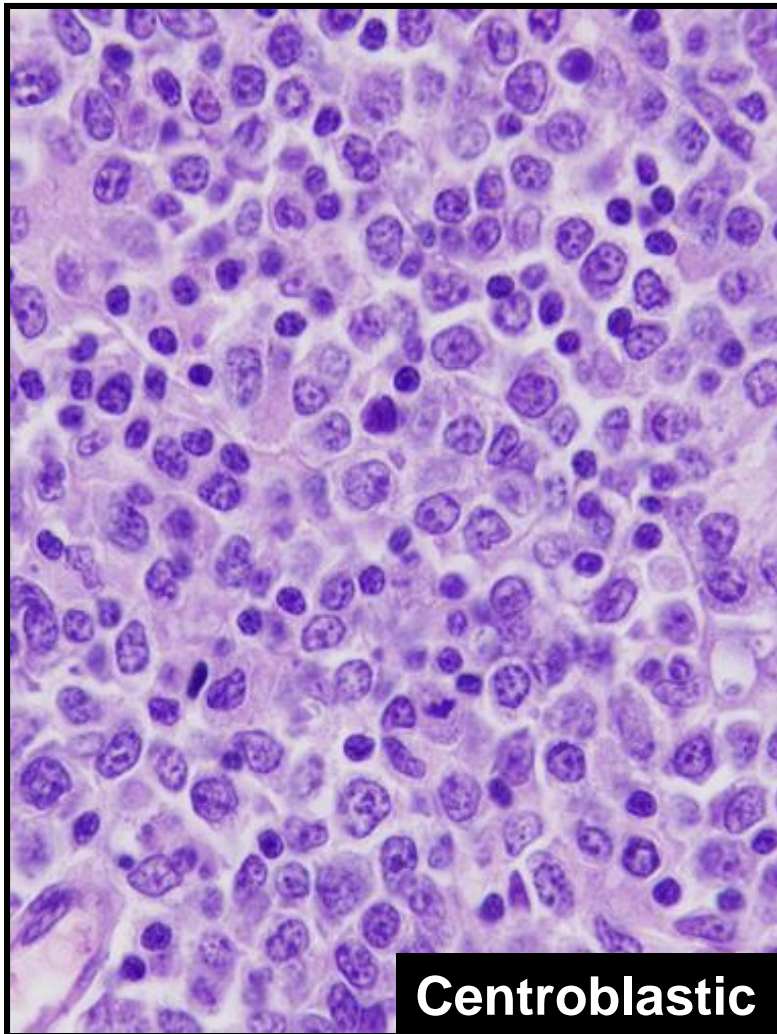
**Table 3. The NCCN-IPI**

<b>NCCN-IPI</b>	<b>Score</b>
<b>Age, y</b>	
>40 to ≤60	1
>60 to ≤75	2
>75	3
<b>LDH, normalized</b>	
>1 to ≤3	1
>3	2
Ann Arbor stage III-IV	1
Extranodal disease*	1
Performance status ≥2	1

\*Disease in bone marrow, CNS, liver/GI tract, or lung.

