

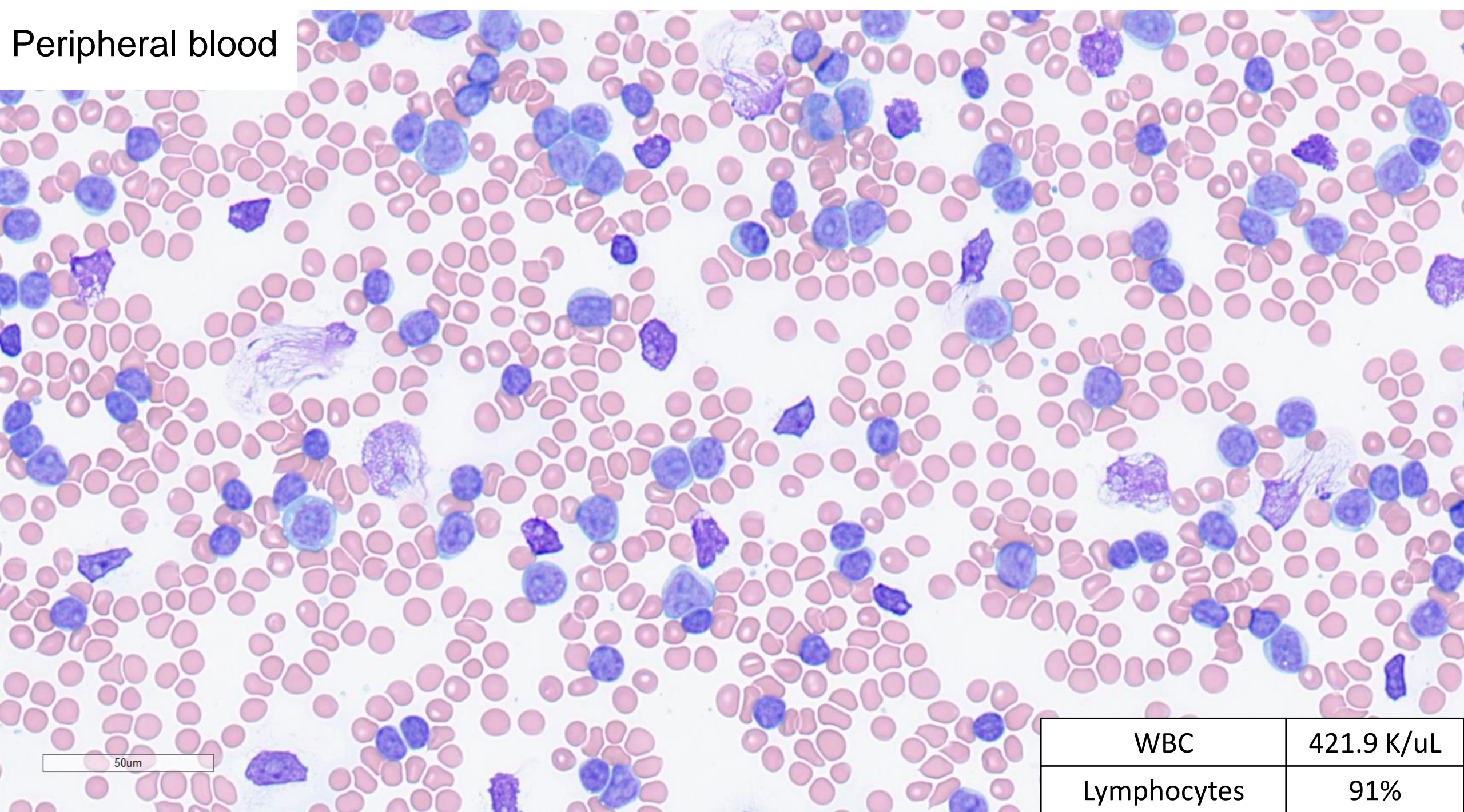
MIHCC Case #2

6/25/2025

Clinical Information

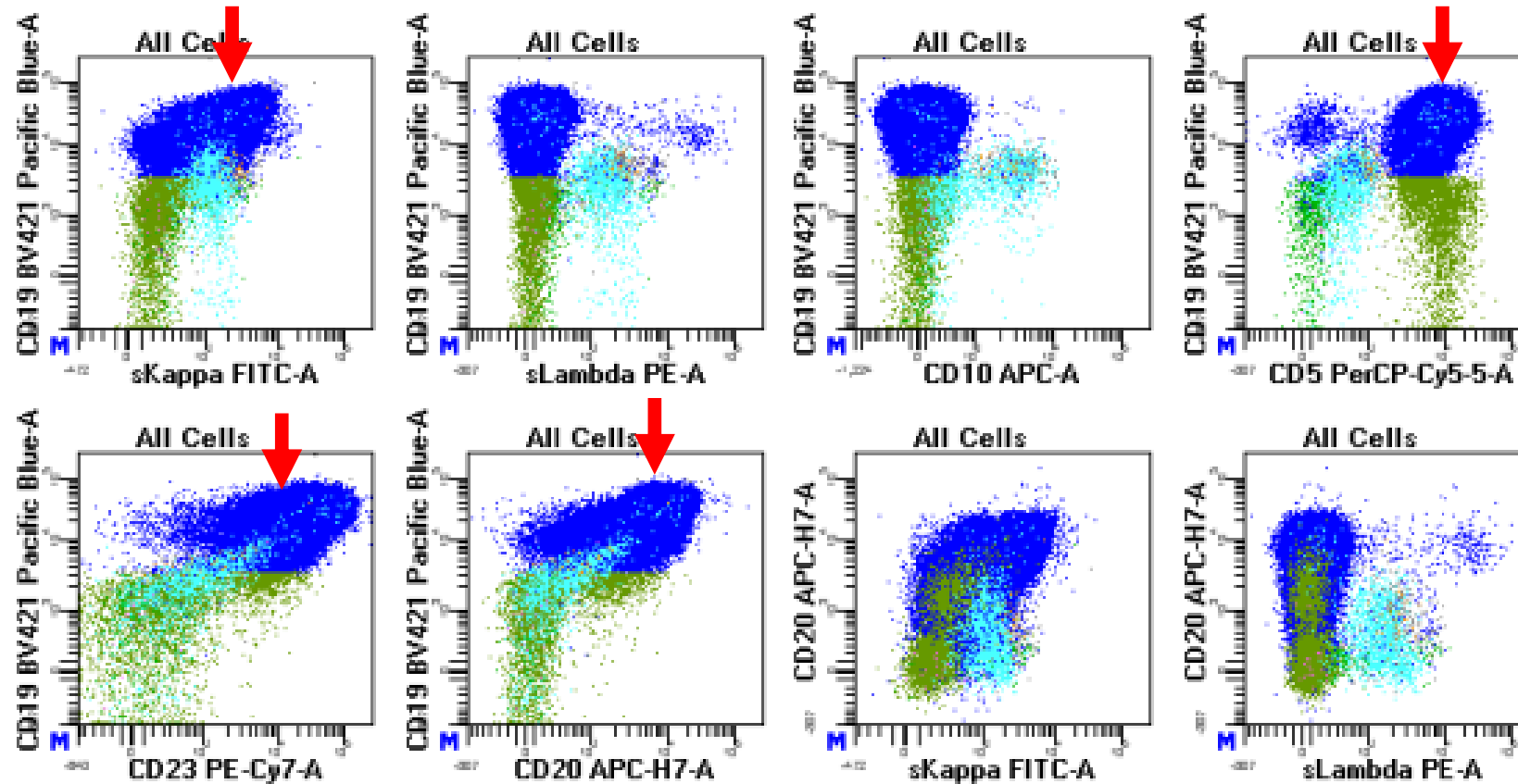
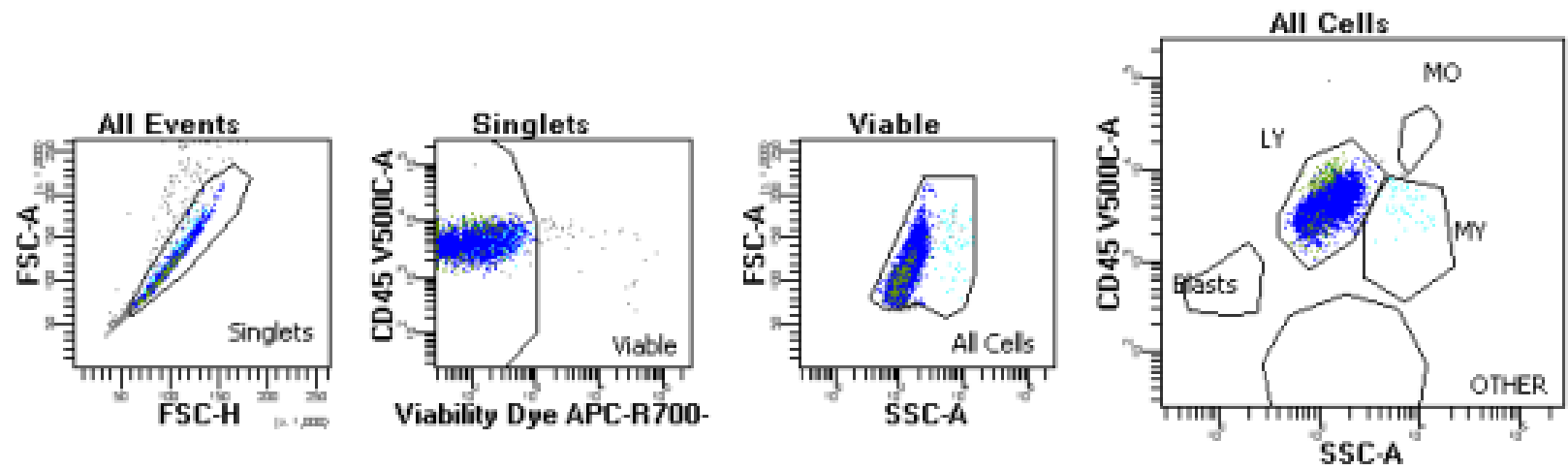
- 68-year-old women with chronic lymphocytic leukemia (CLL), under observation since January 2024.
- In January 2025, she presented with worsening lymphocytosis and lymphadenopathy.

