

Multi-Institutional Hematopathology Interesting Case Conference (MHICC)

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Memorial Sloan Kettering Cancer Center

04/22/2026

Disclosure

I have nothing to disclose

73, F

- weight loss and persistent right leg pain

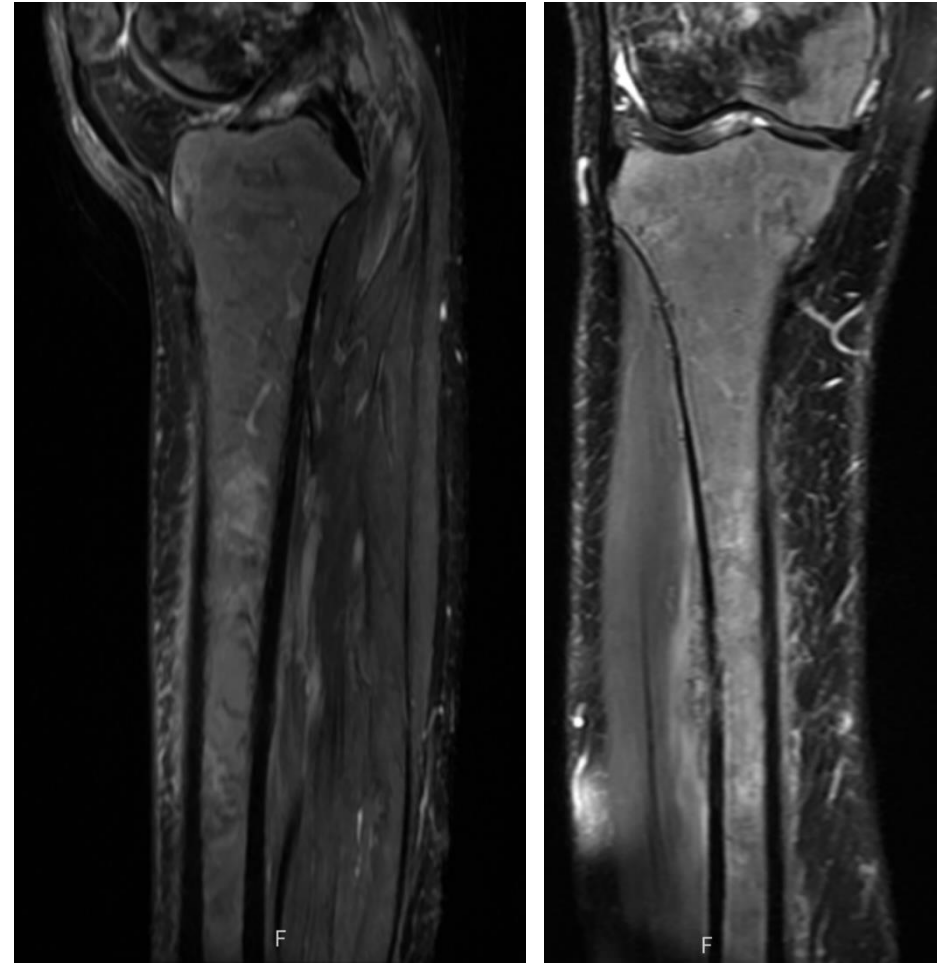
Medical Hx

- Familial Mediterranean Fever
- Osteoarthritis
- Bilateral DCIS, post-lumpectomy (2002)
- CLL (2018), observation





11/12/2025



11/22/2025

Proximal to mid right tibial diaphysis aggressive lesion with cortical erosions and extraosseous soft tissue. Differential diagnosis includes leukemic involvement, primary osseous lesion, and metastasis.