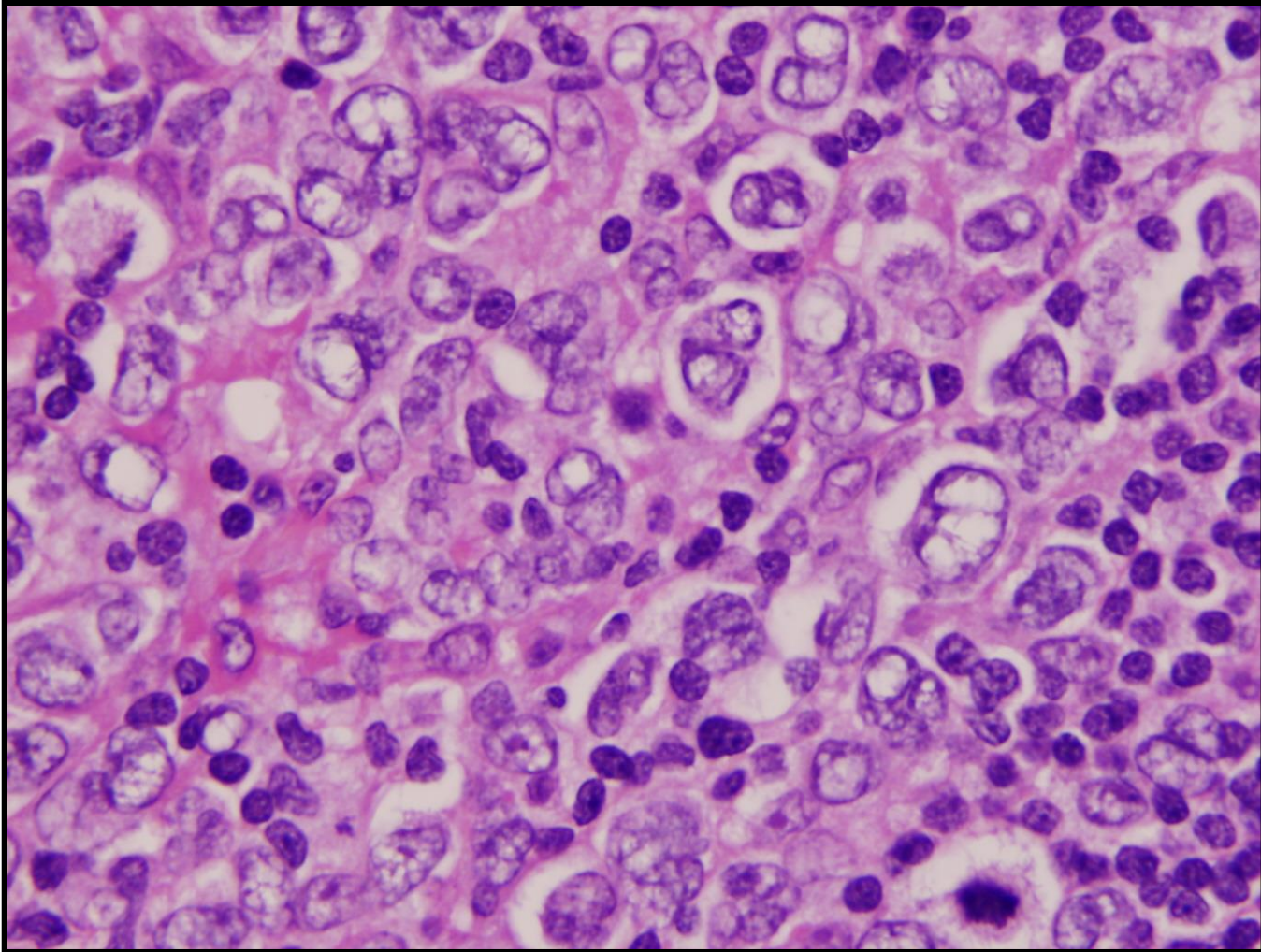
# Diffuse Large B-cell Lymphoma NOS

## Morphologic Variants



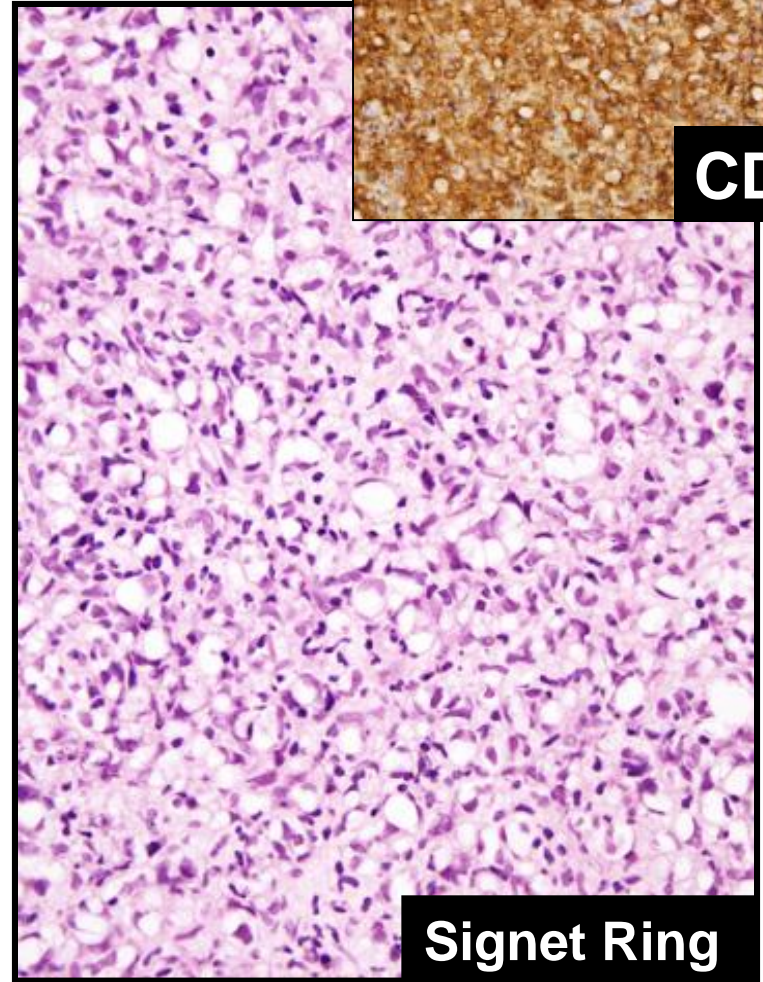
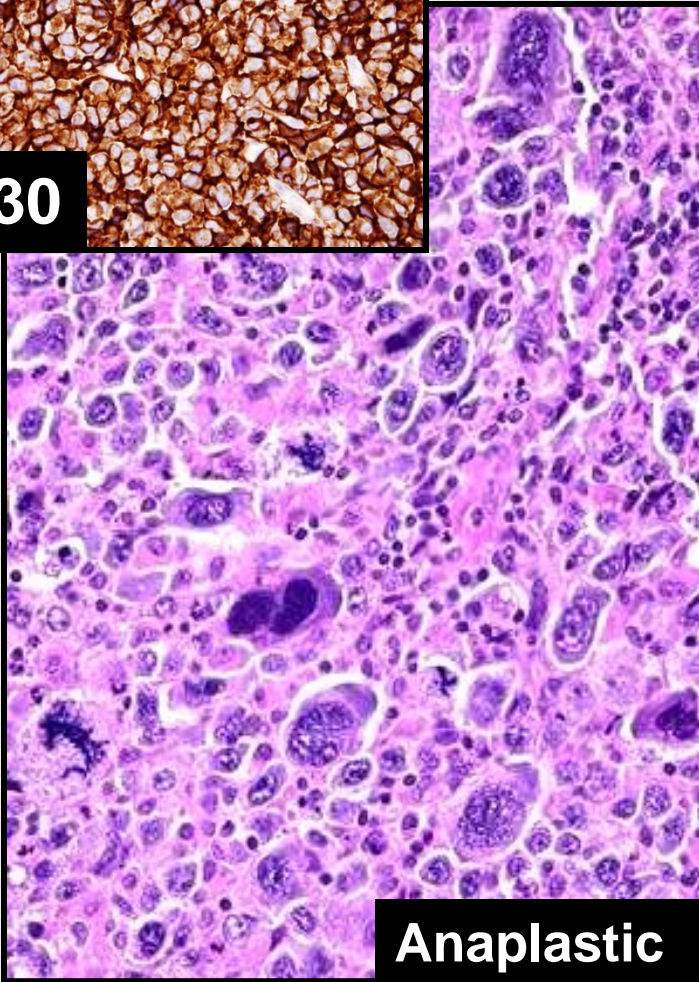
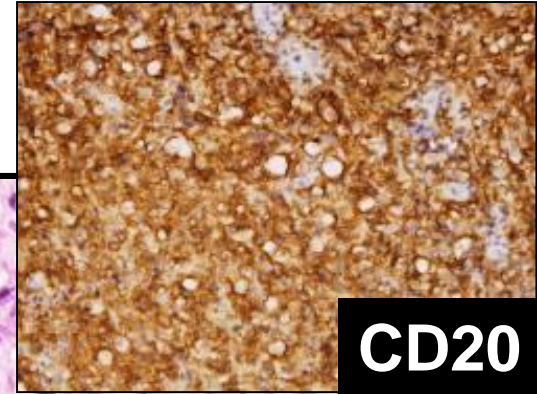
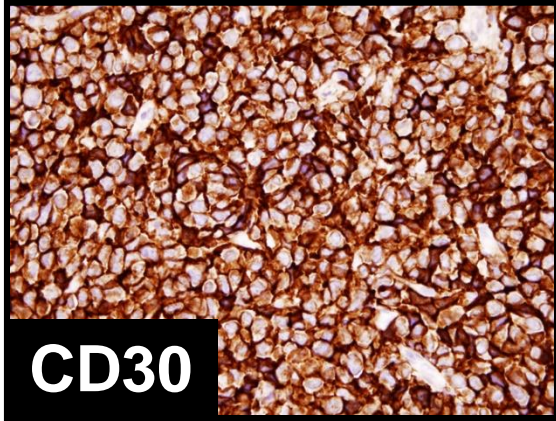
# Diffuse Large B-cell Lymphoma NOS

## Multilobated Variant



# Diffuse Large B-cell Lymphoma NOS

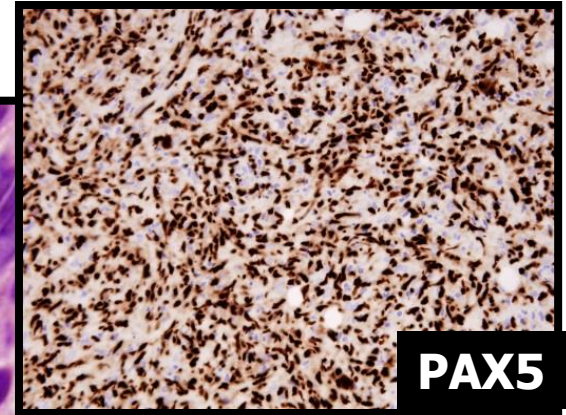
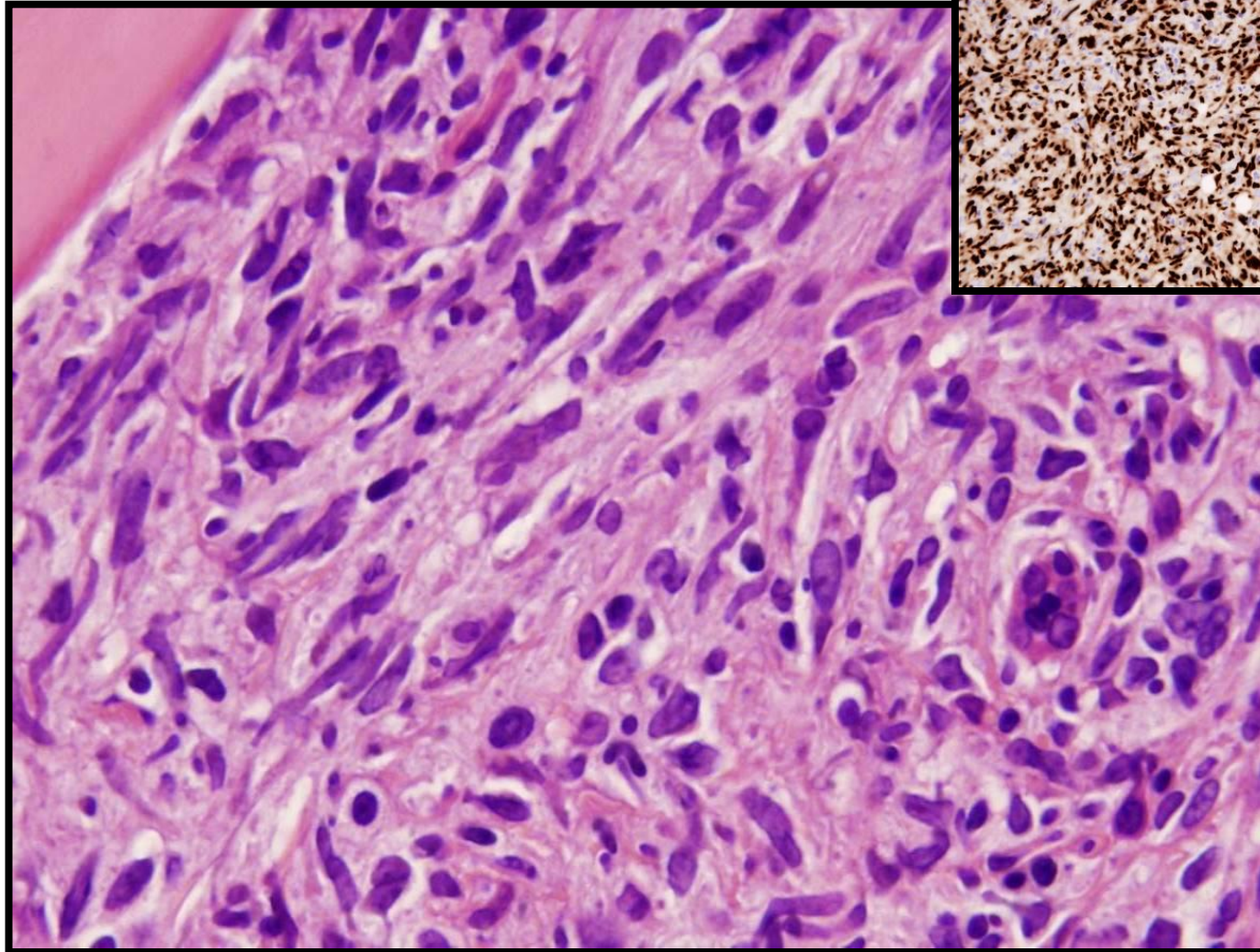
## Morphologic Variants