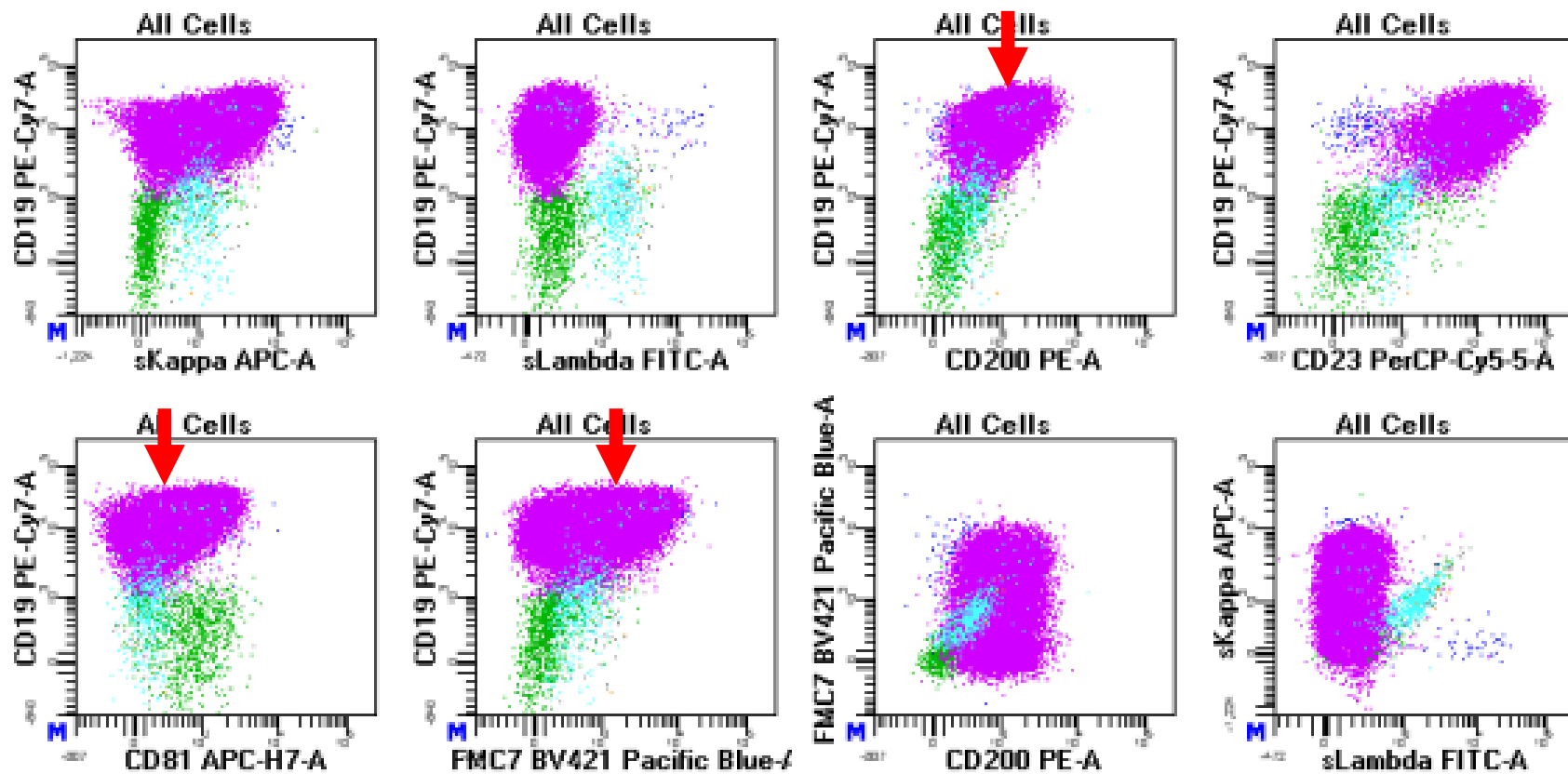
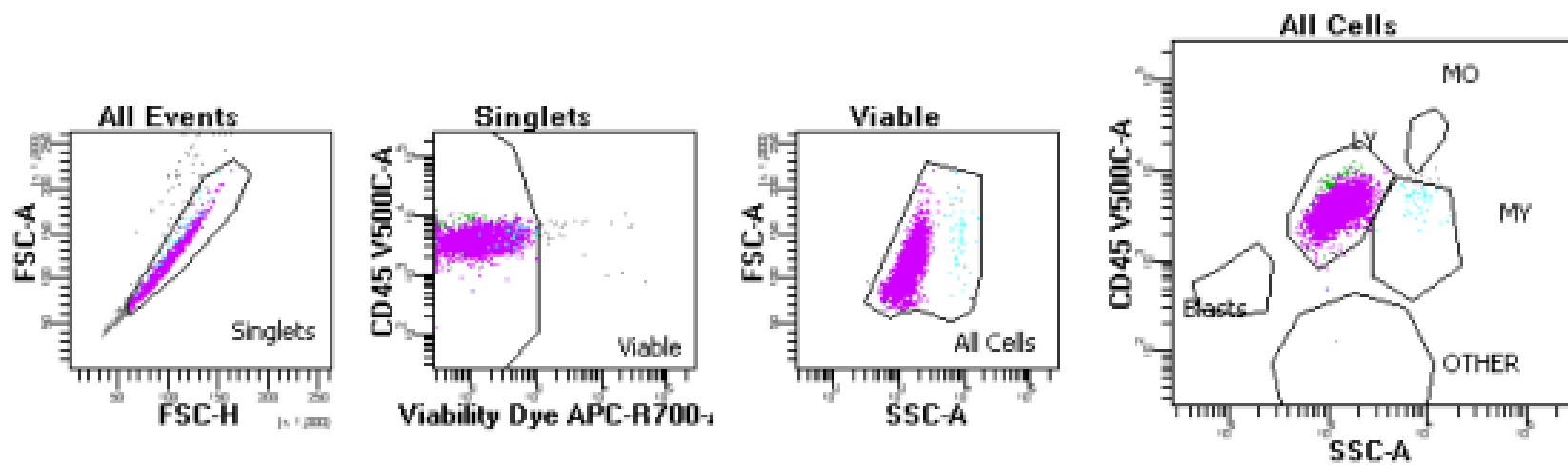
Peripheral blood



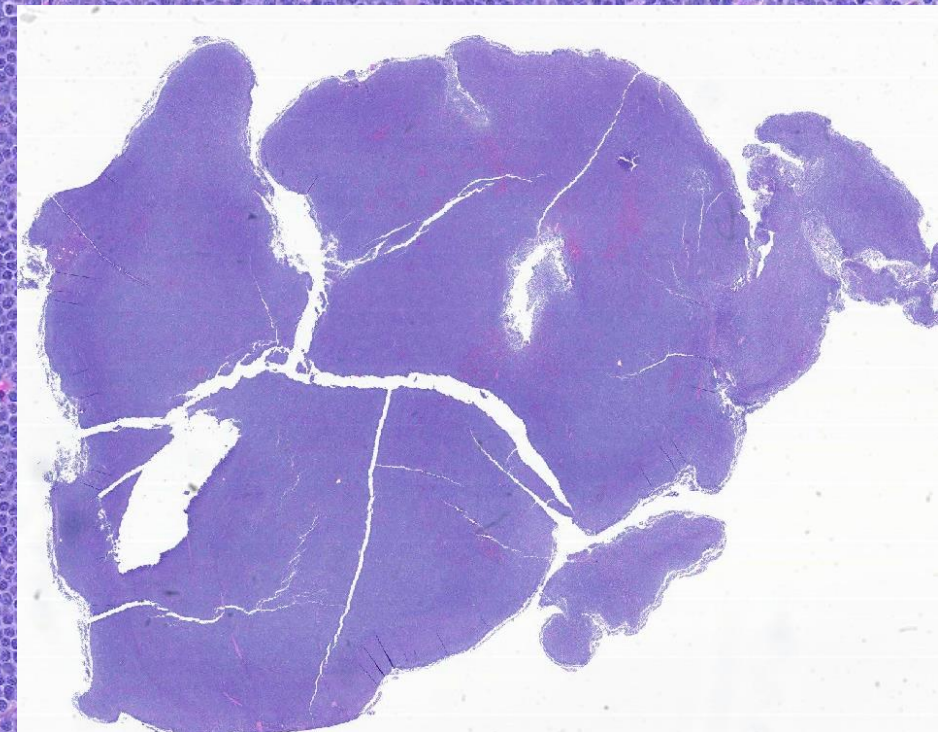
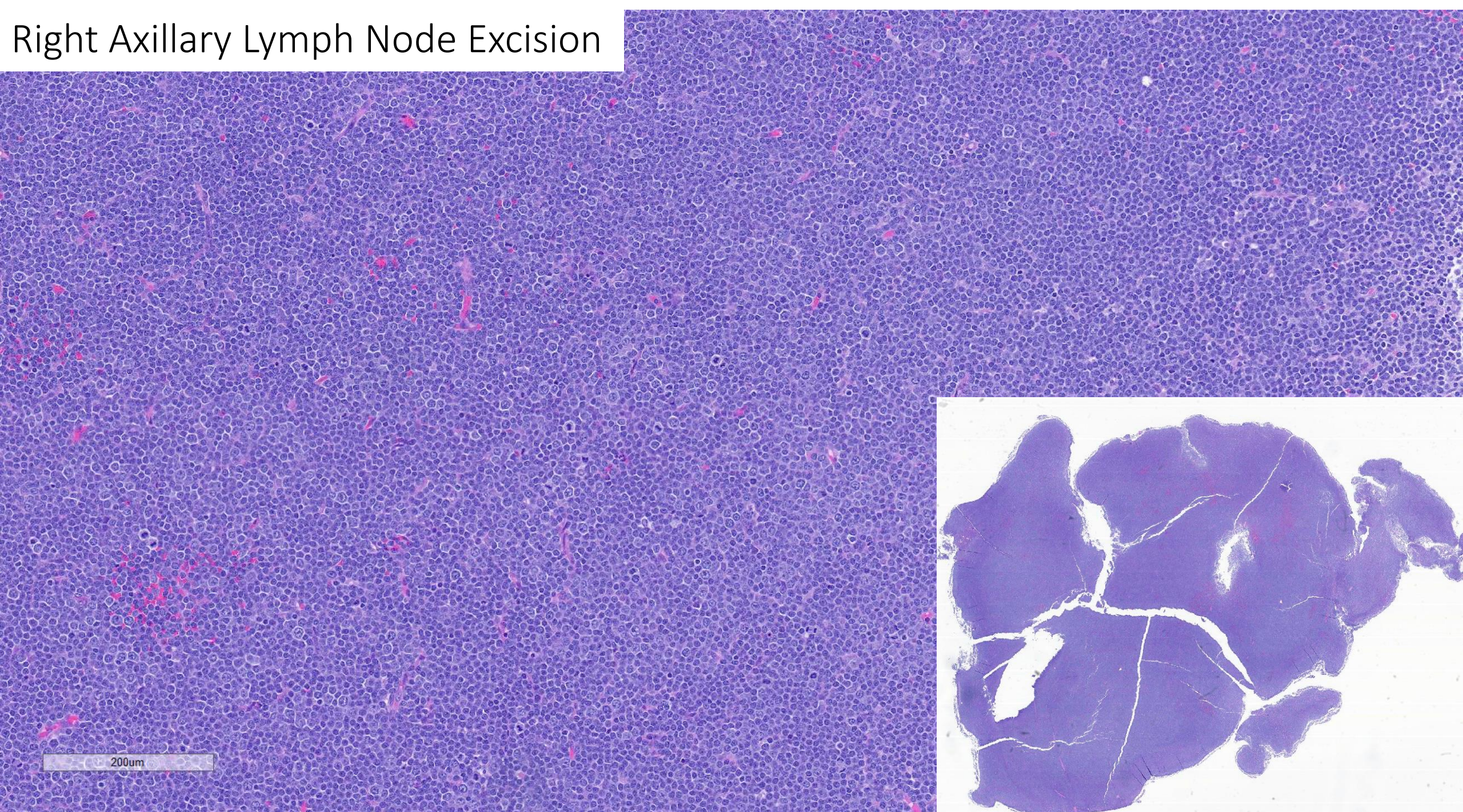
50um