R L



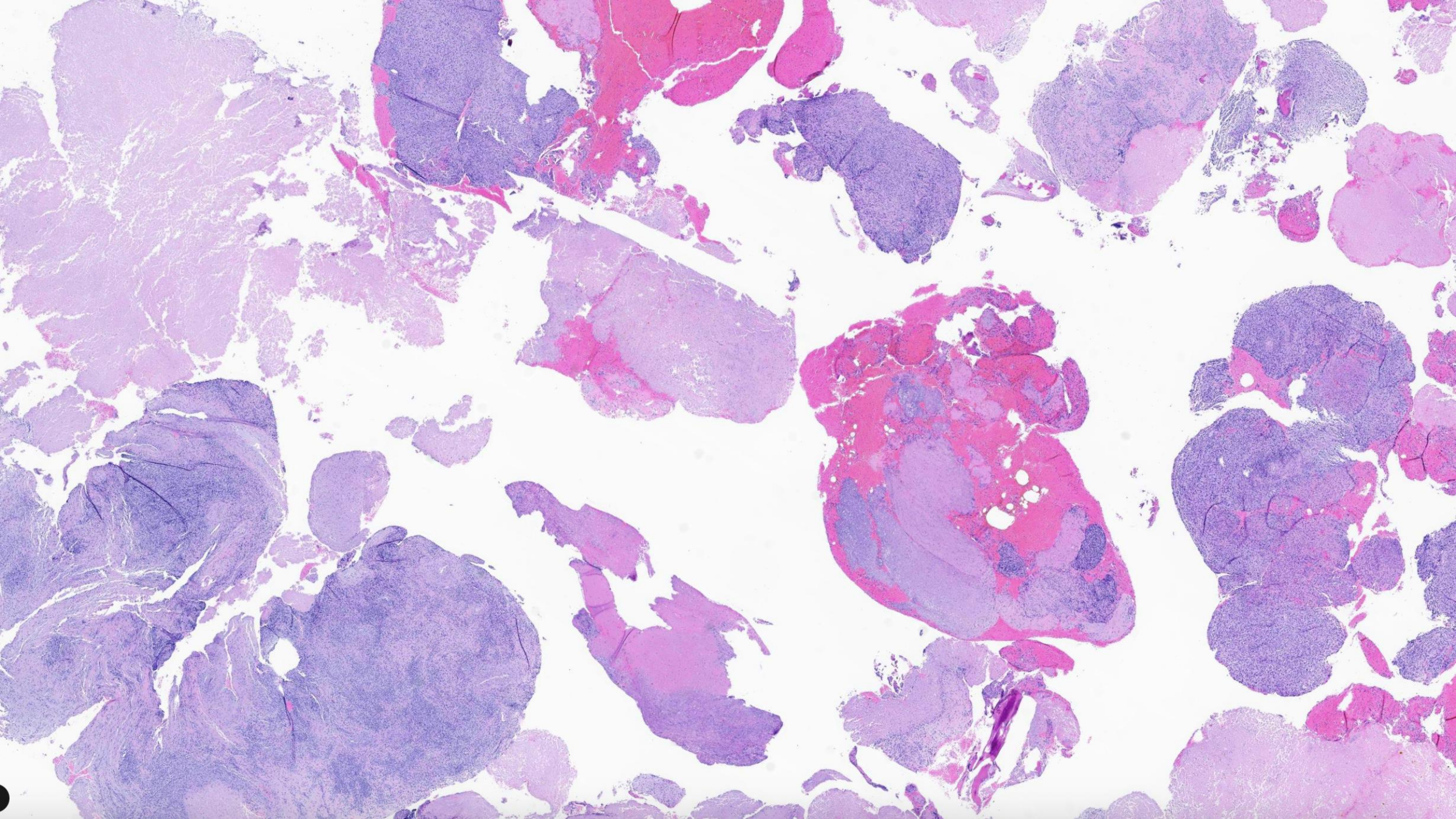
1. Abundant normal sized and slightly enlarged lymph nodes throughout the body with only minimal FDG uptake (SUV <2), consistent with CLL.