# Diffuse Large B-cell Lymphoma NOS

## Spindle Cell Variant



# **Diffuse Large B-cell Lymphoma NOS**

## **Morphologic Variants**

### **Common**

**Centroblastic (~80%)**

**Immunoblastic (~10%)**

**Multilobated (<5%)**

**Anaplastic (<5%)**

### **Rare**

**Sinusoidal**

**Spindled**

**Myxoid**

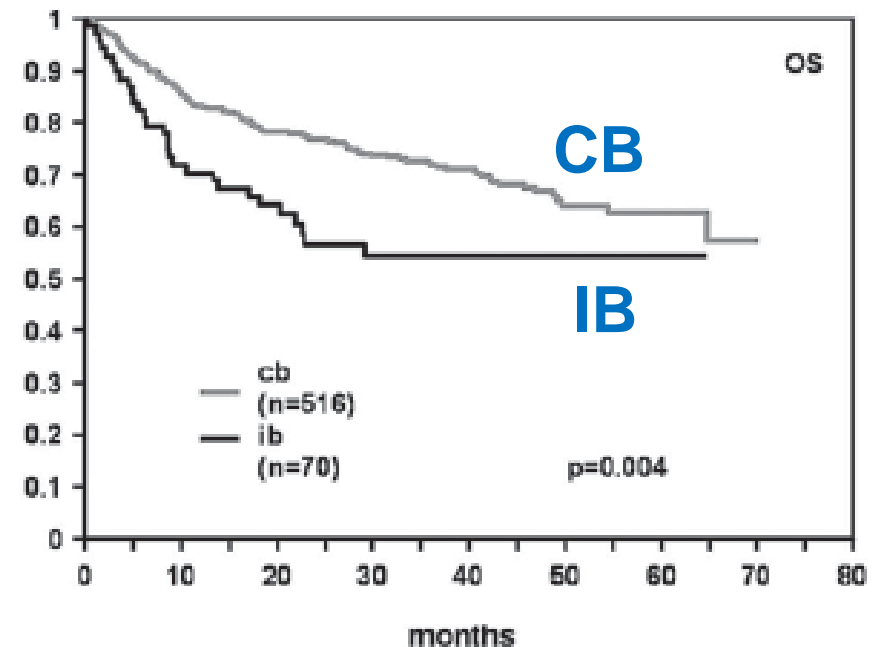
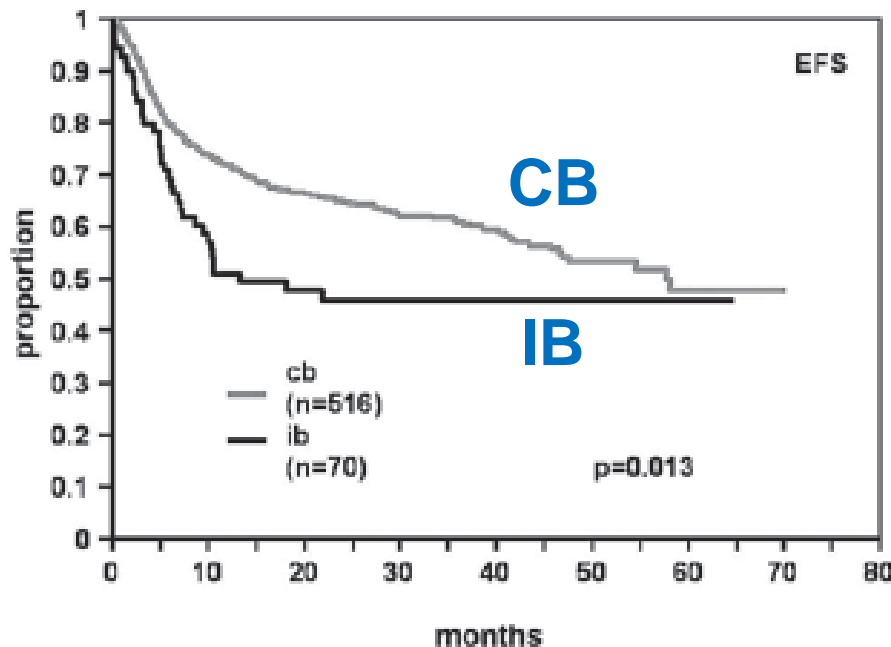
**Signet Ring**

**Rosettes**

**Does morphology correlate with prognosis ?**

# Immunoblastic morphology but not the immunohistochemical GCB/nonGCB classifier predicts outcome in diffuse large B-cell lymphoma in the RICOVER-60 trial of the DSHNHL

German Ott,<sup>1,2</sup> Marita Ziepert,<sup>3</sup> Wolfram Klapper,<sup>4</sup> Heike Horn,<sup>2</sup> Monika Szczepanowski,<sup>4</sup> Heinz-Wolfram Bernd,<sup>5</sup> Christoph Thorns,<sup>5</sup> Alfred C. Feller,<sup>5</sup> Dido Lenze,<sup>6</sup> Michael Hummel,<sup>6</sup> Harald Stein,<sup>6</sup> Hans-Konrad Müller-Hermelink,<sup>1</sup> Matthias Frank,<sup>7</sup> Martin-Leo Hansmann,<sup>7</sup> Thomas F. E. Barth,<sup>8</sup> Peter Möller,<sup>8</sup> Sergio Cogliatti,<sup>9</sup> Michael Pfreundschuh,<sup>10</sup> Norbert Schmitz,<sup>11</sup> Lorenz Trümper,<sup>12</sup> Markus Loeffler,<sup>3</sup> and Andreas Rosenwald<sup>1</sup>



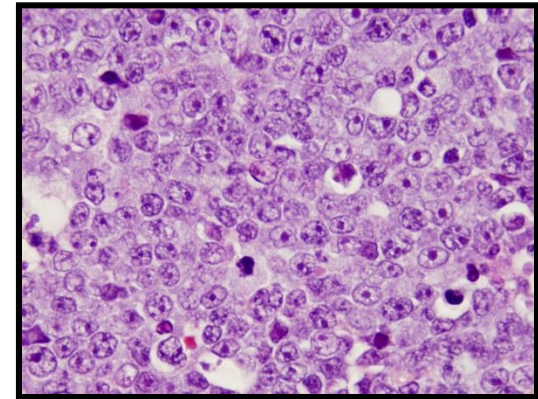
# Diffuse Large B-cell Lymphomas of Immunoblastic Type Are a Major Reservoir for *MYC-IGH* Translocations

Heike Horn, PhD,\* Annette M. Staiger, MSc,\* Matthias Vöhringer, MD,† Ulrich Hay, MD,‡  
Elias Campo, MD,§ Andreas Rosenwald, MD,|| German Ott, MD,\* and M. Michaela Ott, MD¶

**The authors assessed 107 DLBCL using FISH with  
MYC breakapart and *MYC-IGH* fusion probes**

***MYC* translocations detected in**

**13 / 39 (33%) immunoblastic  
5 / 68 (7%) centroblastic**



**All immunoblastic DLBCL with *MYC* translocations  
had *MYC-IGH* fusions**

# **Immunophenotyping of DLBCL**

## **What Is The Purpose ?**

**In the past**

**Diagnosis**

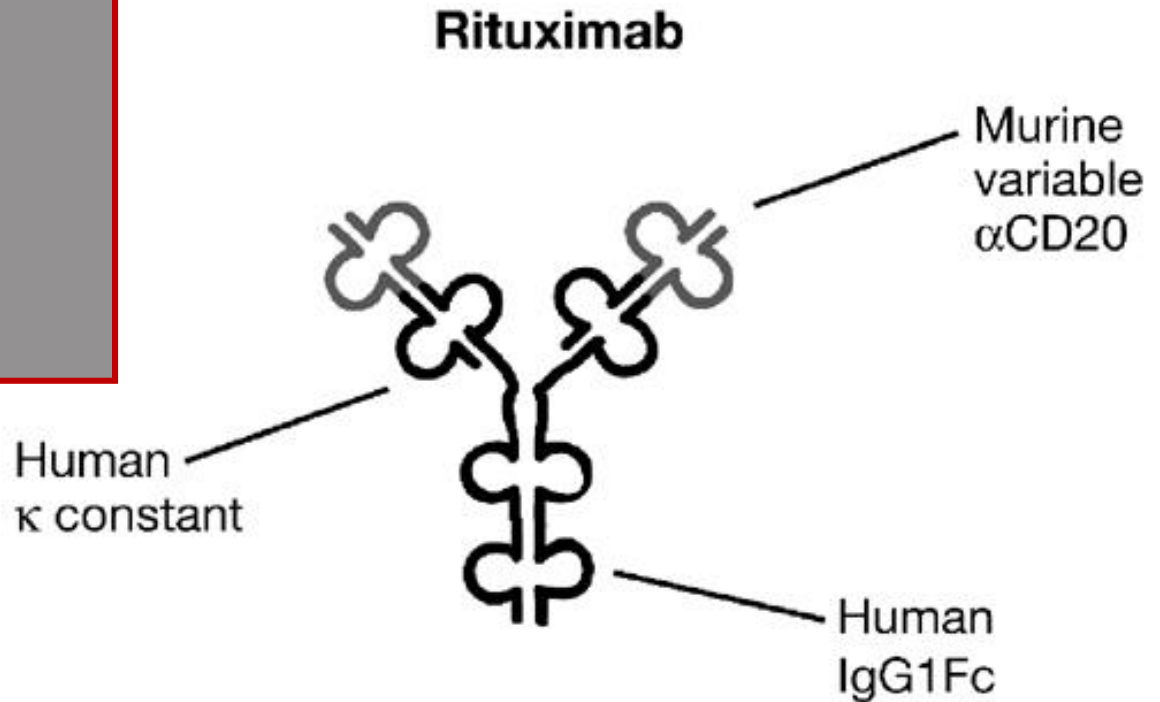
**Currently**

**Diagnosis**

**Prognosis**

**Identifying targets for therapy**

# Monoclonal Antibodies are Being Added to Standard Therapy



**CD20 is used for diagnosis and is a therapeutic target**

# Potential Targets Assessable by IHC

<b>Target</b>	<b>Drug</b>	<b>Pathway</b>
<b>CD19</b>	<b>Tafasitamab</b>	<b>B-cell receptor signaling</b>
<b>CD30</b>	<b>Brentuximab vedotin</b>	<b>NF-κB</b>
<b>CD38</b>	<b>Daratumumab</b>	<b>Cell migration, adhesion, signaling</b>
<b>CD79A</b>	<b>Polatuzumab vedotin</b>	<b>B-cell receptor signaling</b>
<b>BTK</b>	<b>Ibrutinib</b>	<b>B-cell receptor signaling</b>
<b>XPO1</b>	<b>Selinexor</b>	<b>Selective inhibitor of nuclear export</b>
<b>BRAF, MEK</b>	<b>Vemurafinib, cobimetinib</b>	<b>MAP kinase</b>
<b>BCL-2</b>	<b>Venetoclax</b>	<b>Apoptosis</b>
<b>PD-L1/L2</b>	<b>Nivolumab, others</b>	<b>Checkpoint inhibitors</b>