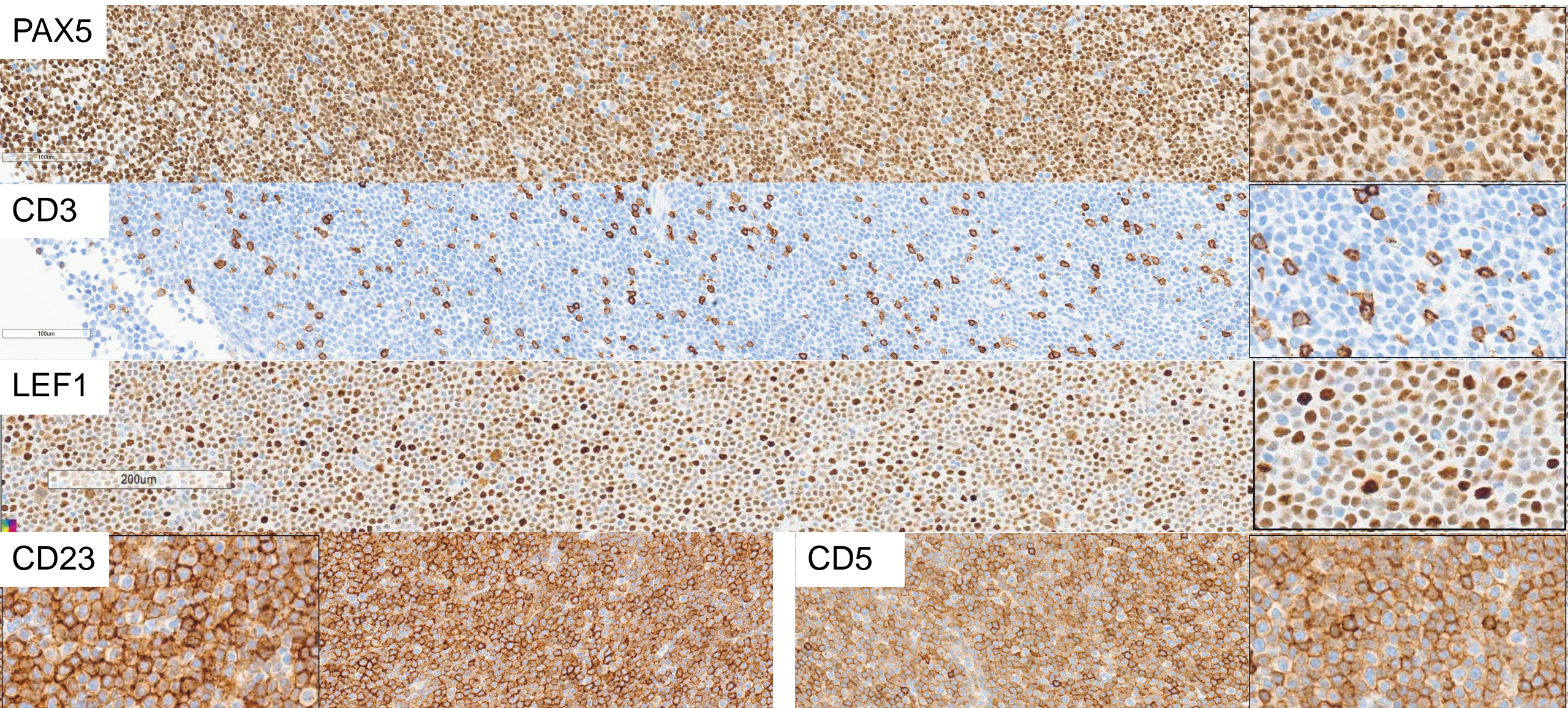
WBC	421.9 K/uL
Lymphocytes	91%





Right Axillary Lymph Node Excision



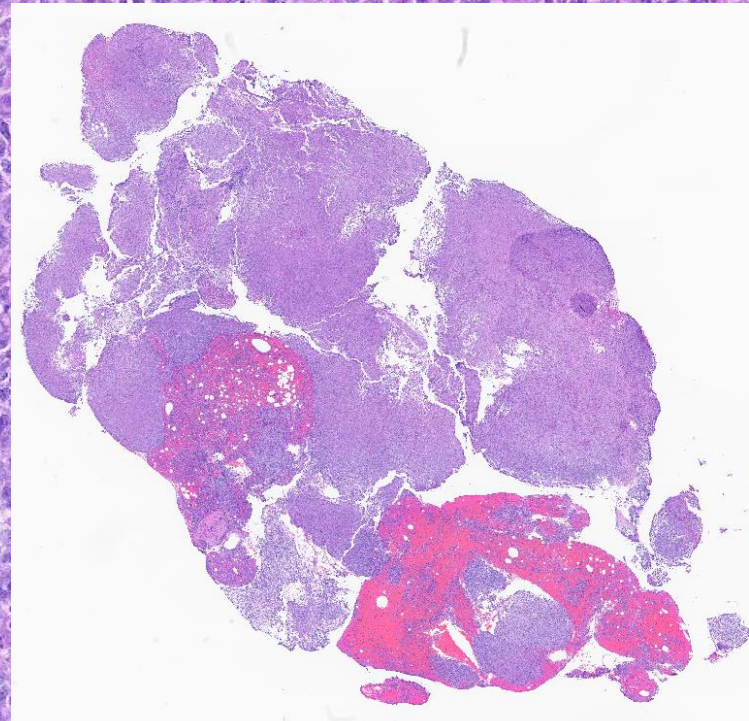
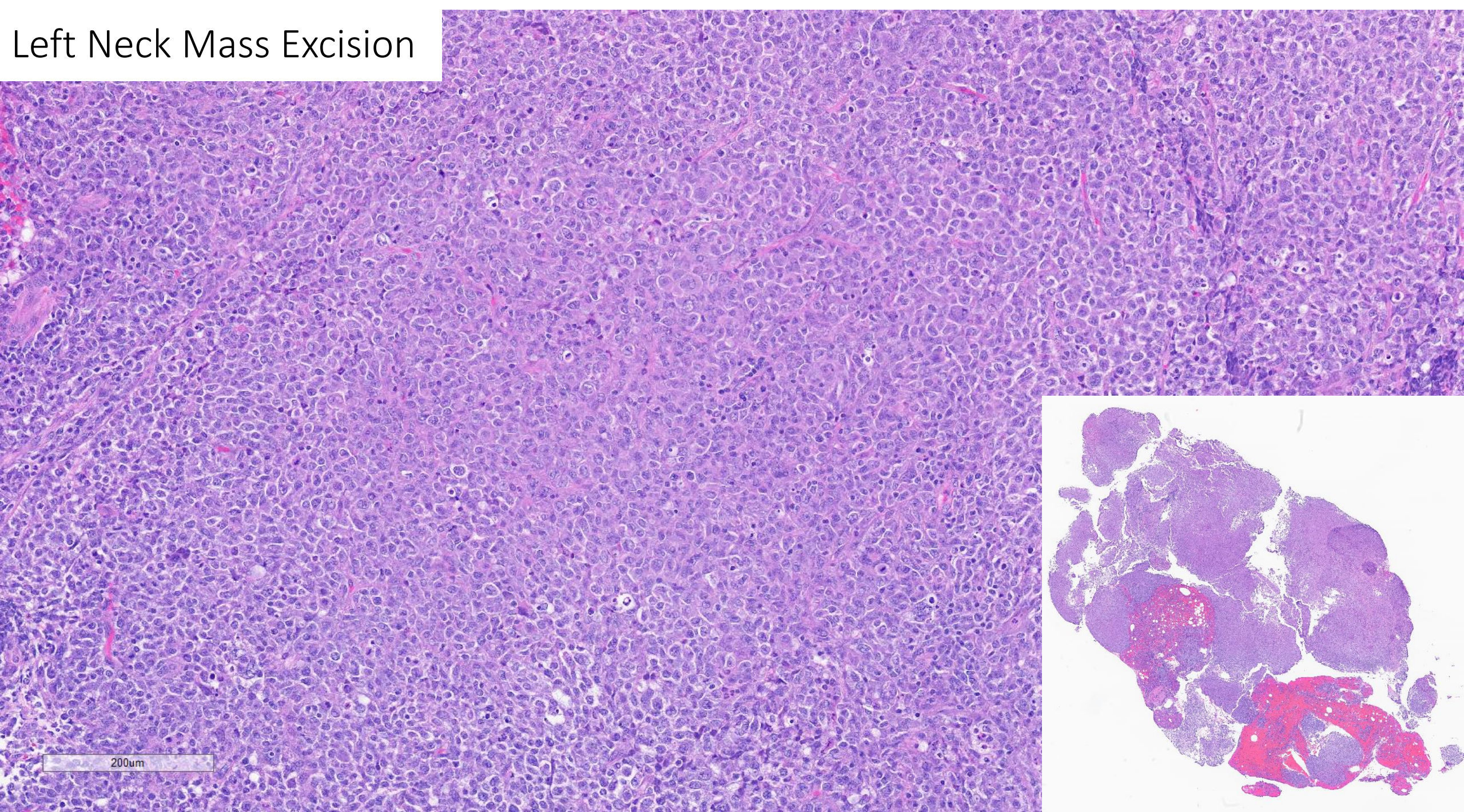


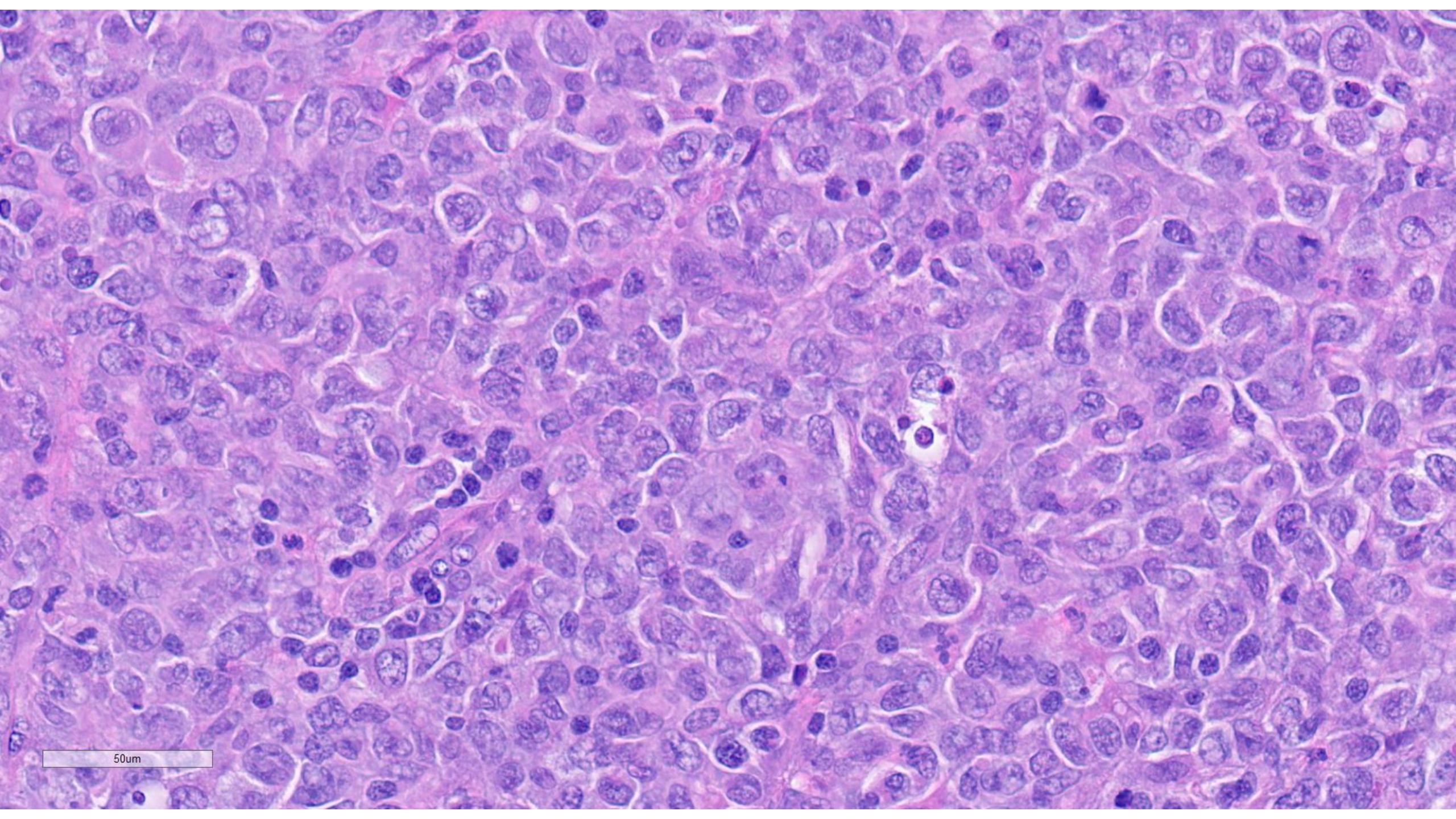
Diagnosis: Chronic lymphocytic leukemia/small lymphocytic lymphoma (CLL/SLL)