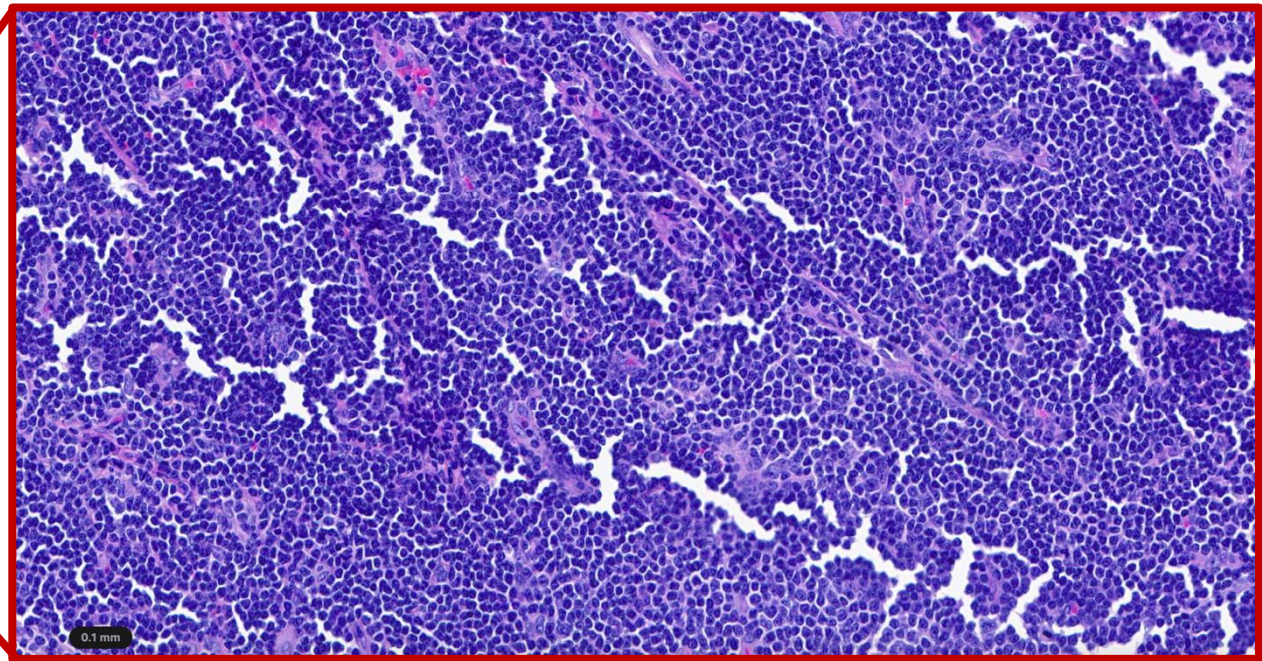
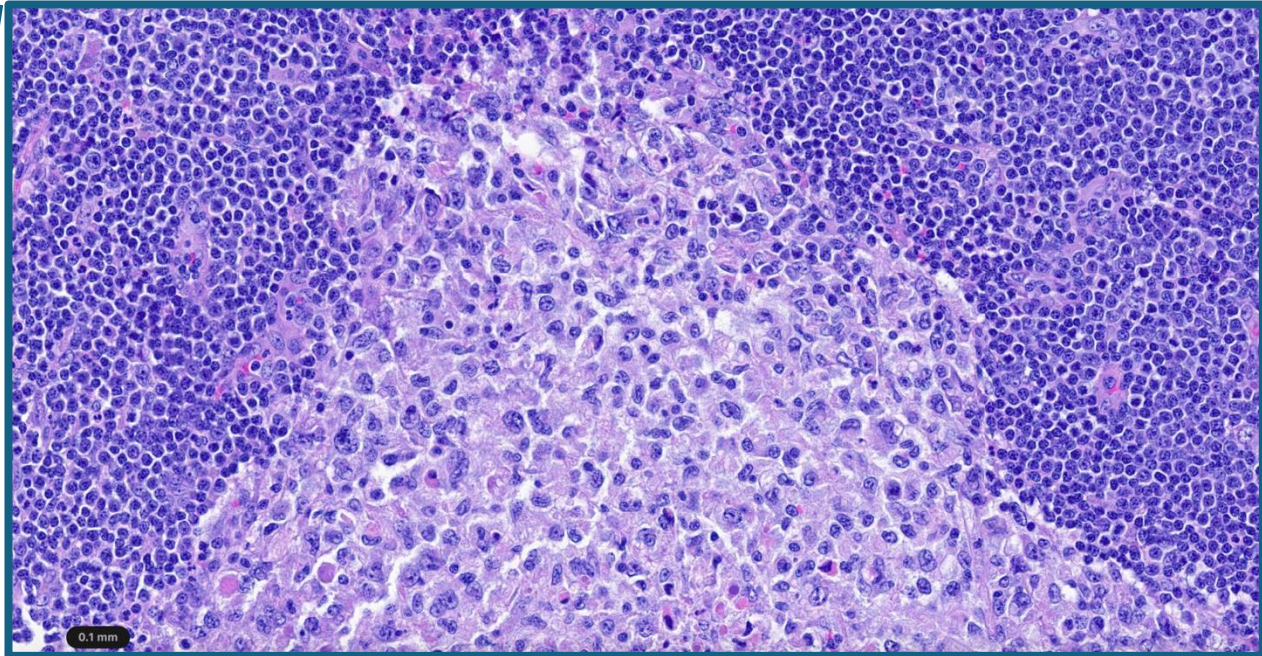