# Common Translocations in DLBCL

**t(3;14)(q27;q32); *BCL6::IGH*** **~25%**

**BCL6 also partners with other genes**

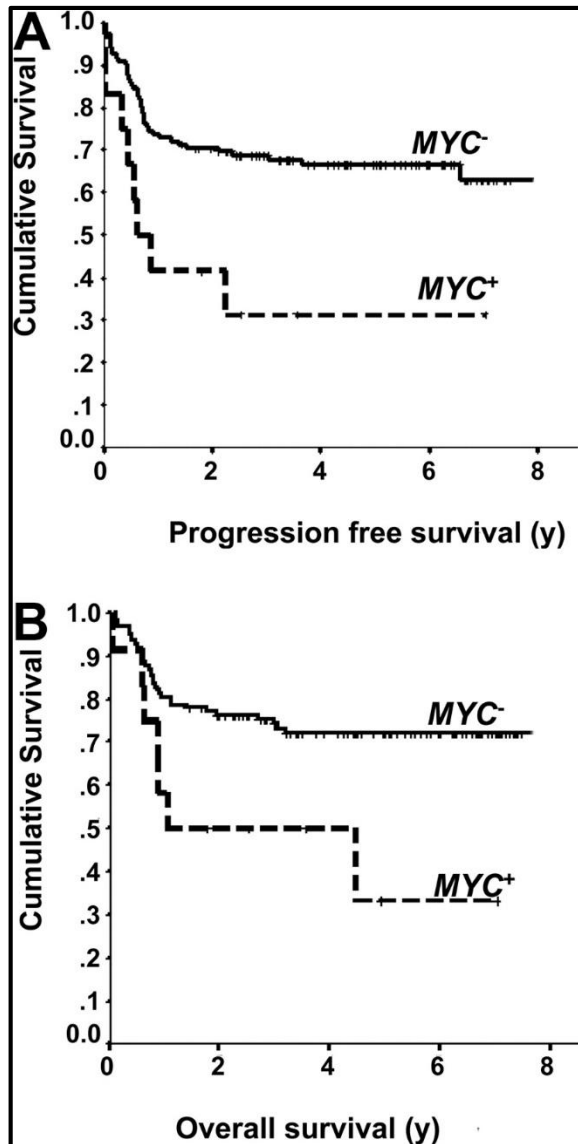
**t(14;18)(q32;q21); *IGH::BCL2*** **~20%**

**t(8;14)(q24;q32); *MYC::IGH*** **~10%**

**MYC also partners with other genes**



# MYC Rearrangment is Prognostic in DLBCL



**t(8;14)(q24;q32) - *IGH* (80%)**  
**t(8;22)(q24;q11) - *IGλ* (15%)**  
**t(2;8)(p11;q24) - *IGκ* (5%)**

## Diagnostic tests

Conventional cytogenetics

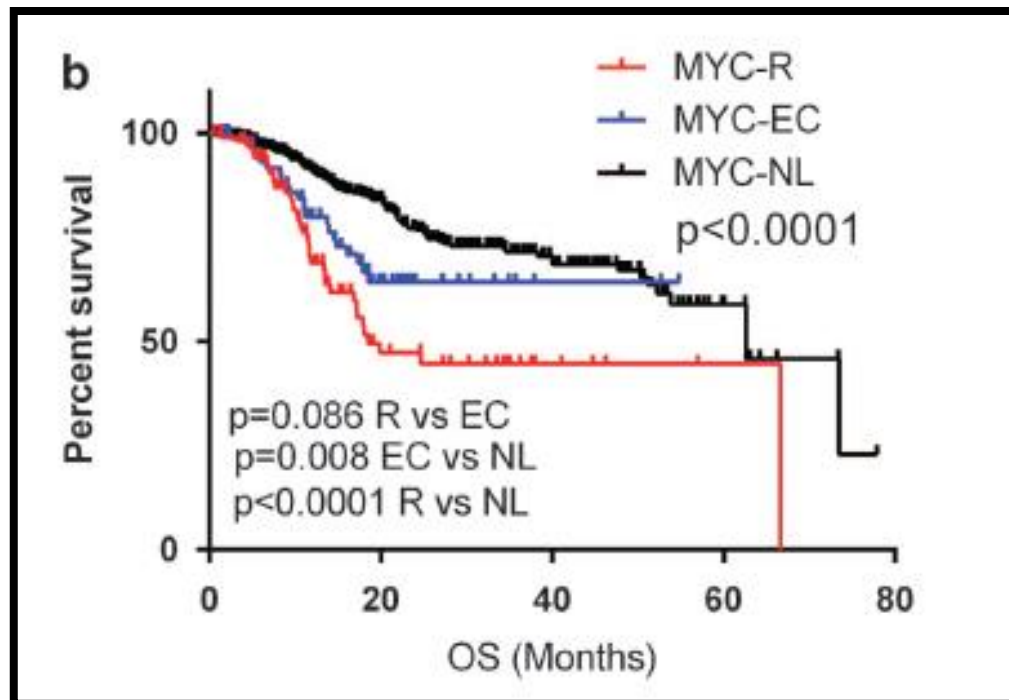
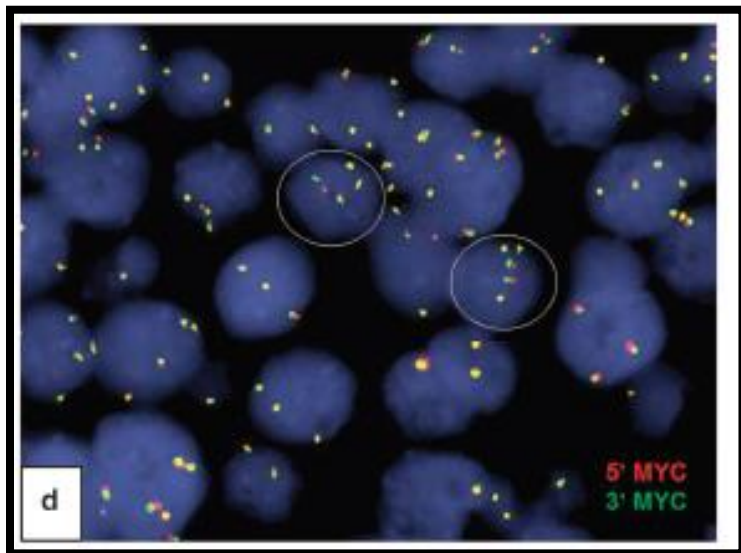
Need viable cells

## FISH

*IGH* and *MYC* probes

*MYC* breakapart probe

# MYC Extra Copies by FISH Predict Poorer Prognosis in DLBCL



Andres Quesada, MD

Increased *MYC* copy number is an independent prognostic factor in patients with diffuse large B-cell lymphoma

Andrés E Quesada<sup>1</sup>, L Jeffrey Medeiros<sup>1</sup>, Parth A Desai<sup>1</sup>, Pei Lin<sup>1</sup>, Jason R Westin<sup>2</sup>, Huda M Hawsawi<sup>1</sup>, Peng Wei<sup>3</sup>, Guilin Tang<sup>1</sup>, Adam C Seegmiller<sup>4</sup>, Nishitha M Reddy<sup>5</sup>, C Cameron Yin<sup>1</sup>, Wei Wang<sup>1</sup>, Jie Xu<sup>1</sup>, Roberto N Miranda<sup>1</sup>, Zhuang Zuo<sup>1</sup> and Shaoying Li<sup>1</sup>