Clinical Information

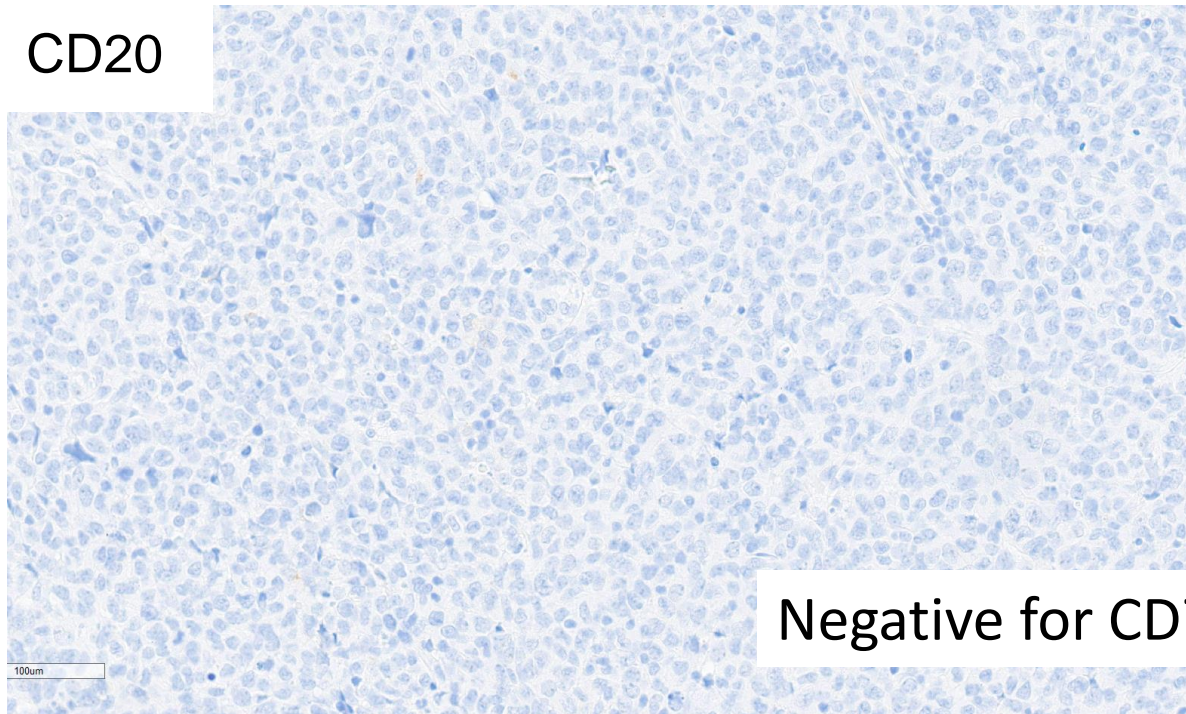
- The patient was treated with rituximab, zanubrutinib, venetoclax, and obinutuzumab.
- Subsequent peripheral blood analysis showed normalization of white blood cell and lymphocyte counts, with a marked reduction in abnormal monotypic B-cell populations.
- In March 2025, the patient developed worsening abdominal pain. PET-CT revealed new FDG-avid lymphadenopathy and multiple lesions in the liver and spleen. A left neck mass excision was subsequently performed.

