

# Classification of Leukemias and Lymphomas: Increasing Role of Molecular Testing

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# Objectives

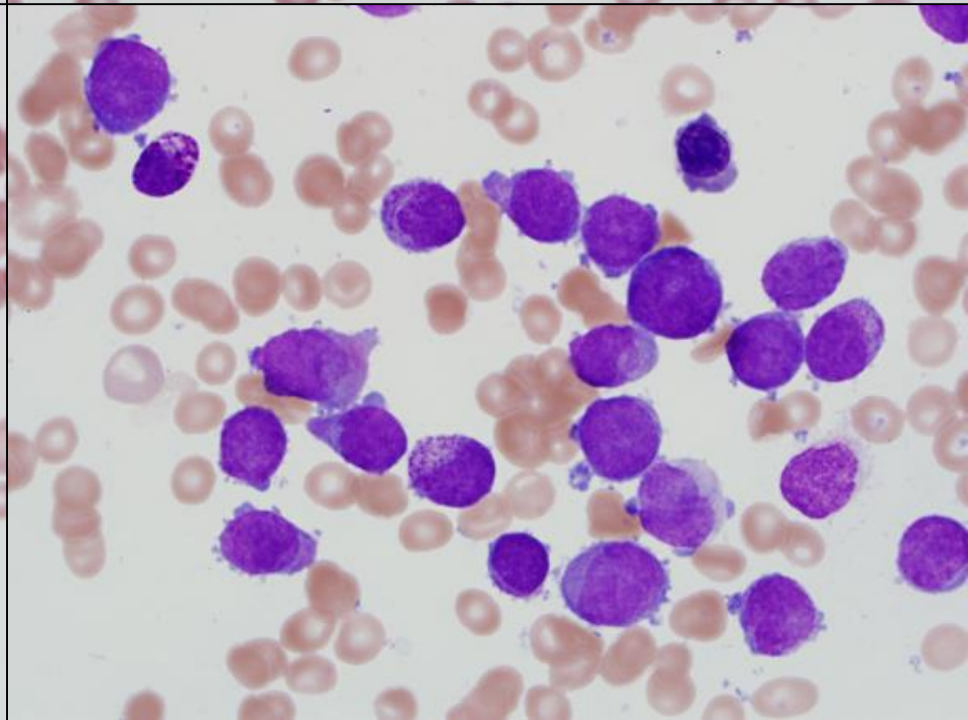
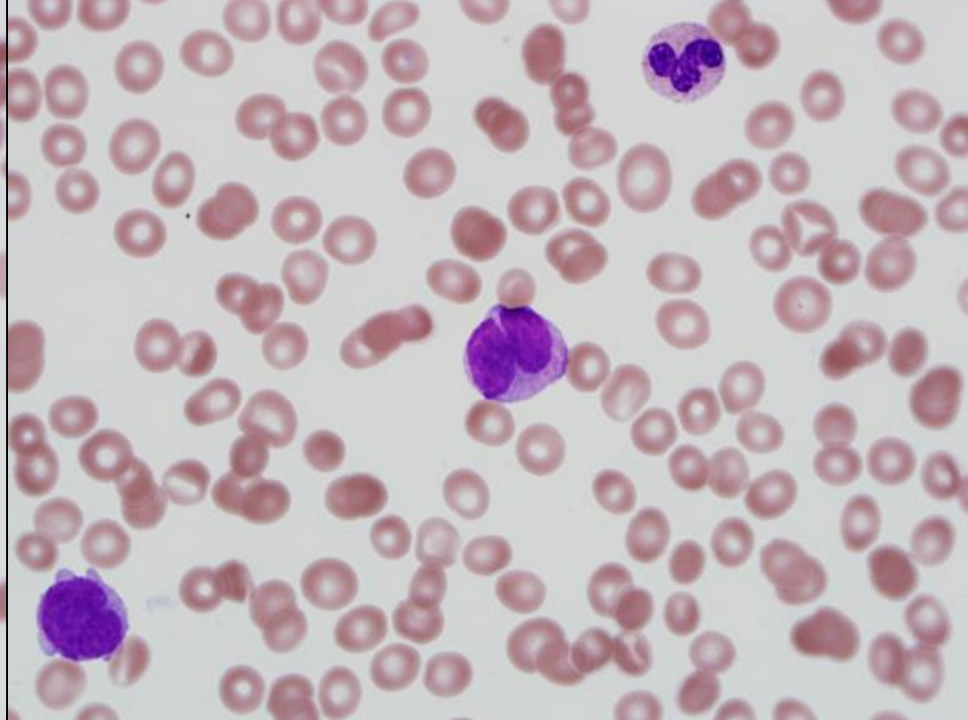
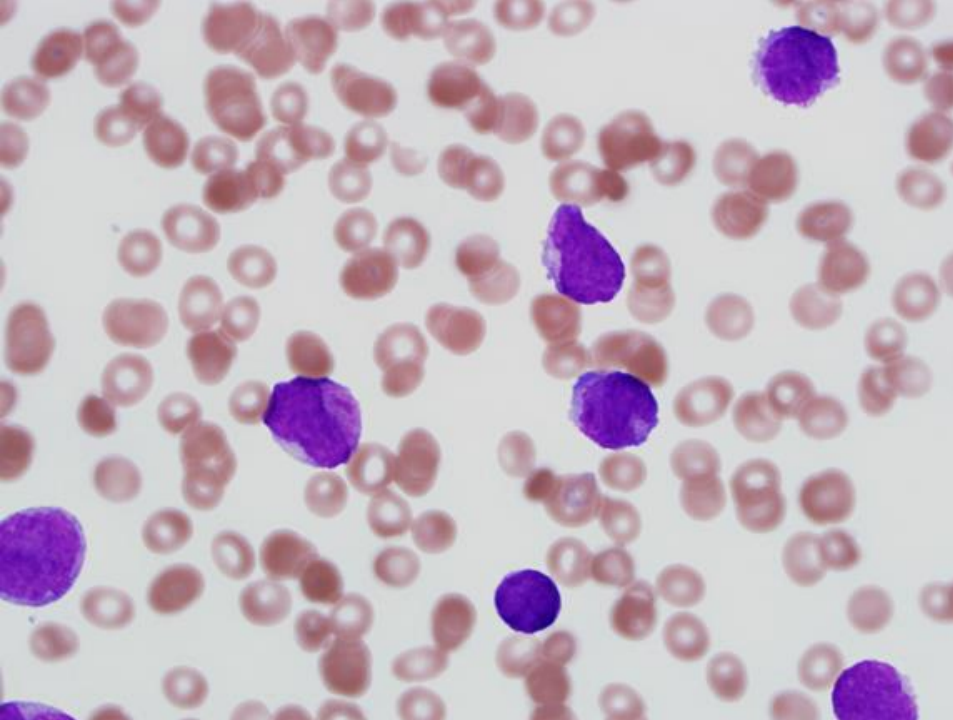
- Discuss how molecular and cytogenetic findings are used to:
  - More precisely classify leukemia and lymphoma.
  - Add prognostic information
    - May evolve into classification in the future.
- Discuss clonal hematopoiesis and “pre-MDS.”
- Include a mix of cases where molecular and/or cytogenetic data were used as above.

# Genetically Defined Hematologic Malignancies

- CML was the first
  - Diagnosis requires t(9;22)
- Acute leukemia diagnosis: recurrent cytogenetic changes and mutations.
  - Diagnose AML with <20% blasts.
- Lymphoma: Limited role now, but expect it to increase.

# Acute Myeloid Leukemia with Recurrent Genetic Abnormalities

- \*AML with t(8;21)(q22;q22), *RUNX1-RUNX1T1 (CBFA/ETO)*
- \*AML with inv(16)(p13q22) or t(16;16)(p13;q22), *CBFB-MYH11*
- \*APL with t(15;17)(q22;q11-12), *PML-RARA*
- AML with t(9;11)(p22;q23), *MLLT3-MLL and other balanced translocations of 11q23 (MLL)*
- AML with t(6;9)(p23;q34), *DEK-NUP214*
- AML with inv(3)(q21;q26.2) or t(3;3)(p13;q13), *GATA2, MECOM*
- AML with t(1;22)(p13;q13), *RBM15-MKL1*



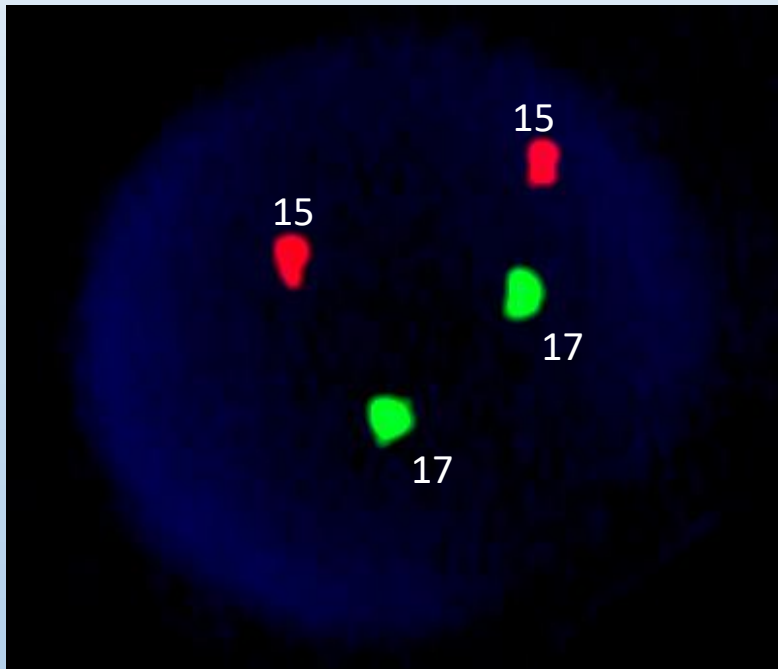
# Morphology Highly Suggestive of Acute Promyelocytic Leukemia

- Patients can present with DIC.
  - Thrombocytopenia, schistocytes.
- Emergent diagnosis required.
- Patients respond well to all-trans retinoic acid (ATRA).
- Treatment regimen distinct from other AMLs: ATRA, arsenic.
- Good prognosis.

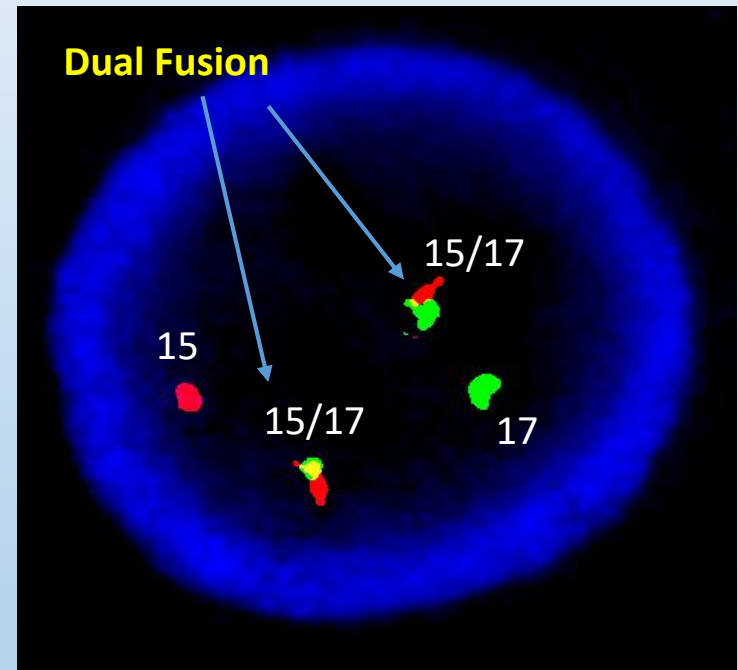


# t(15;17) *PML-RARA* fusion is diagnostic of APL

Normal



Abnormal



- FISH (or RT-PCR) is recommended at diagnosis for quick turn-around time.
- RT-PCR for *PML-RARA* fusion product for disease monitoring.

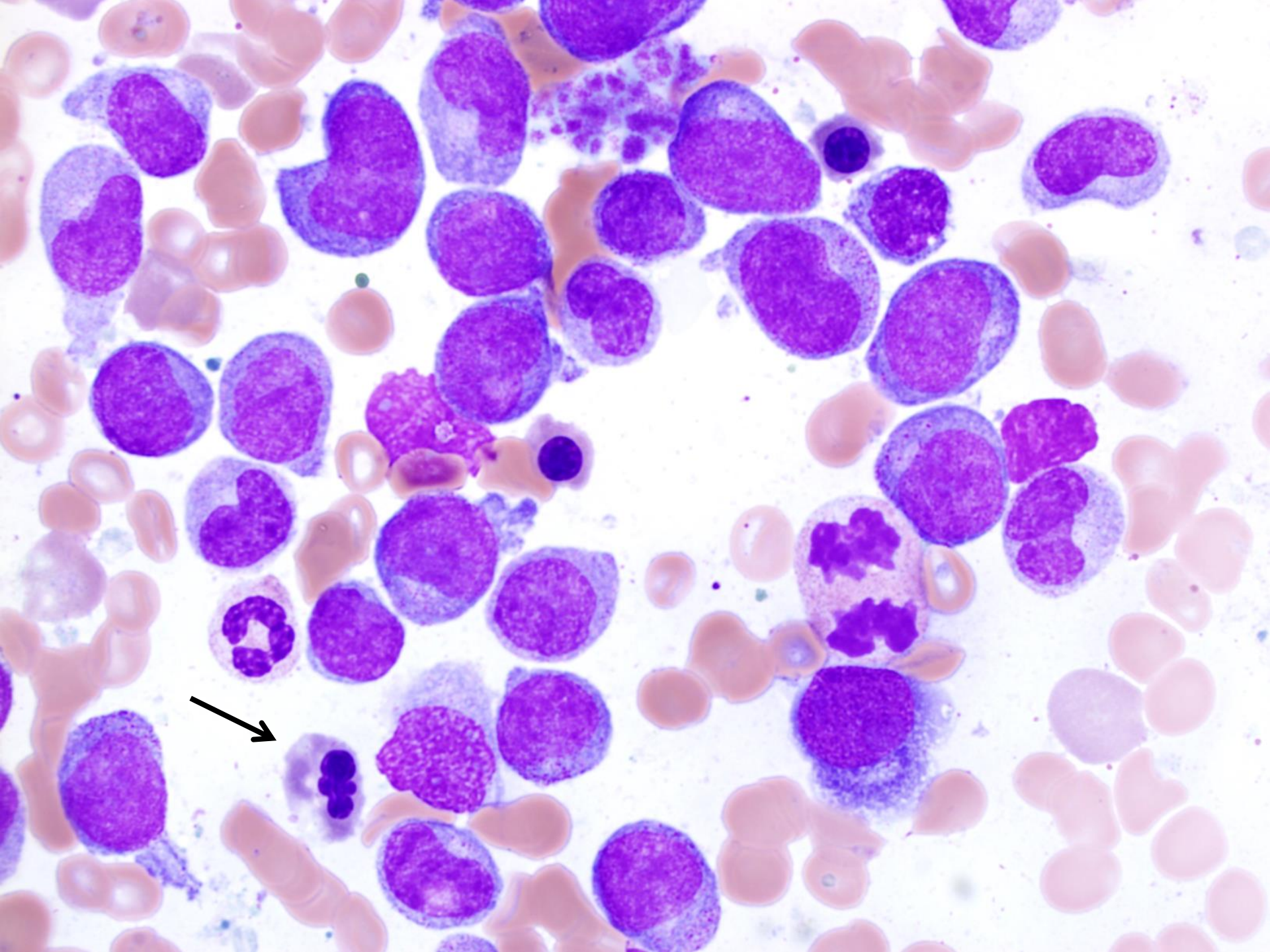
# Cytogenetically Normal AMLs: Mutation Studies for Classification

- Mutations
  - *NPM1* (nucleophosmin)
  - *CEBPA* (CCAAT enhancer binding protein alpha)
- WHO 2017 Categories:
  - AML with mutated *NPM1*
  - AML with biallelic mutations in *CEBPA*
  - AML with mutated *RUNX1* (*provisional*)



# Case:

- 54-year-old woman noticed increased bleeding while brushing her teeth.
- CBC:
  - WBC: 7.2 k/ $\mu$ L with occasional circulating blasts
  - Hgb: 10.1 g/dL
  - HCT: 30.7%
  - PLT: 97 k/ $\mu$ L
- Bone marrow evaluation.



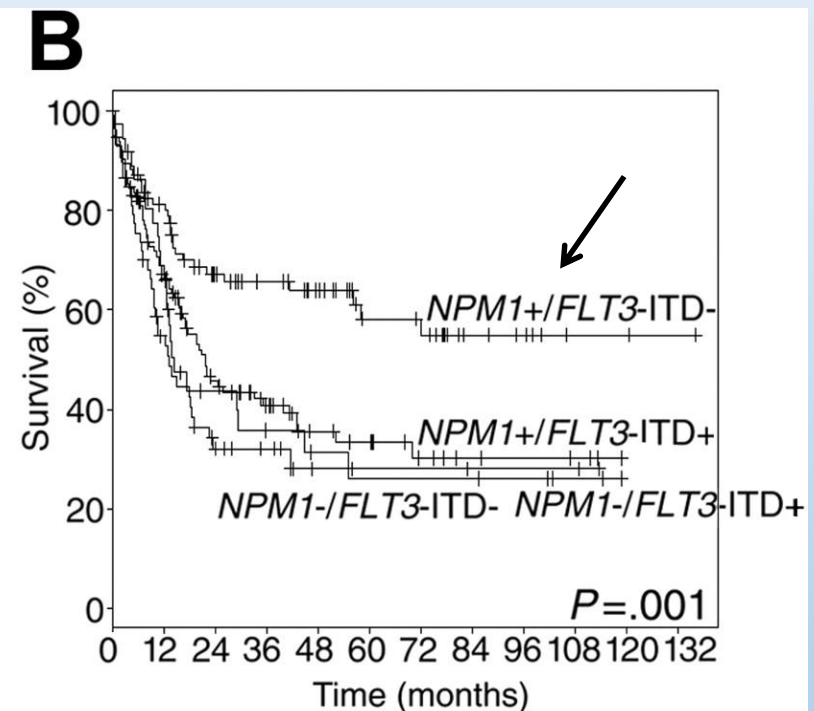
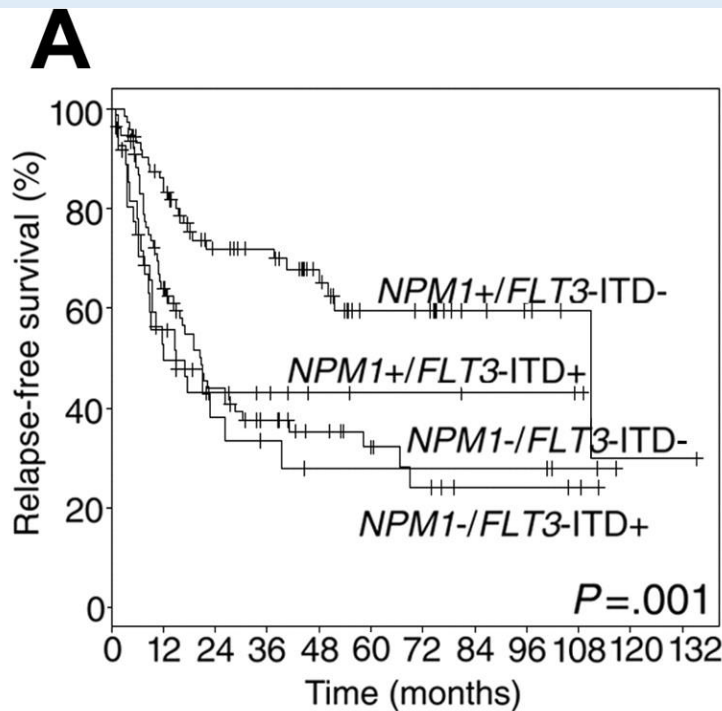
# Hematopathology Work-up

- Blasts represented 57% of the bone marrow nucleated cells by differential count.
- Flow cytometry identified a prominent blast population which expressed CD13, CD33, CD117, CD34, CD7, myeloperoxidase and HLA-DR.
- Diagnosis: acute myeloid leukemia.
- 1-4 days later, normal AML FISH panel.
  - APL t(15;17) result first.
- 7 days later, normal karyotype: 46,XX [20].
- 14 days later, NGS panel results: *NPM1* mutation, no *FLT3* or *CEBPA* mutations.