# Diffuse Large B-cell Lymphoma

## Gene Expression Profiling Using DNA Microarrays



**Ash Alizadeh, MD, PhD**

**Lymphochip with 17,856 cDNA clones**

**12,069 Germinal center B-cell genes**

**2,338 B-cell NHL genes**

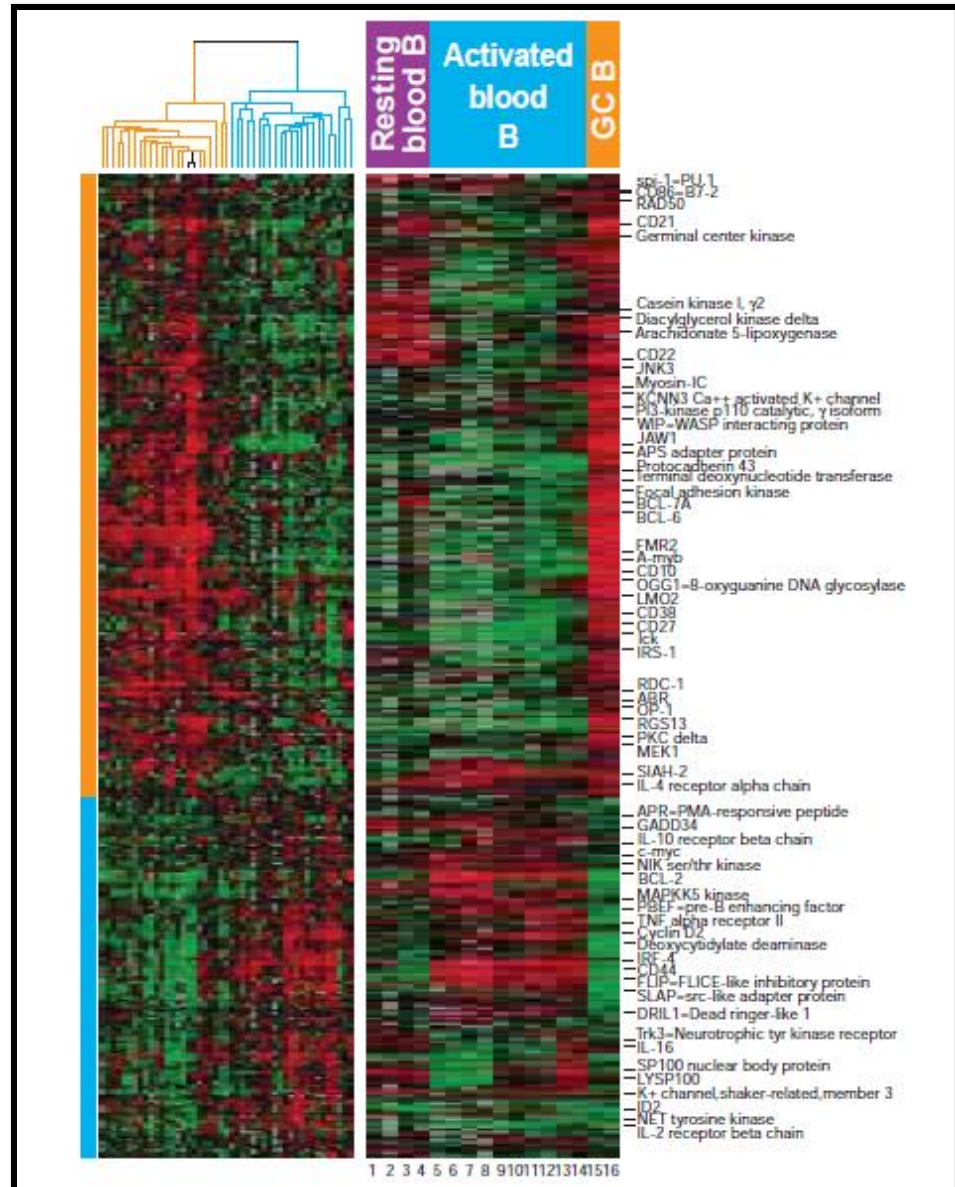
**3,186 Activated lymphocyte genes**



**Louis Staudt, MD, PhD**

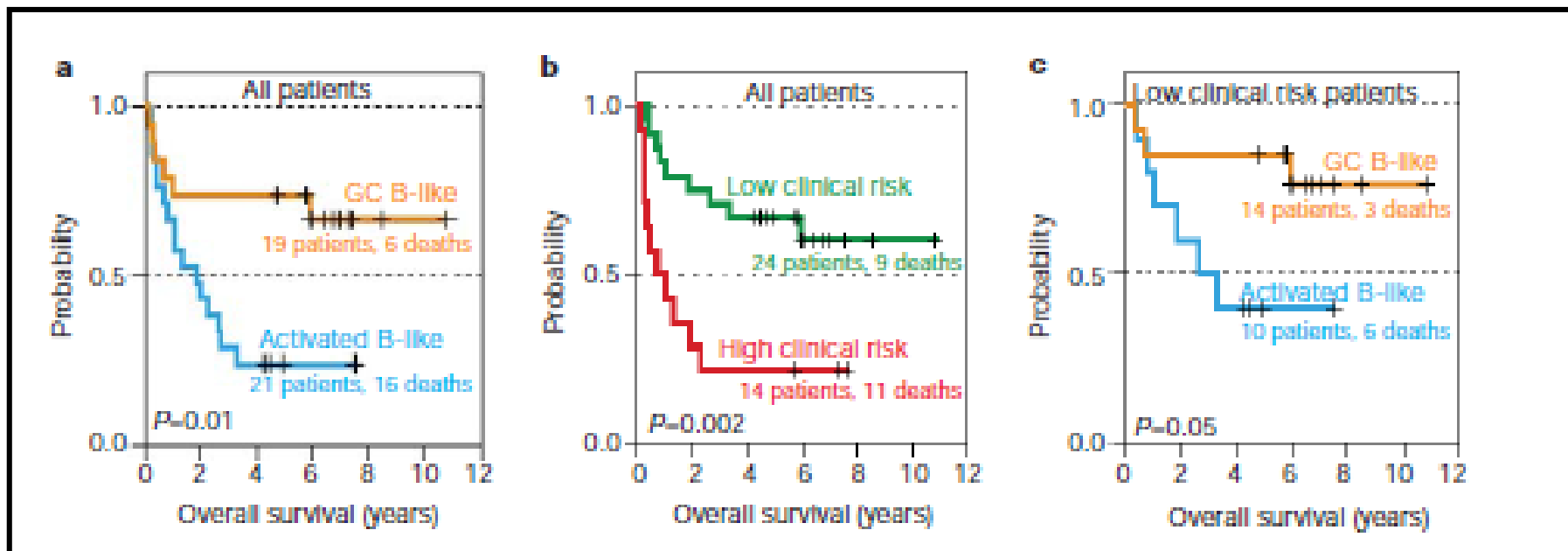
# Diffuse Large B-cell Lymphoma

GCB  
ABC



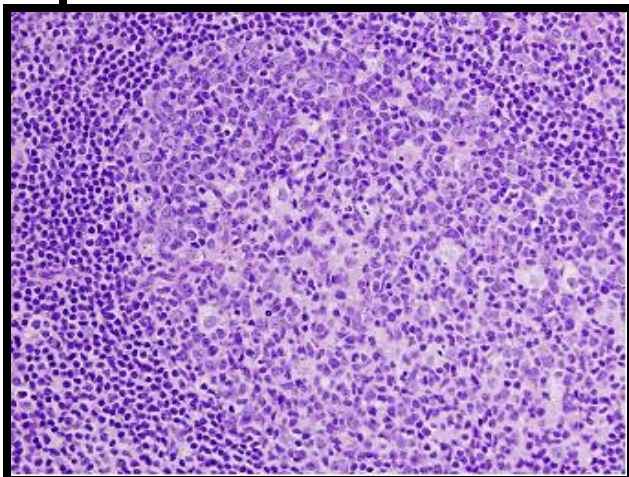
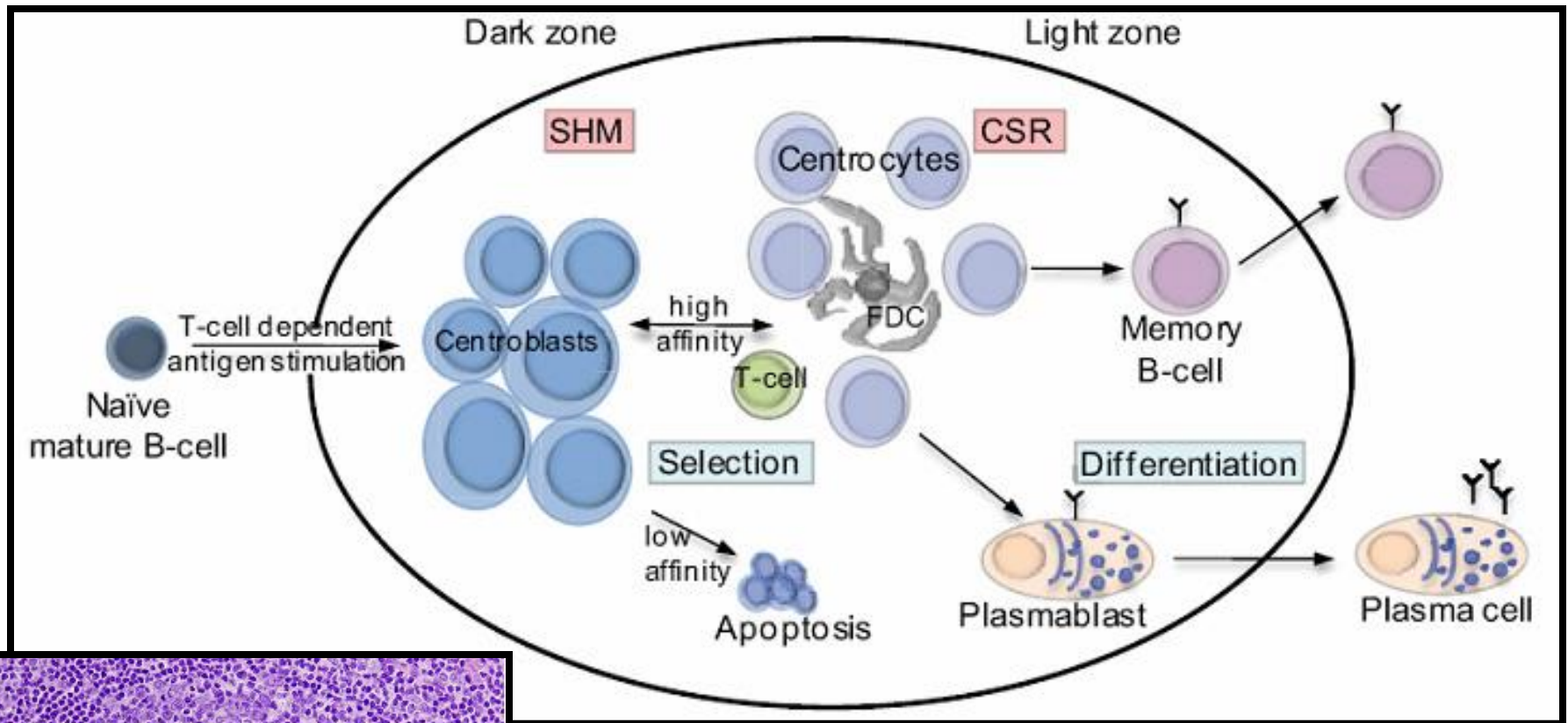
# Diffuse Large B-cell Lymphoma

