

# MIHCC

June 25, 2025

Case 3

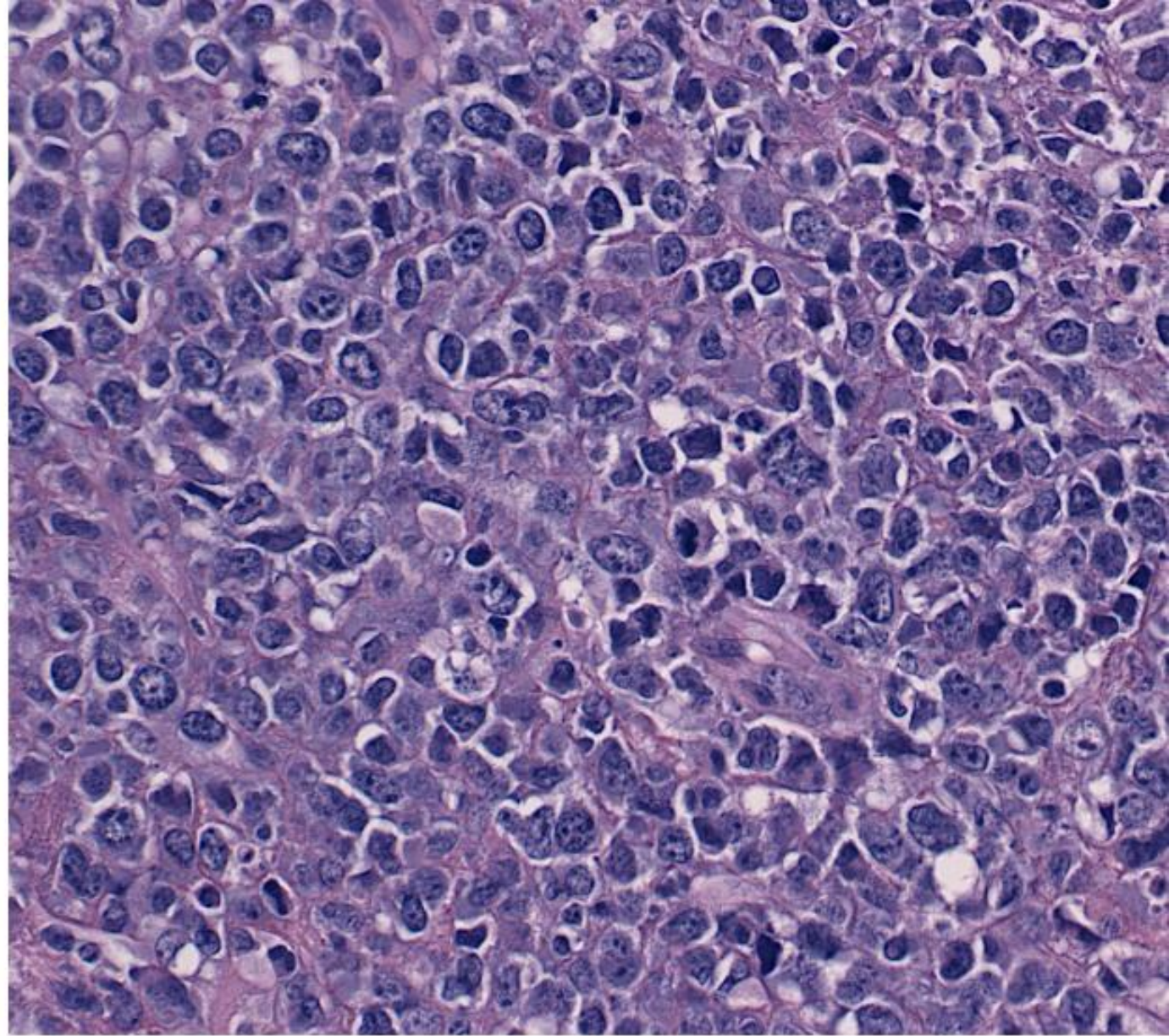
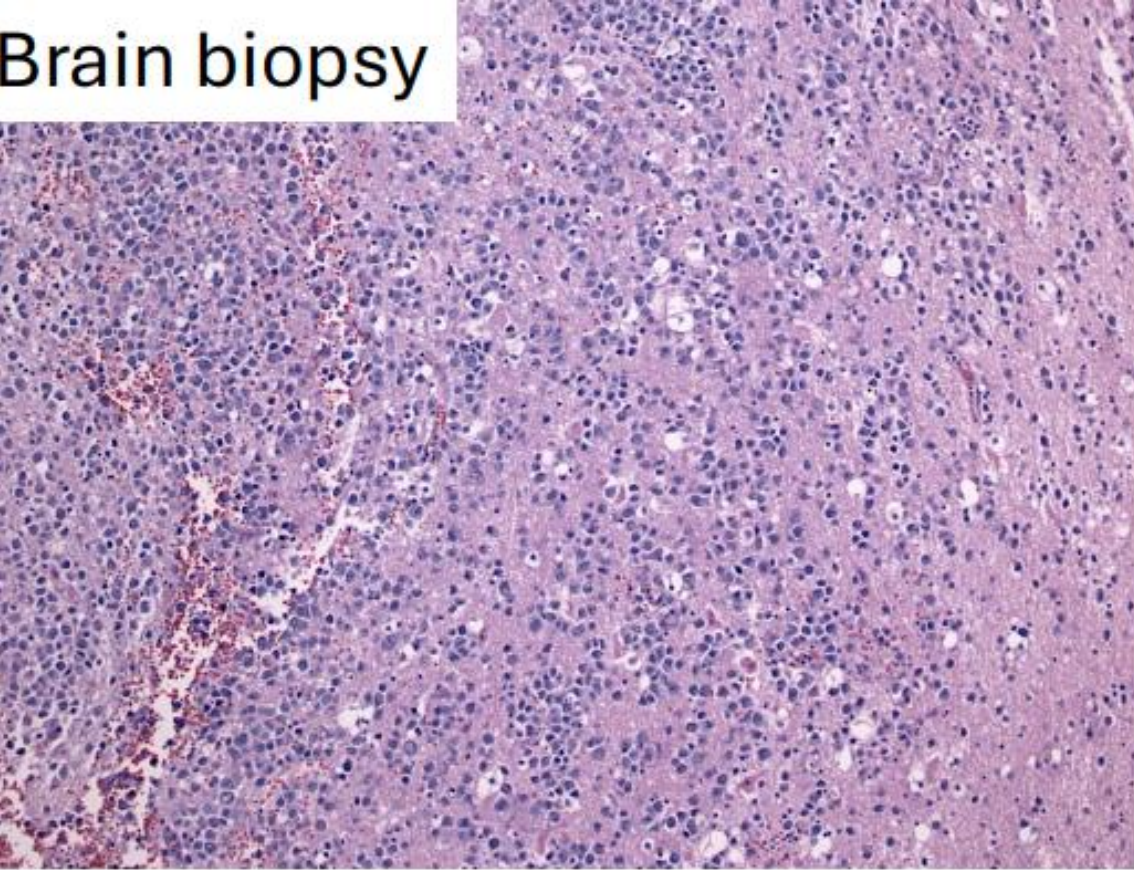
# Clinical History

- June 2019: A 61-year-old man with a history of bipolar disorder and alcohol abuse presents with declining mental status over the past month.
- MRI showed multiple intracerebral lesions concerning for either metastatic disease or lymphoma.