# Diagnosis: AML with mutated *NPM1*

- Original pathology report is amended to include precise classification.
- Significant dysplasia not associated with worse prognosis in this setting.
- Monitoring option: *NPM1* mutation by quantitative RT-PCR to detect minimal residual disease (MRD).
- Prognostic information.

# *NPM1* mutated, *FLT3* Wild Type Identifies a Subgroup with a Better Prognosis



Number at risk:

<i>NPM1</i> +/ <i>FLT3</i> -ITD-	74	58	40	36	26	15	13	6	4	2	1	1
<i>NPM1</i> -/ <i>FLT3</i> -ITD-	78	47	28	20	14	9	6	3	3	2	0	0
<i>NPM1</i> +/ <i>FLT3</i> -ITD+	37	16	10	8	4	3	3	2	2	1	0	0
<i>NPM1</i> -/ <i>FLT3</i> -ITD+	28	15	8	6	4	4	4	4	4	2	0	0

<i>NPM1</i> +/ <i>FLT3</i> -ITD-	86	66	45	38	30	19	17	8	6	2	2	1
<i>NPM1</i> +/ <i>FLT3</i> -ITD+	117	71	42	29	18	14	8	5	4	3	0	0
<i>NPM1</i> -/ <i>FLT3</i> -ITD-	59	27	14	10	4	3	3	2	2	2	0	0
<i>NPM1</i> -/ <i>FLT3</i> -ITD+	38	24	11	8	6	5	5	5	4	2	0	0



# AML Summary

- Morphology still the first and most important step.
  - Most cases still diagnosed based on blast percentage.
- Cytogenetics are critical.
  - Define AML in absence of increased blasts.
  - Provide prognostic information.
- Molecular testing.
  - Specific classification and prognostic information.
- Amount of testing required depends on the clinical situation.

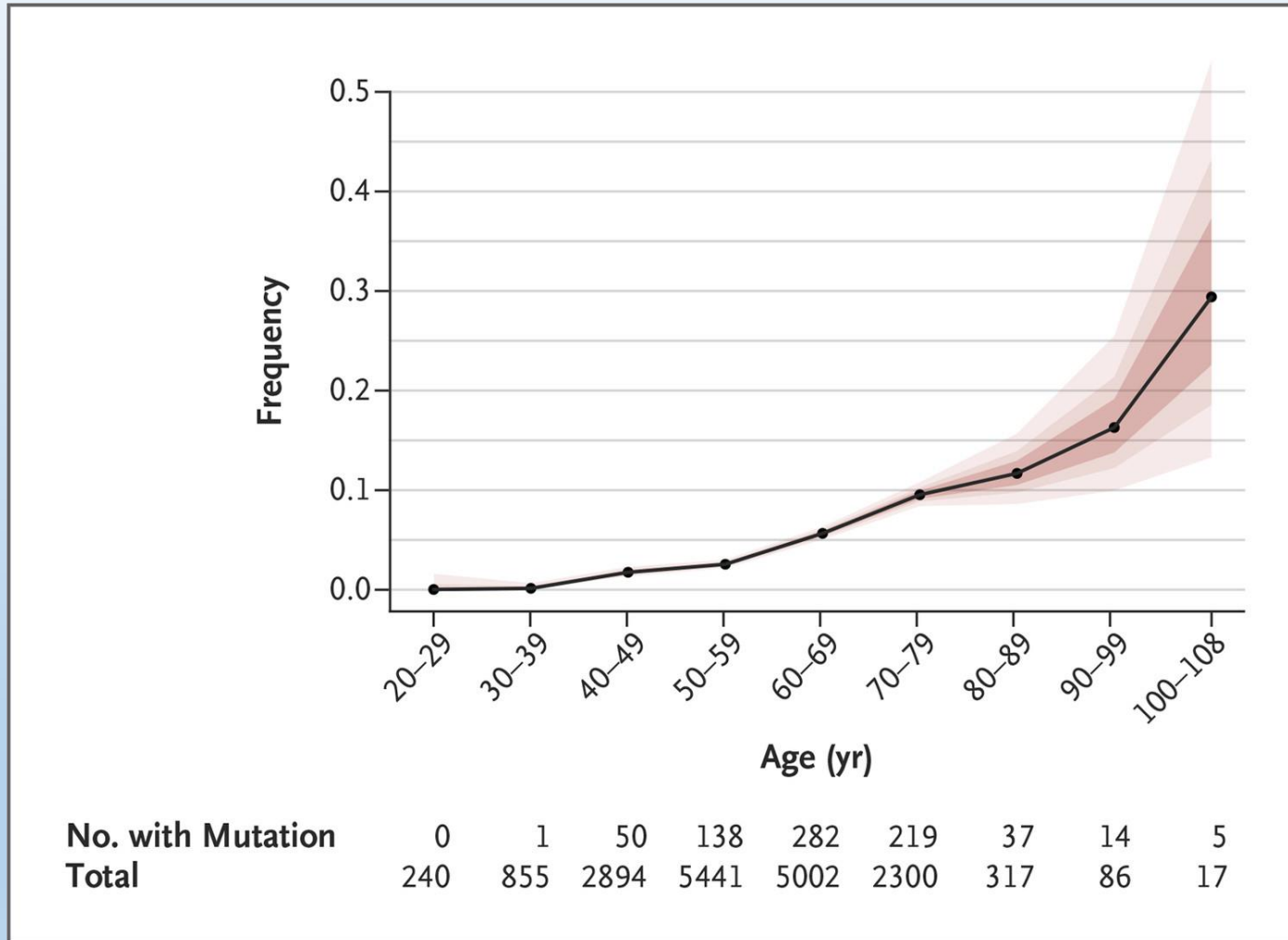
# Clonal Hematopoiesis +/- Cytopenias



# Clonal Hematopoiesis

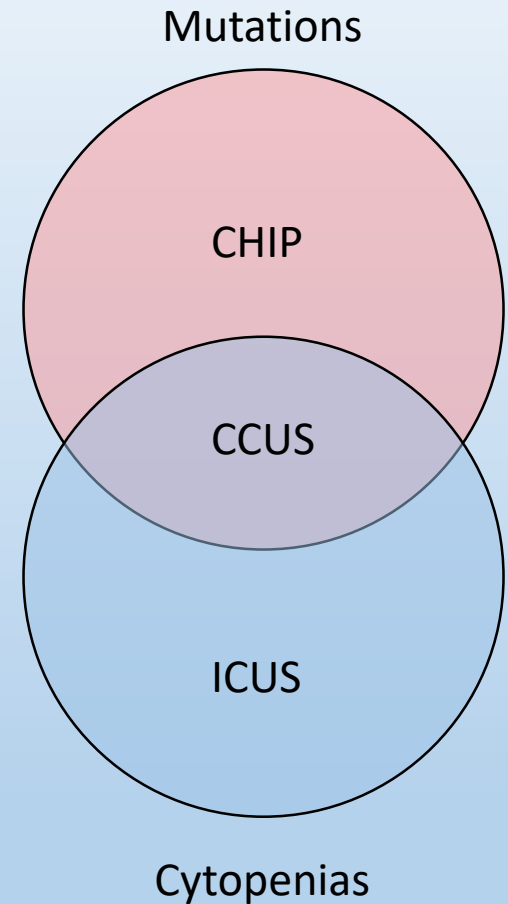
- Mutations in leukemia-associated driver genes are commonly detected in apparently healthy older people with normal blood counts.
  - Predominantly *DNMT3A*, *TET2*, and *ASXL1*
- Mutations are present in a clone and confer a survival advantage.
  - Would not be detectable in a single cell.
- Associated with risk of developing myeloid neoplasms (sometimes lymphoid) or cardiovascular events.

# Prevalence of Somatic Mutations Increases with Age

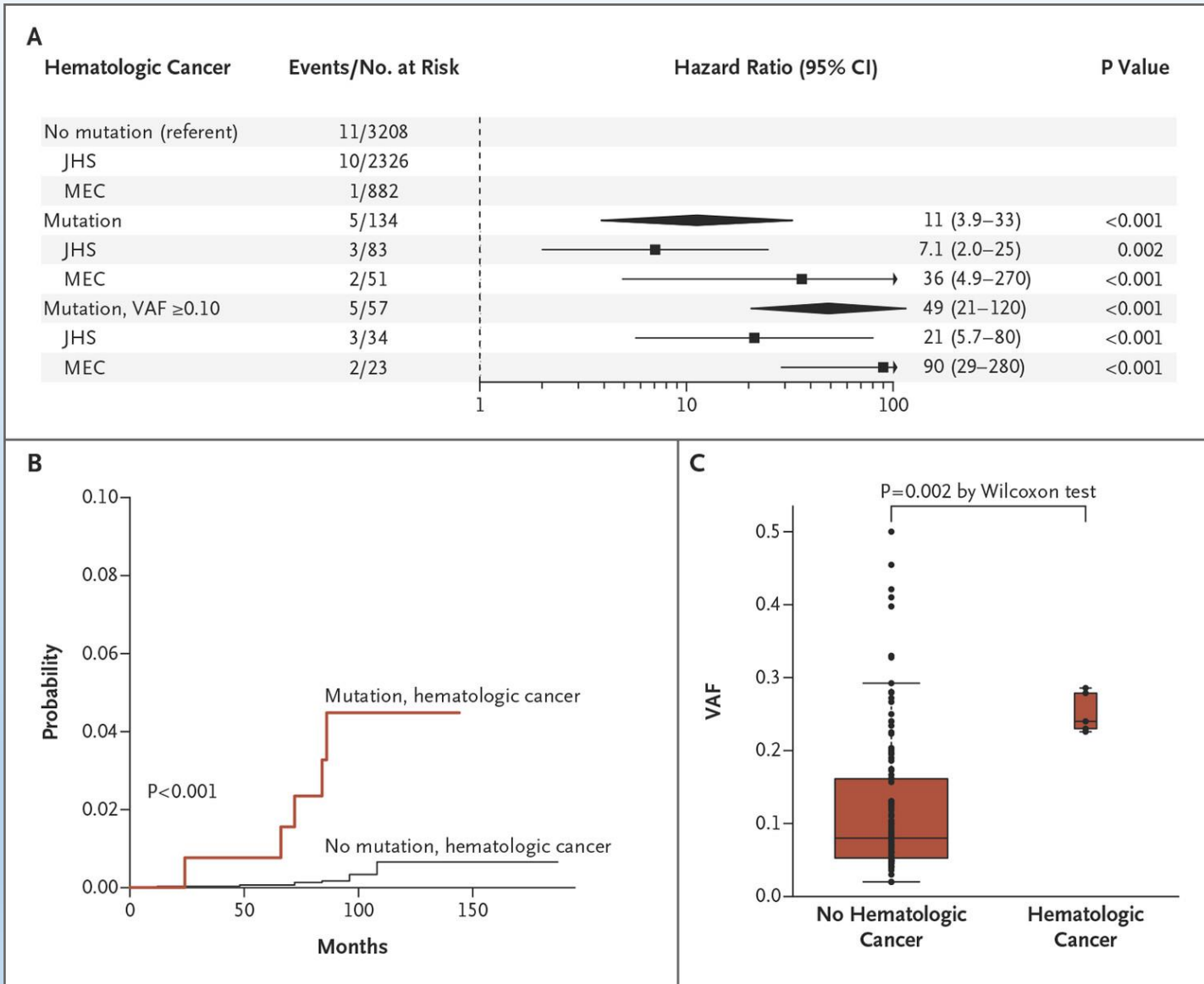


# Terminology

- CHIP: Clonal Hematopoiesis of Indeterminate Potential.
  - By definition, patients lack cytopenias.
- CCUS: Clonal Cytopenias of Undetermined Significance.
- ICUS: Idiopathic Cytopenias of Undetermined Significance (no mutation detected).
- Clonality  $\neq$  Malignancy
- MDS: Fulfills WHO 2017 criteria.

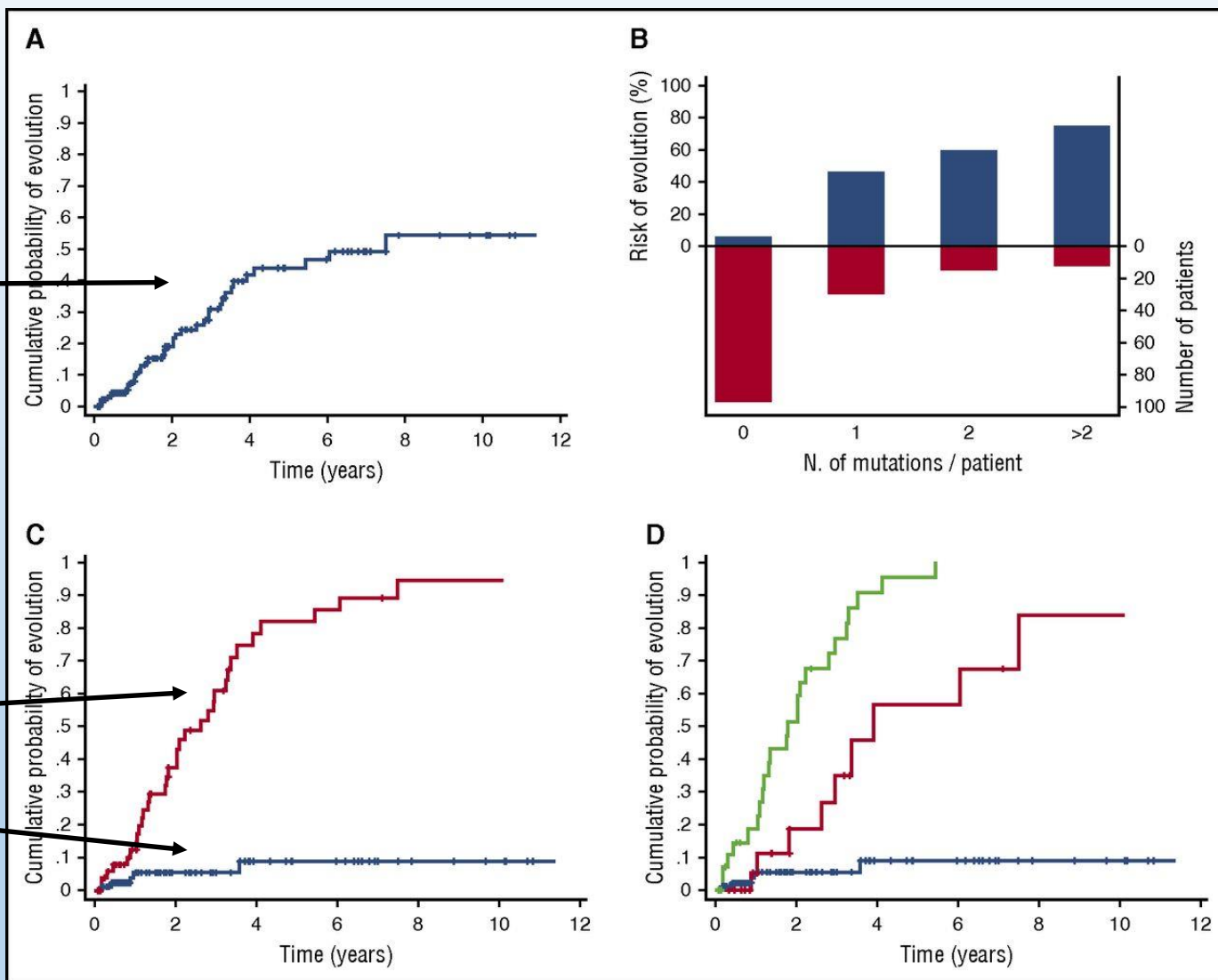


# Risk of Hematologic Cancers is Related to Mutant Variant Allele Frequency



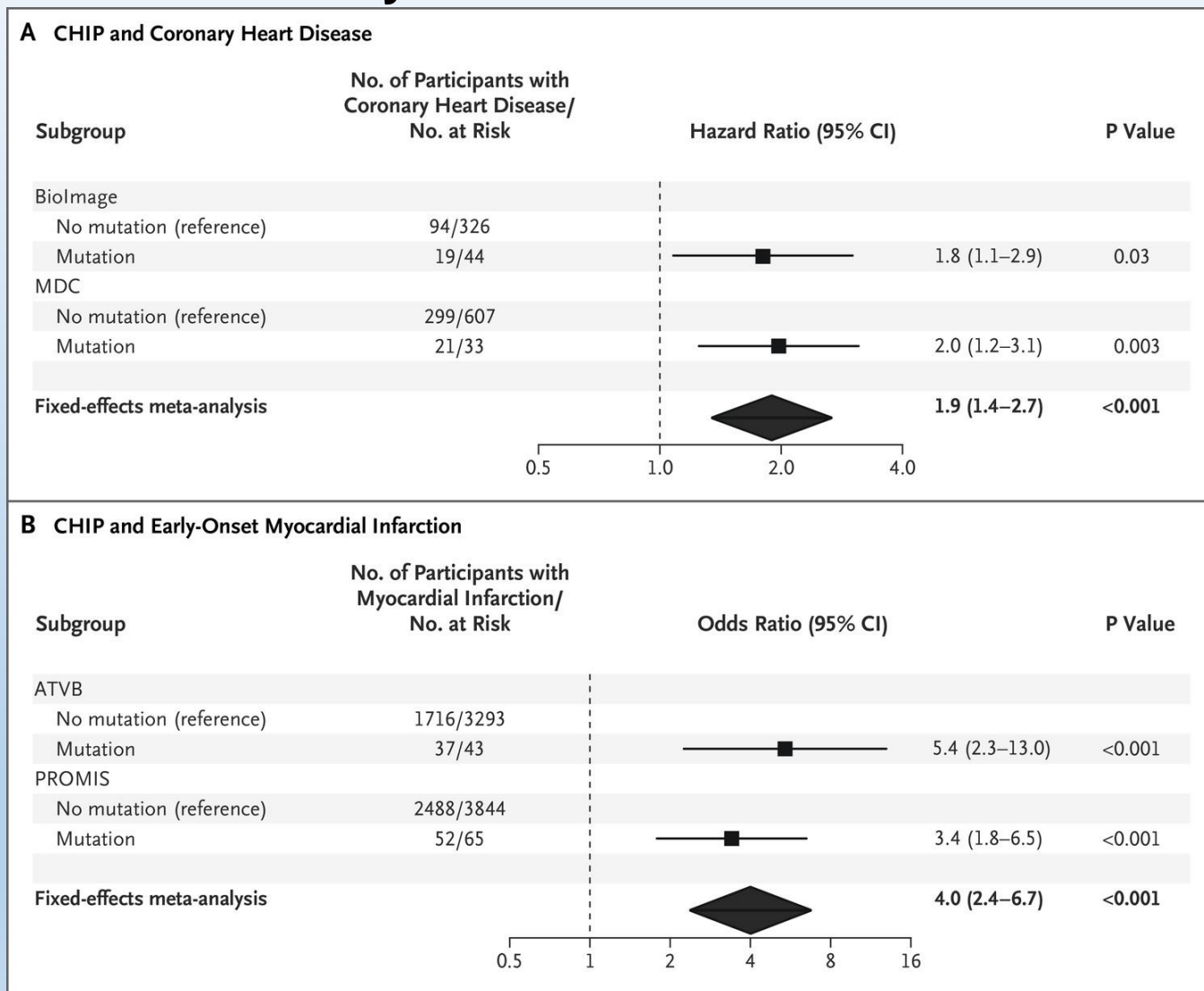
# Probability of progression to a myeloid neoplasm in patients with cytopenias is higher when mutation(s) are detected

All patients with cytopenias



Luca Malcovati et al. Blood 2017;129:3371-3378

# Association between Clonal Hematopoiesis of Indeterminate Potential (CHIP) and Coronary Heart Disease and Early-Onset Myocardial Infarction.