Left Neck Mass Excision

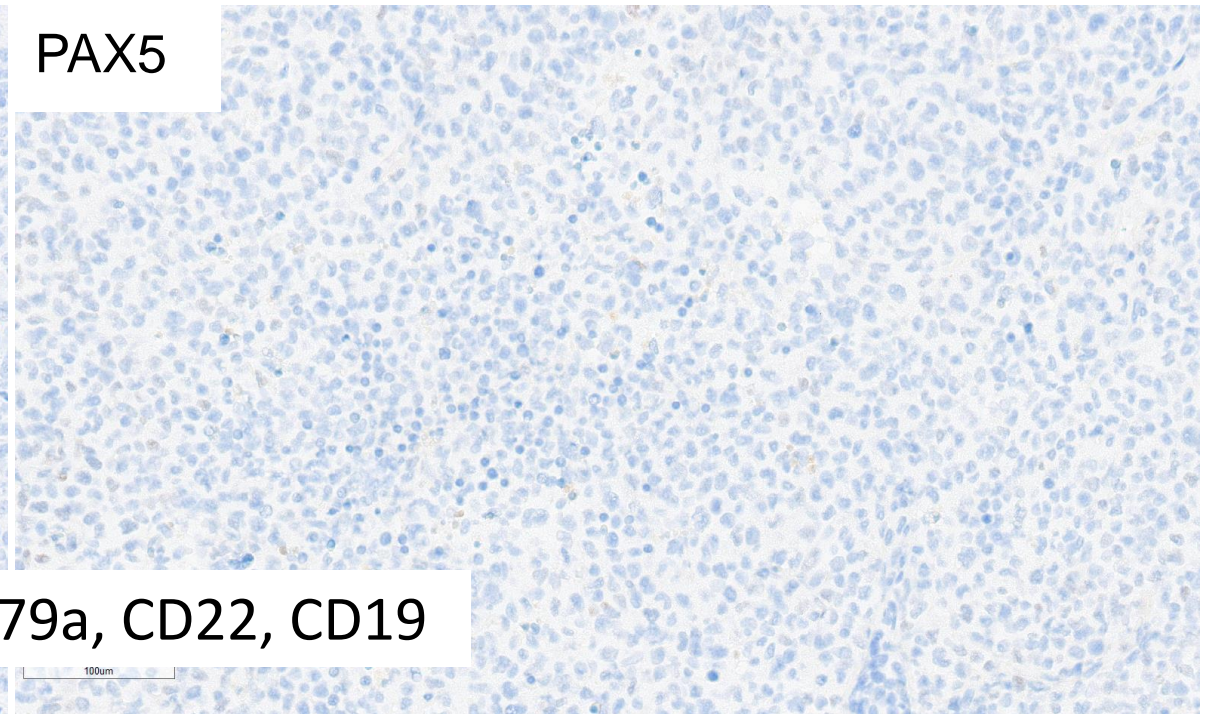




CD20

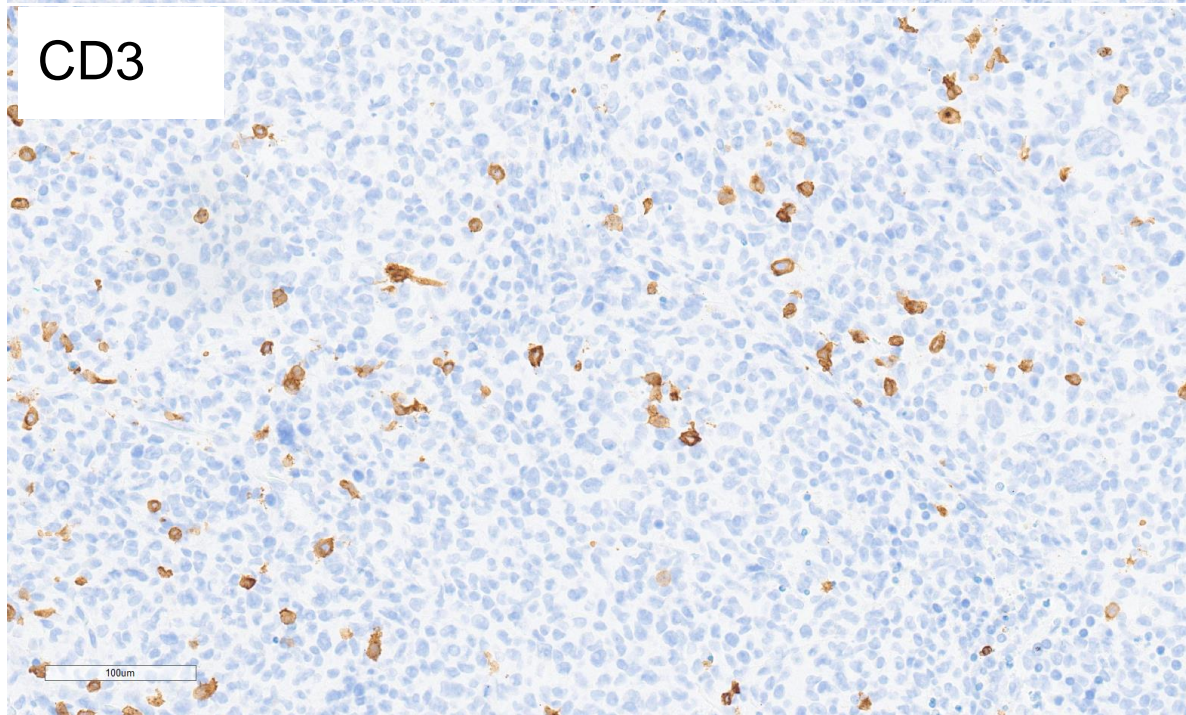


PAX5

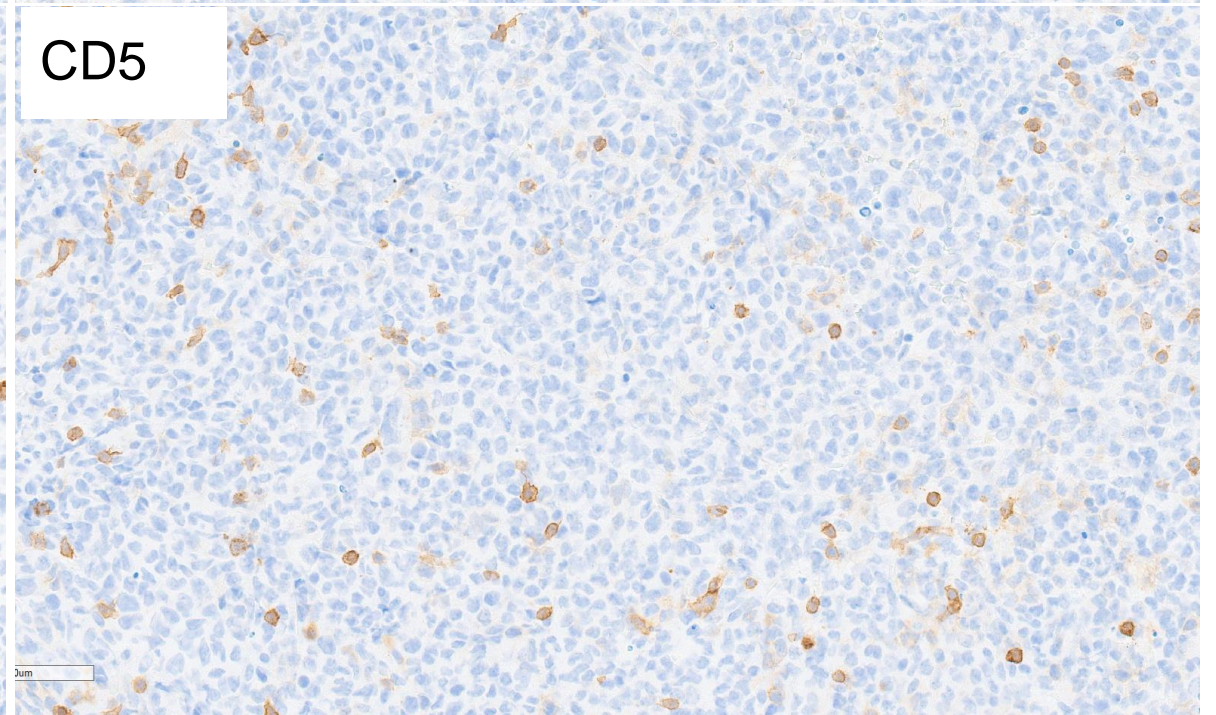


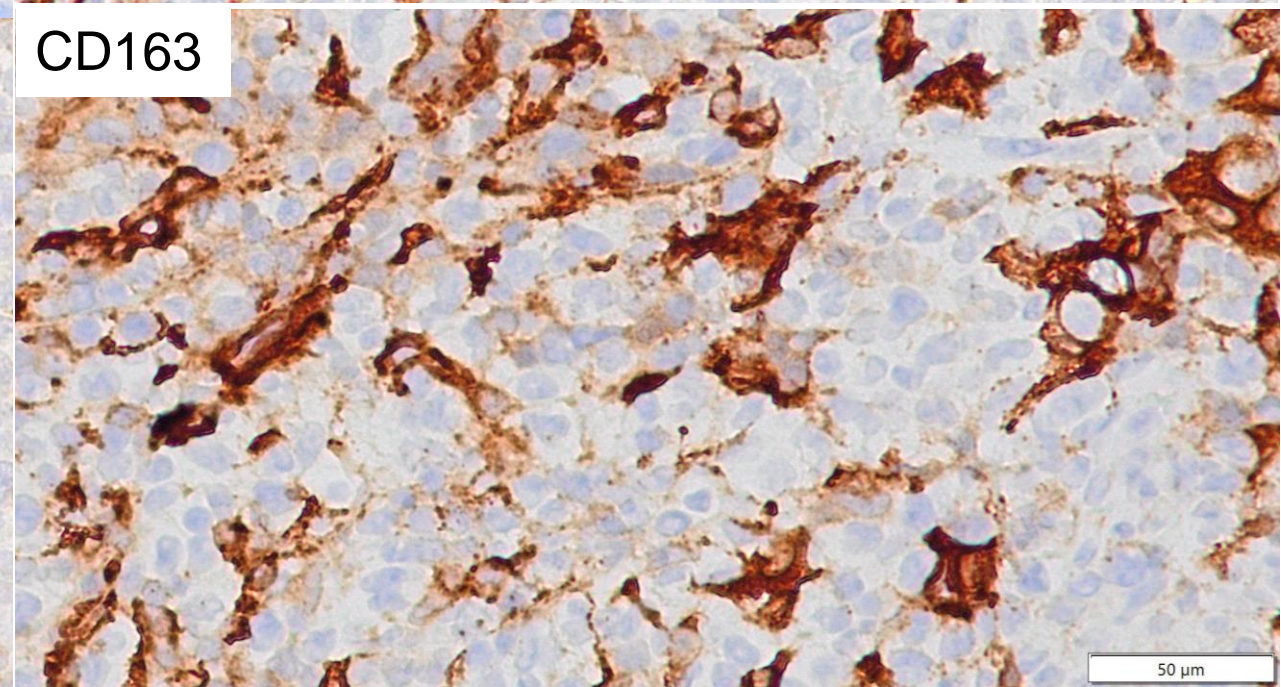
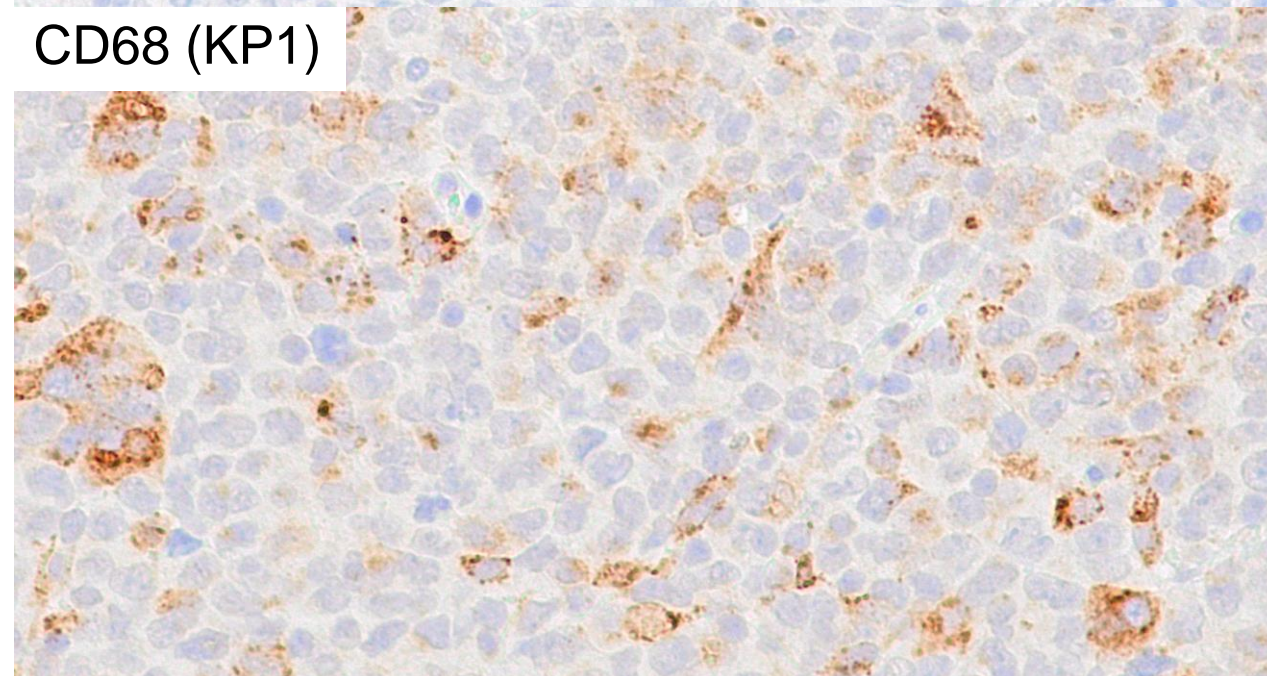
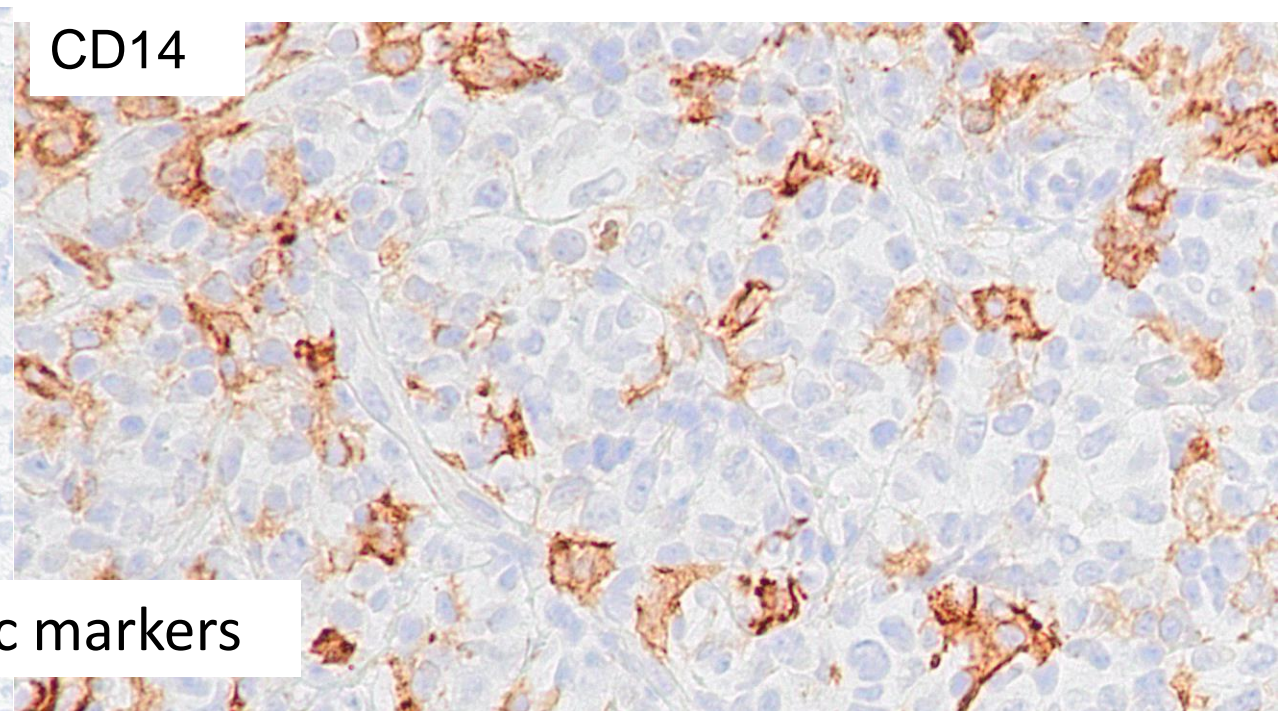
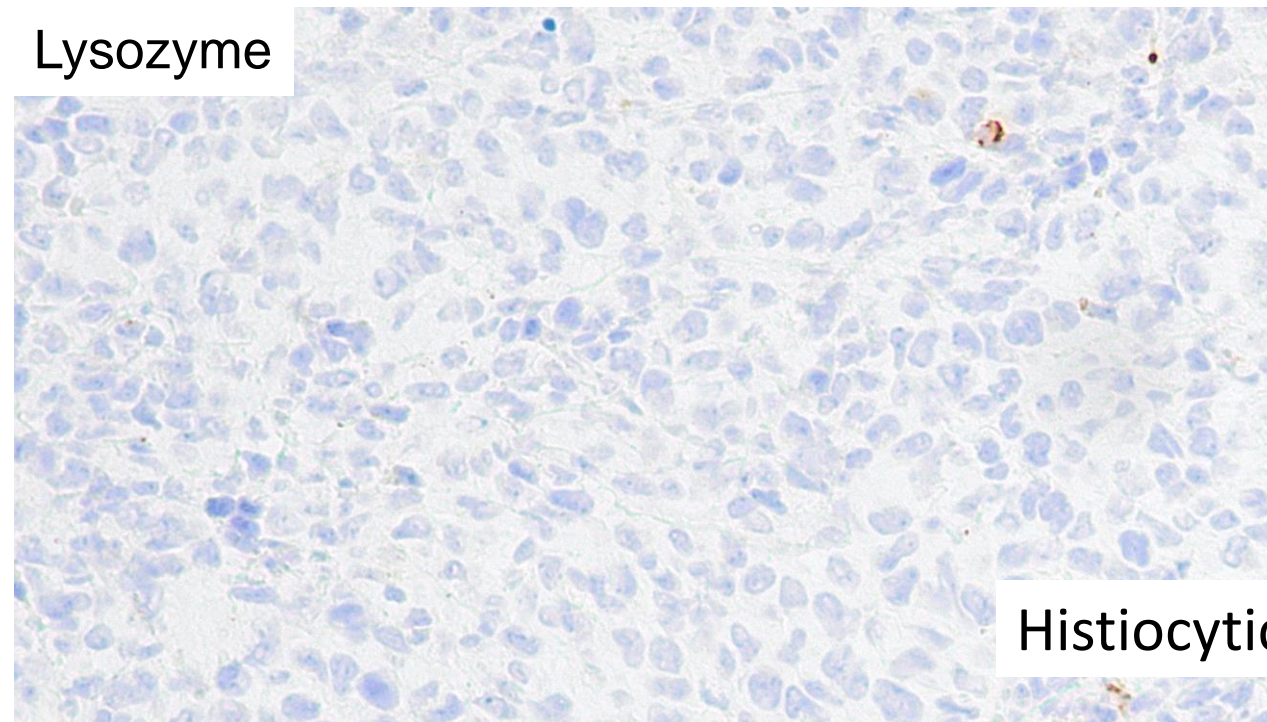
Negative for CD79a, CD22, CD19