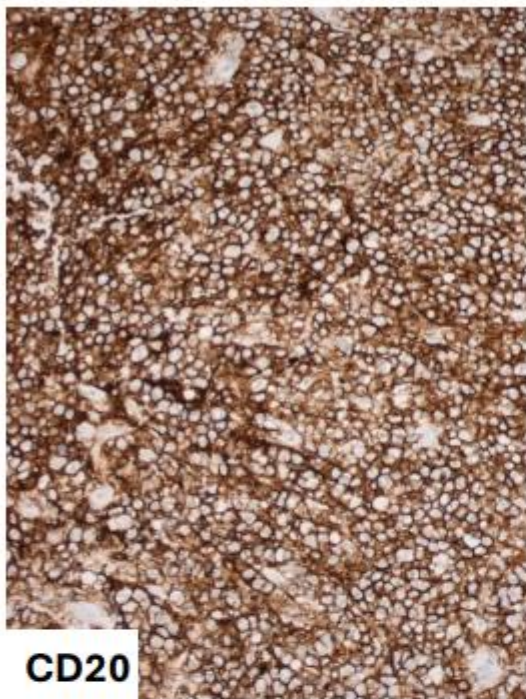




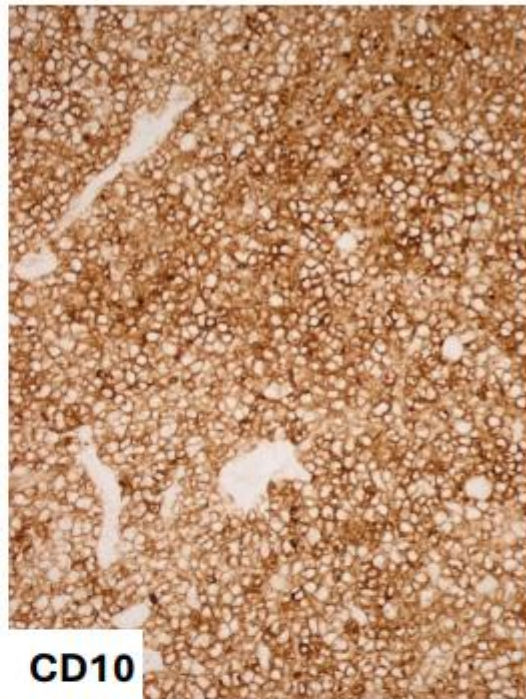
Brain biopsy



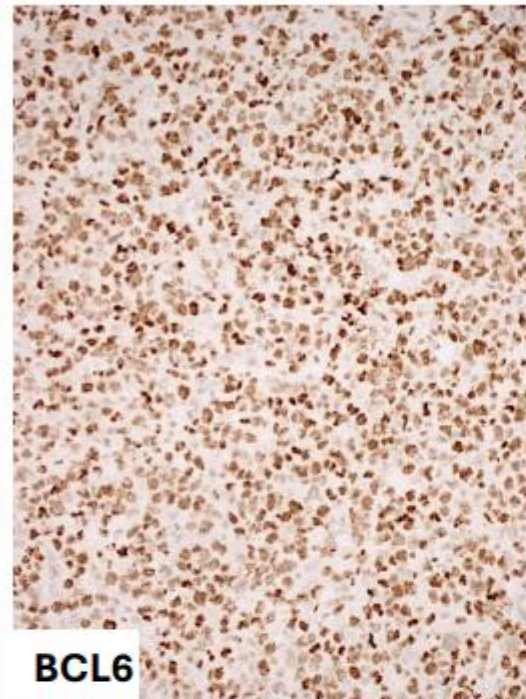




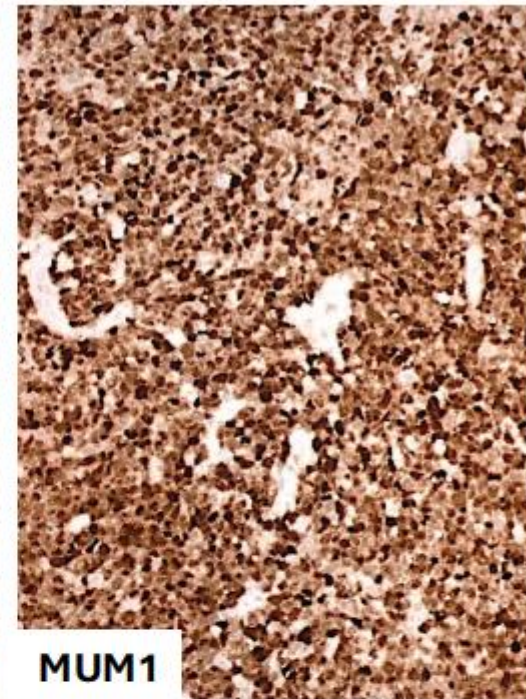
**CD20**



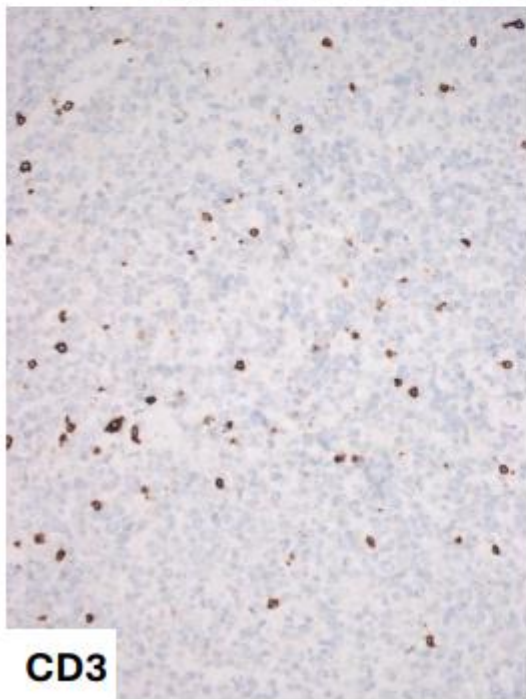
**CD10**



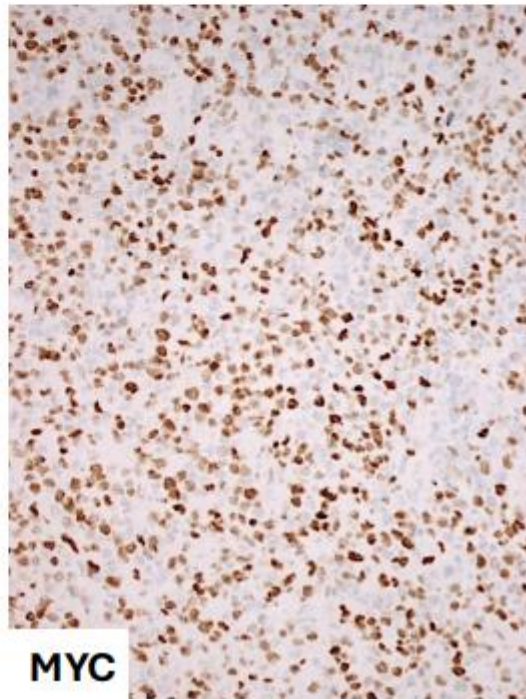
**BCL6**



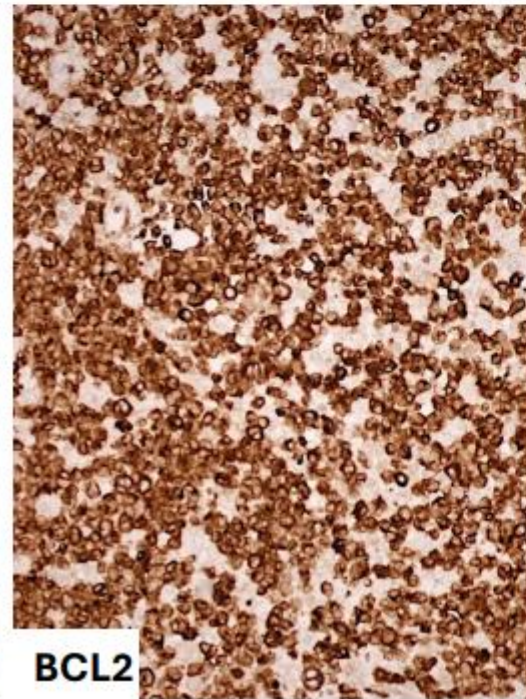
**MUM1**



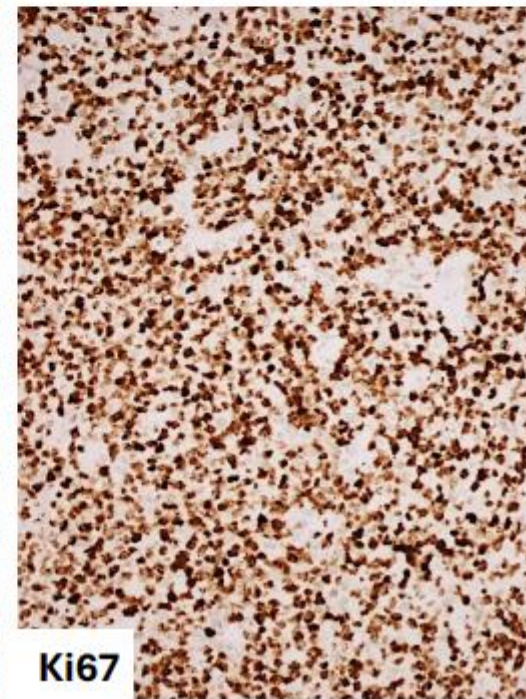
**CD3**



**MYC**



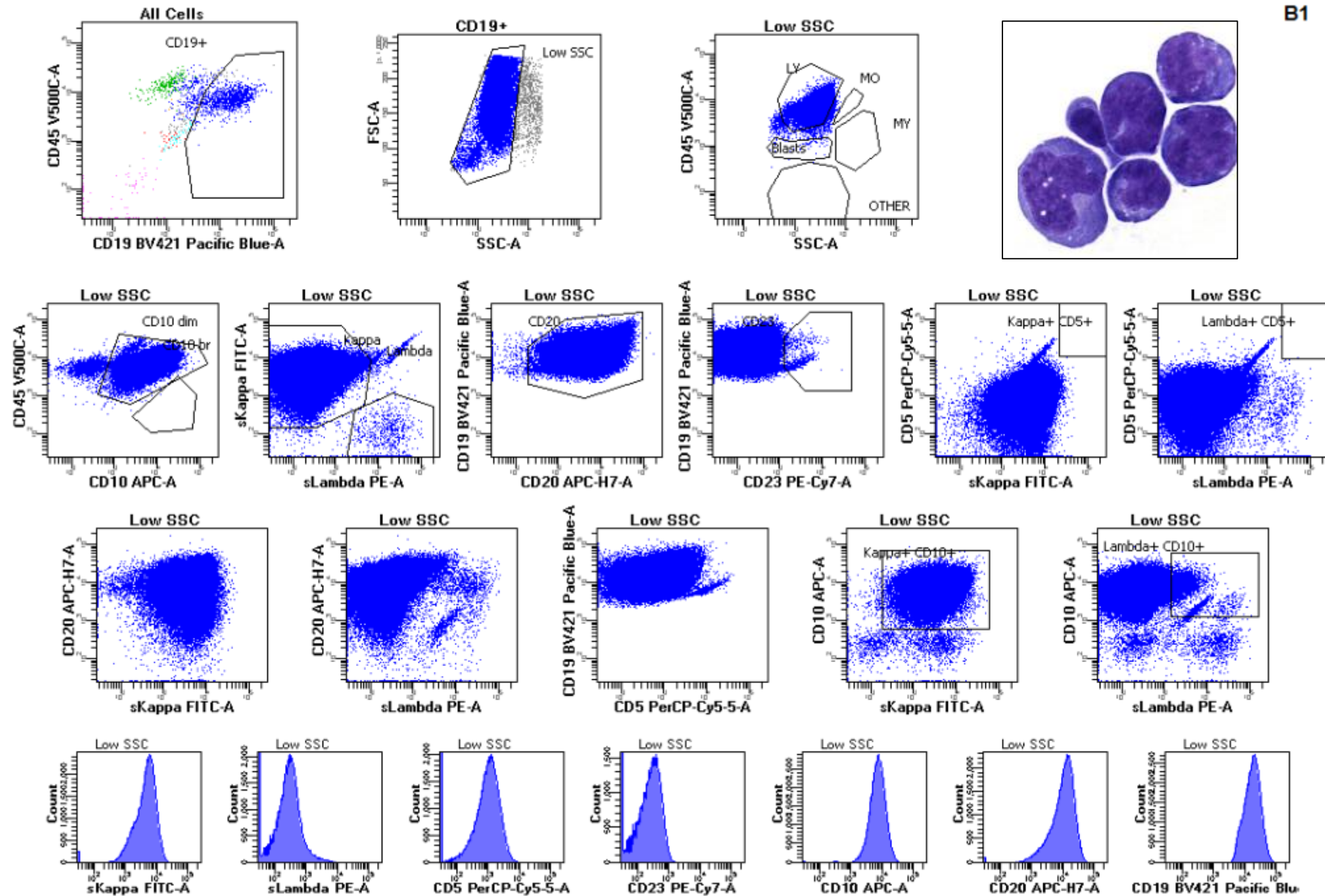
**BCL2**



**Ki67**



B1



# FISH Studies

nuc ish(BCL6x3~4)[64/200],(MYCx3~4)[65/200]/(D8Z1x2~4,MYCx3~4,IGHx3~6)[165/200],  
(IGHx3~6,BCL2x2~4)[166/200]

**Interphase FISH assays ruled out the presence of BCL6, MYC, MYC-IGH and IGH-BCL2 gene rearrangements in this sample.**

**FINAL DIAGNOSIS:**

**A & B. Brain tumor:**

**Diffuse large B-cell lymphoma, in brain. See note.**

Note: The immunohistochemical profile of this DLBCL is consistent with a DLBCL of the germinal center B-cell subtype (BCL6+, CD10+, MUM1+, Hans' classifier). Differential diagnoses include a primary diffuse large B-cell lymphoma of the CNS versus CNS involvement by a systemic DLBCL. CD10 is expressed infrequently (<10%) by primary CNS DLBCL, but its expression is more frequent in systemic DLBCL. Thus, the