# NGS Sequencing Panels

- Include many myeloid driver mutations including those most common in CHIP.

## ARUP's Myeloid NGS Panel:

Genes – *ASXL1*, *ASXL2*, *BCOR*, *BCORL1*, *BRAF*, *CALR*, *CBL*, *CEBPA*, *CSF3R*, *DNMT1*, *DNMT3A*, *EED*, *ELANE*, *ETNK1*, *ETV6*, *EZH2*, *FAM5C*, *FLT3*, *GATA1*, *GATA2*, *HNRNPK*, *IDH1*, *IDH2*, *JAK2*, *JAK3*, *KDM6A*, *KIT*, *KRAS*, *LUC7L2*, *MAP2K1*, *MLL*, *MPL*, *NOTCH1*, *NPM1*, *NRAS*, *NSD1*, *PHF6*, *PRPF40B*, *PRPF8*, *PTPN11*, *RAD21*, *RUNX1*, *SETBP1*, *SF1*, *SF3A1*, *SF3B1*, *SMC1A*, *SMC3*, *SRSF2*, *STAG2*, *SUZ12*, *TET2*, *TP53*, *U2AF1*, *U2AF2*, *WT1*, *ZRSR2*



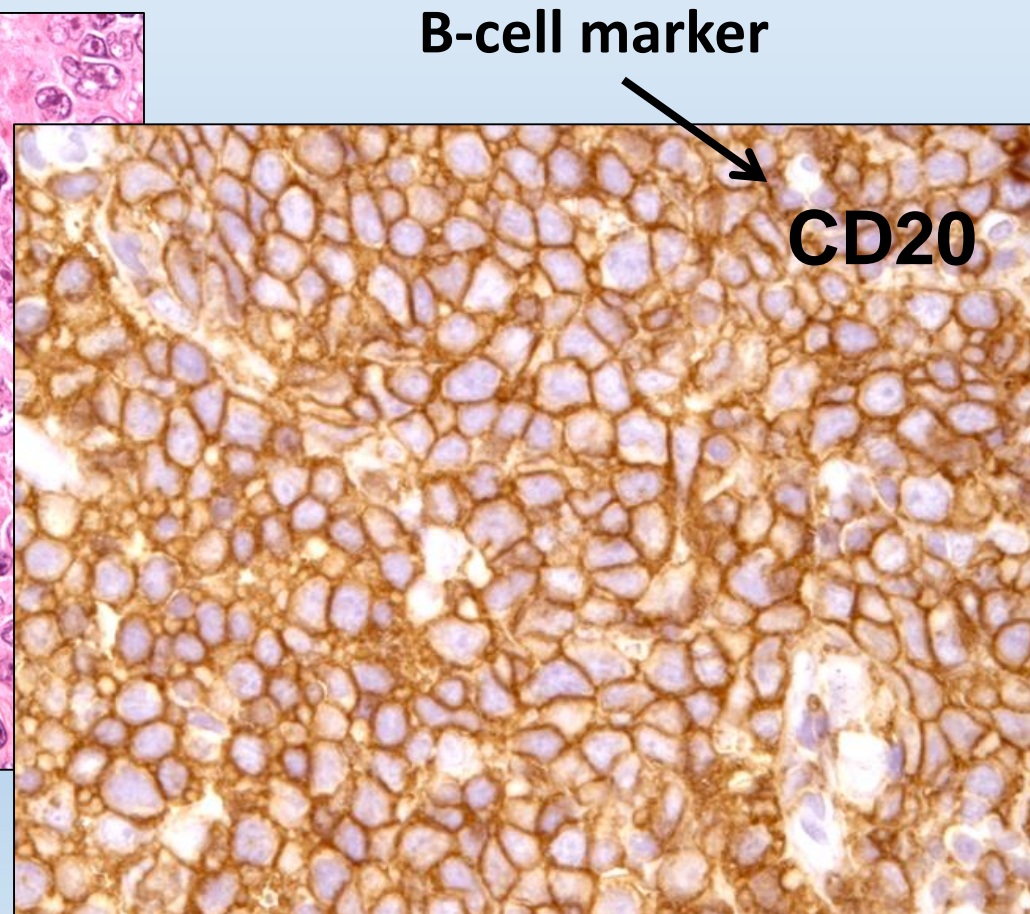
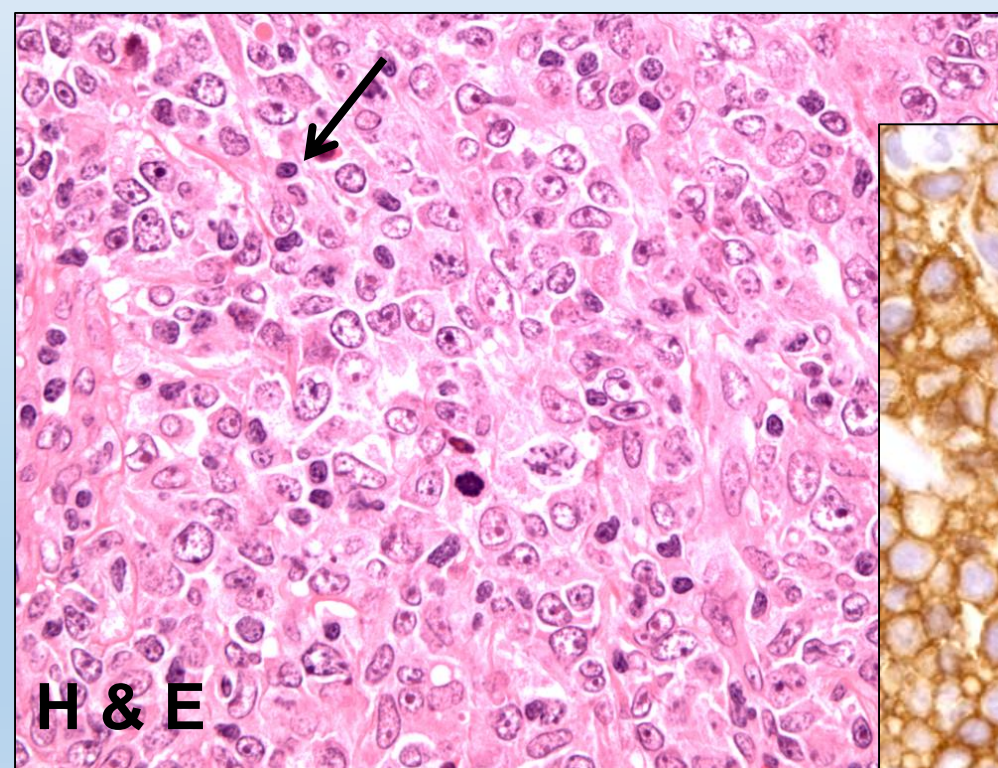
# CHIP and Clonal Cytopenias: Key Points

- CHIP: when you run NGS on normal older patients.
- ICUS: Cytopenias without (detected) mutations
- CCUS: MDS without dysplasia.
  - Similar to cytogenetically-defined MDS.
  - Likely considered as such in future classification.
- Cardiovascular disease is much more common than MDS, so this is the bigger risk for patients with CHIP.
  - Likely involves clonal monocyte/macrophages and increased local inflammation in atherosclerotic plaques.
  - Anti-inflammatory anti-IL-1beta monoclonal antibody decreased recurrent events after MI.
  - Anti-inflammatory therapy for patients with CHIP?
  - Fasting lipid panel...and NGS myeloid panel?

# Genetic Testing in Lymphoma

# Diffuse Large B-cell Lymphoma

- Ancillary testing required for diagnosis: CD20.



# Diffuse Large B-cell Lymphoma

## Ancillary Testing

- Useful for for sub-classification and/or prognostic information
  - EBER
  - FISH for *MYC*, *BCL2*, *BCL6* translocations
  - MIB-1
  - GC vs. non-GC subtyping
  - *MYC*, *BCL2* protein expression

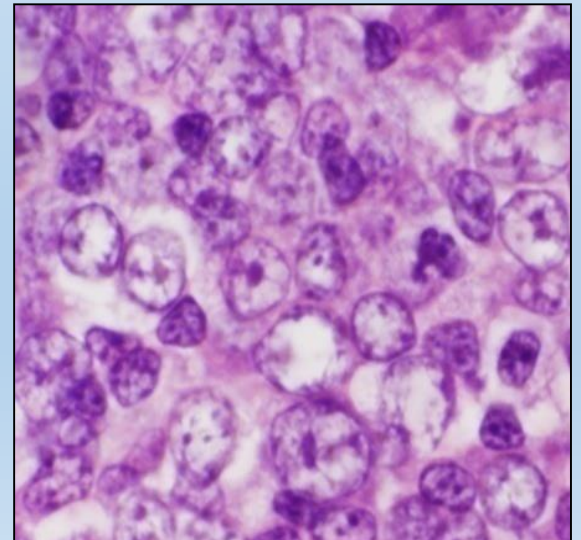
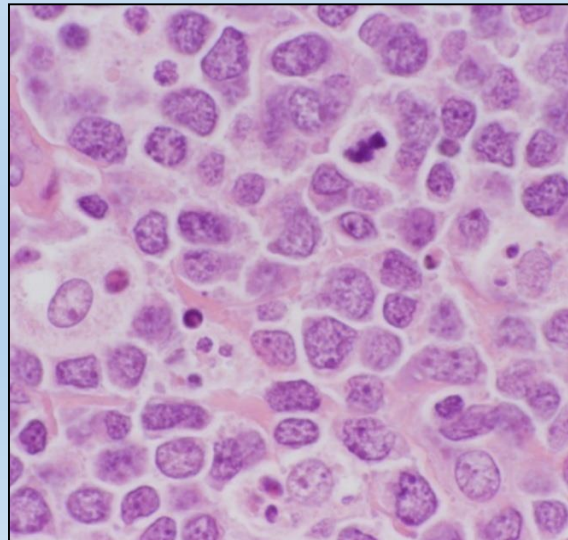
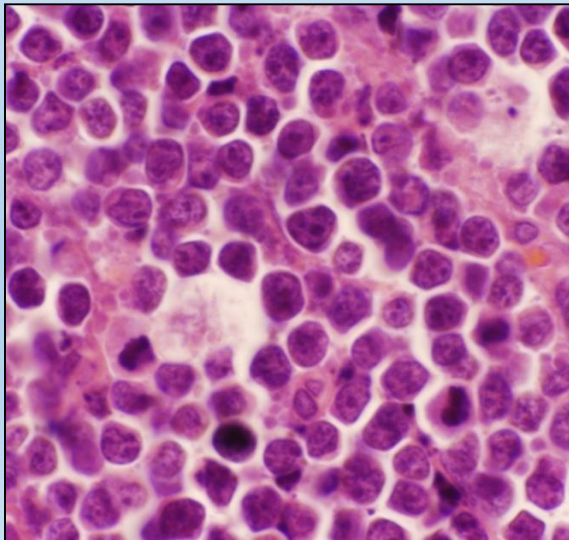
# *MYC* and *BCL2* Rearrangements and Protein Expression: Inform Prognosis and Guide Therapy

- Diffuse large B-cell lymphoma, NOS
- Double-expresser (DE) DLBCL, NOS
  - Expresses *MYC* (>40%) and *BCL2* (>50%) protein
- High grade B-cell lymphoma double hit (HGBL-DH), 4-6% of DLBCL.
  - *MYC/BCL2*, 80% (includes 20% triple hit).
  - *MYC/BCL6*, 20%.

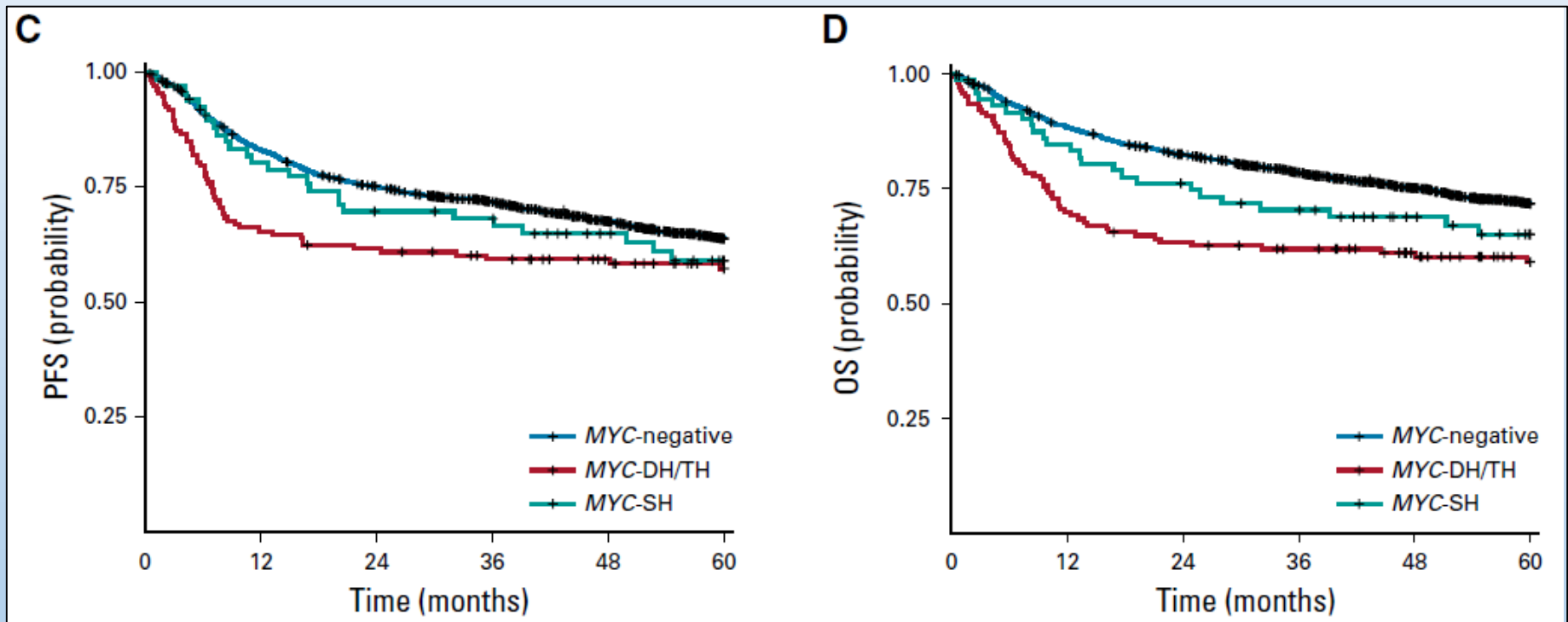


# High-Grade B-cell Lymphoma with *MYC* and *BCL2* and/or *BCL6* Rearrangements (WHO 2017)

- Aggressive presentation, often disseminated (PB, BM, CSF).
- Can resemble BL with increased pleomorphism and/or atypical immunophenotype or genetic features.
- *MYC* complex karyotype is common.

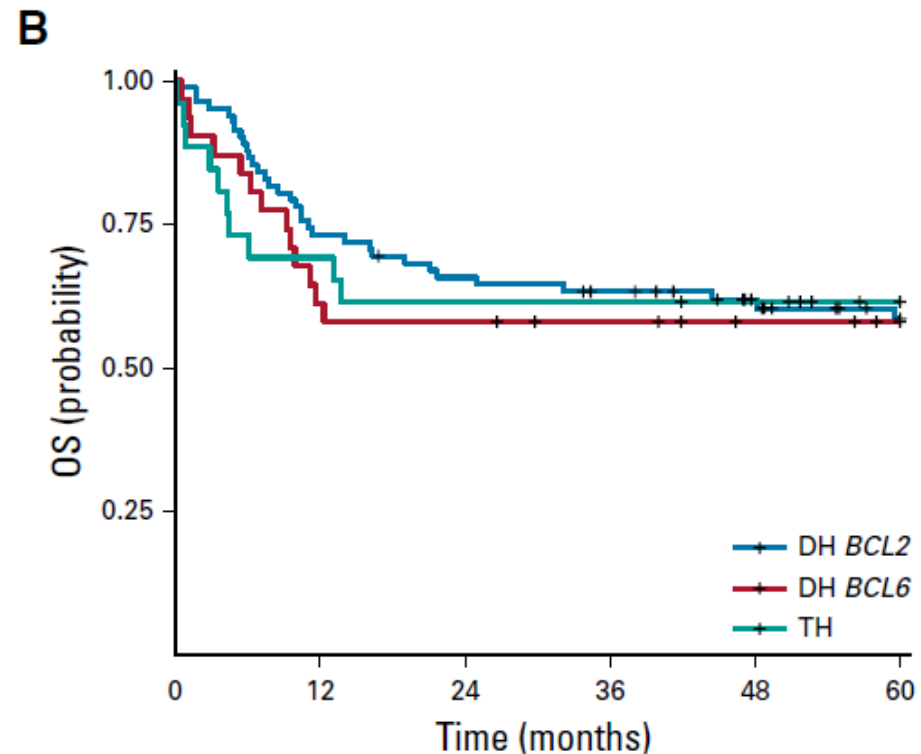
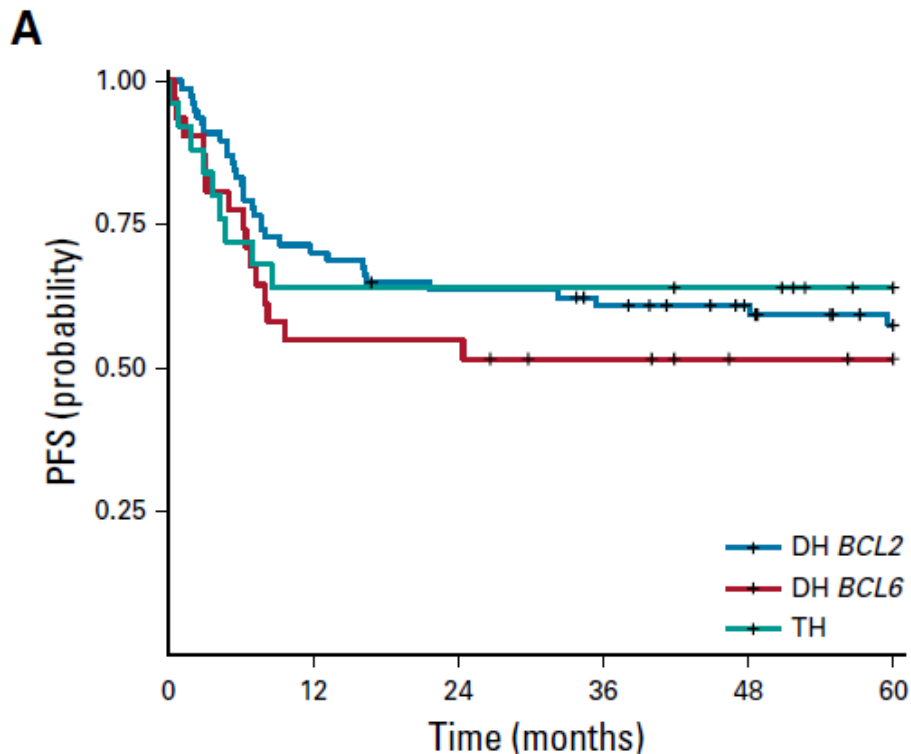