CD3



CD5





CD33

CD56

Myeloid/monocytic markers

CD4

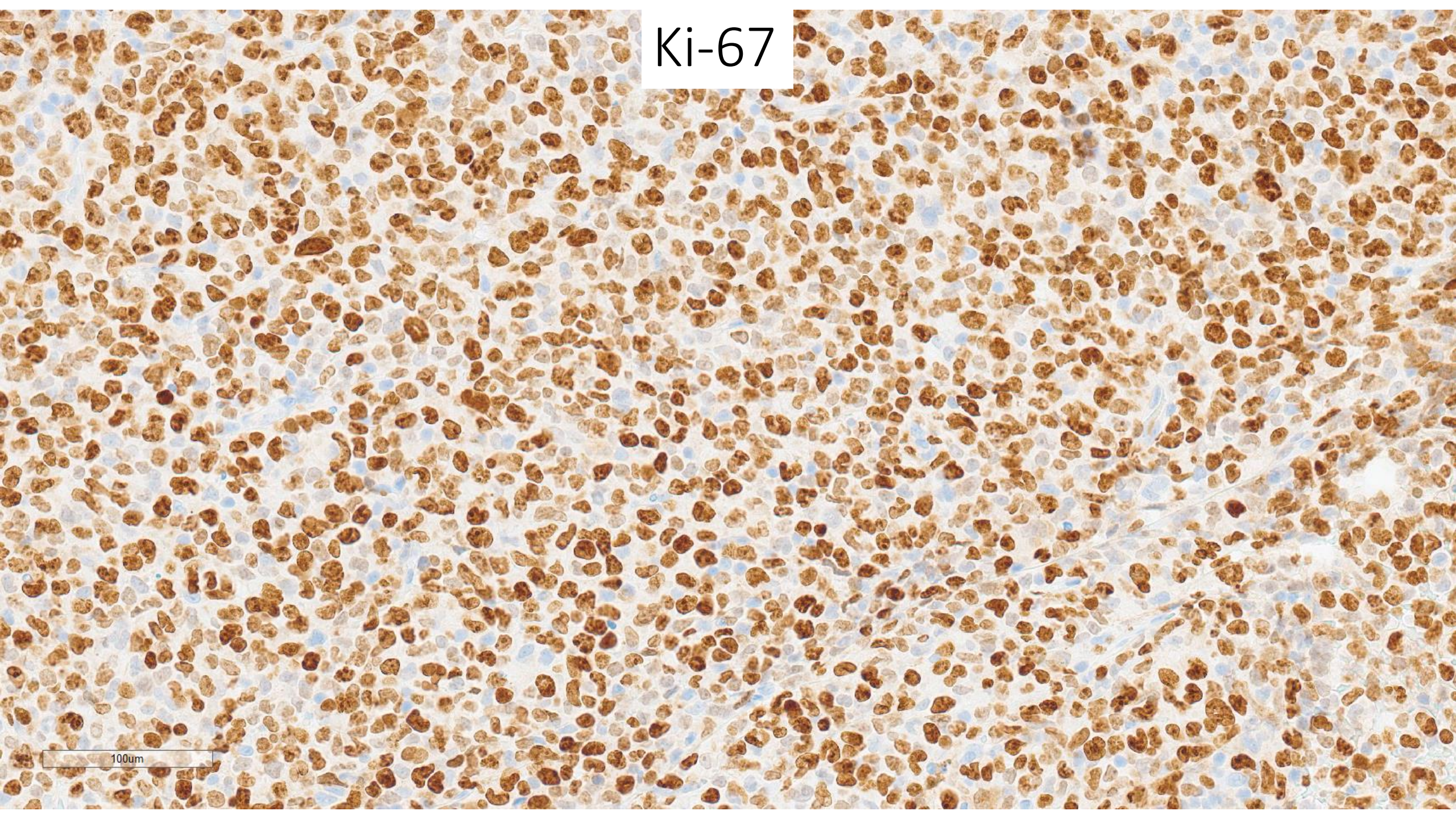
HLA-DR

c-Myc

100um

Ki-67

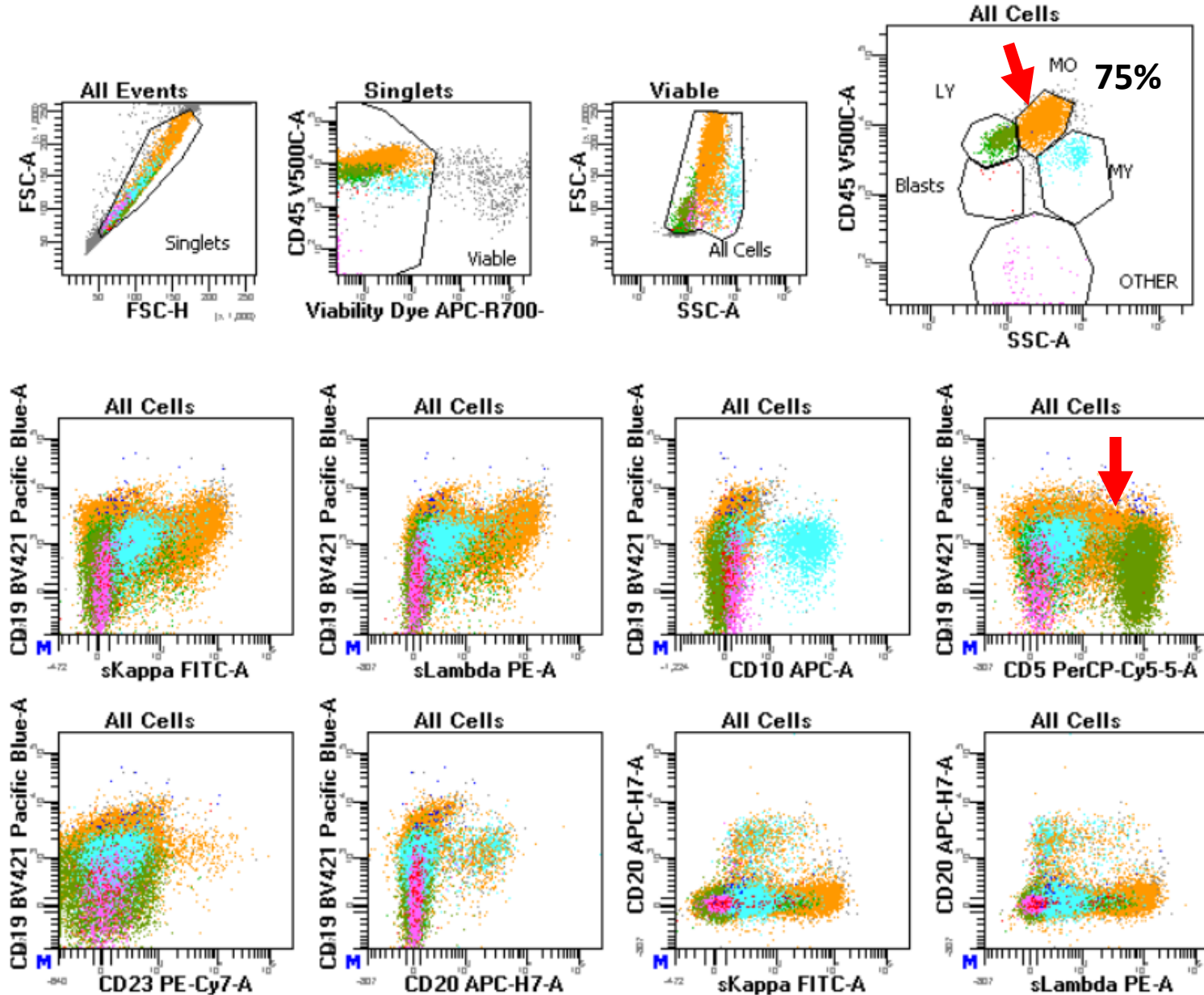
100um



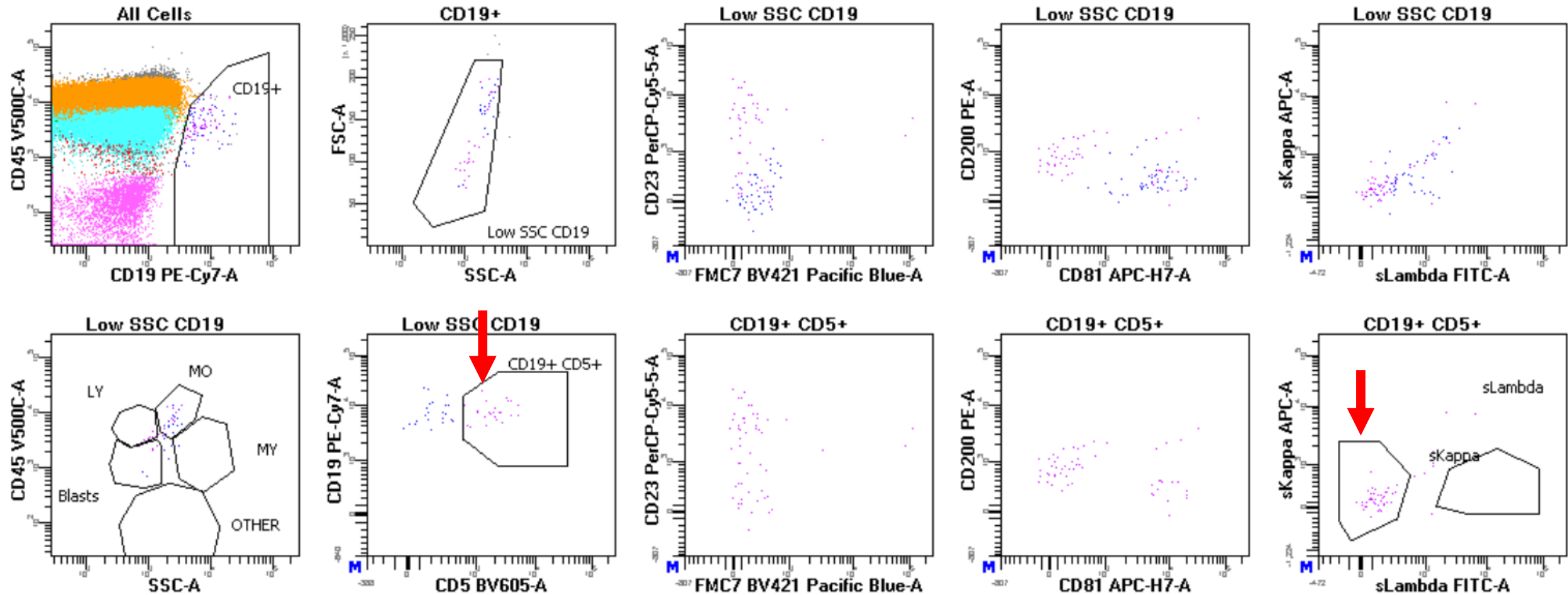
Summary of immunohistochemistry stains

Positive	Negative
HLA-DR, CD33, CD56, CD4, fascin, CD31, c-Myc, LEF1, CD38 (weak/partial), CD5 (small subset/partial), BCL2 (small subset/weak), and BCL1 (partial/weak)	CD3, CD19, CD20, CD22, Pax-5, CD79a, MUM1, CD30, EMA, CD21, CD23, CXCL13, D2-40, CD34, CD117, MPO, Lysozyme, CD15, CD11c, CD14, CD68/KP1, CD68/PGM1, CD163, S100, CD1a, Langerin, PD-L1, CD123, p53, mutant NPM1, Kappa ISH, Lambda ISH, EBER ISH

The specimen is predominantly composed of cells with elevated forward and side scatter properties localized within the conventional monocyte gate.



Minute kappa-monotypic B cell population



Diagnosis

- Involvement by a myeloid sarcoma.
- Transdifferentiation of the patient's CLL/SLL into a myeloid sarcoma?