strong CD10 positivity seen in this DLBCL raises the possibility of CNS dissemination by a systemic DLBCL. Therefore, clinical correlation, including thorough evaluation for a systemic DLBCL, is recommended.

June 2019: Work-up showed no other sites of disease

Treated with 6 cycles of R-MVP

(rituximab, methotrexate, procarbazine, vincristine)

Summer/Fall 2019: Interim/end of treatment MRIs showed decreased disease with improved mental status



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Imaging showed recurrent disease

CSF flow cytometry: small population of kappa-restricted, CD10+ B cells



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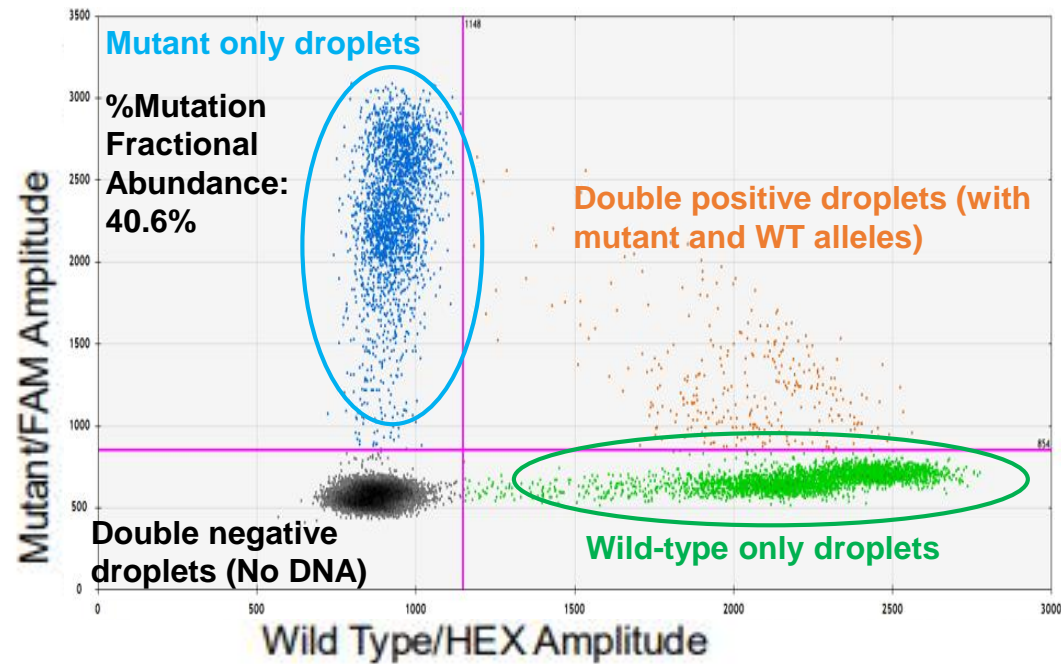
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**Molecular testing for *MYD88* mutation was done**

# MYD88 Testing

(Quantitative droplet digital PCR)



$$\begin{array}{l} \% \text{ Mutation} \\ \text{Fractional} \\ \text{Abundance} \end{array} = \frac{\text{Mutant Droplets}}{\text{Mutant Droplets} + \text{Wild Type Droplets}} \times 100$$



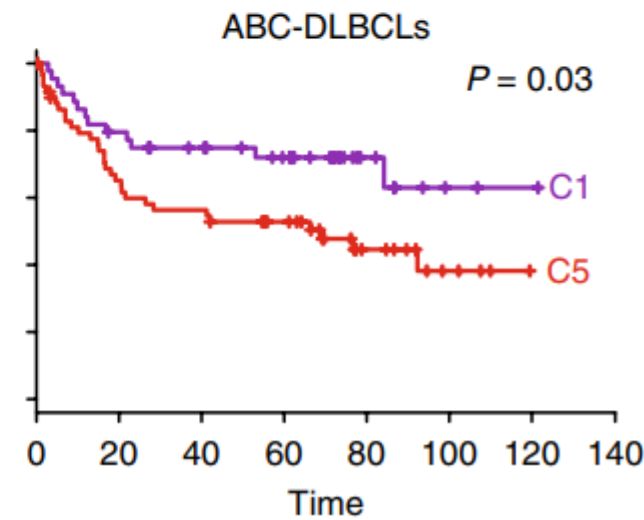
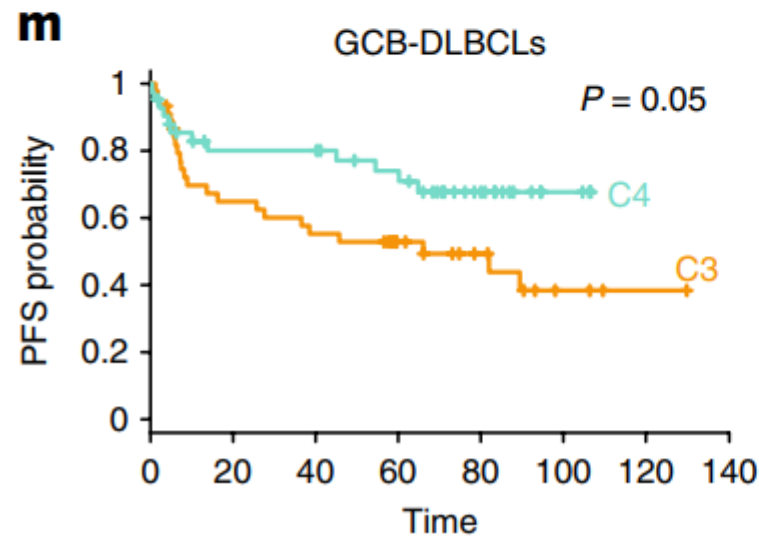
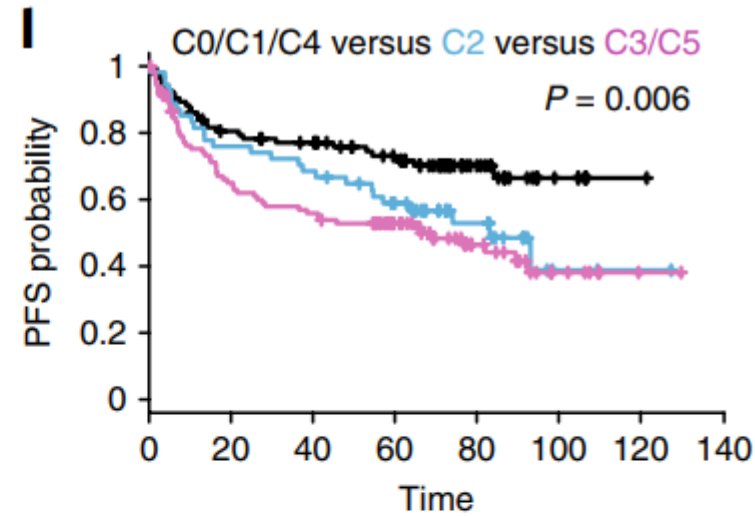
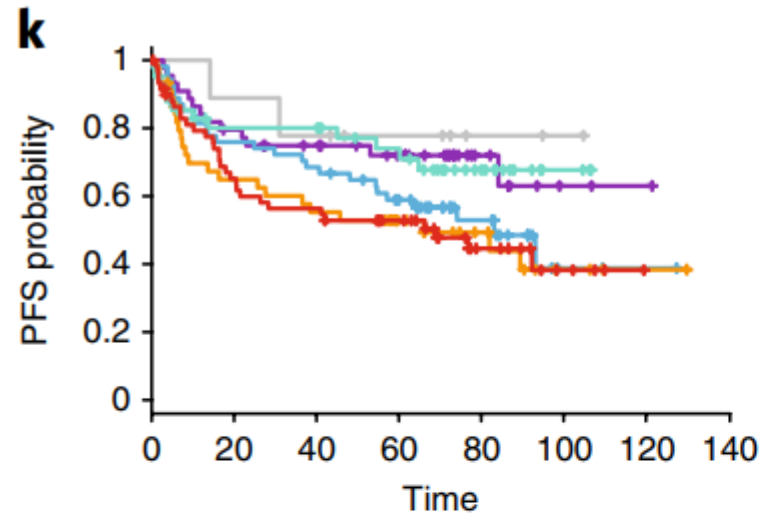
# Harvard classification