# MYC DH/TH Have Worse Outcomes than MYC-N or MYC-SH

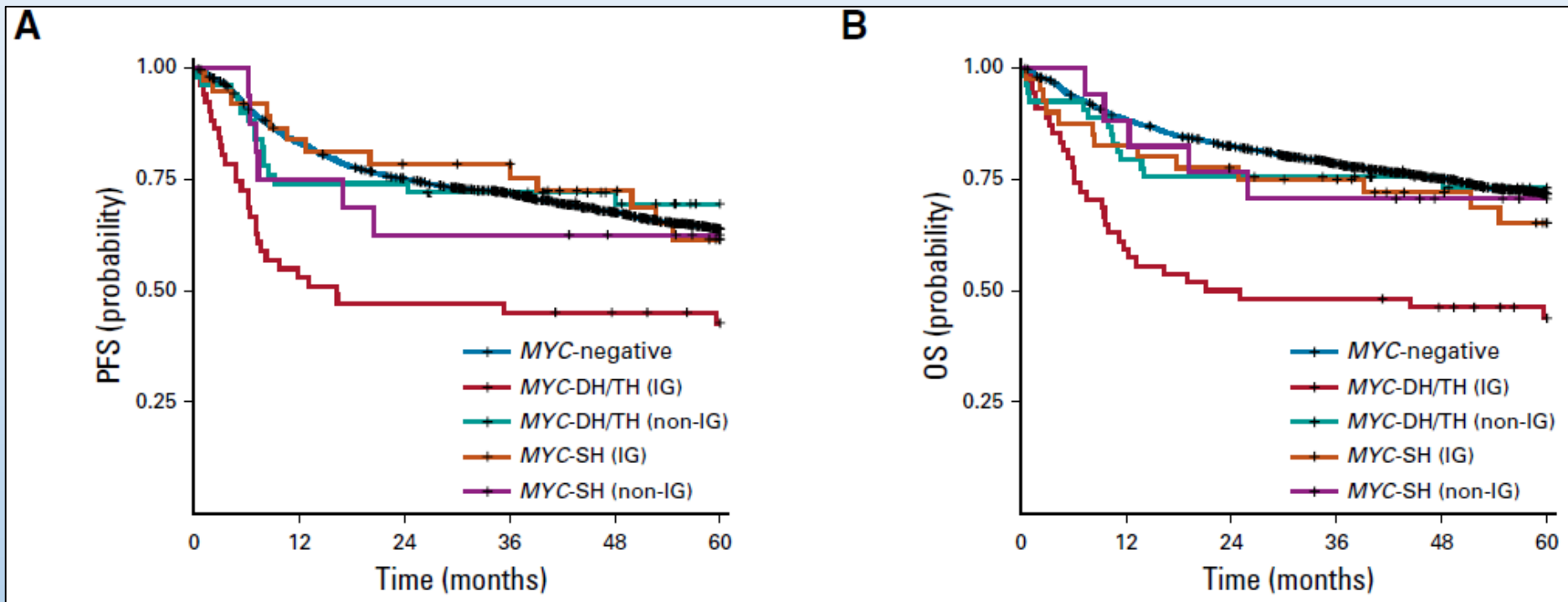




# No Significant Difference Between MYC/BCL2, MYC/BCL6, or TH

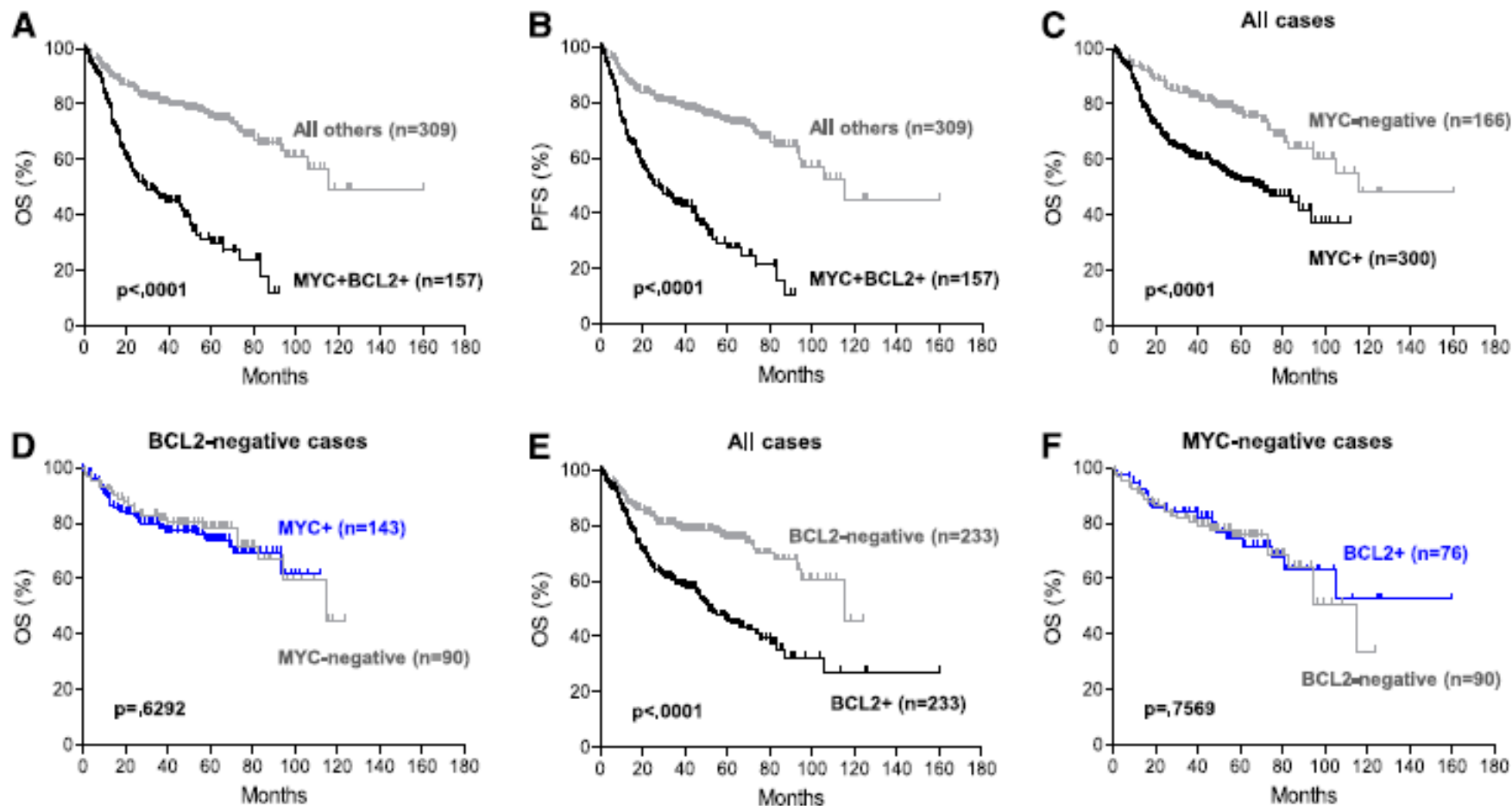


# Partner Matters: DH/TH with non-IG MYC Partners Don't Do Worse



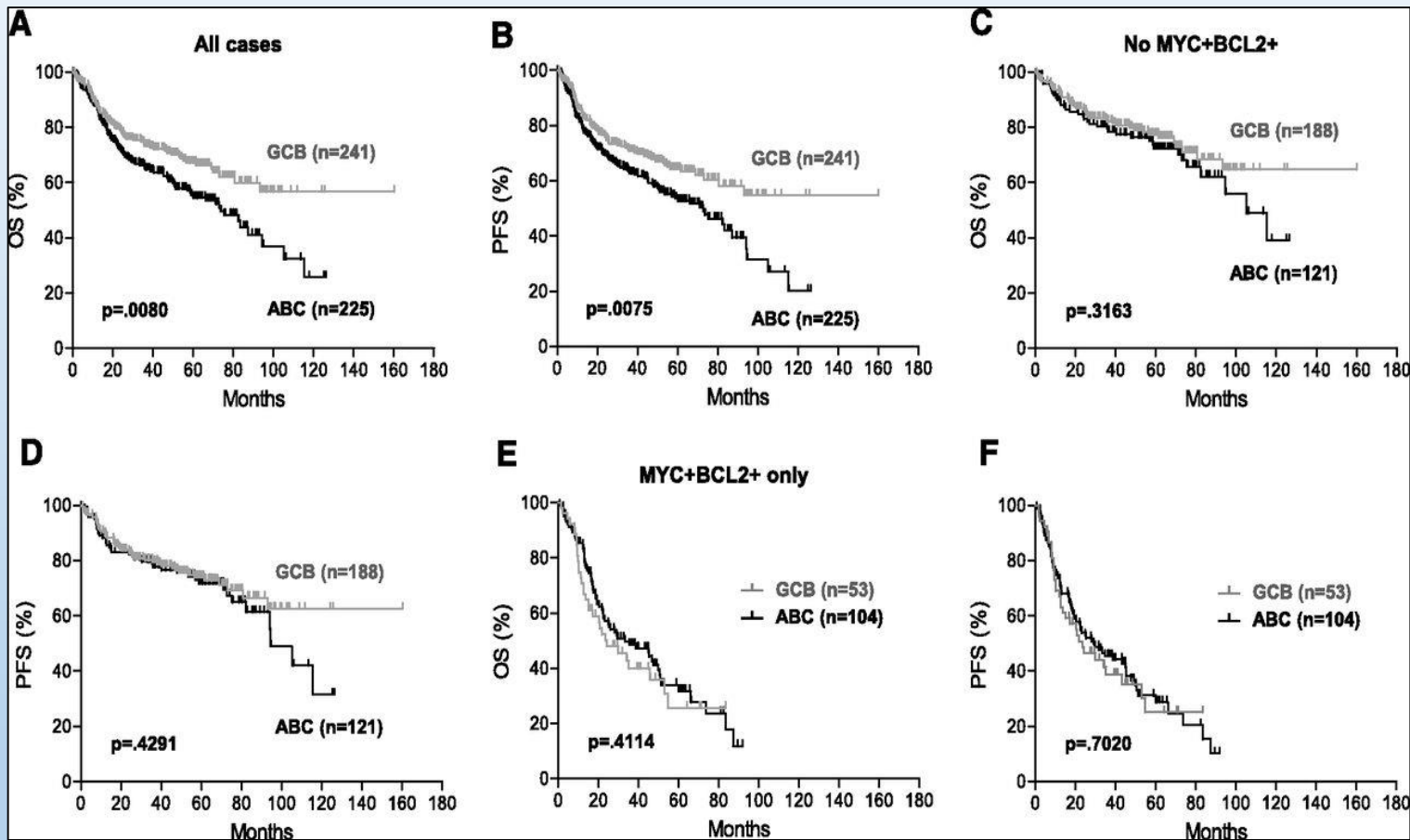
- Suggests that MYC break-apart should be followed by FISH for MYC-IGH, IGK-MYC, MYC-IGL.
- IG promoters/enhancers drive the highest MYC Expression.

MYC/BCL2 Co-expression has adverse effect on survival.  
Neither MYC nor BCL2 expression alone impacts survival.



**Figure 1. Prognostic impact of MYC/BCL2 coexpression in DLBCL.** (A-B) OS (A) and PFS (B) of patients with DLBCL with MYC/BCL2 coexpression (MYC<sup>+</sup>BCL2<sup>+</sup>) in the training set. (C-D) OS of patients with MYC<sup>+</sup> DLBCL in the presence (C) or absence (D) of BCL2 coexpression in the training set. (E-F) OS of patients with BCL2<sup>+</sup> DLBCL in the presence (E) or absence (F) of MYC coexpression in the training set.

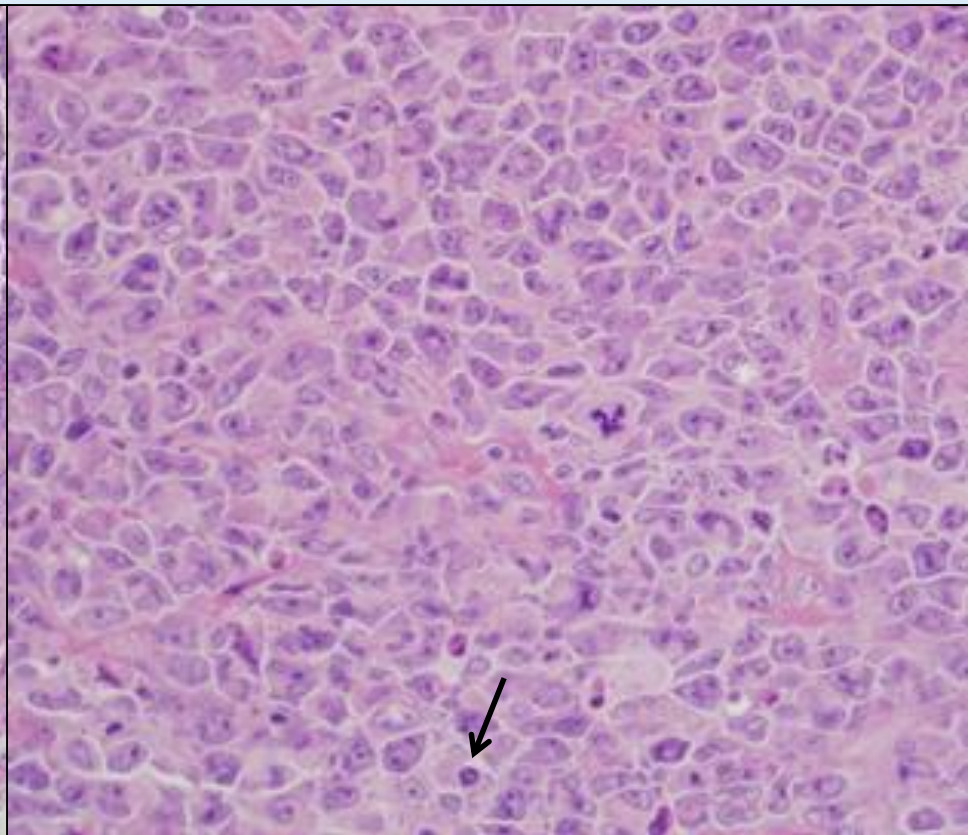
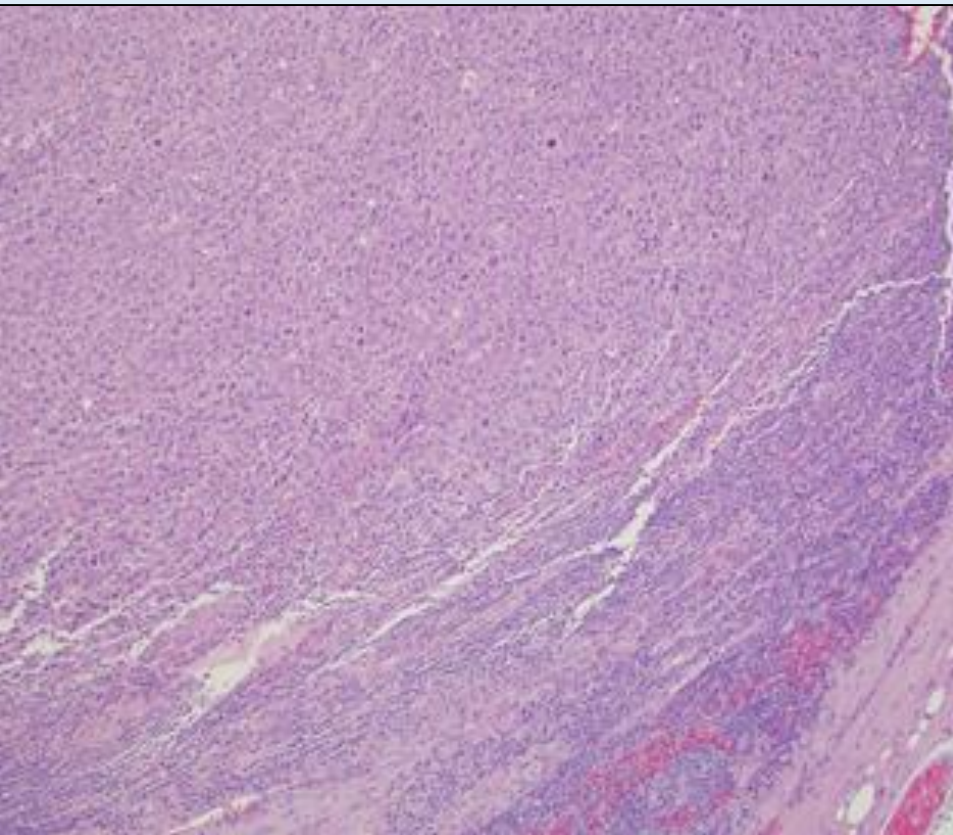
## MYC/BCL2 coexpression contributes to the inferior prognosis of ABC-DLBCL.