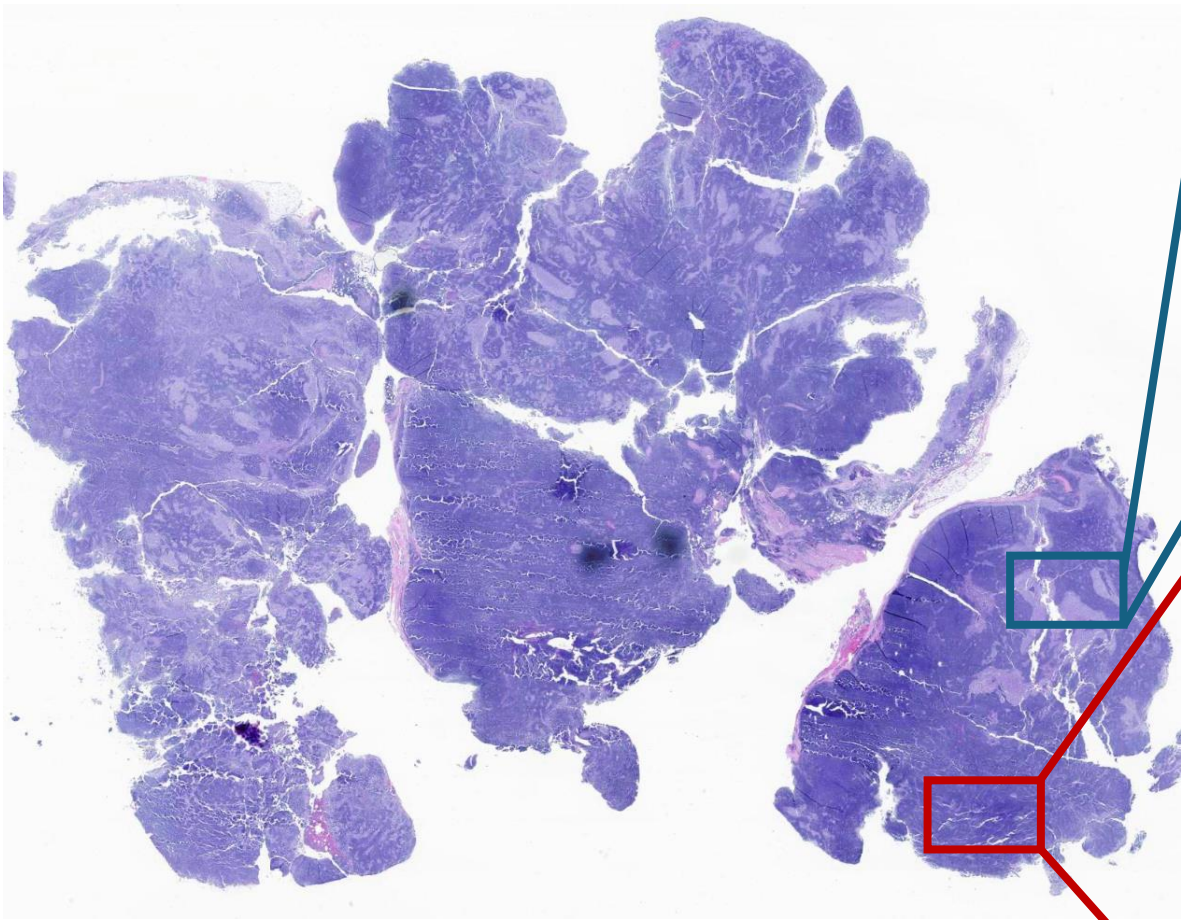
3. Prominent homogeneous marrow uptake is suspicious for CLL involvement.