## Harvard study

- Retrospective
- WES, copy number changes and translocation
- R-CHOP like treatment

<i>BCL6</i> SV with mutations in <i>NOTCH2</i> pathway	Biallelic inactivation of <i>TP53</i> , 17p loss, <i>CDKN2A</i> loss and associated genomic instability	<i>BCL2</i> SV, mutations and/or loss of <i>PTEN</i> , alterations of epigenetic enzymes	Alterations in <i>JAK/STAT</i> , <i>BRAF</i> pathways and multiple histones	<i>BCL2</i> copy gain, <i>MYD88</i> , <i>CD79B</i> mutations
C1	C2	C3	C4	C5
Mutations in NF- $\kappa$ B pathway members	Loss of RB1 → increased levels of E2F targets	Mutations in chromatin modifiers	Mutations in H1 linker histones	Associated with extranodal disease
Mutations associated with immune escape	More driver somatic copy number alterations	2 mechanisms of PTEN inactivation	Primarily GCB	Aberrant somatic hypermutation
Majority ABC	GCB and ABC	95% GCB		96% ABC

# Genetic subtypes and outcome

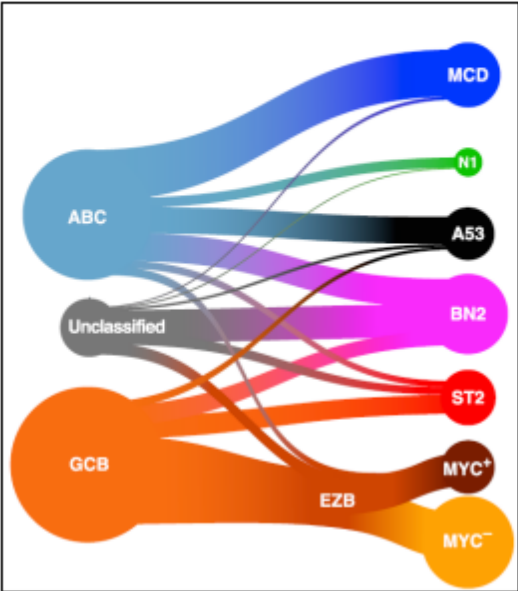


C1: ABC  
C2: Both  
C3: GCB  
C4: GCB  
C5: ABC

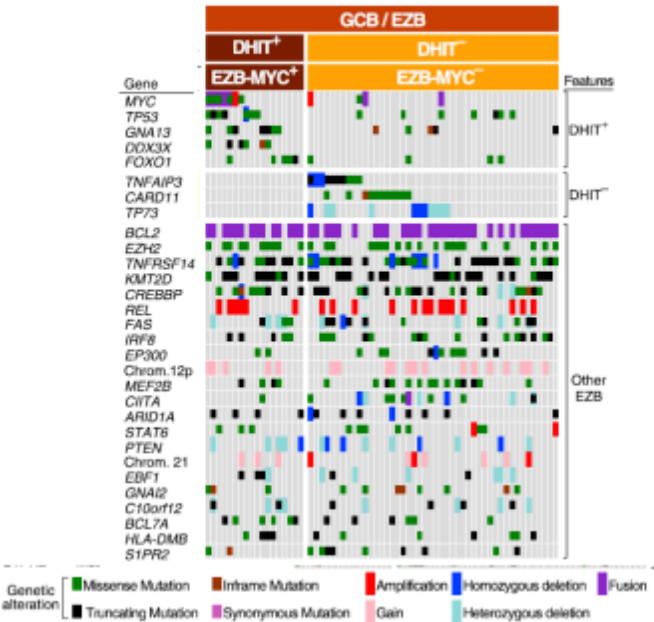
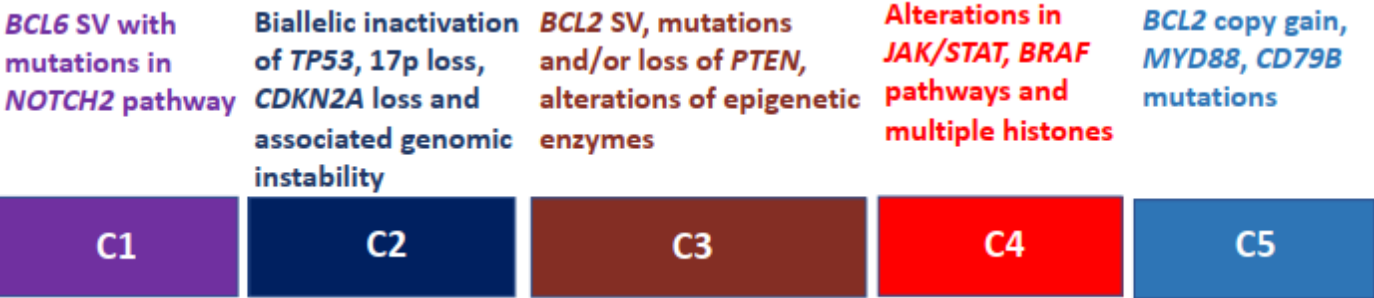


# LymphGen

- 6 genetic subtypes of DLBCL were proposed:
  - MCD (*MYD88*L265P and *CD79B* mutations)
  - BN2 (*BCL6* translocations and *NOTCH2* mutations)
  - N1 (*NOTCH1* mutations)
  - EZB (*BCL2* translocations and *EZH2* mutations)
  - A53 (aneuploid with *TP53* inactivation)
  - ST2 (*SGK1* and *TET2* mutations)