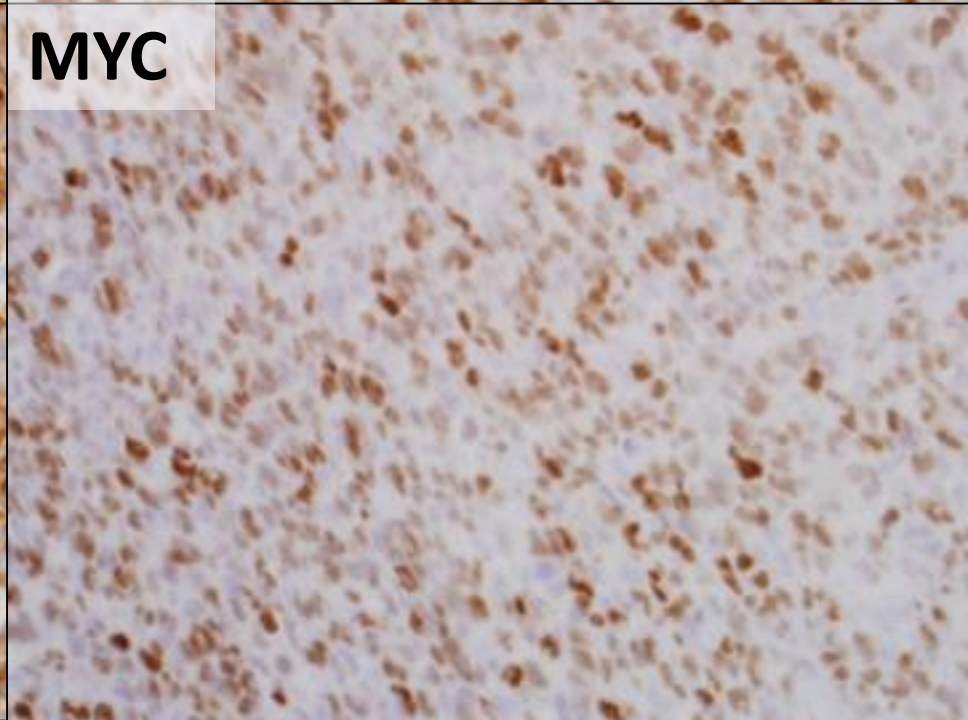
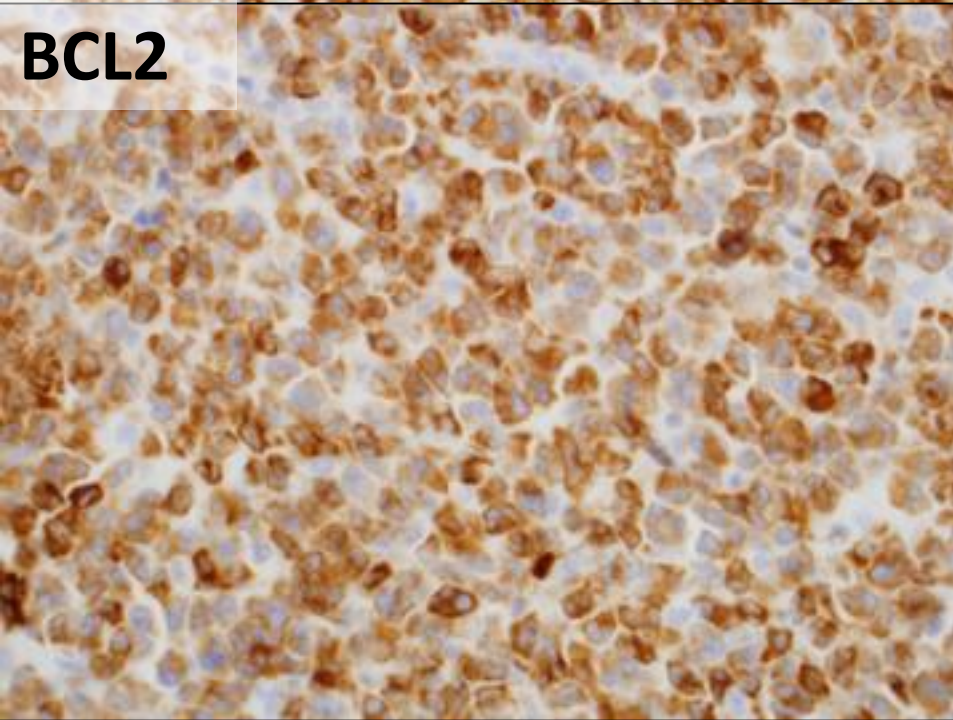
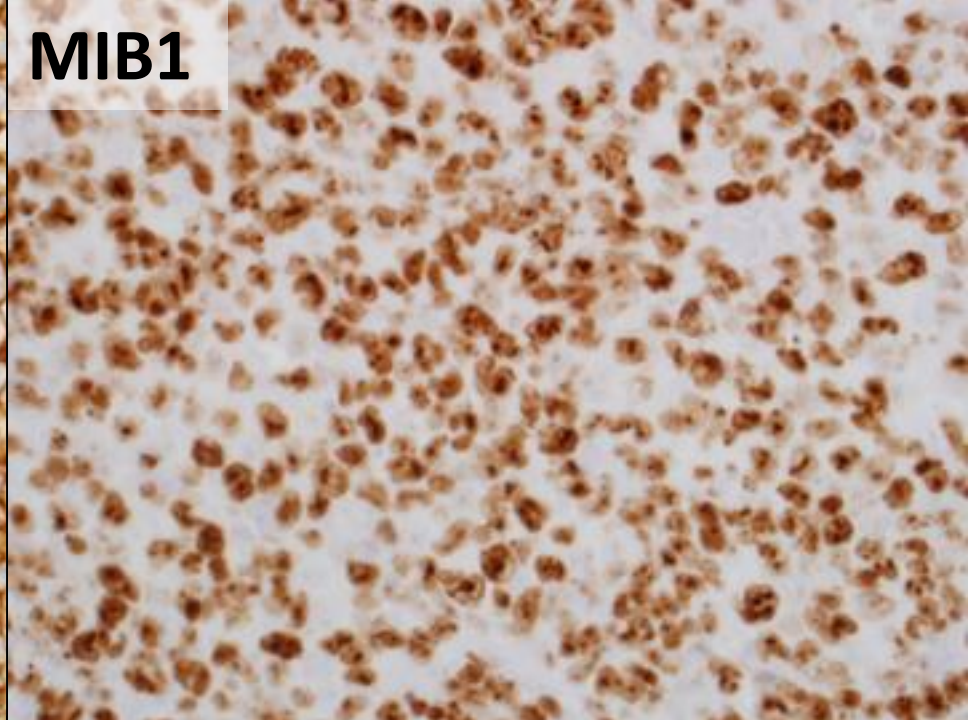
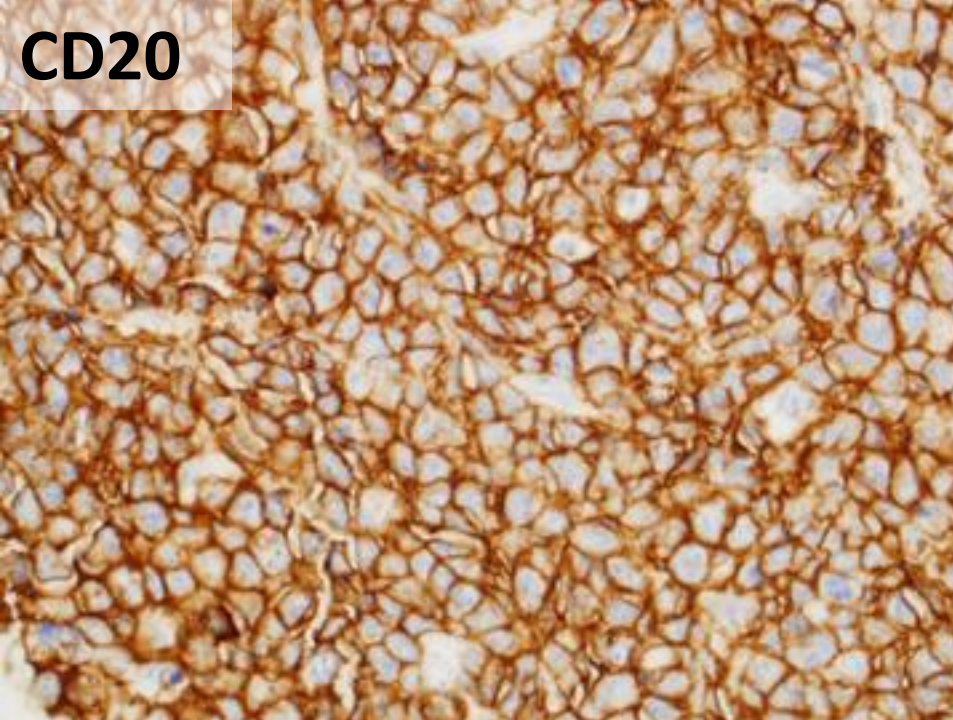
Hu S et al. Blood 2013;121:4021-4031

79-year-old woman with recent breast cancer diagnosis and enlarged lymph nodes.

- Right neck mass.







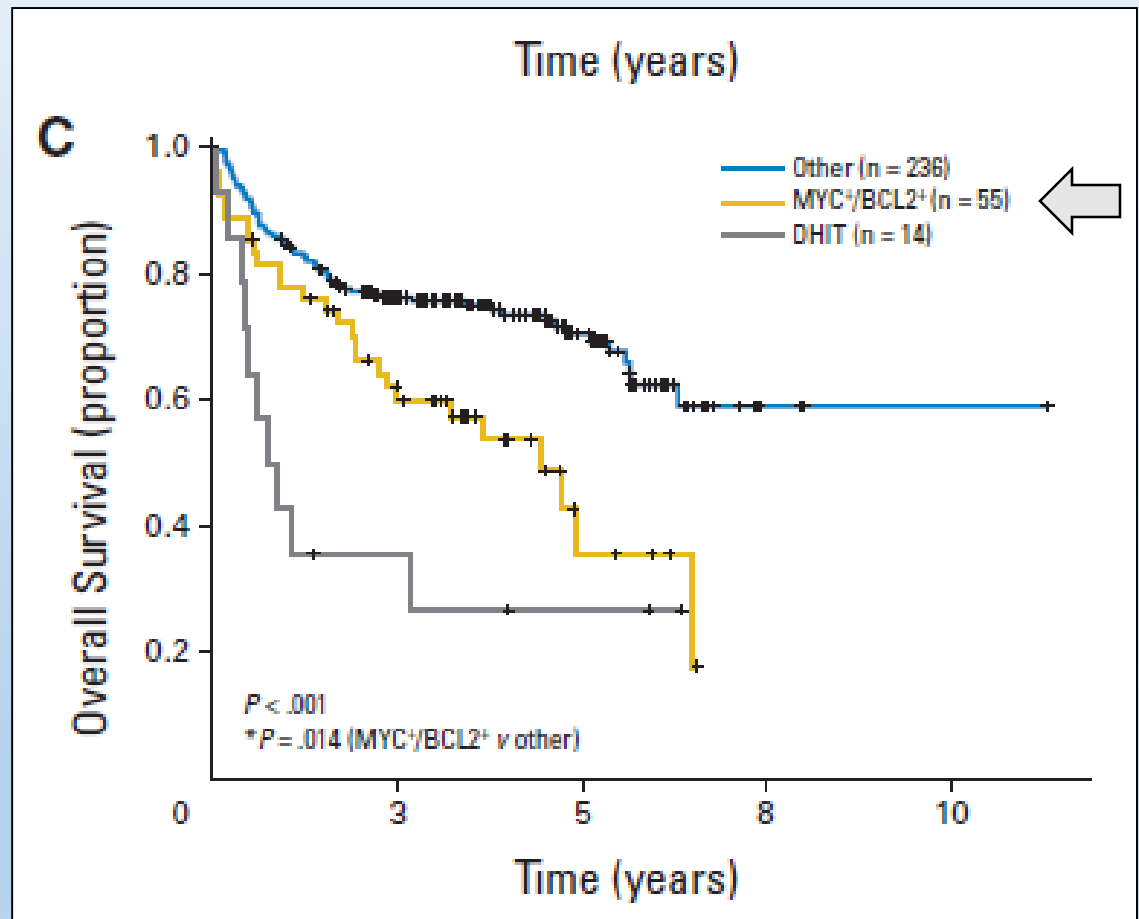
# Results of FISH and Final Classification

- *MYC* breakapart probe assay negative.
  - *MYC* rearrangement excluded.
  - *MYC/BCL2* double hit lymphoma excluded.
- Classification:
  - Diffuse large B-cell lymphoma, activated B-cell subtype, with co-expression of *MYC* and *BCL2*.
- Prognosis: bad, but not as bad as *MYC/BCL2* double hit.



# Prognosis and Treatment

- Consider DA-EPOCH-R rather than R-CHOP.



# DLBCL Prognostic Testing Strategy

De novo DLBCL (excludes relapse, PTLD, transformation?)

DLBCL, NOS

*MYC* FISH, If +,  
*BCL2*, *BCL6*

*MYC* and *BCL2*  
Immunohistochemistry  
CD10, *BCL6*, and MUM1  
for Hans COO

Clinical and/or morphology  
suggests DLBCL subtype:

- TCHRBCL
- EBV+ DLBCL of elderly
- Primary mediastinal (PMBL)
- Primary CNS
- Primary cutaneous leg type

Yes

No

No

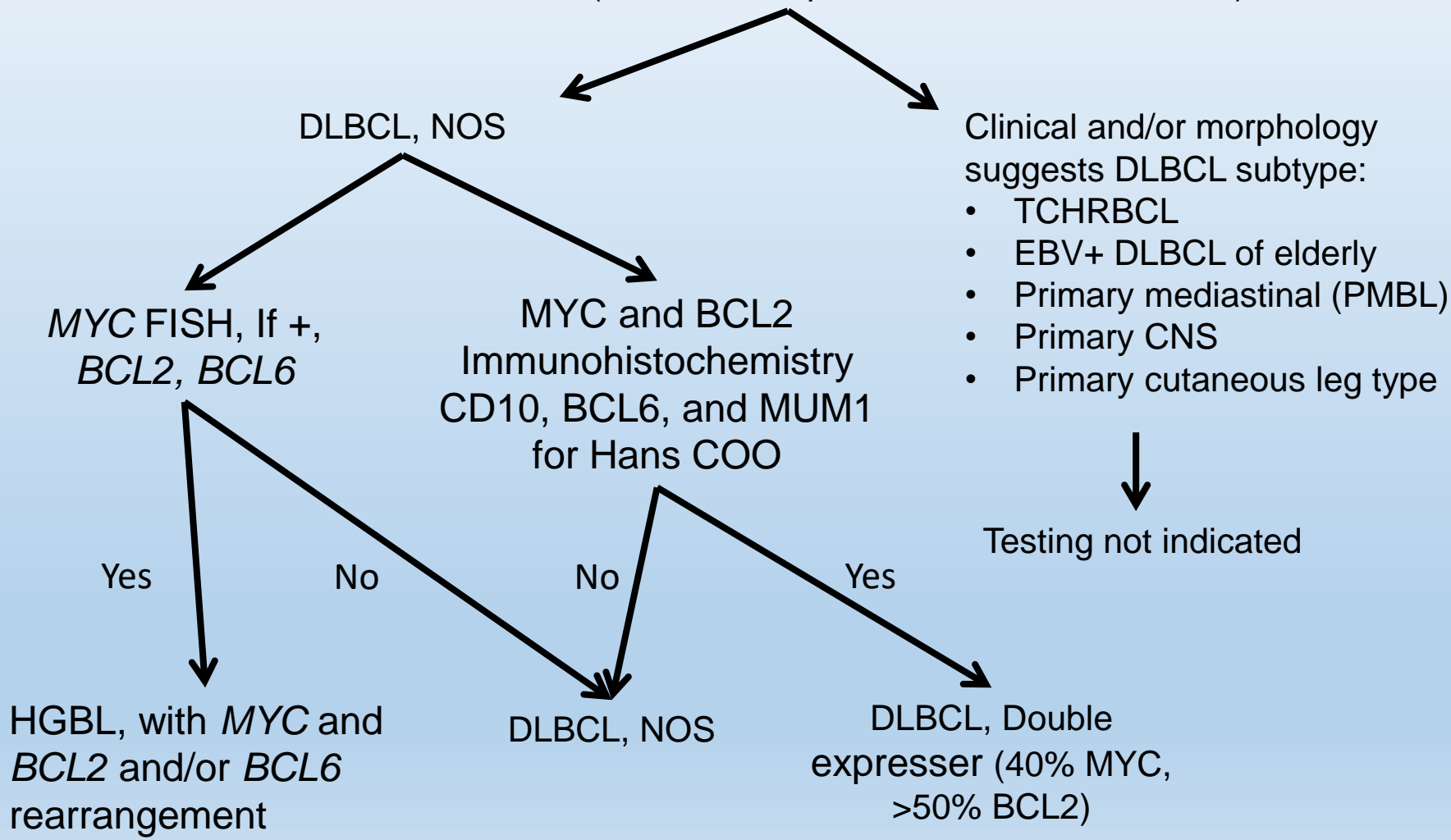
Yes

Testing not indicated

HGBL, with *MYC* and  
*BCL2* and/or *BCL6*  
rearrangement

DLBCL, NOS

DLBCL, Double  
expresser (40% *MYC*,  
>50% *BCL2*)