RESULTS OF GENE REARRANGEMENT ANALYSIS

SPECIMEN: A. Peripheral Blood for Molecular **CLL/SLL**

RESULTS:

Ig Heavy Chain: Monoclonal

Ig Kappa Light Chain: Monoclonal

INTERPRETATION: For IGH, a monoclonal pattern was identified (149bp). For IGK, a monoclonal rearrangement was detected (152bp, 200bp in tube A).

SPECIMEN: A. Left Neck Mass for Molecular **Myeloid sarcoma**

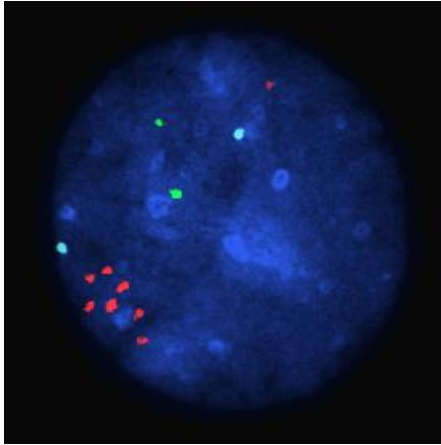
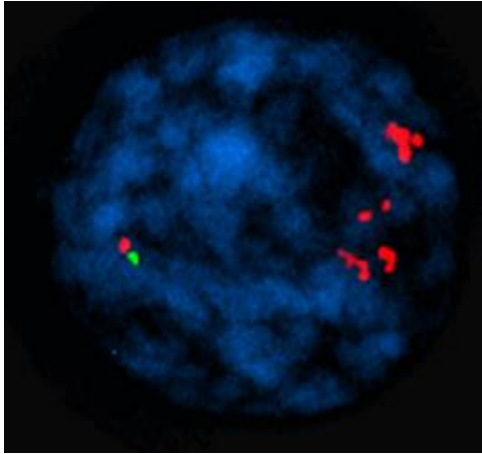
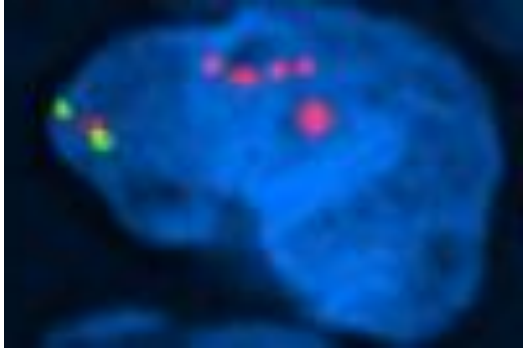
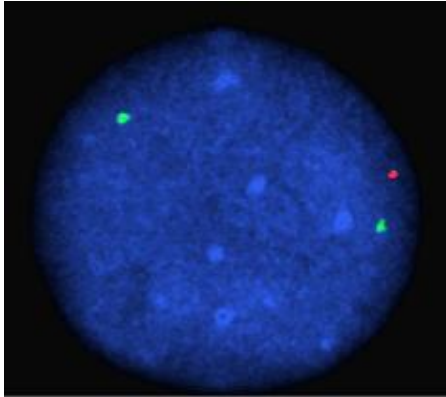
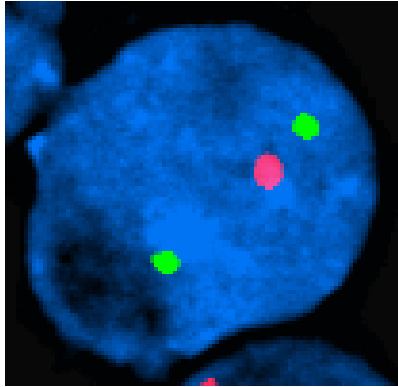
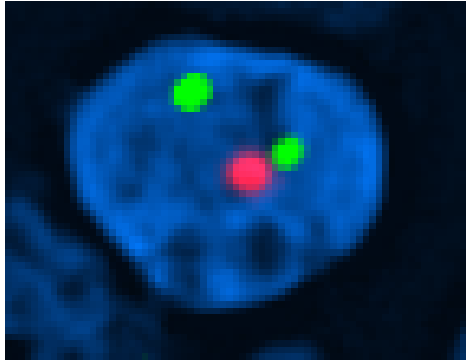
RESULTS:

Ig Heavy Chain: Monoclonal

Ig Kappa Light Chain: Monoclonal

INTERPRETATION: For IGH, a monoclonal pattern was identified (149bp). For IGK, a monoclonal rearrangement was detected (152bp + 200bp in tube A).

FISH positive for MYC amplification and 7q deletion

	Peripheral blood CLL/SLL	Right axillary lymph node CLL/SLL	Left neck mass Myeloid sarcoma
MYC amplification			
7q deletion			

Bone marrow biopsy

Diagnosis:

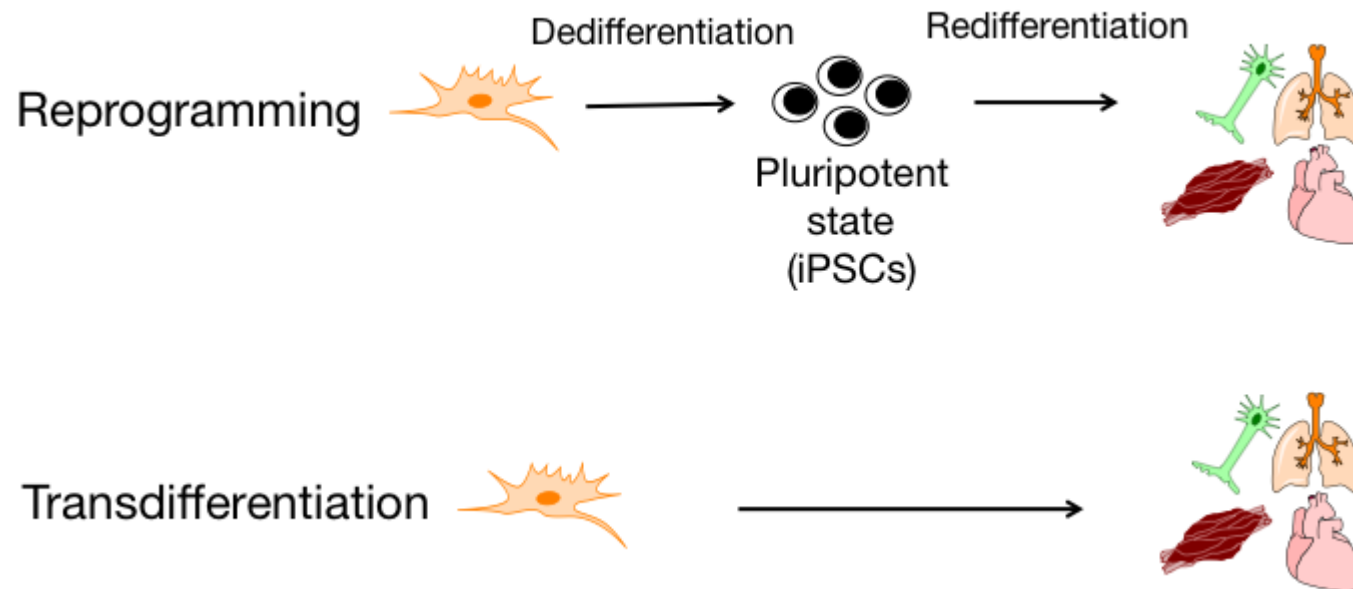
- Chronic lymphocytic leukemia/small lymphocytic lymphoma, residual focal bone marrow involvement (~5% of marrow cellularity).

Cytogenetics:

- 46,XX[20]
- FISH AML/MDS panels: Normal

Transdifferentiation

- Transdifferentiation refers to a process in which a mature cell type switches lineage identity into a phenotypically and functionally distinct lineage without reverting to a pluripotent state.

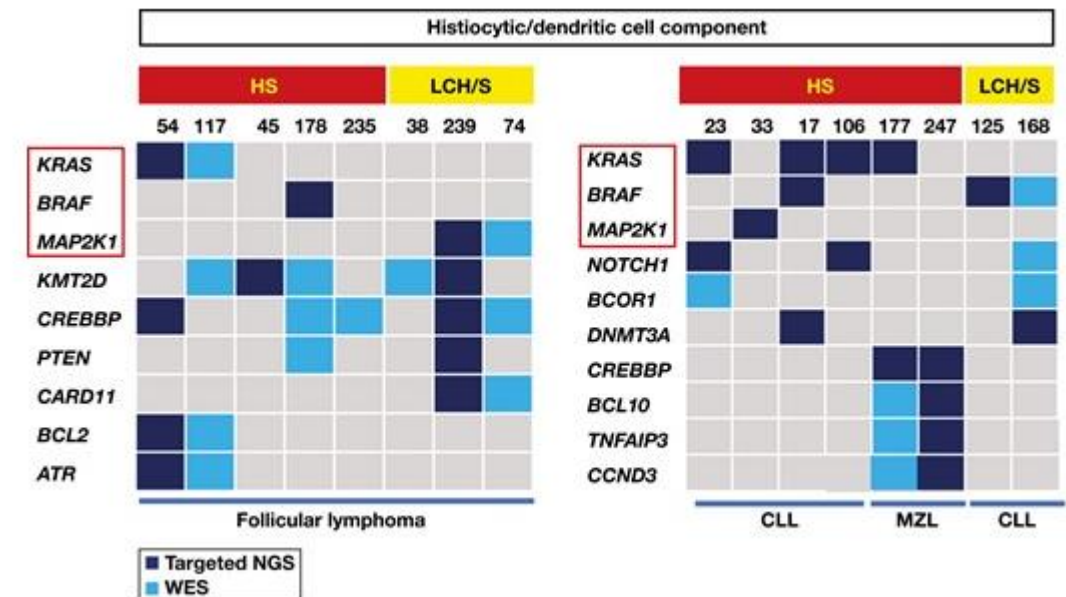


- In hematologic malignancies, transdifferentiation is most notably observed in low-grade B-cell lymphomas (follicular lymphoma, CLL/SLL, and marginal zone lymphoma) that transdifferentiate into histiocytic/dendritic cell neoplasm.
- These cases often occur in the setting of disease progression or relapse and may follow targeted therapies such as rituximab or BTK inhibitors.
- To the best of our knowledge, transformation of CLL/SLL into a lesion most consistent with myeloid sarcoma has not been previously reported in the literature.

Mutational profile

- A high frequency of mutations in the RAS/MAPK pathway was found in secondary histiocytic/dendritic neoplasms associated with diverse lymphoid malignancies.

	FL						BCL	CLL	B-ALL	TCL	T-ALL			
IG				PC						ND	PC	PC	-	PC
TRG	ND	ND	ND	ND	ND	PC	PC	ND		ND				
IGH/BCL2	*	*	*	*	*	*	-	ND	ND	ND	ND	ND	ND	ND
BRAF														
KRAS														
MAP2K1														
NRAS														
NF1														



Egan C. et al. Mod Pathol. 2021 Feb;34(2):336-347.

Xiao W. et al. Am J Clin Pathol. 2023 Jun 1;159(6):522-537.

Genomic Findings	
IA	IB
No variants reported.	No variants reported.