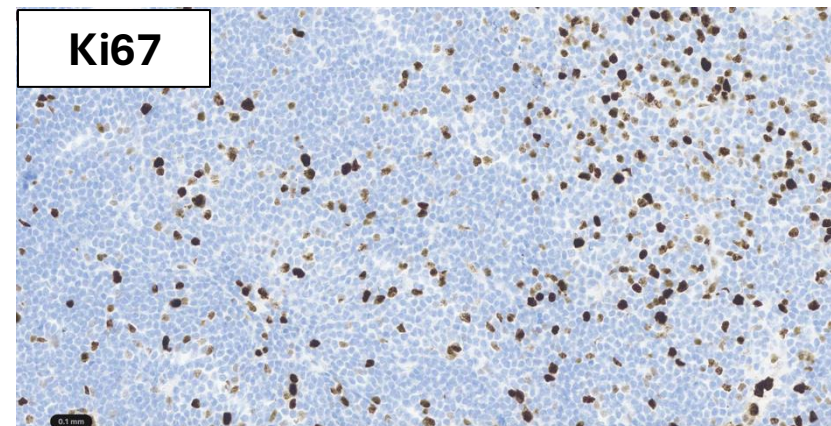
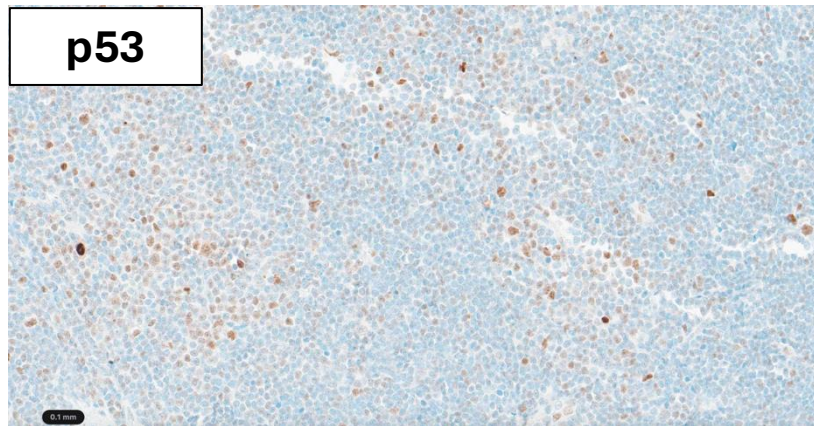
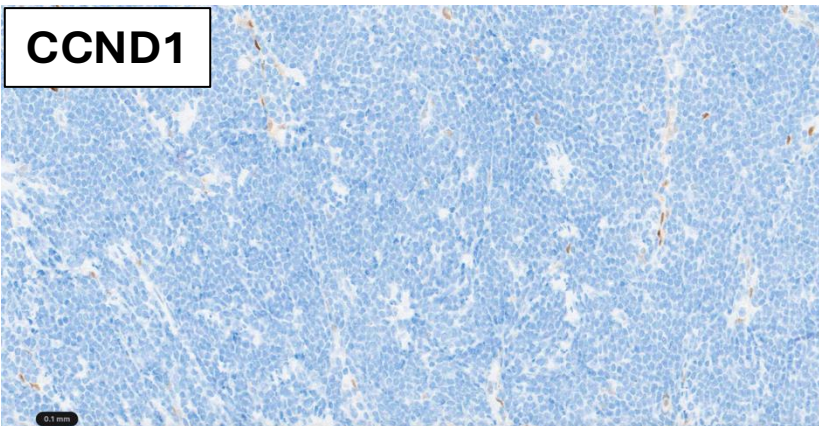
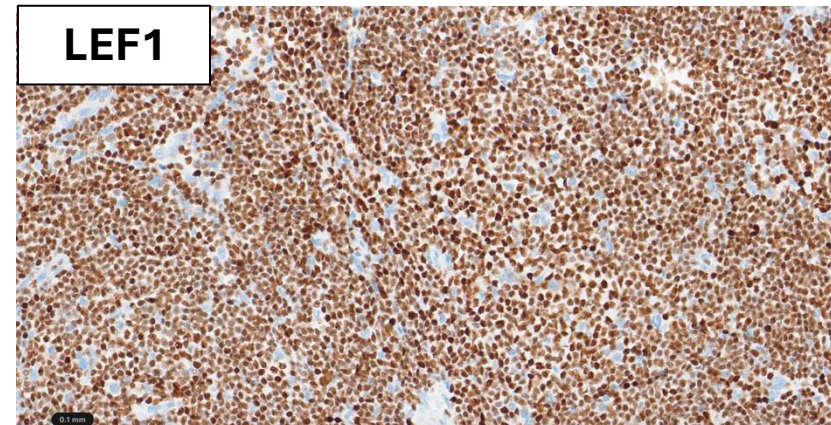
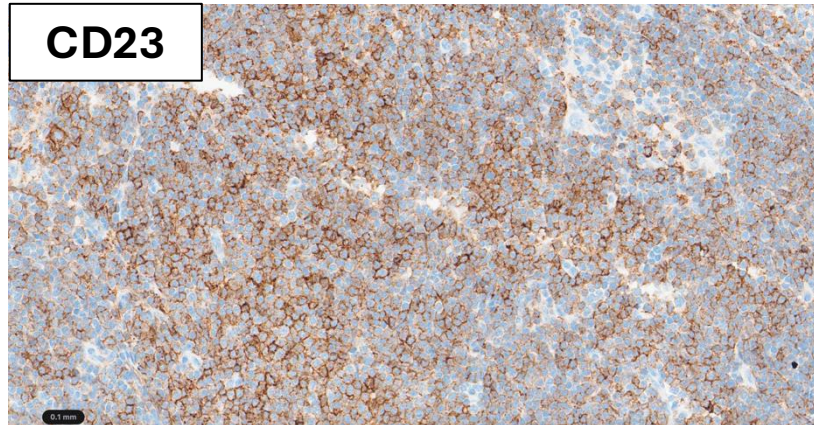
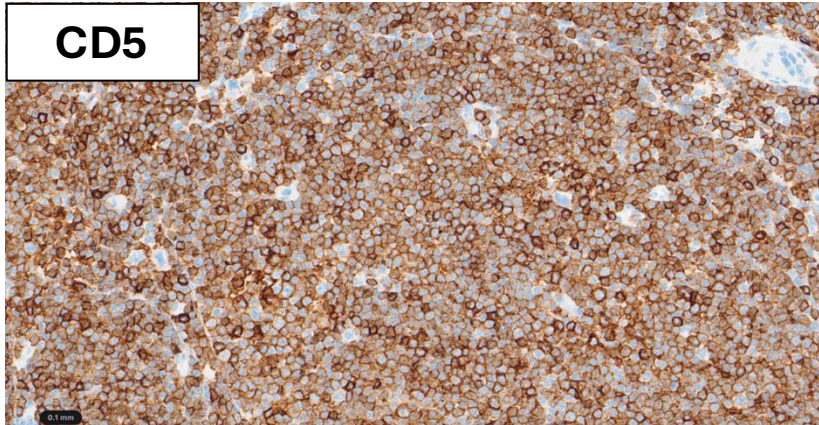
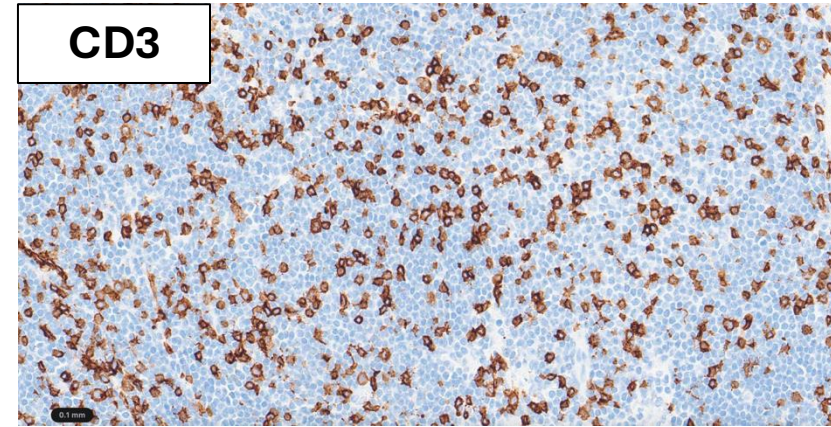