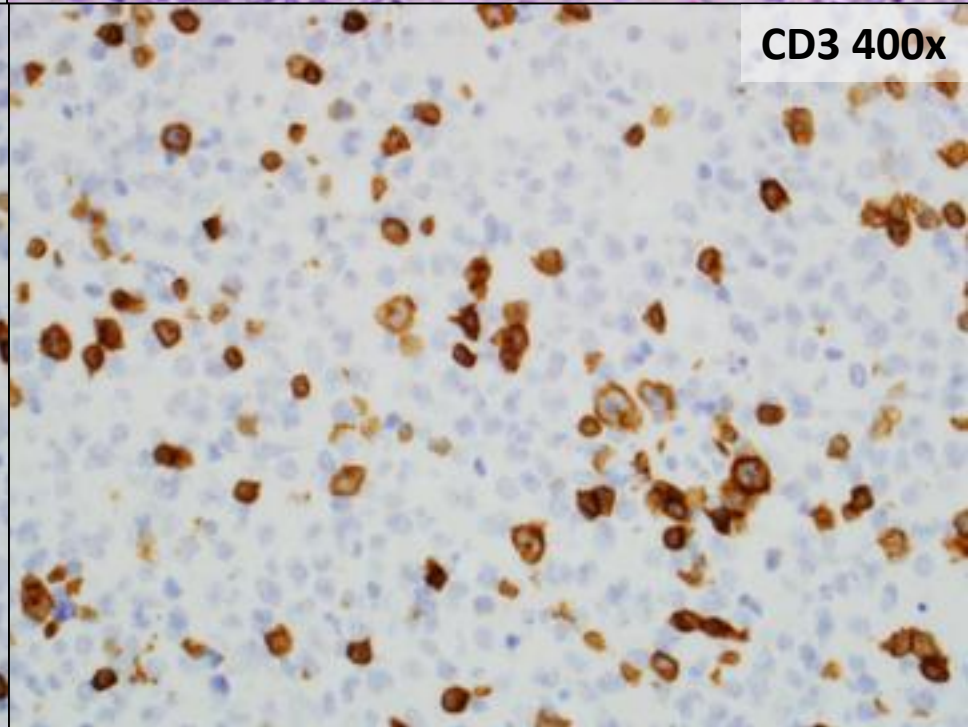
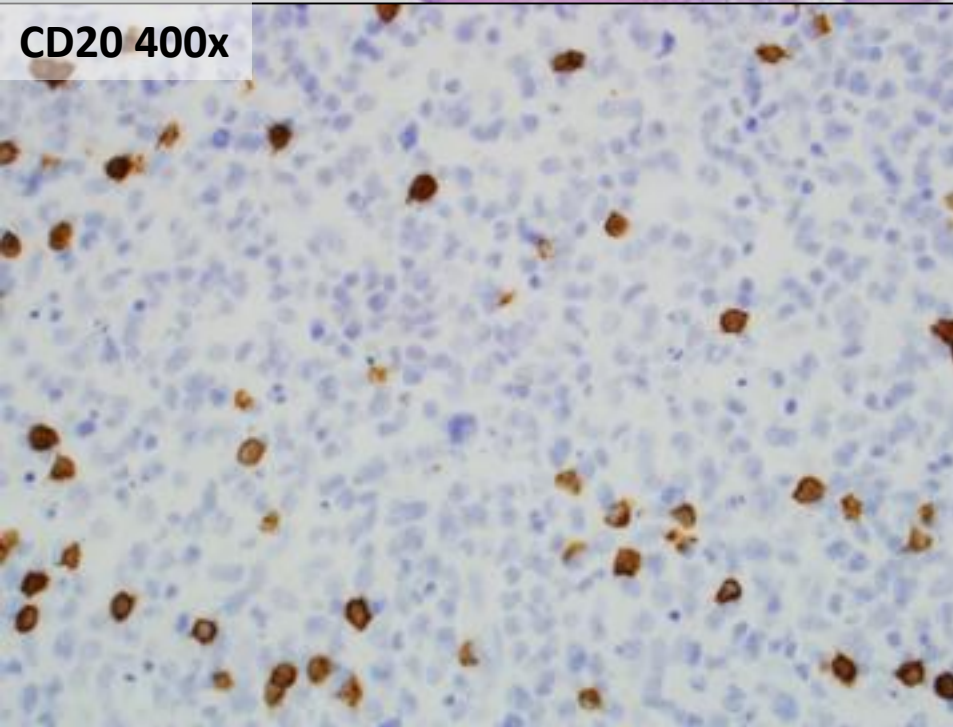
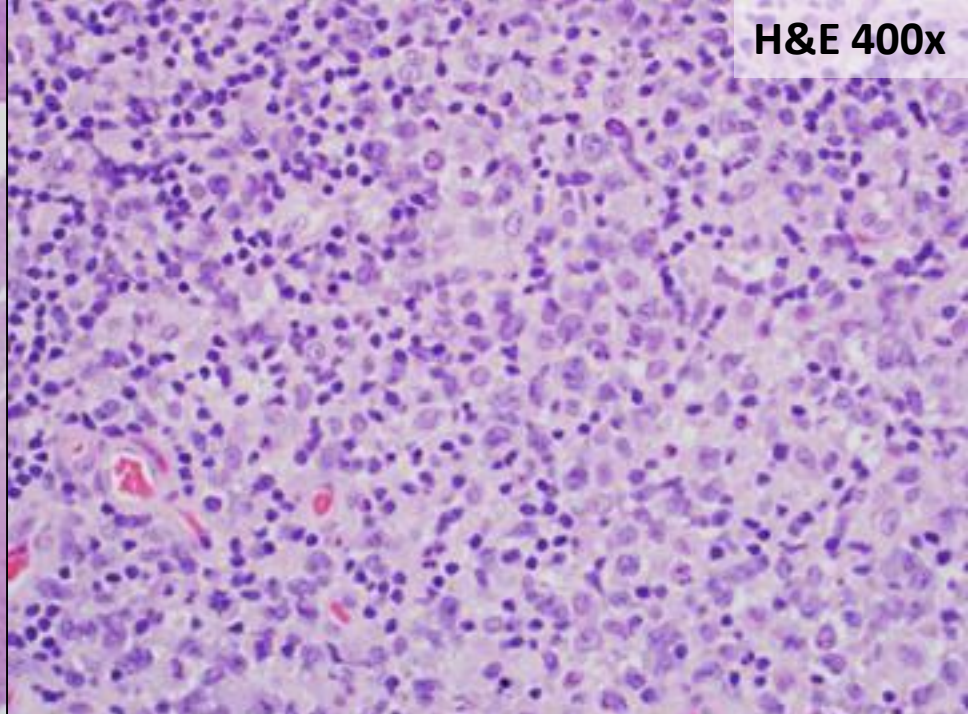
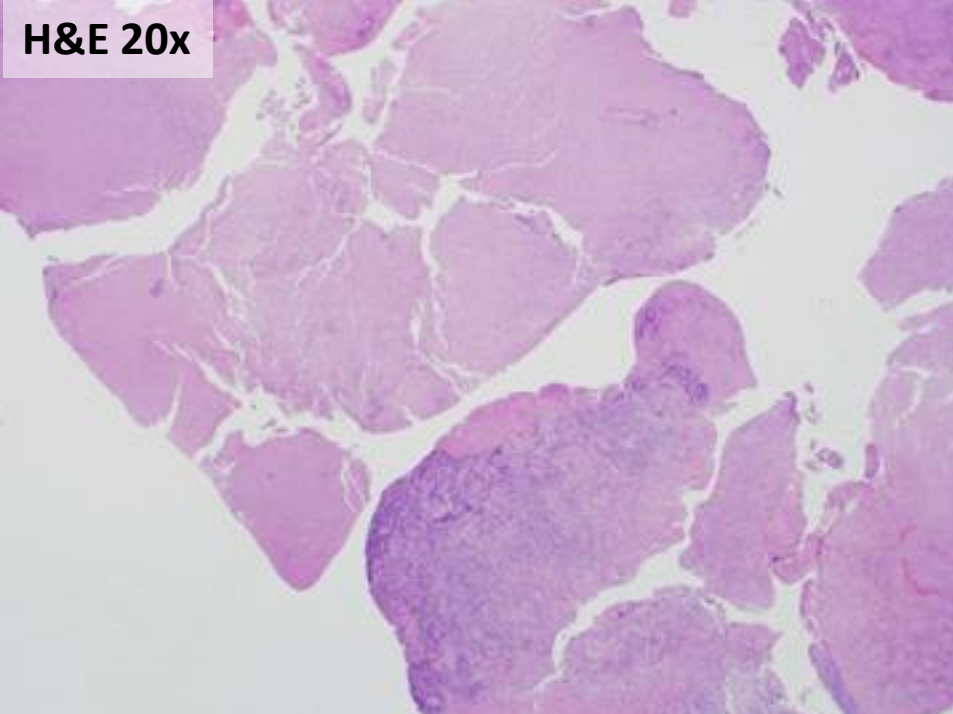
# When should FISH be performed?

- FISH for *MYC* on most DLBCLs.
  - If positive, follow with *BCL2* and *BCL6* FISH.
- Clinical context should be considered.
  - Will it change clinical approach/therapy?

# DLBCL Conclusions

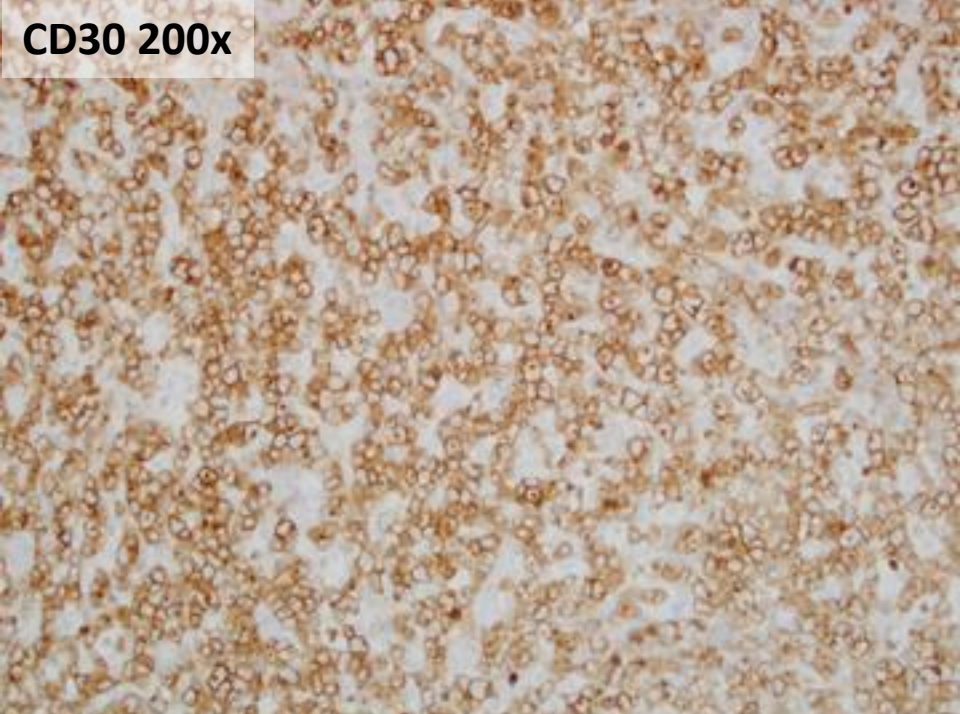
- Diagnosis of DLBCL requires only morphology and immunophenotype.
- Diagnosing or excluding the WHO 2017 category HGBL, with *MYC+BCL2* +/- *BCL6* rearrangement requires FISH.
  - A genetically-defined lymphoma.
- Testing should be performed when results will affect patient care.

Case: 9-year-old boy with a  
mediastinal mass

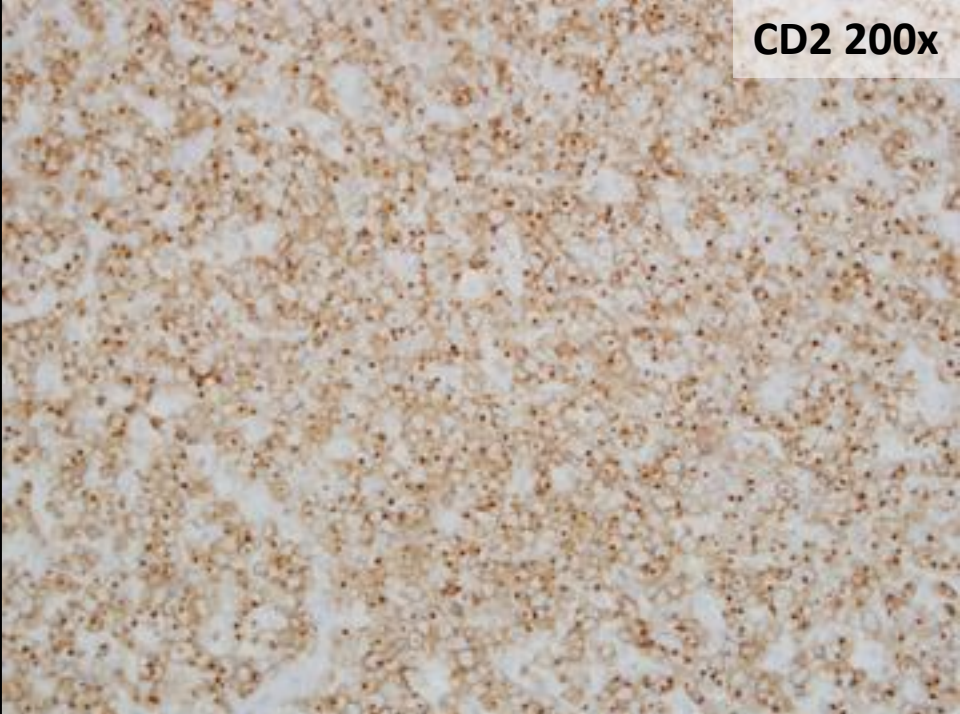




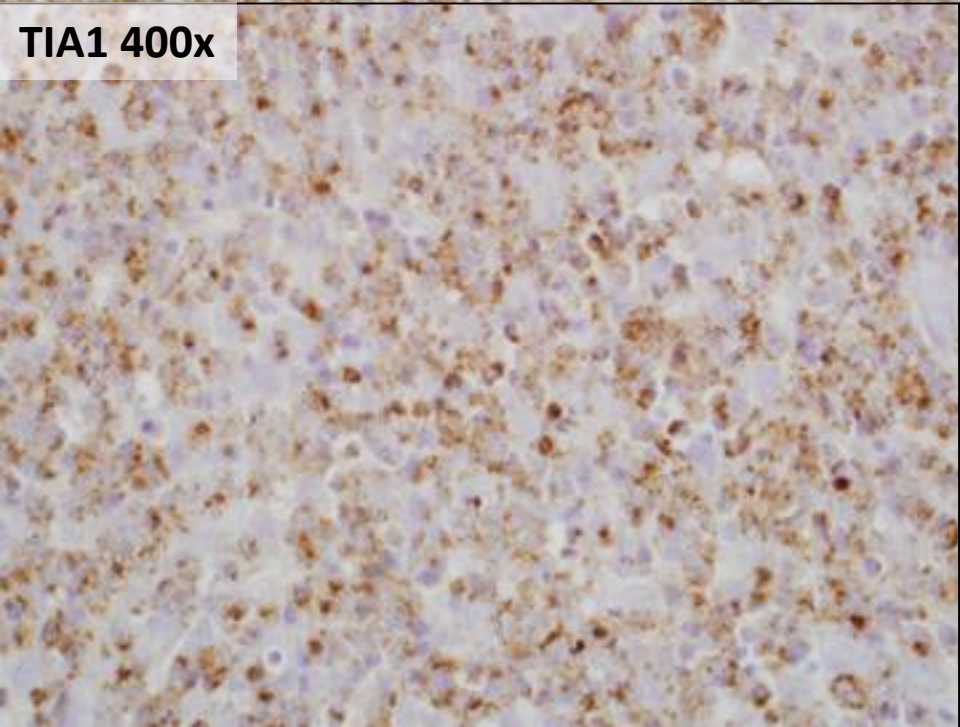
**CD30 200x**



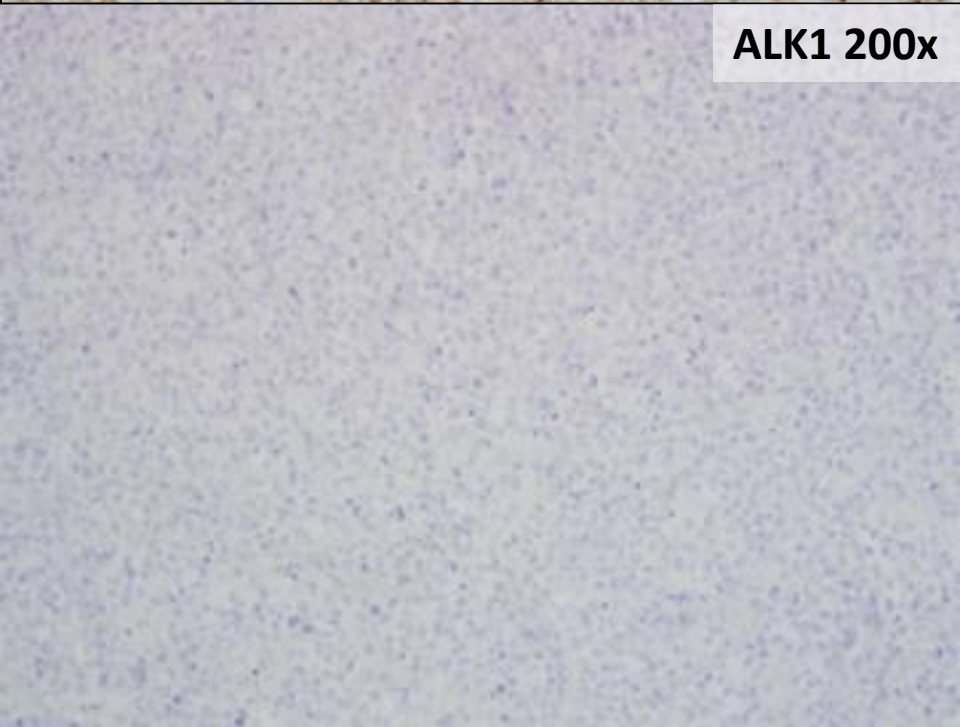
**CD2 200x**



**TIA1 400x**



**ALK1 200x**



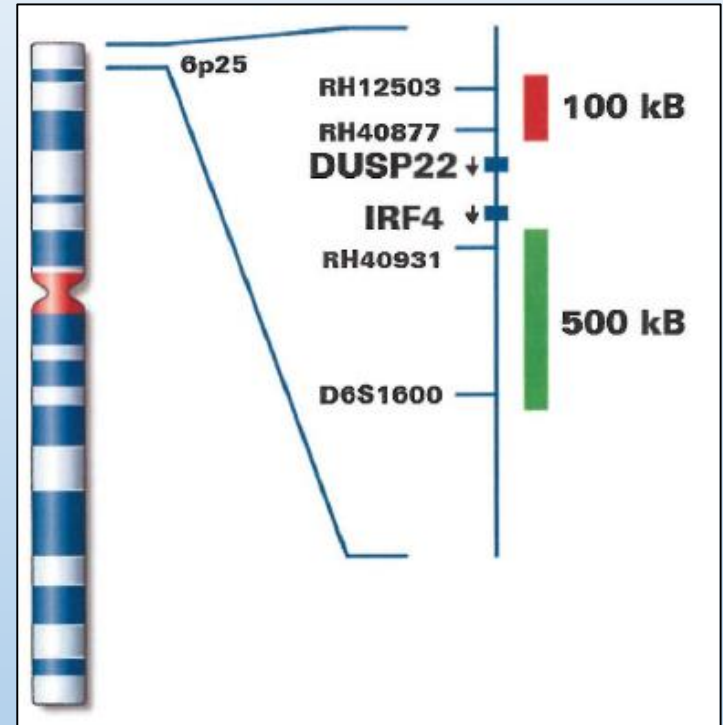
# Immunophenotype and Diagnosis

- Positive for CD2, CD30 (strong and diffuse), TIA1.
- Positive: CD45, CD4 (weak), CD7 (subset).
- Negative: CD20, PAX5, CD15, CD8, CD5.
  
- Diagnosis: Anaplastic Large Cell Lymphoma (ALCL), ALK1 negative.
- Other considerations:
  - Hodgkin: morphology; CD15-, PAX5-, CD3+
  - PTCL, NOS: CD30 too strong and diffuse.

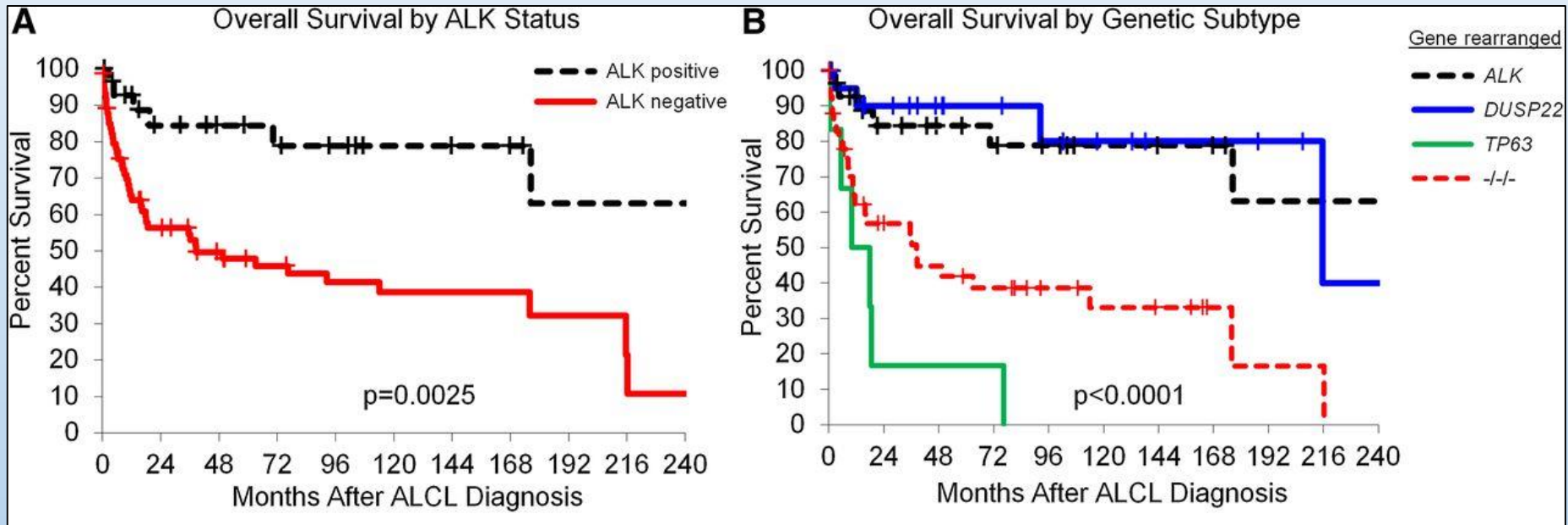


# Additional Studies (I thought we were done?)

- FISH NEGATIVE for rearrangement of *DUSP22/IRF4*.
- Immunohistochemical stain for p63 is NEGATIVE in tumor cells.