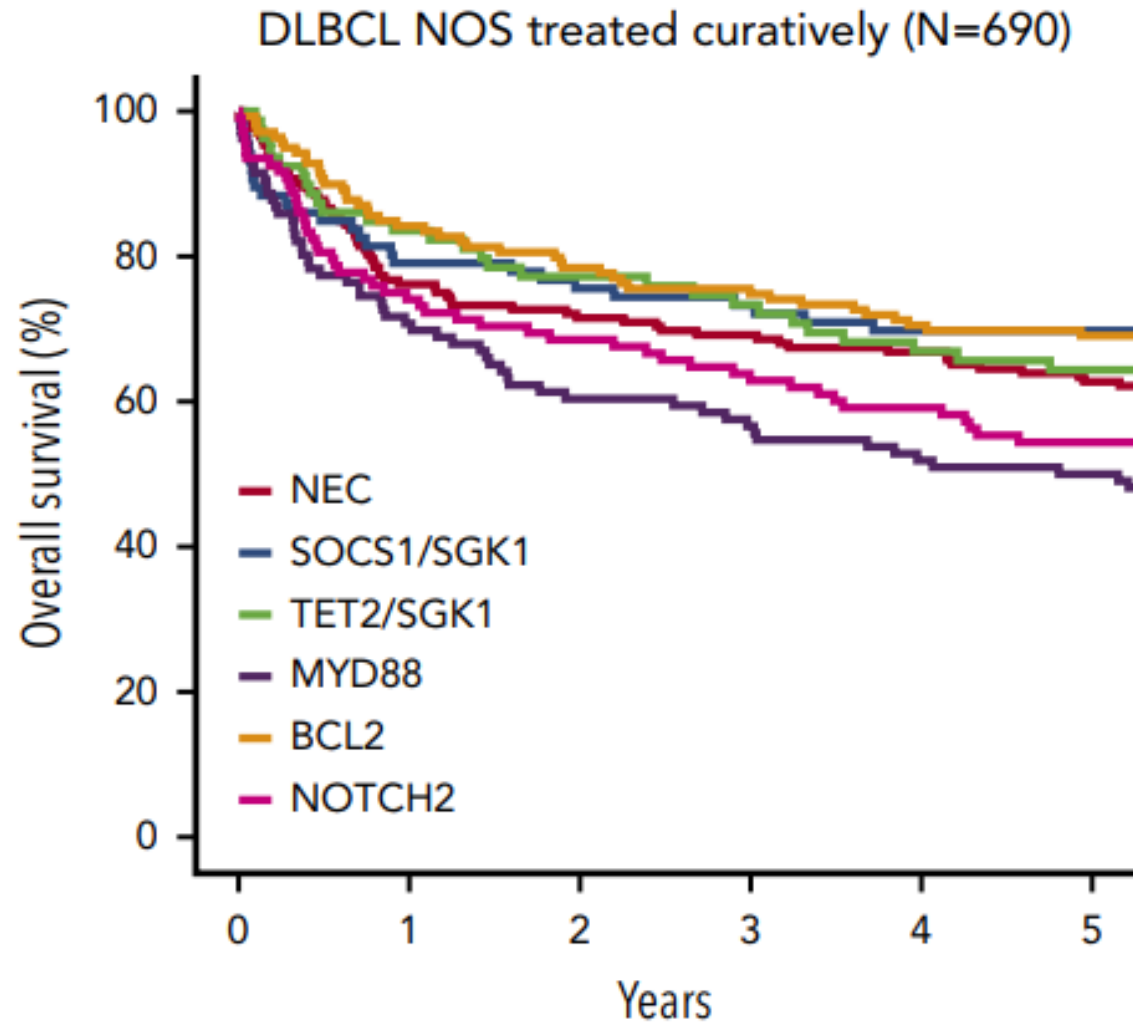
## Harvard classification



# UK Haematological Malignancy Research Network (HMRN) classification

UK HMRN classification					
<ul style="list-style-type: none"> <li>Prospective</li> <li>Based predominantly on target panel mutation analysis; copy number changes are considered for a small number of genes-&gt; <b>cannot reliably identify aneuploidy</b></li> <li>Modified classification used the presence of <i>MYC</i> hotspot and <i>NOTCH1</i> PEST domain mutations</li> <li>Including a subset patients with R-CHOP full treatment</li> </ul>					
<i>NOTCH1</i> mutations	<i>NOTCH2, BCL10, TNFAIP3, CCND3, SPEN, TMEM30A FAS, and CD70</i> mutations	<i>EZH2, BCL2, CREBBP, TNFRSF14, KMT2D and MEF2B</i> mutations	<i>SOCS1, CD83, SGK1, NFKB1A, HIST1H1E and STAT3</i> mutations	<i>TET2, SGK1, KLHL6, ZFP36L1, BRAF, MAP2K1 and KRAS</i> mutations	<i>MYD88L265P, PIM1, CD79B, ETV6</i> mutations and loss of <i>CDKN2A</i>
NOTCH1	NOTCH2	BCL2	SOCS1/SGK1	TET2/SGK1	MYD88
Not elsewhere classified (NEC)	Correlation with BCL6 rearrangement	Majority with t(14;18) BCL2 translocation	Genes are known targets of somatic hypermutation	Mutations of the ERK pathway	Signatures associated with ABC, IRF4, and MYC
	Biologically similar to marginal zone lymphoma	Predominantly GCB	Biologically similar to PMBCL	Predominantly GCB	
	Mixture of ABC, GCB, unclassified DLBCL		Predominantly GCB		

# Genetic subtypes and outcome

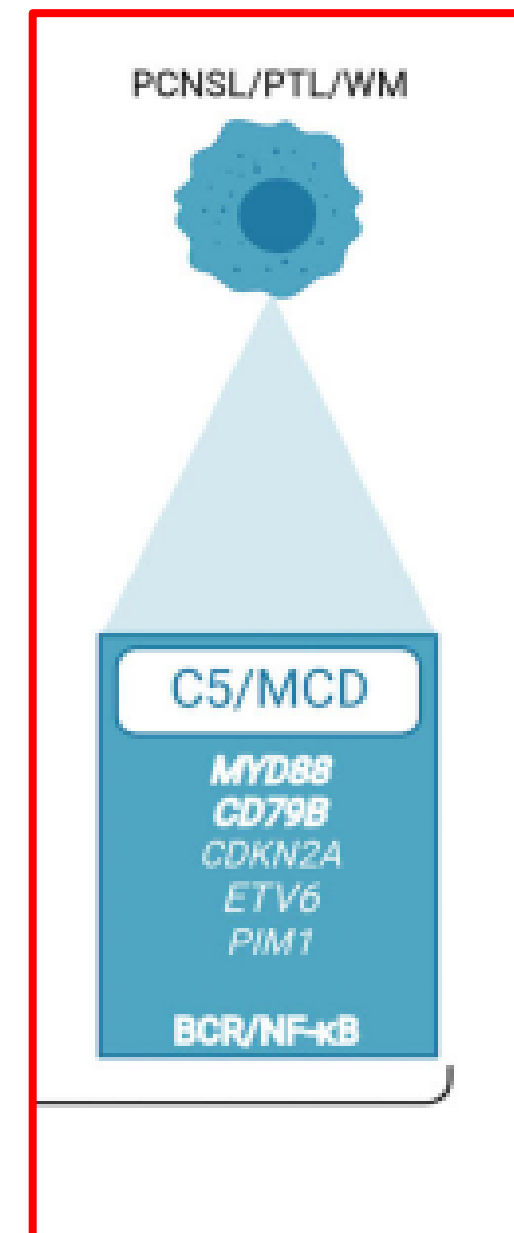
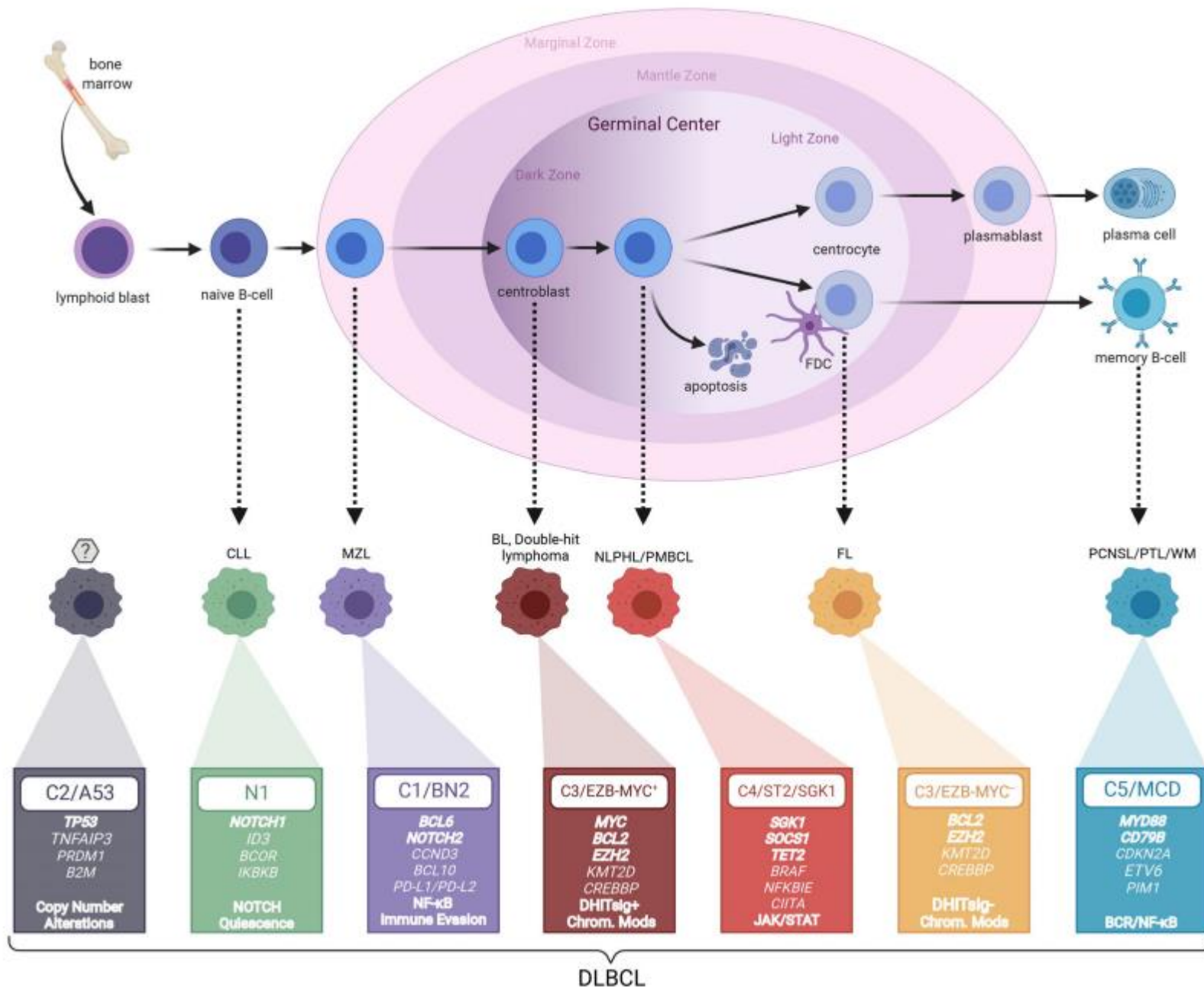


Better outcomes in  
SOCS1/SGK1, BCL2, and  
TET2/SGK1 clusters  
(**GCB**)