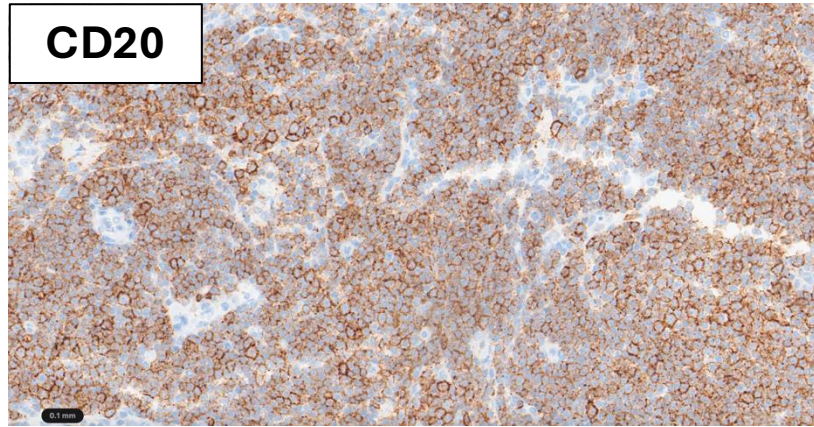
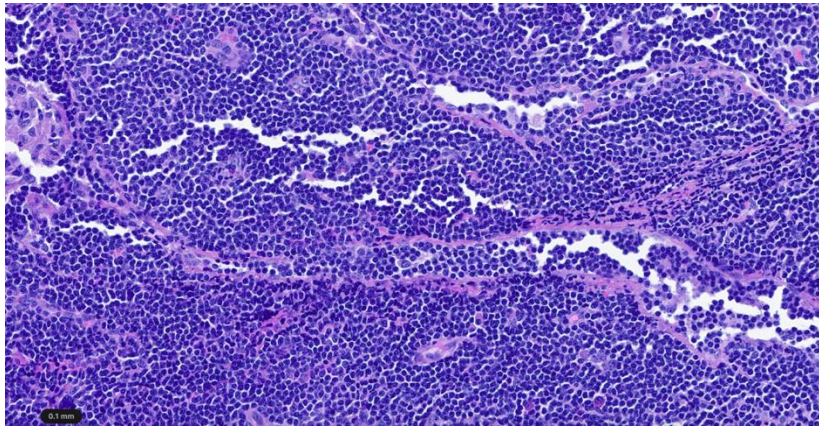
A few areas of more prominent uptake, in particular in the right tibia (SUV 15.4) with destruction of lateral cortex and in the medial condyle of the right humerus (SUV 4.6) .