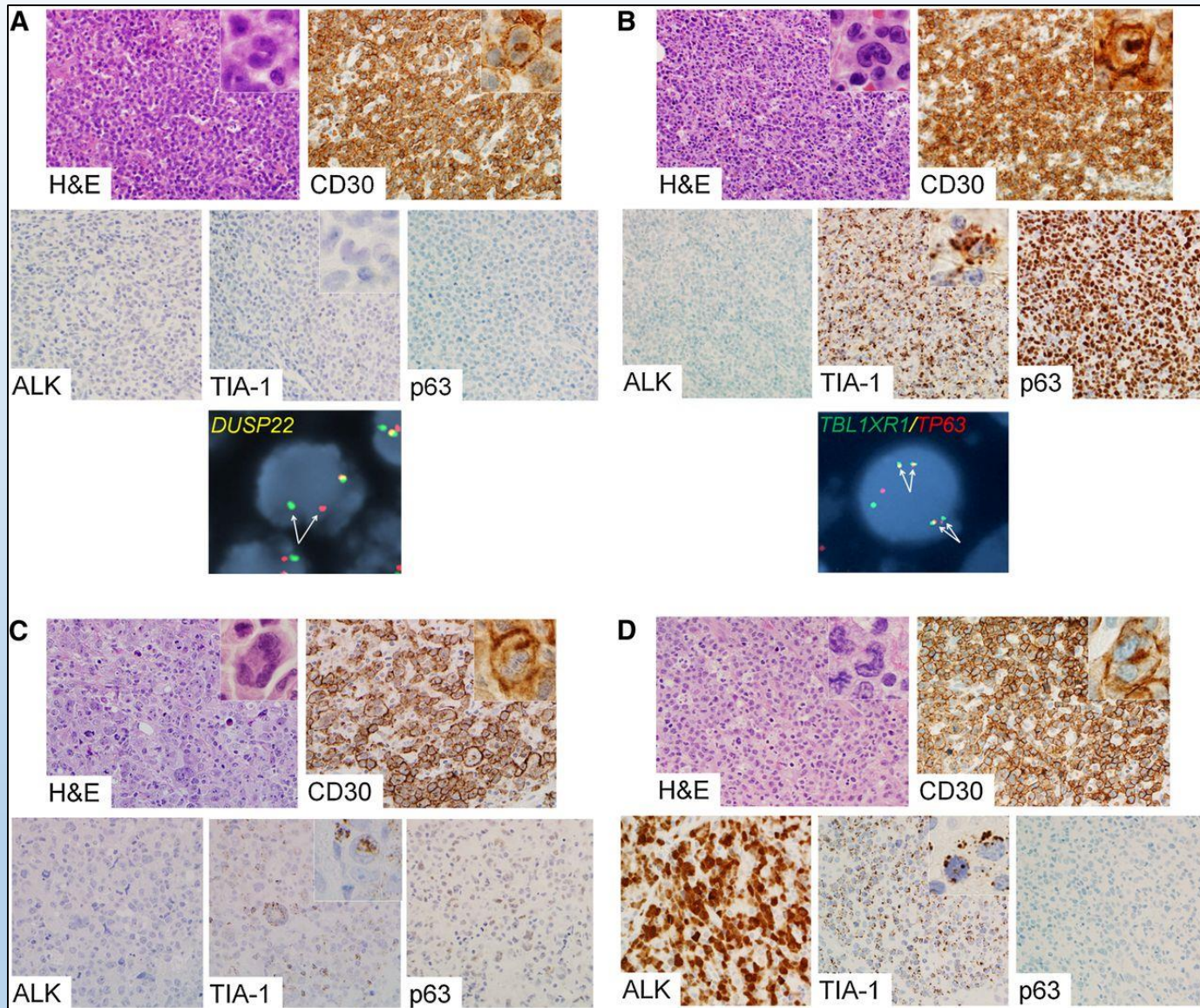
# IRF4/DUSP22 Rearranged ALK- ALCL Shows Outcomes Similar to ALK+ ALCL



Edgardo R. Parrilla Castellar et al. Blood 2014;124:1473-1480



# Genetic subtypes of ALCL



Edgardo R. Parrilla Castellar et al. Blood 2014;124:1473-1480

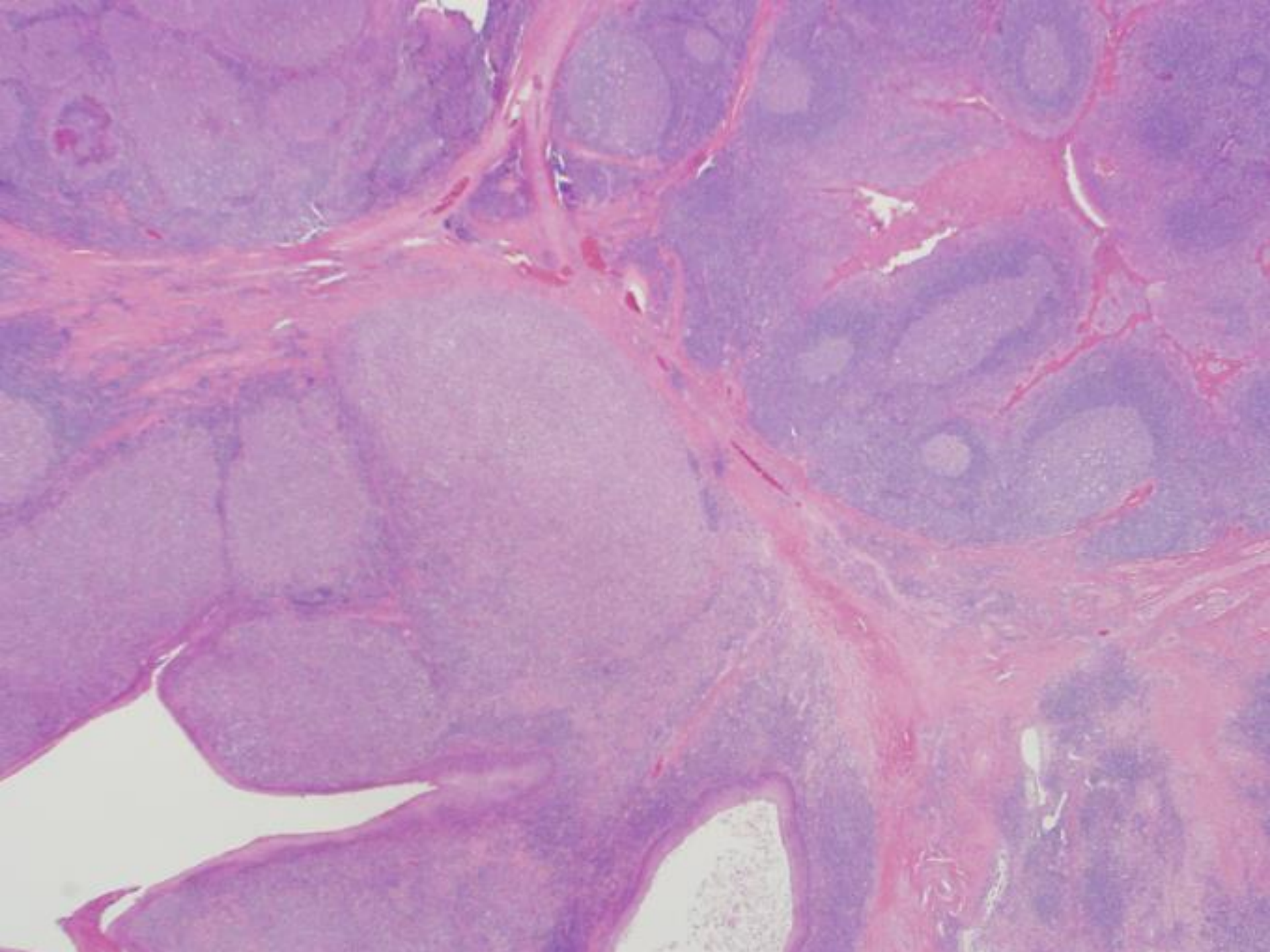
# Approach to ALCL

- ALK+ ALCL: Beware of morphologic variants, which can show varying amounts of CD30 expression and large, atypical cells.
  - Common pattern (60%): sheets of large, atypical cells.
  - Lymphohistiocytic pattern (10%): large, atypical cells can hide among histiocytes but cluster around vessels.
  - Small cell (5-10%): most cells are small to medium-sized, large cells cluster around vessels.
  - Hodgkin-like pattern (3%): Resembles NS-cHL.
- ALK negative ALCL
  - Diffuse or sinusoidal growth pattern.
  - Resembles common pattern of ALK+ ALCL with strong, diffuse CD30 (no variants recognized).
- p63 expressed in all cases with TP63 rearrangement, but also in some cases without rearrangement.
  - Expression did not have prognostic significance.
  - Negative p63 has good negative predictive value.

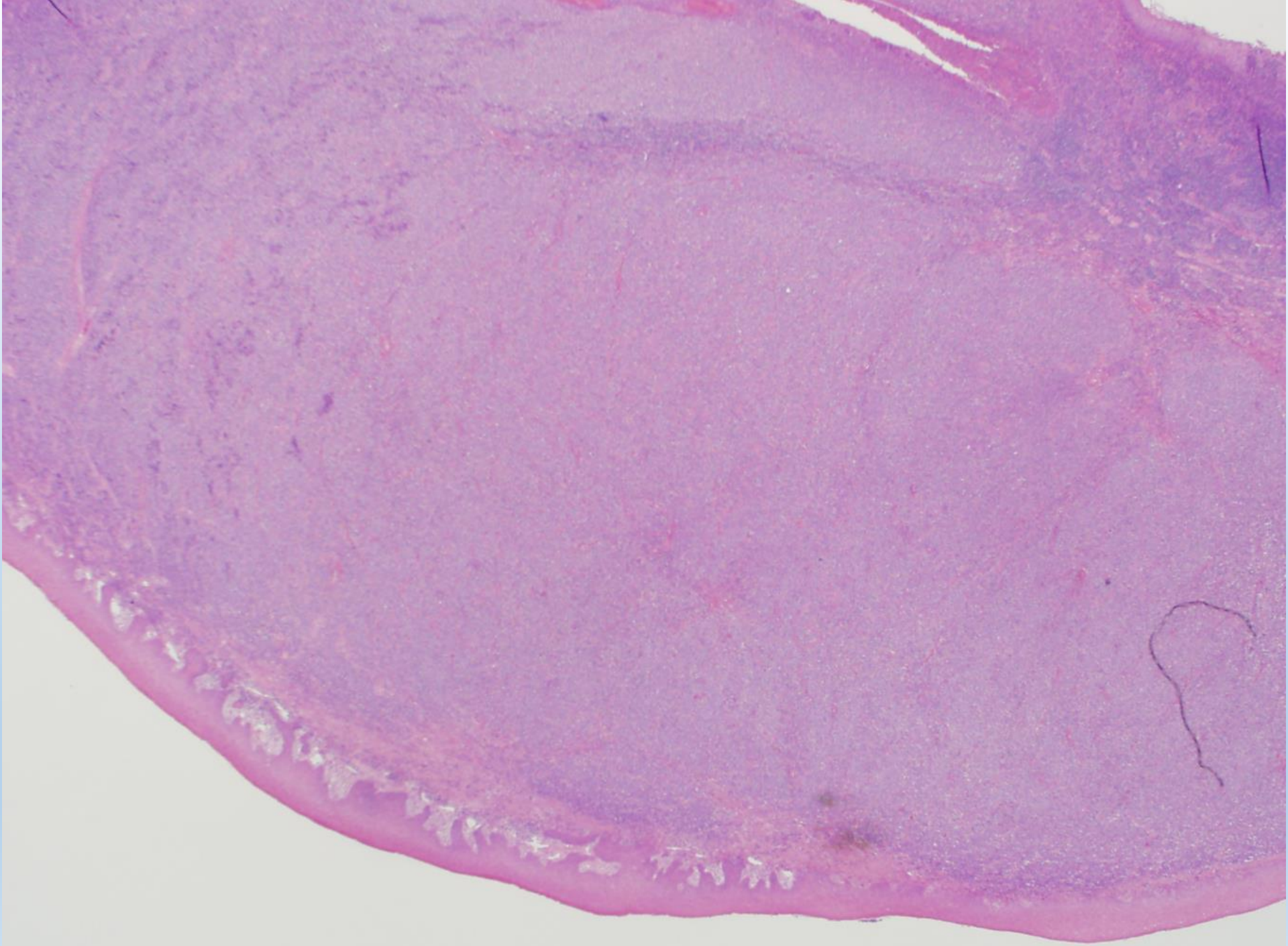


# 17-year-old girl with enlarged tonsils

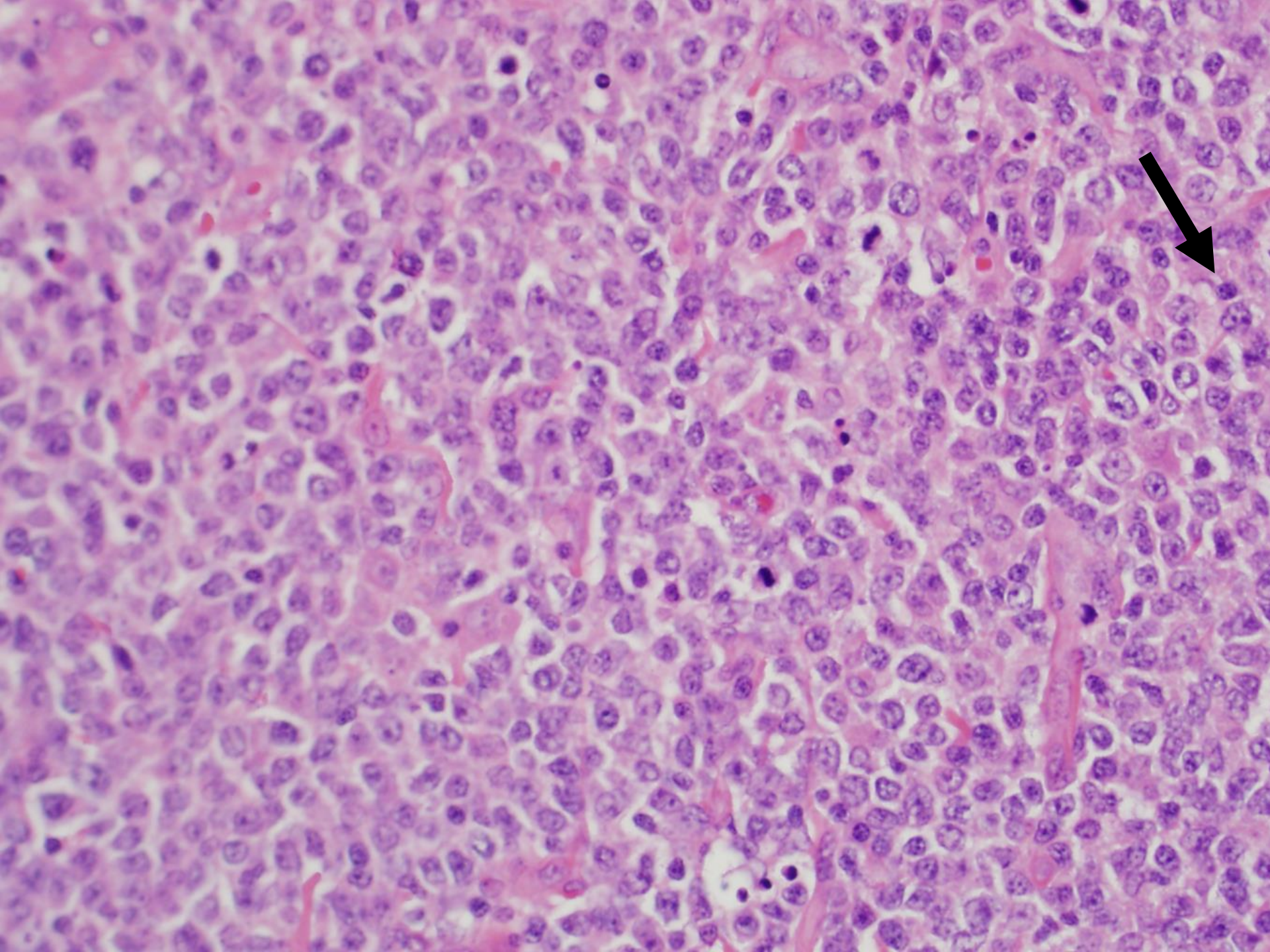
- Original pathology report:
- R tonsil: malignant lymphoma, favor high grade.
- L tonsil: follicular hyperplasia.
- “The overall features favor a high-grade lymphoma.”
- Implications: High-grade implies Burkitt lymphoma in this age group, could include double-hit lymphomas in older adults.
- Intensive chemotherapy regimen required.