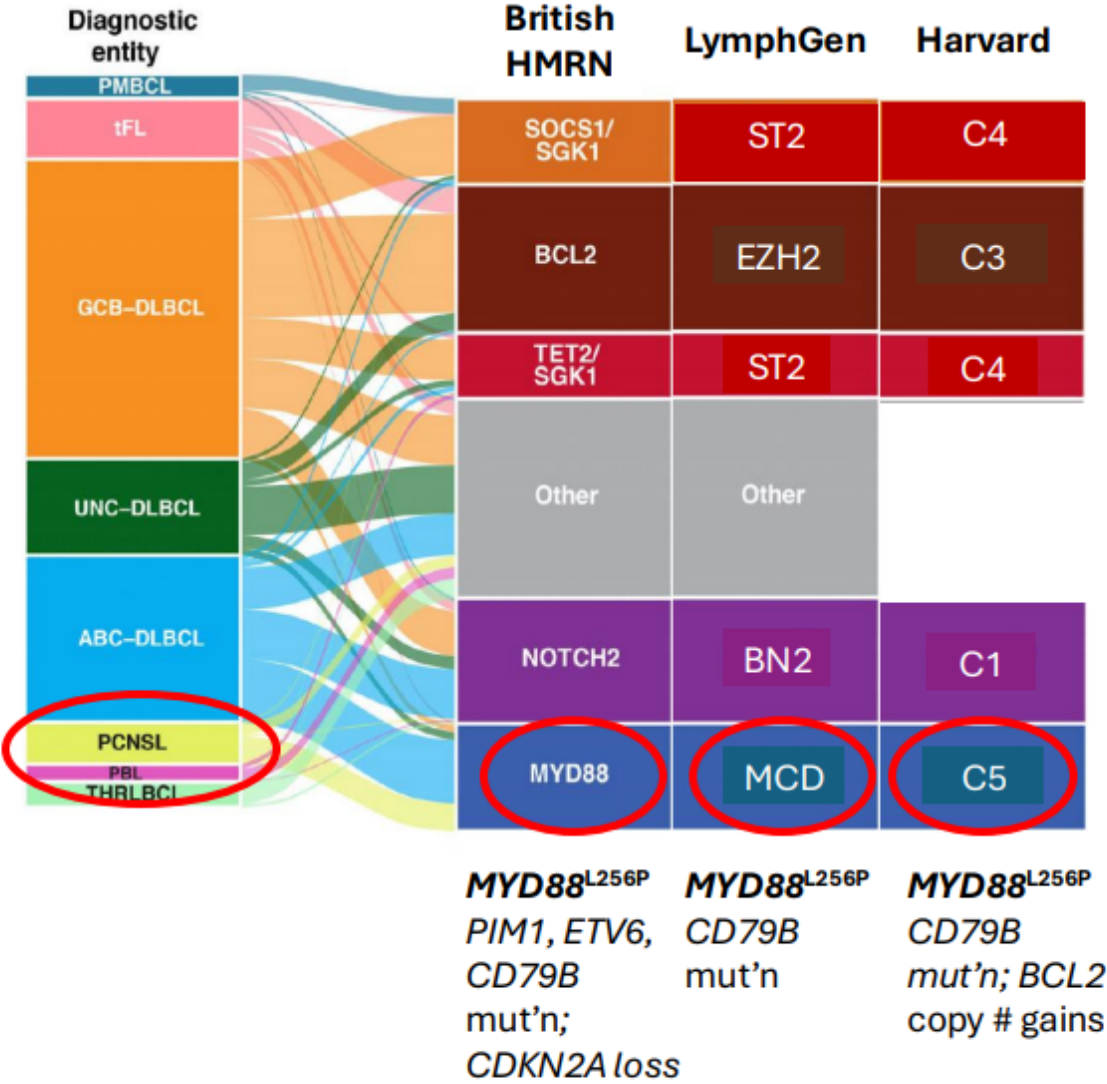
Intermediate survival for  
NEC and NOTCH2  
clusters (**mix**)

Poor outcome in for  
MYD88 cluster (**ABC**)

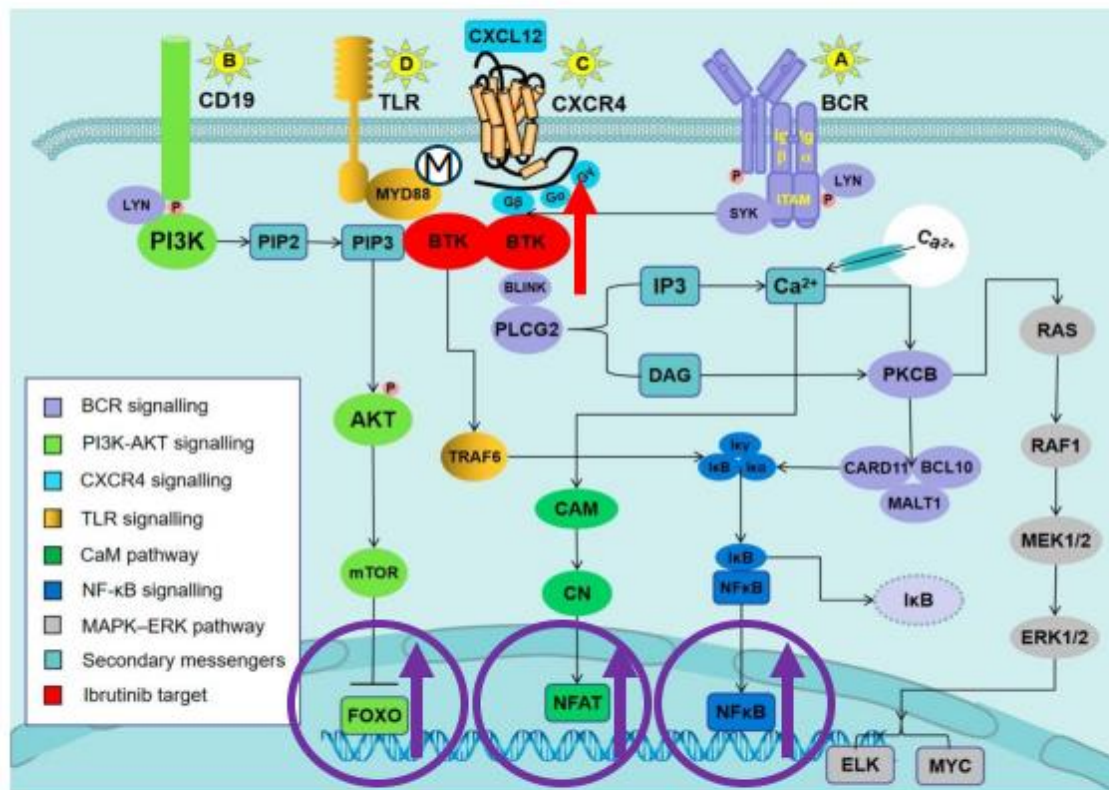
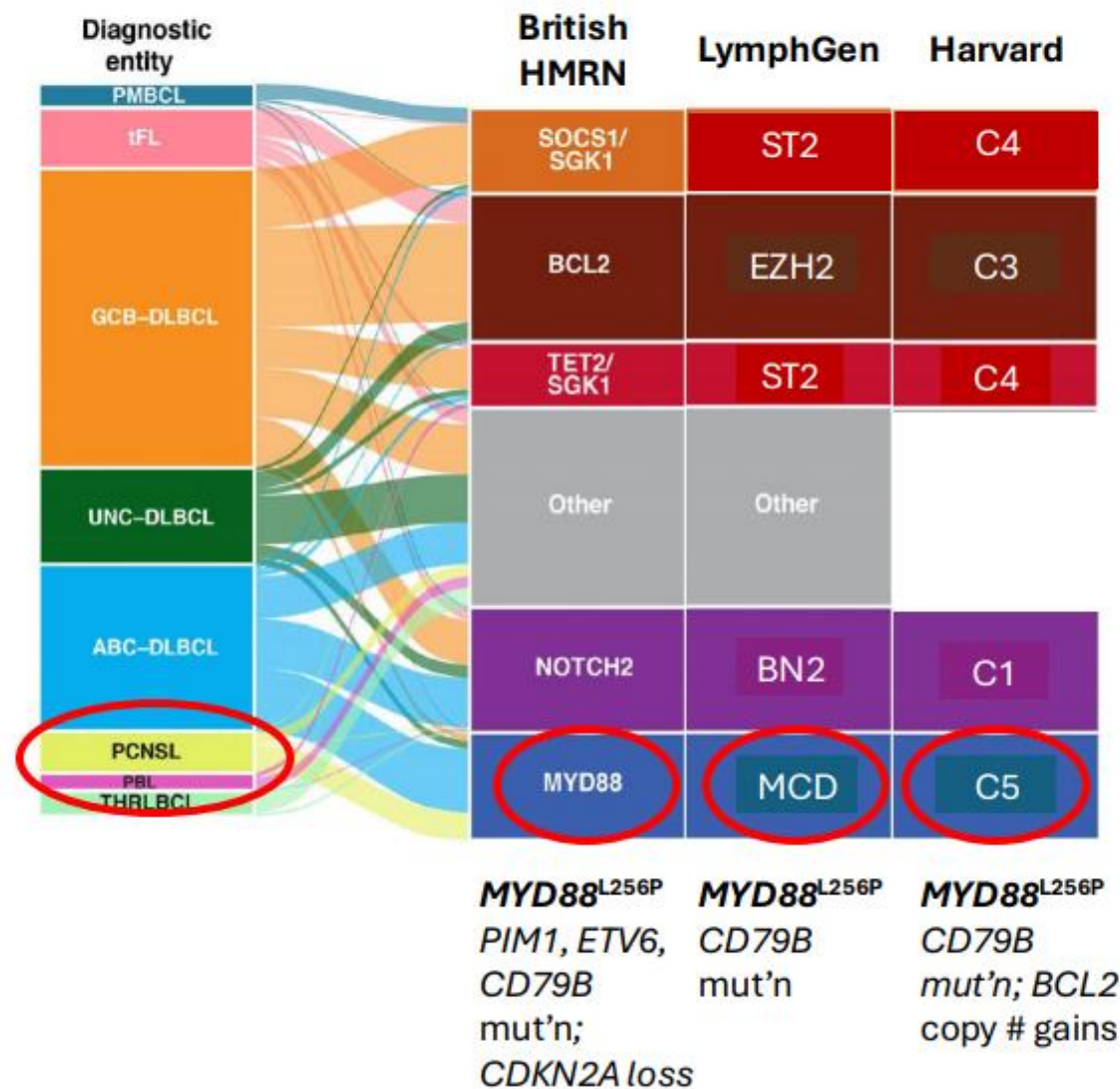