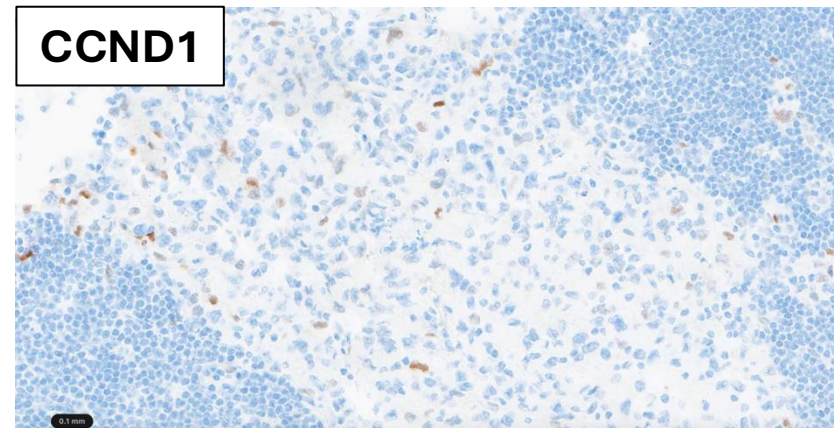
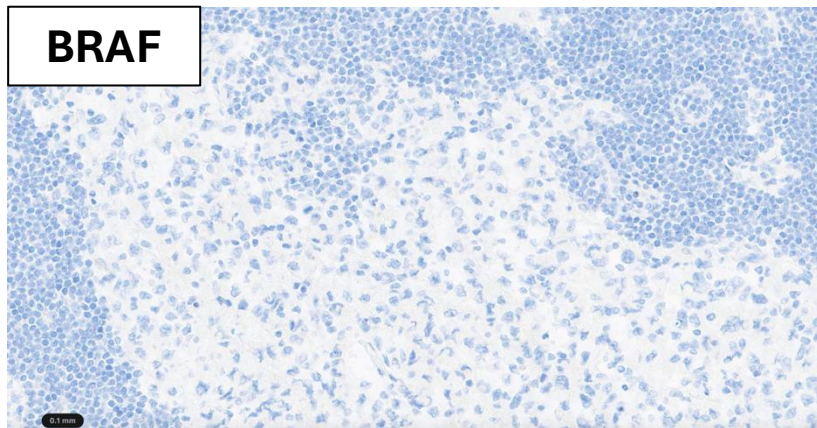
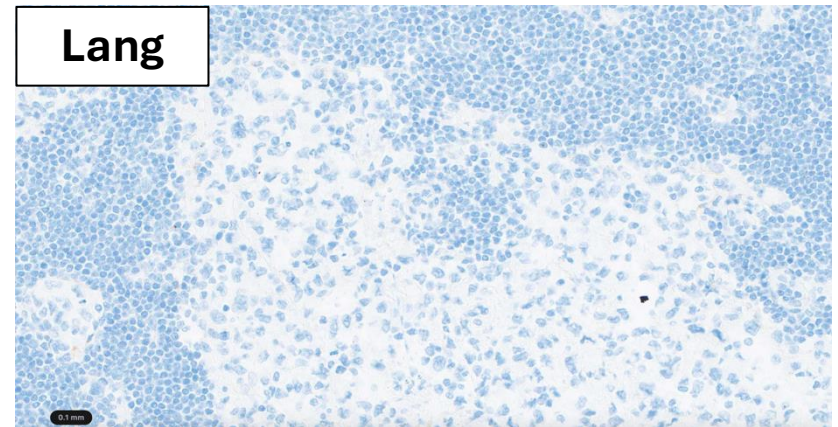
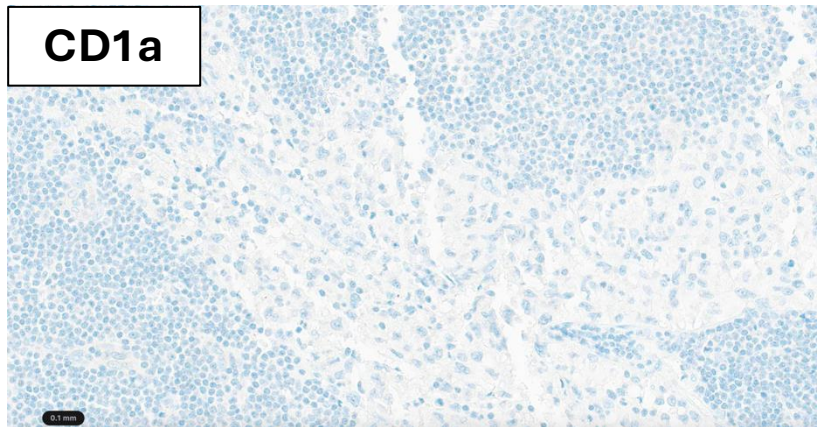
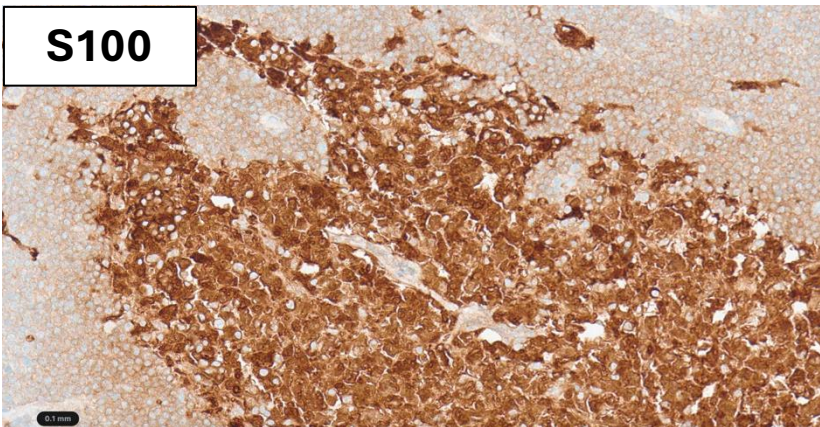
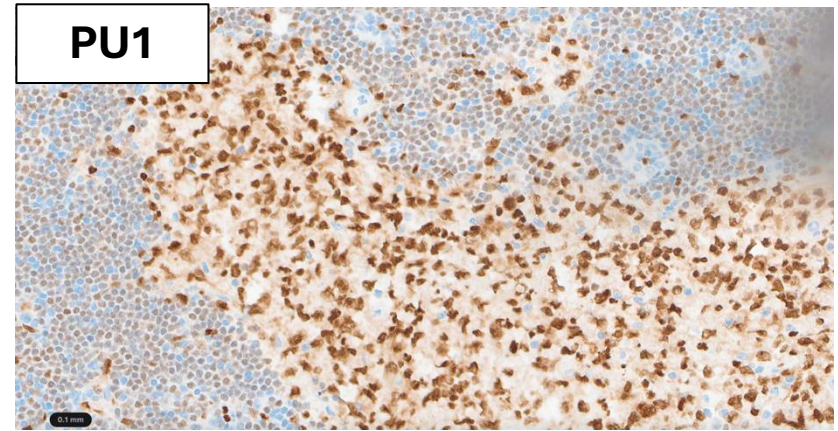