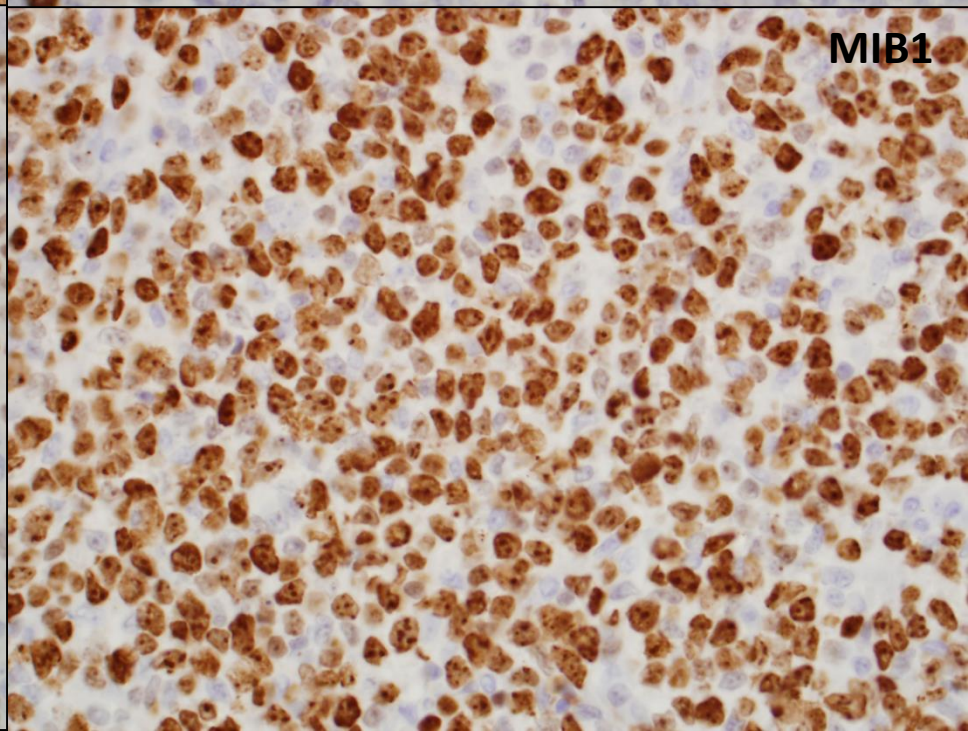
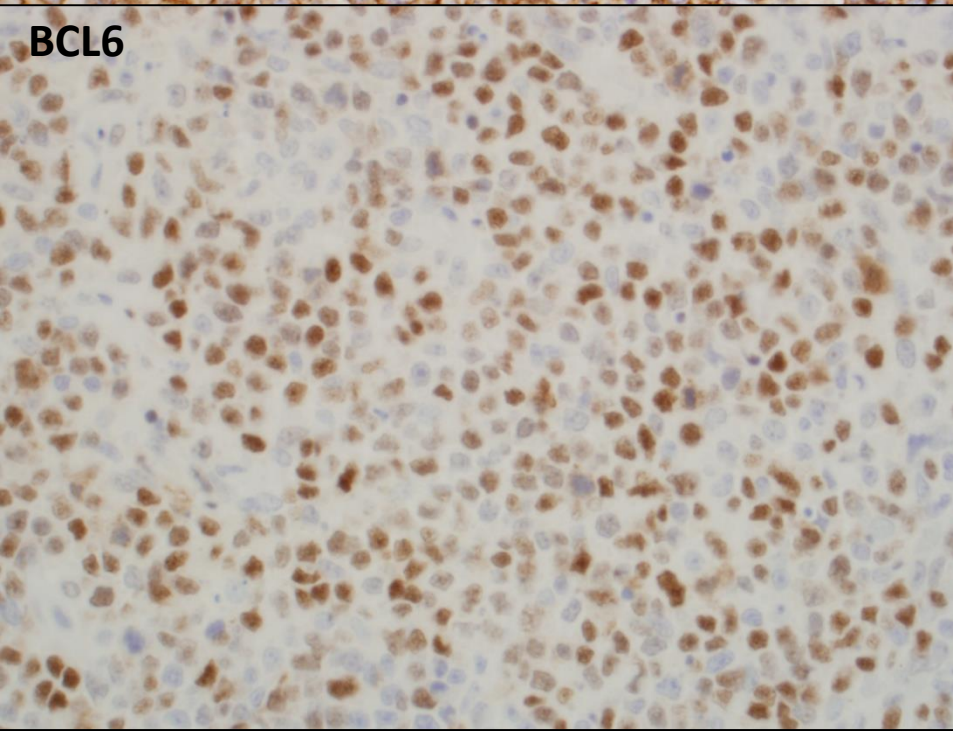
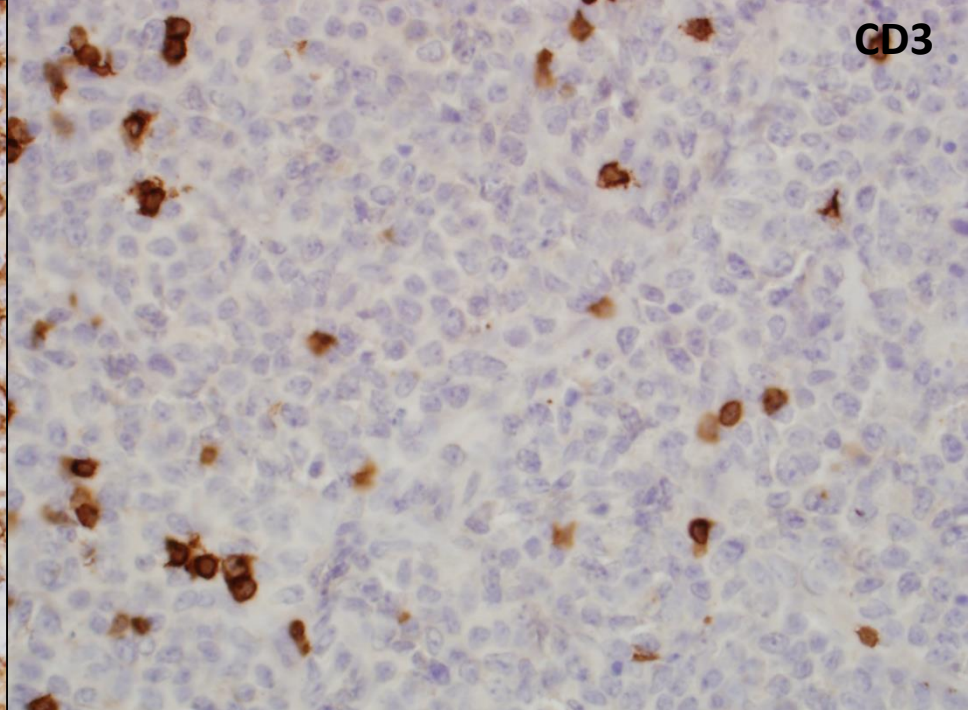
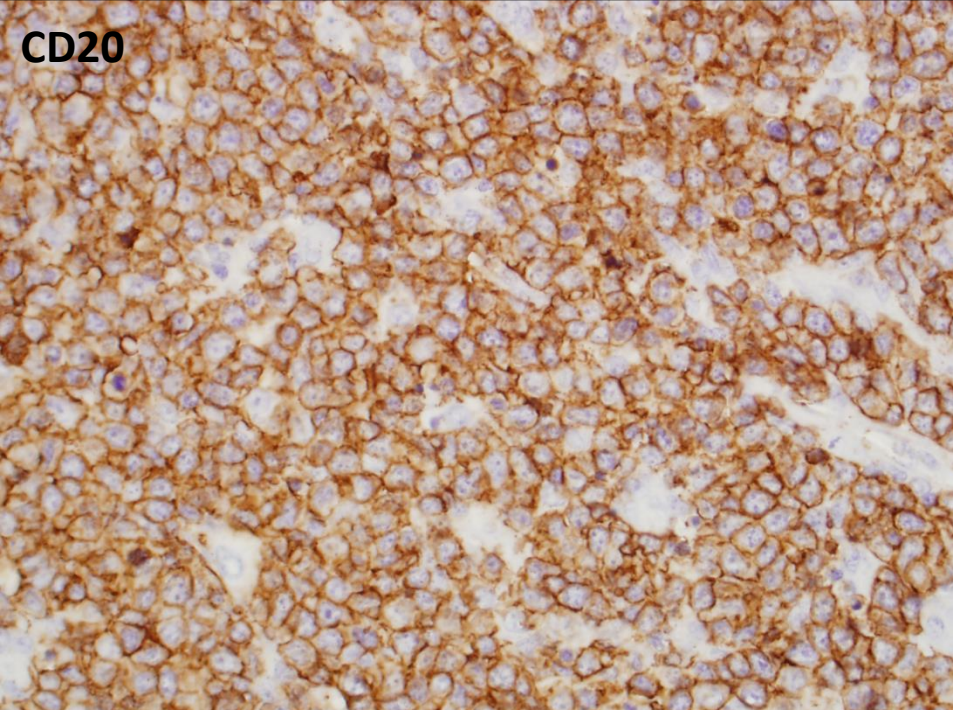






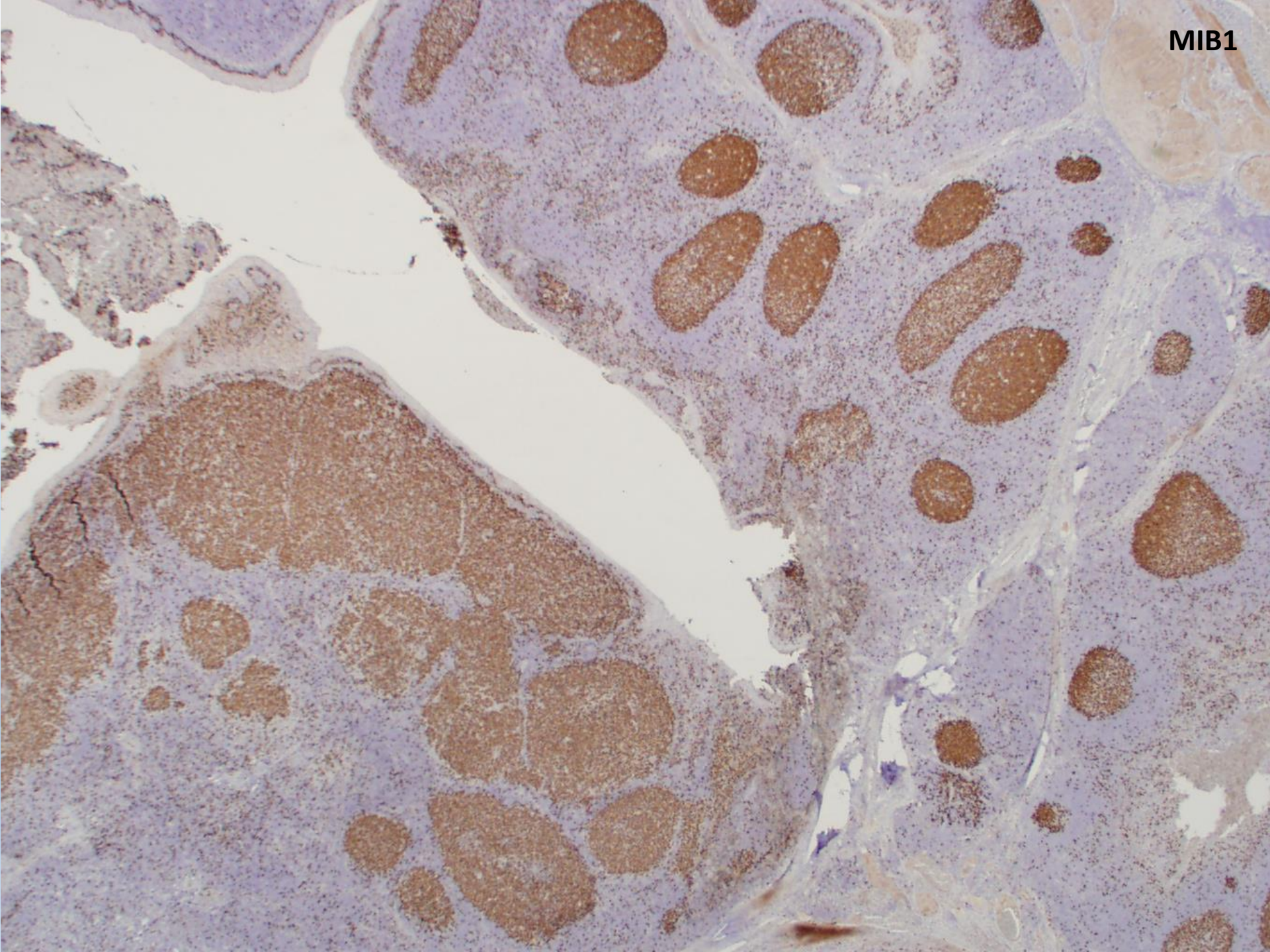






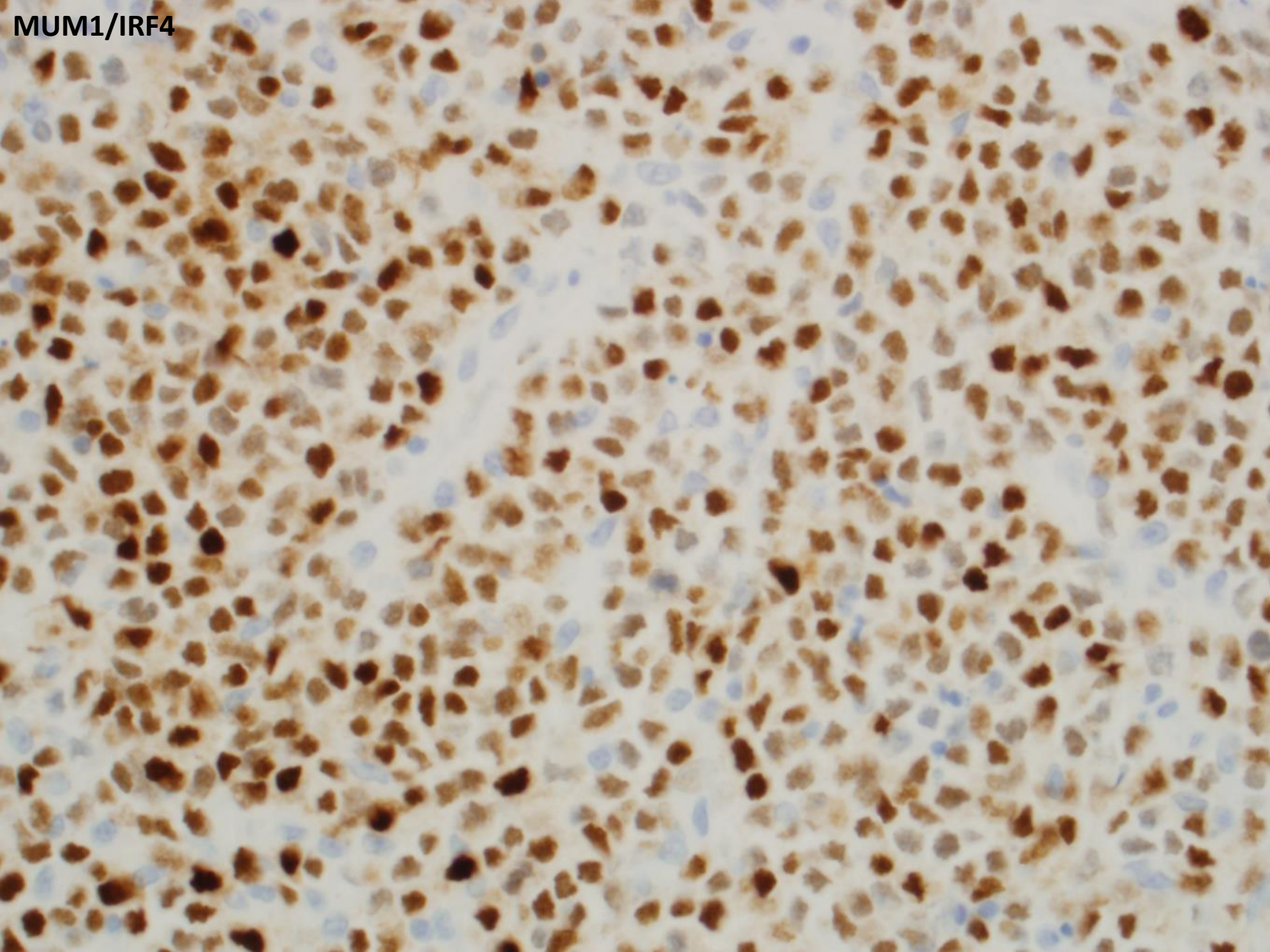


MIB1





MUM1/IRF4



# Differential Diagnosis

- Burkitt lymphoma: Excluded by morphology
  - Lacks tingible body macrophages, too pleomorphic.
- Follicular lymphoma, grade 3.
- Diffuse large B-cell lymphoma.
- Large B-cell lymphoma with *IRF4* rearrangement.
  - New entity in WHO 2017.
- FISH results:
  - FISH for *MYC* and *BCL6* and *BCL2* negative.
  - FISH for *IRF4/DUSP22* positive.

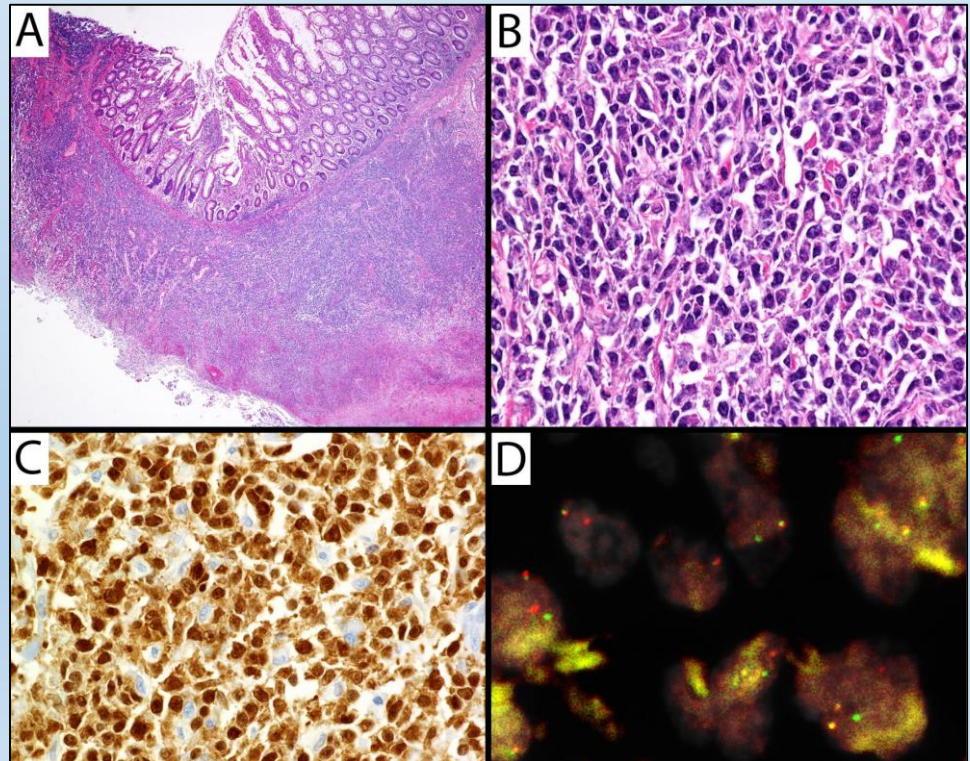


# Large B-cell lymphoma with *IRF4* rearrangement.

- Localized in head and neck.
- Median age 12 (range 4-79).
- Morphologically fit into DLBCL, follicular lymphoma grade 3, pediatric type follicular lymphoma.
- Positive for BCL6 and IRF4/MUM1.
- Good outcome after chemotherapy.
  - Less intensive therapy than Burkitt lymphoma.
- In the appropriate clinical context, FISH for *IRF4/DUSP22* should be performed.

# Overall uncommon (<1%), but more common in younger patients

- We studied 32 patients from Children's Oncology Group protocols.
- FISH for *IRF4/DUSP22* positive in 2/32 cases (6%).
  - One in tonsils.
  - One in ileum.



# Genetically Defined Lymphomas

- ALK+ ALCL
- Mantle cell lymphoma
- HGBL, with *MYC+BCL2* +/- *BCL6* rearrangement
- Large B-cell lymphoma with *IRF4* rearrangement
- Lymphomas with highly characteristic genetic changes that do not define them:
  - Burkitt lymphoma: (*IG-MYC*)
  - Follicular lymphoma: t(14;18)
  - Lymphoplasmacytic lymphoma: *MYD88* mutation.
  - Hairy cell leukemia (BRAF V600E).

# Summary

- Some entities are now defined in the WHO Classification by genetic (mutations, translocations) features.
  - More in AML, ALL
  - Emerging in lymphoma
- When clinically indicated, consider additional testing for precise classification.