## GEP Shows 2 Types that Predict Prognosis



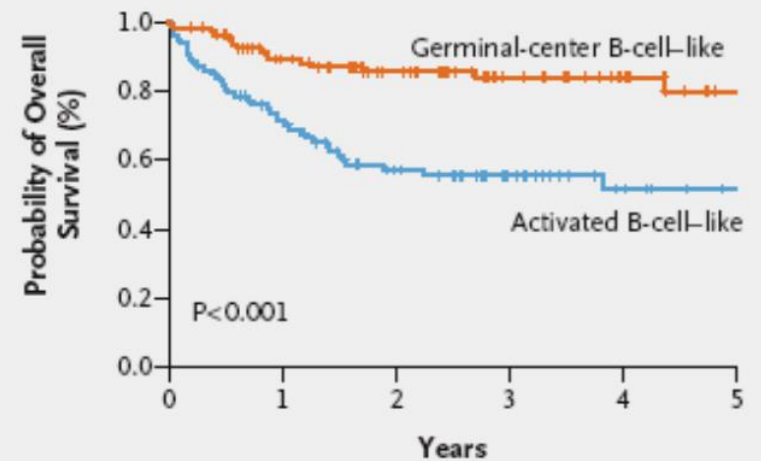
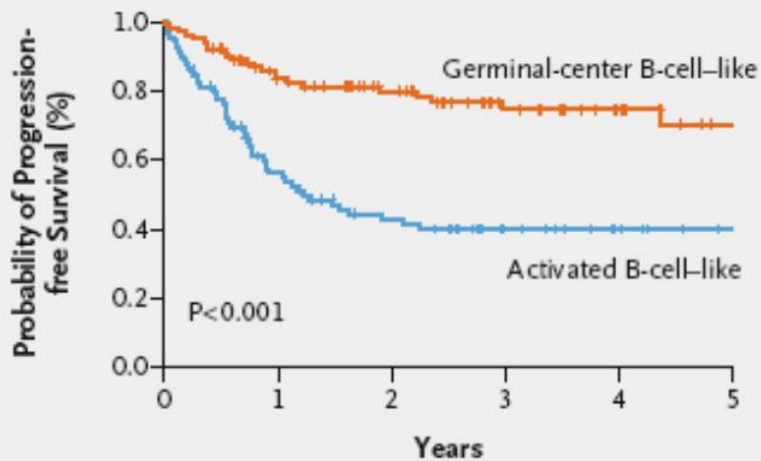
**CHOP Therapy**

# Germinal Center Reaction



# Diffuse Large B-cell Lymphoma

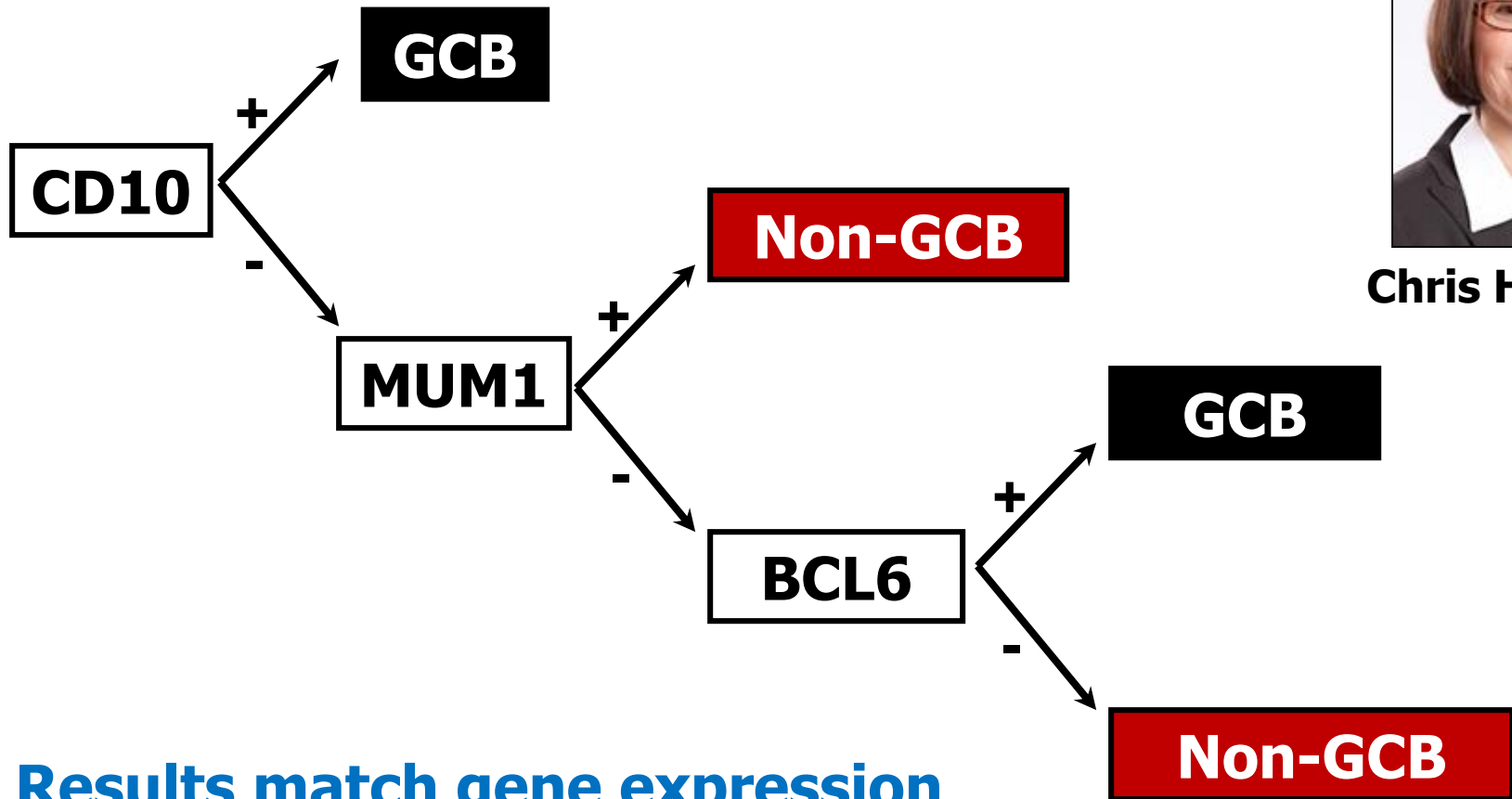
## GEP is Valid for R-CHOP Treated Patients



### No. at Risk

Germinal-center B-cell-like	107	82	61	39	27	15	101	74	56	35	24	14
Activated B-cell-like	93	60	38	23	11	6	90	45	30	17	10	5

# Can Immunohistochemistry be used as a Surrogate for GEP in DLBCL?



Chris Hans, MD

Results match gene expression profile in ~80% of cases



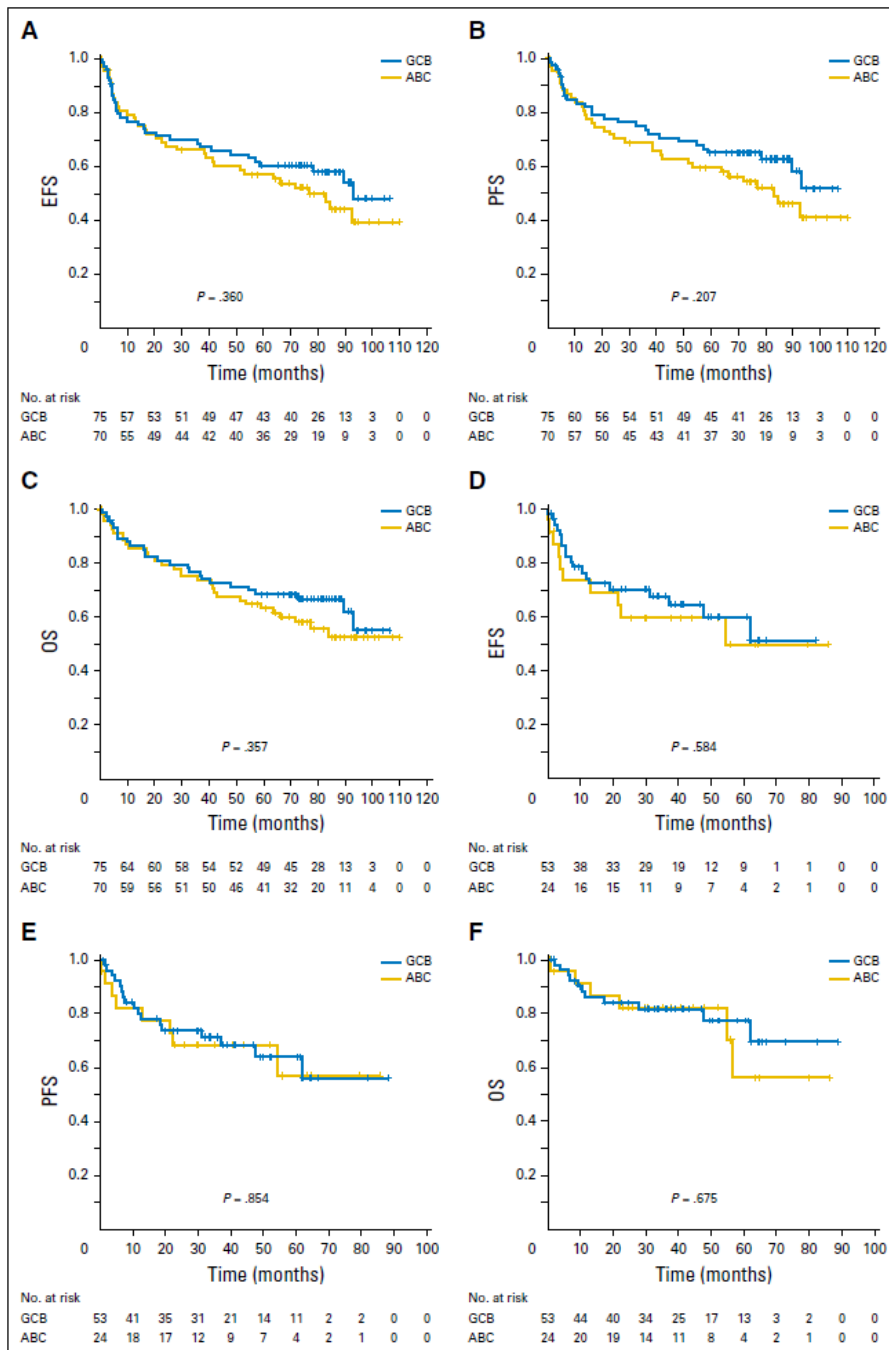
Clinical Impact of the Cell-of-Origin Classification and the MYC/BCL2 Dual Expresser Status in Diffuse Large B-Cell Lymphoma Treated Within Prospective Clinical Trials of the German High-Grade Non-Hodgkin's Lymphoma Study Group