Genomic Findings

IA	IB	IIC		IID	
No variants reported.	No variants reported.	NRAS	p.G12D c.35G>A VAF 41.4%	FBXW7	p.R361P c.1082G>C
		KRAS	p.G13D c.38G>A VAF 53.3%	CDKN2B	Copy number loss in <i>CDKN2B</i> (1 copy)
		0 Clinical Trials		CDKN2A	Copy number loss in <i>CDKN2A</i> (0 copies)
				ATM	Copy number loss in <i>ATM</i> (1 copy)
				MYC	Copy number gain in <i>MYC</i> (5 copies)
				0 Clinical Trials	

Proposed mechanism

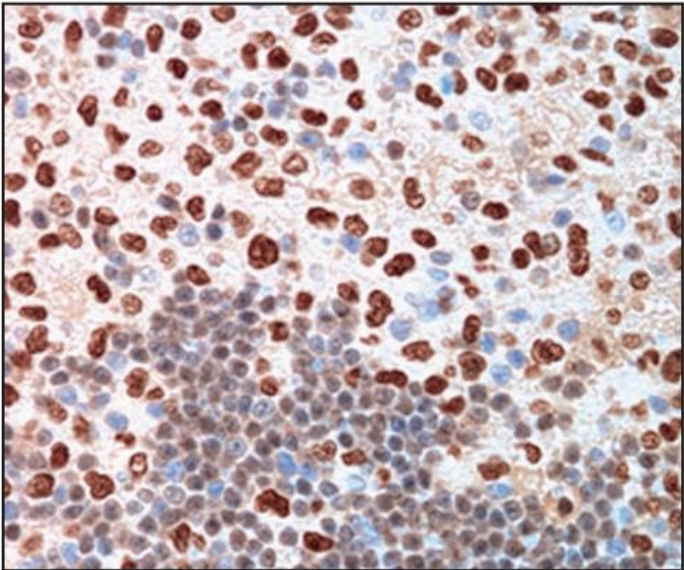
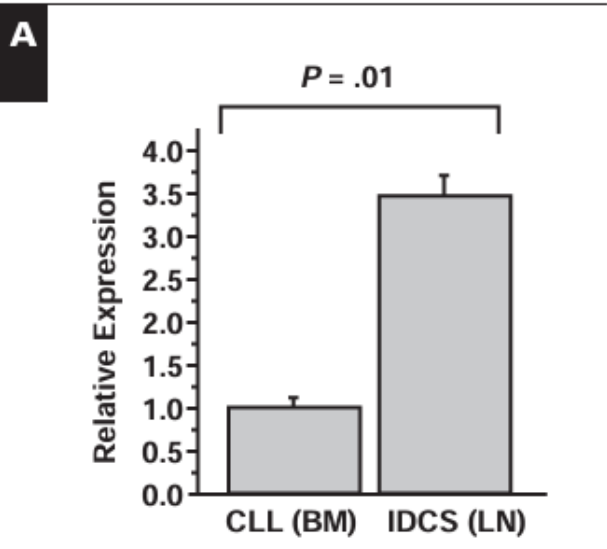
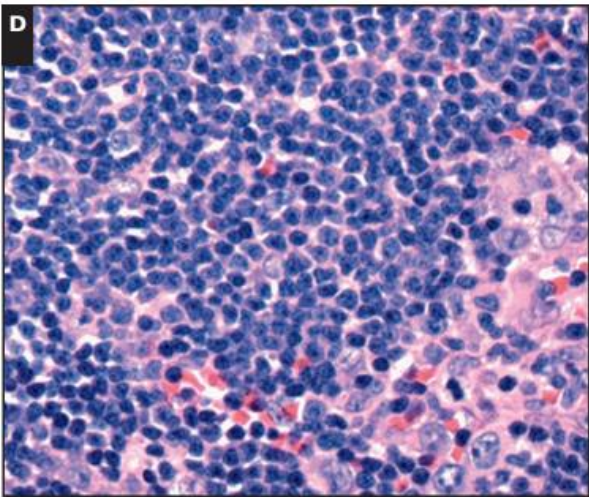
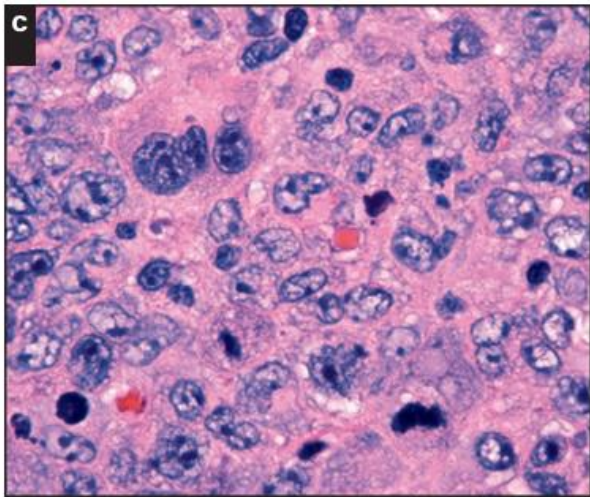
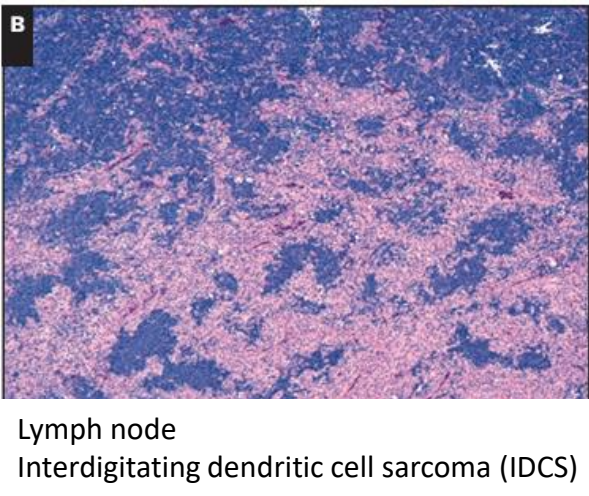
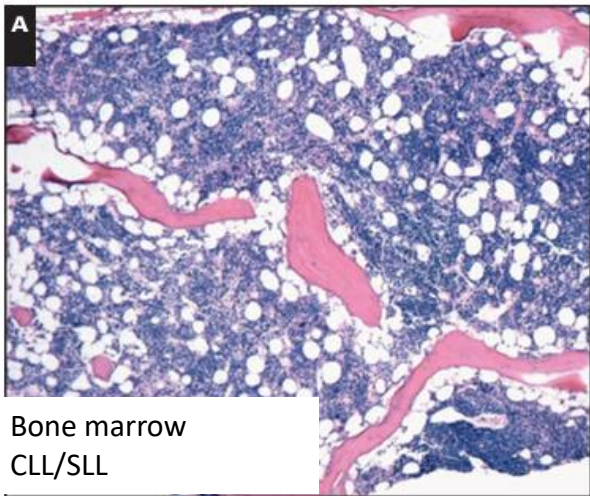
- Transdifferentiation is driven by disruption of lineage-specifying transcription factors (e.g., PAX5) and induction of myeloid-associated regulators like PU.1 and C/EBP α .
- In murine models, the deletion of *Pax5* in B cells results in reprogramming toward macrophage lineages.

Transformation of chronic lymphocytic leukemia/small lymphocytic lymphoma to interdigitating dendritic cell sarcoma: evidence for transdifferentiation of the lymphoma clone

Cory R Fraser¹, Wei Wang, Mario Gomez, Taotao Zhang, Susan Mathew, Richard R Furman, Daniel M Knowles, Attilio Orazi, Wayne Tam

Affiliations + expand

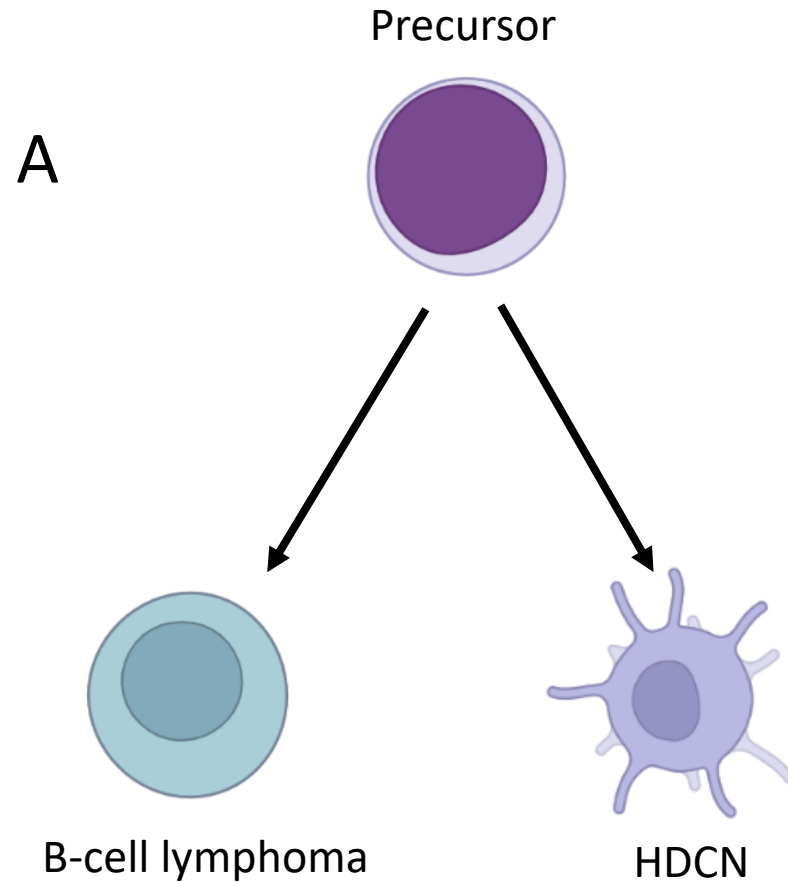
PMID: 19926586 DOI: 10.1309/AJCPWQ0I0DGXBMHO



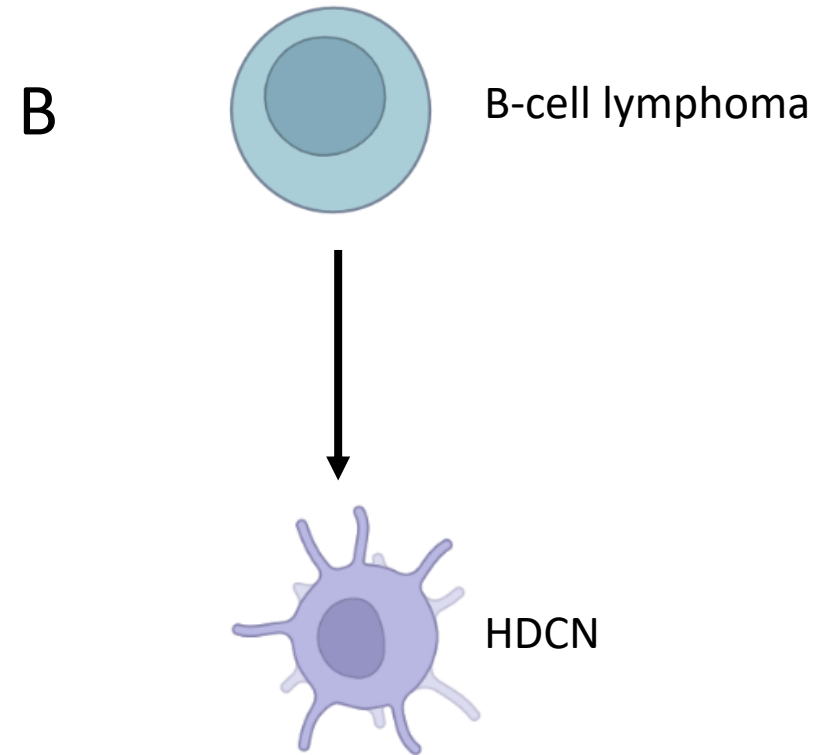
PU.1 RT-PCR

High expression of PU.1 in IDCS cells compared with the weak expression in the adjacent CLL cells

Patterns of clonal evolution



A, Divergent clonal evolution.



B, Linear evolution.

Prognostic and therapeutic implications

- These transformed neoplasms are typically clinically aggressive, with limited response to conventional therapies. Median survival is often less than two years following transformation.
- Treatment strategies are not standardized and may rely on phenotype-directed regimens.
- Emerging therapies targeting lineage plasticity, epigenetic modifiers, and tumor microenvironmental signaling are currently under investigation.



Thank you!