# MYD88 Testing



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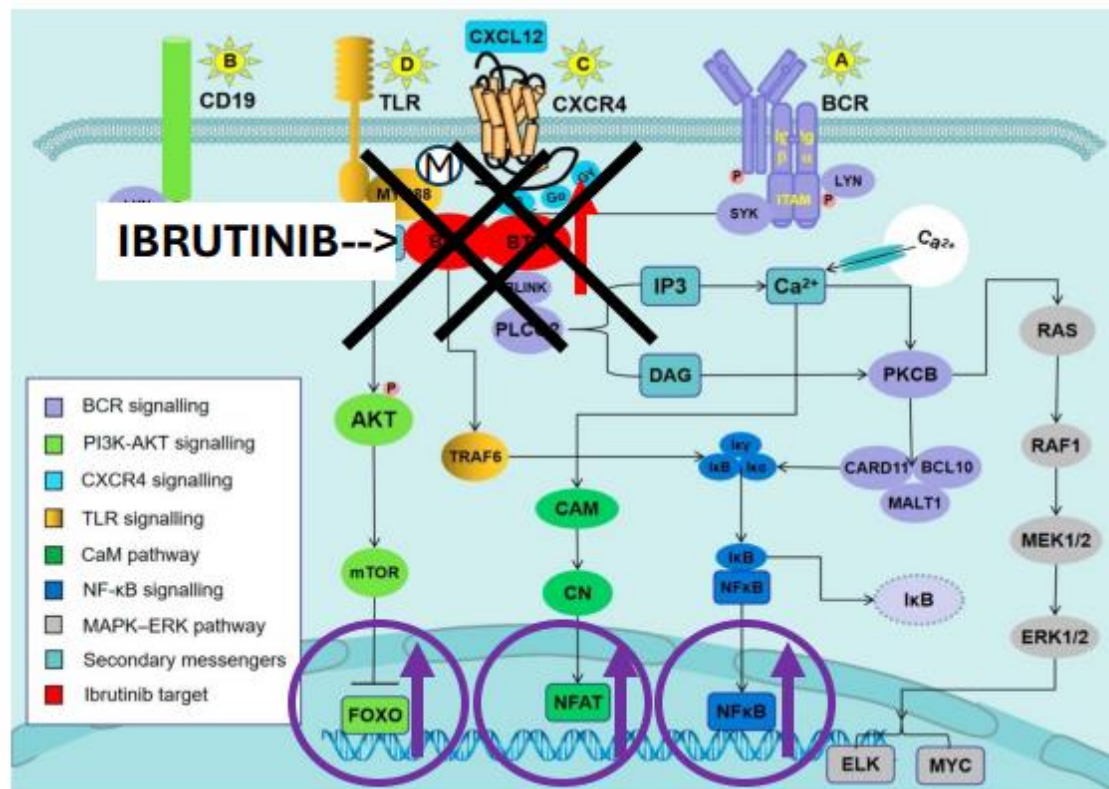
# MYD88 Testing

Diagnostic entity	British HMRN	LymphGen	Harvard
PMBCL			
tFL			
GCB-DLBCL	SOCS1/SGK1	ST2	C4
	BCL2	EZH2	C3
	TET2/SGK1	ST2	C4
UNC-DLBCL	Other	Other	
ABC-DLBCL	NOTCH2	BN2	C1
PCNSL	MYD88	MCD	C5
PBL			
THRLBCL			

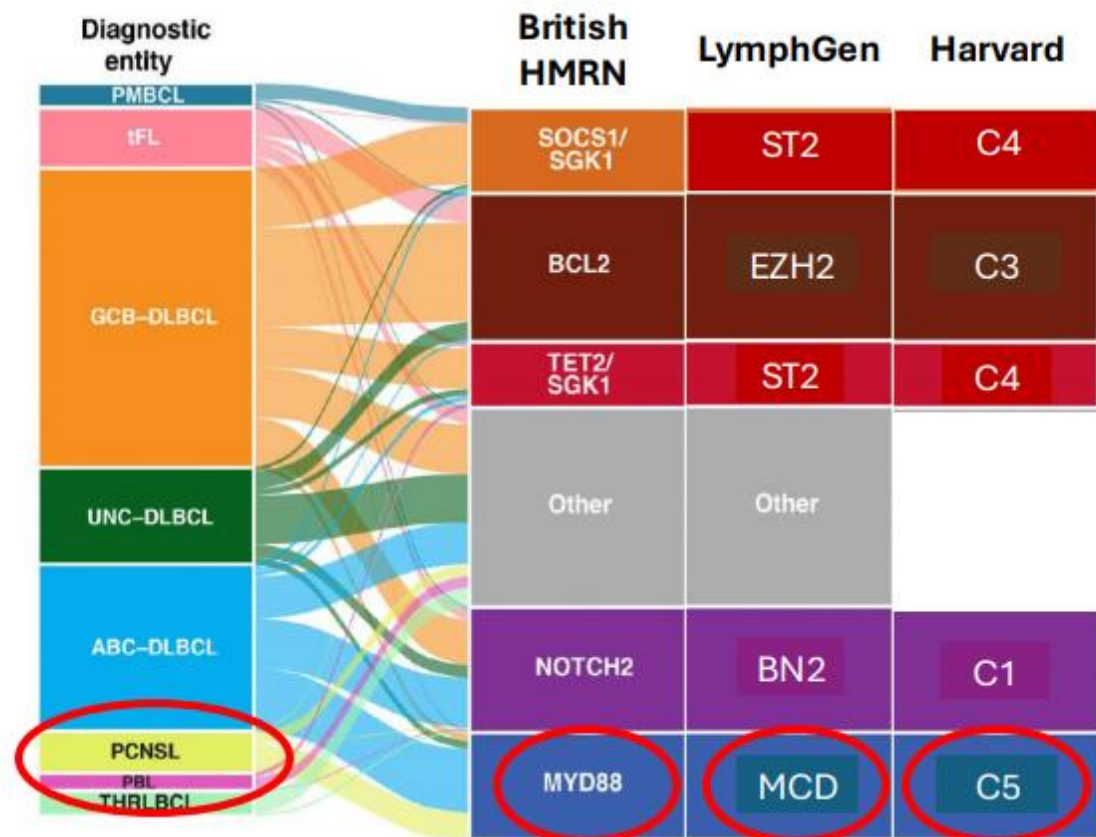
**MYD88<sup>L256P</sup>**  
*PIM1, ETV6,*  
*CD79B*  
*mut'n;*  
*CDKN2A loss*