Annette M. Staiger, Marita Ziepert, Heike Horn, David W. Scott, Thomas F.E. Barth, Heinz-Wolfram Bernd, Alfred C. Feller, Wolfram Klapper, Monika Szczepanowski, Michael Hummel, Harald Stein, Dido Lenz, Martin-Léo Hansmann, Sylvia Hartmann, Peter Möller, Sergio Cogliatti, Georg Lenz, Lorenz Trümper, Markus Löffler, Norbert Schmitz, Michael Pfrendschuh, Andreas Rosenwald, and German Ott for the German High-Grade Lymphoma Study Group

**A,C,E. RICOVER-60 trial**  
**B,D,F. R-MegaCHOEP trial**

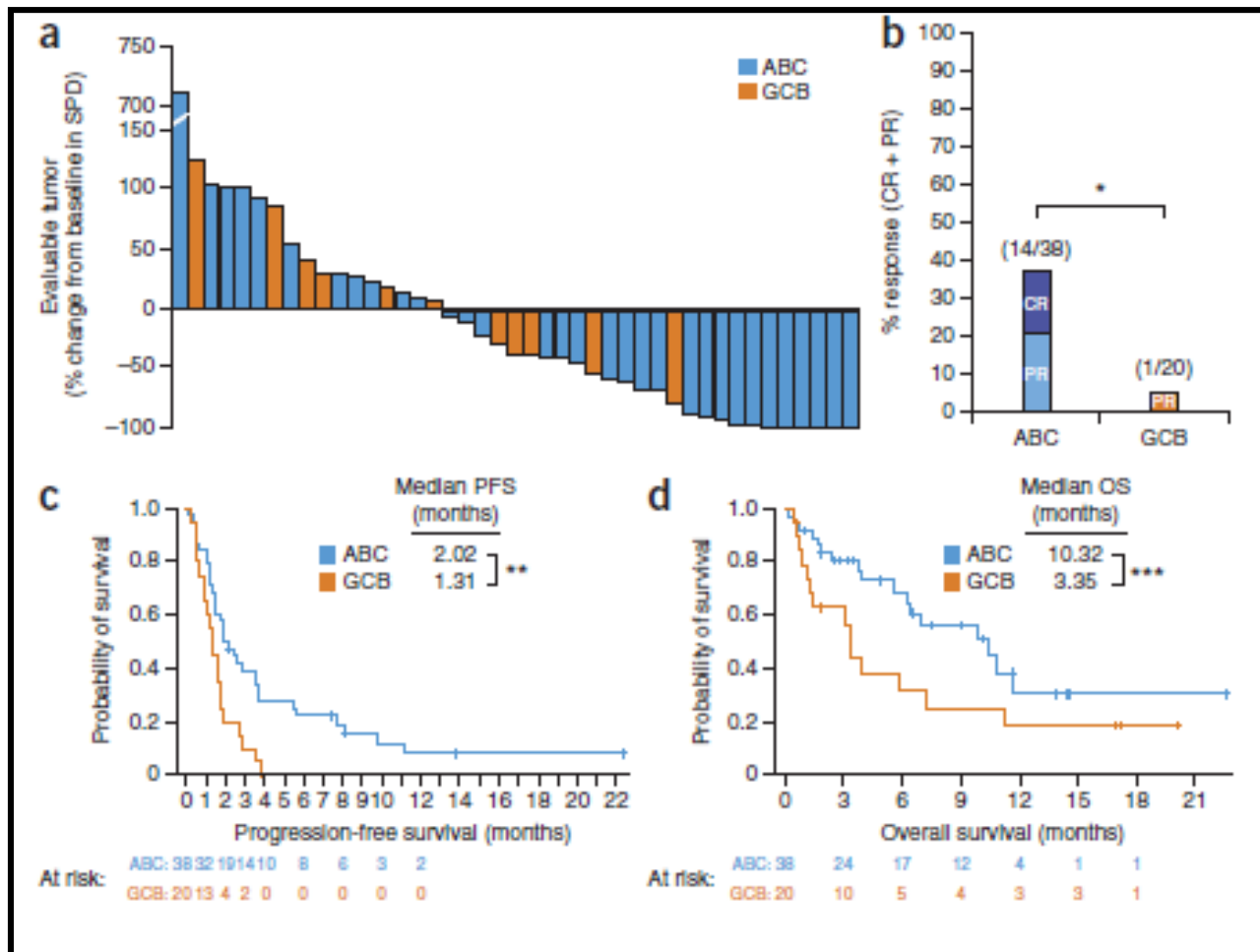
Cell-of-origin classification did not correlate with prognosis

**J Clin Oncol 35:2515, 2017**



# R-CHOP+Ibrutinib for DLBCL

## Impact of GCB versus ABC



# Mutations in Pathways Involved in DLBCL

## B-cell receptor signaling

**CD79A, CD79B, CARD11**

## Toll-like receptor signaling

**MYD88**

**NF- $\kappa$ B**

## Lymphocyte differentiation

**TNFAIP3/A20, TRAF3, BIRC3, IKK $\beta$**

## DNA repair and transcriptional regulation

**p53**

## Lymphocyte activation

**STAT6, BCL10**

## DNA methylation

**EZH2, MLL2**

## DNA acetylation

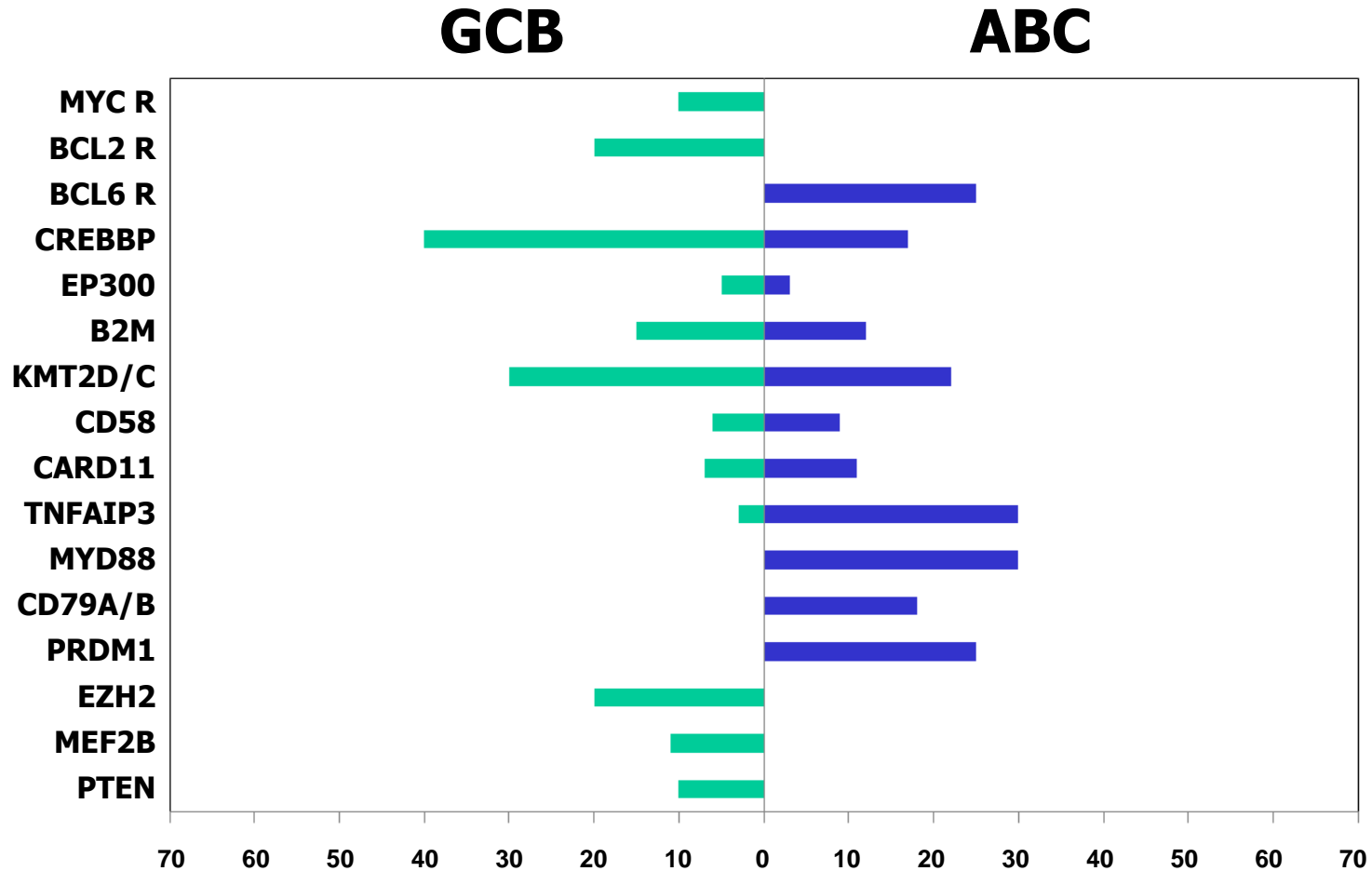
**CREBBP, MEF2B**

## Immune surveillance

**$\beta$ 2M, CD58**

# Diffuse Large B-cell Lymphoma, NOS

Rearrangements and mutations correlate with COO



Frequency of Mutations

# **Diffuse Large B-cell Lymphoma, NOS**

## **3 Major Studies on Molecular Subgroups**

**Chapuy B, et al. Molecular subtypes of diffuse large B cell lymphoma are associated with distinct pathogenic mechanisms and outcomes. Nat Med 2018; 24:679-690.  
(Harvard)**