**MYD88<sup>L256P</sup>**  
*CD79B*  
*mut'n*

**MYD88<sup>L256P</sup>**  
*CD79B*  
*mut'n; BCL2*  
*copy # gains*



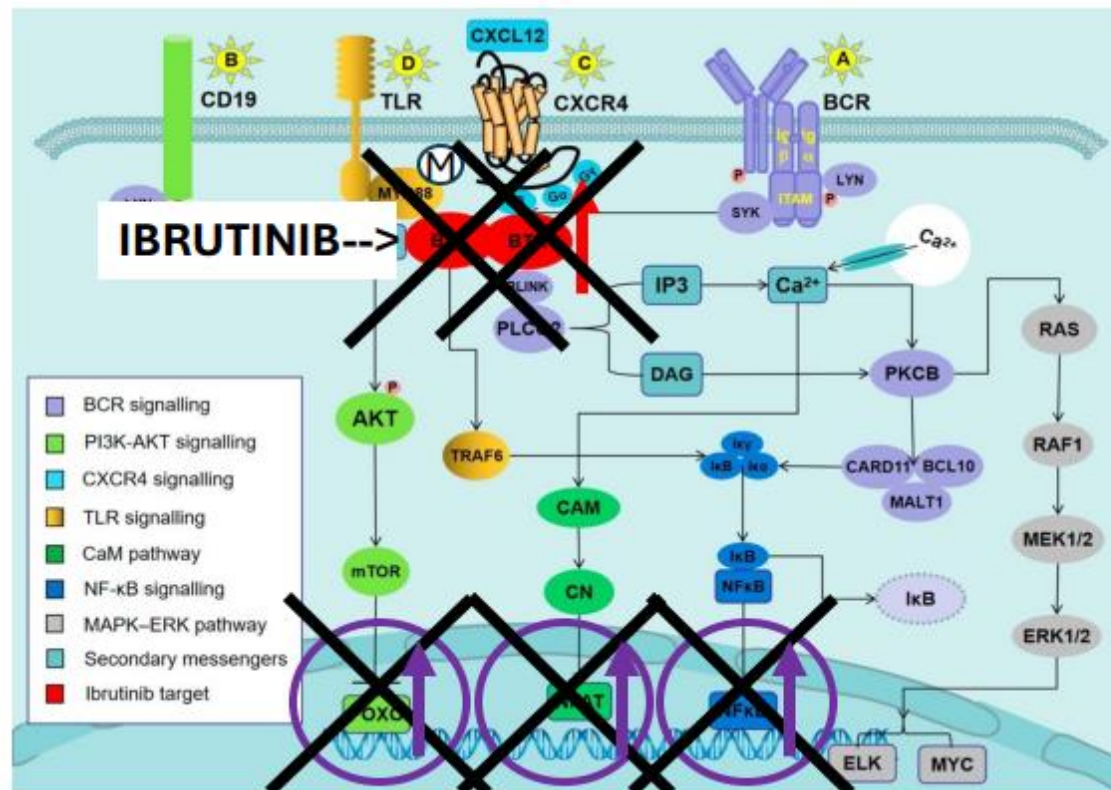
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**MYD88<sup>L256P</sup>**  
CD79B  
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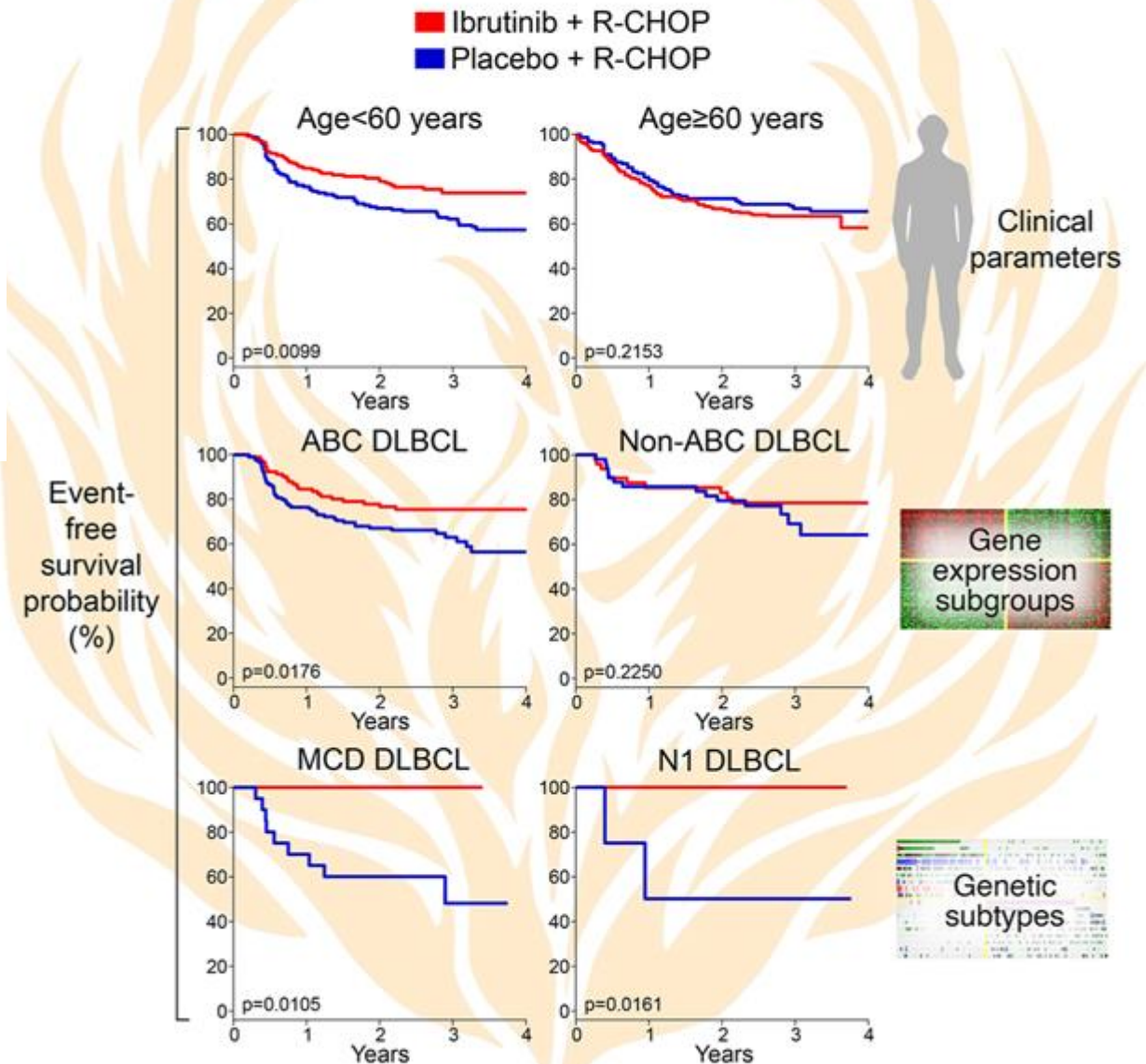


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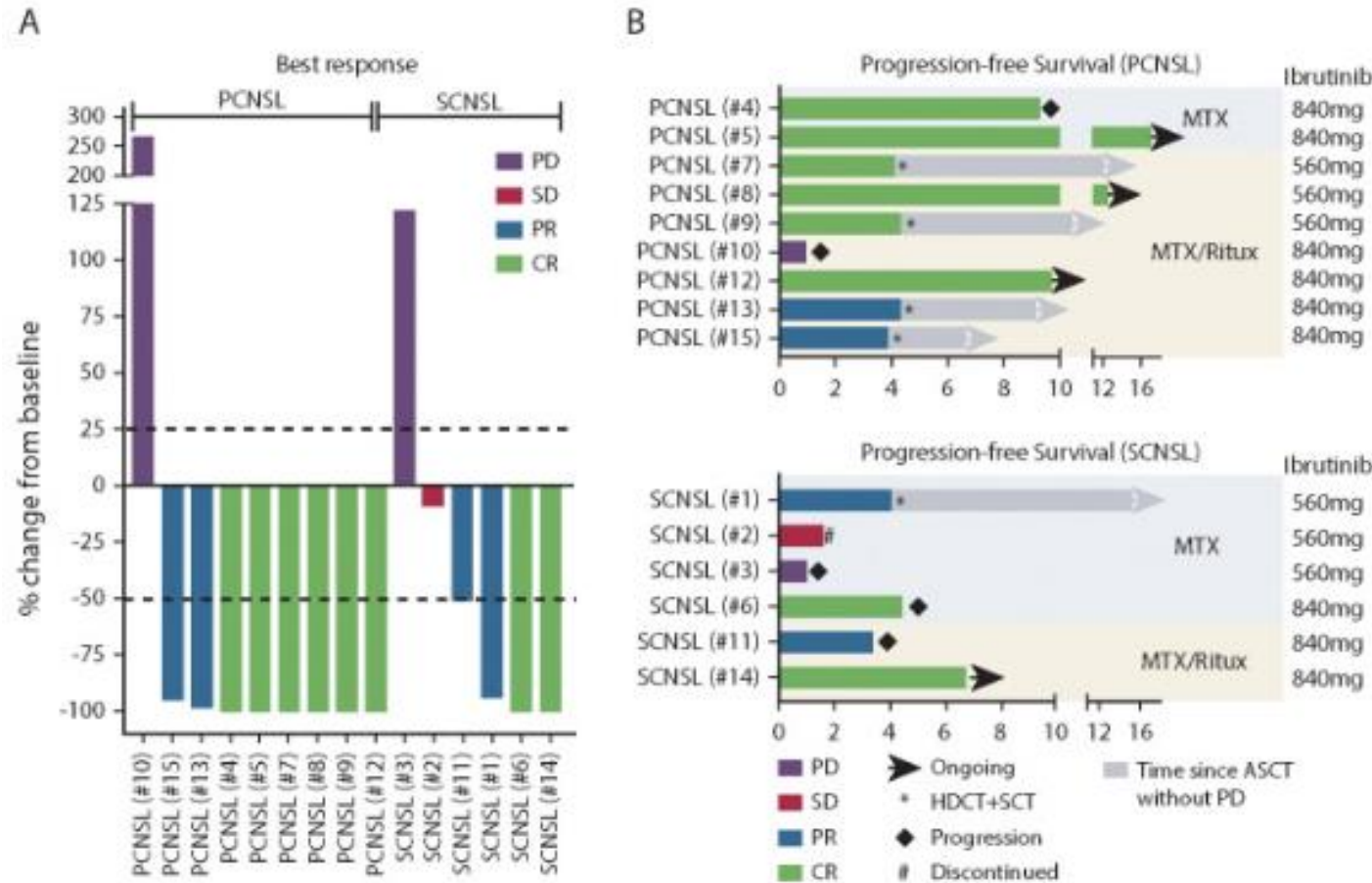
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Two genetic subtypes of DLBCL – MCD and N1 – have 100% survival when treated with the BTK inhibitor ibrutinib plus R-CHOP chemotherapy but ≤50% survival when treated with R-CHOP alone.

## Phoenix Phase III Clinical Trial in Previously Untreated Non-GCB Diffuse Large B Cell Lymphoma



# Phase 1b trial of an ibrutinib-based combination therapy in recurrent/refractory CNS lymphoma



- Explored the sequential combination of ibrutinib with high-dose methotrexate (HD-MTX) and rituximab in patients with R/R CNS lymphoma (both primary and secondary)
- Eight of 9 (89%) primary CNS lymphoma patients and 4 of 6 (67%) secondary CNS lymphoma patients responded to ibrutinib-based combination therapy.



Patient started on TEDDI-R + brain XRT

(temozolomide, etoposide, doxorubicin, dexamethasone, ibrutinib, and rituximab)

April 2021: MRI showed some response; no clinical progression of disease

Patient started on TEDDI-R + brain XRT

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September 2021: Admitted with bilateral PEs

MRI showed no evidence of CNS lymphoma

PET-CT showed no evidence of extra-CNS disease

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**Last follow-up: No evidence of disease**

Thank you