**Wright GW, et al. A probabilistic classification tool for genetic subtypes of diffuse large B cell lymphoma with therapeutic implications. Cancer Cell 2020; 37:551-568.  
(LymphGen NIH)**

**Lacy SE, et al. Targeted sequencing in DLBCL, molecular subtypes, and outcomes: a Haematological Malignancy Research Network report. Blood 2020; 135:1759-1771.**

# Comparison of Three Systems

LymphGen	Modified HMRN	Harvard	Main gene mutations	COO	Outcome	Related Lymphoma
<b>MCD</b>	<b>MYD88</b>	<b>C5</b>	<i>MYD88<sup>L265P</sup>, CD79B, PIM1</i>	ABC	Poor	Primary CNS Lymphoma, Primary Testicular Lymphoma
<b>EZB</b>	<b>BCL2</b>	<b>C3</b>	<i>BCL2, EZH2, CREBBP, KMT2D</i>	GCB	Good	Follicular Lymphoma
<b>EZB-MYC+</b>	<b>BCL2-MYC</b>				Poor	Double-hit Lymphoma
<b>BN2</b>	<b>NOTCH2</b>	<b>C1</b>	<i>NOTCH2, BCL10, SPEN, CD70, BCL6</i>	ABC, GCB, UC	Intermediate/Good	Marginal Zone Lymphoma
<b>ST2</b>	<b>TET2/SGK1</b>	<b>C4</b>	<i>TET2, SGK1, KLHL6, BRAF, MAP2K1, KRAS</i>	GCB	Good	Nodular Lymphocyte Predominant Hodgkin Lymphoma
	<b>SOCS1/SGK1</b>		<i>SOCS1, SGK1, CD83, NFKBIA, HIST1H1E, STAT3</i>	GCB	Very Good	Primary Mediastinal B-Cell Lymphoma
<b>N1</b>	<b>NOTCH1</b>		<i>NOTCH1, ID3</i>	ABC	Poor	Chronic Lymphocytic Leukaemia
<b>A53</b>		<b>C2</b>	<i>TP53, aneuploidy</i>	Mixed	Intermediate	
<b>Other</b>	<b>NEC</b>	<b>C0</b>		ABC, GCB, UC	Intermediate	

Wright

Chapuy

Courtesy of Daniel Hodson, MD  
Cambridge, UK

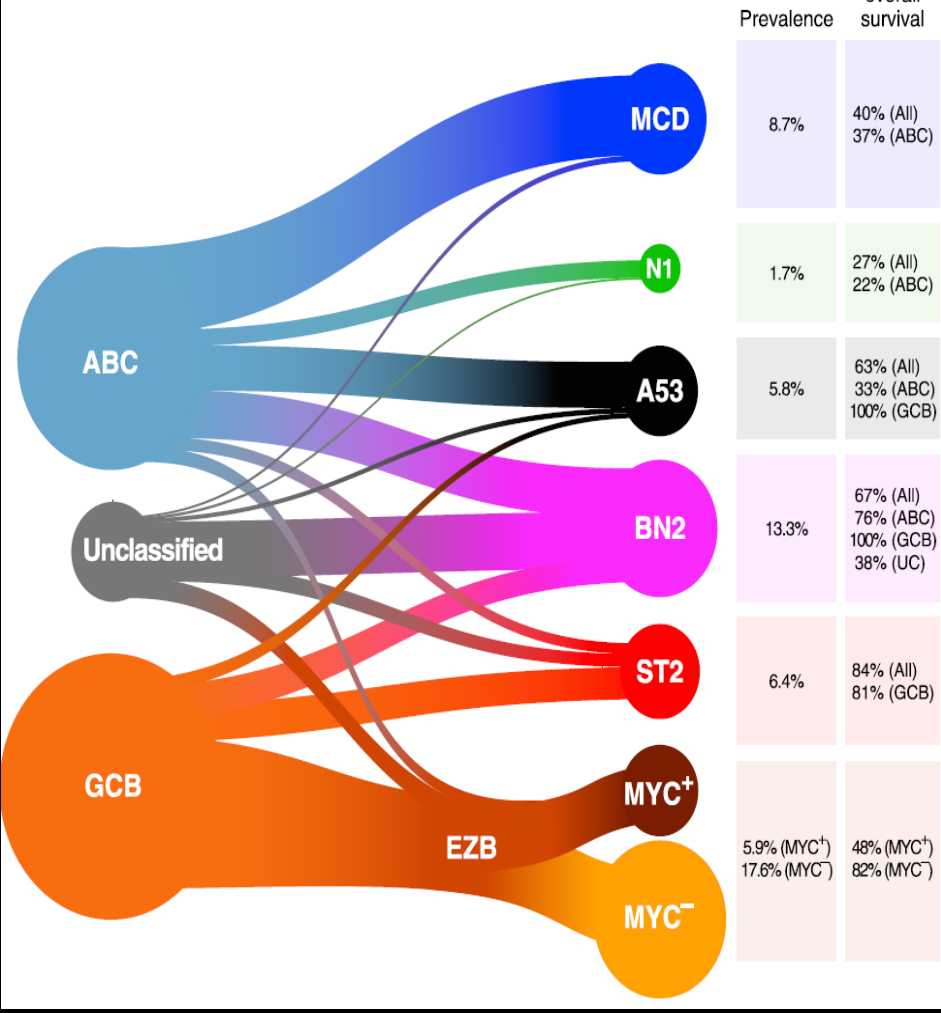
**A Probabilistic Classification Tool  
for Genetic Subtypes of Diffuse Large B Cell  
Lymphoma with Therapeutic Implications**

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**LymphGen Classifier**

**DLBCL Subgroups**

**Cancer Cell 37: 551, 2020**



**MCD**  
*MYD88 + CD79B* mutations

**N1**  
*NOTCH1* pathway

**A53**  
Aneuploidy + TP53 mutations

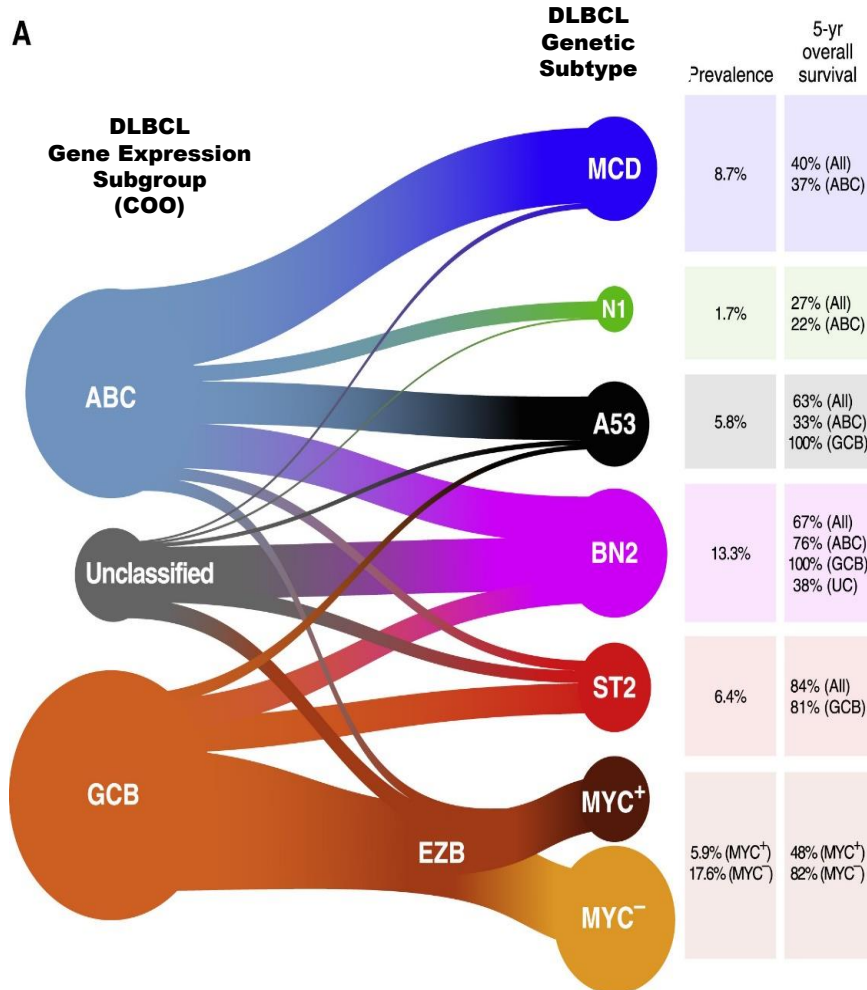
**BN2**  
*BCL6* fusions + *NOTCH2* mutations

**ST2**  
*SGK1* and *TET2* mutations

**EZB**  
*EZH2* mutations +  
*BCL2* translocations

# DLBCL Genetic Subtypes

## Implications for Pathogenesis and Therapy



## Potential drug targets

**MCD**

**BTK, PI3K, BCL2, JAK**

**N1**

**A53**

**BN2**

**BTK, PI3K, BCL2**

**ST2**

**PI3K, JAK**

**EZB**

**EZH2, PI3K, BCL2**



# Take Home Points

**The traditional cell-of-origin model (GCB vs ABC) is **not** sufficiently granular to predict prognosis or to plan therapy**

**For now, we will need to keep using this model, but only until a better, more practicable system becomes available**

**A new model may not lead to optimal therapy currently, but it will lead to design of clinical trials and evaluation of therapies**

**However, this new system needs to be practical**