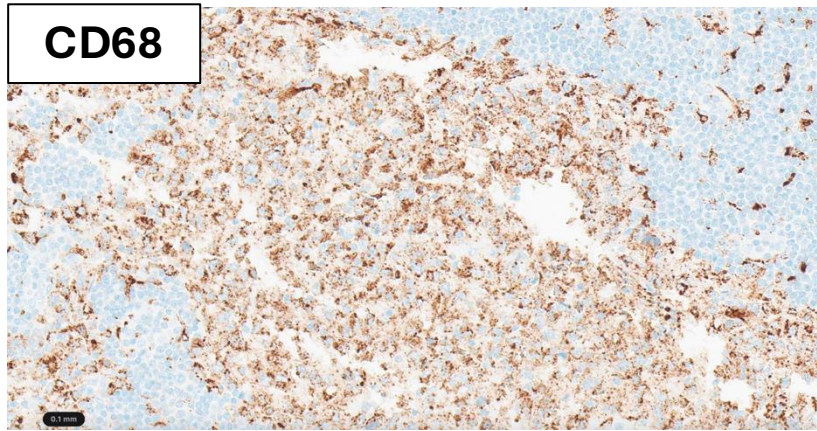
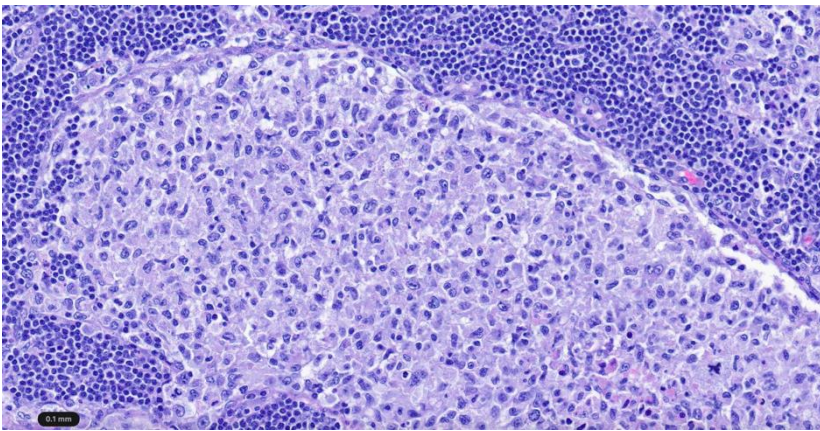
Since no FDG avid primary tumor is identified, these are possibly sites of transformed lymphoma; biopsy is recommended.

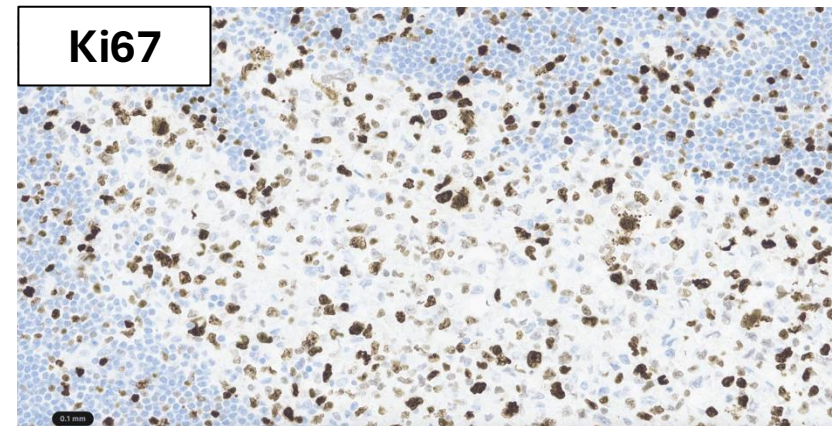
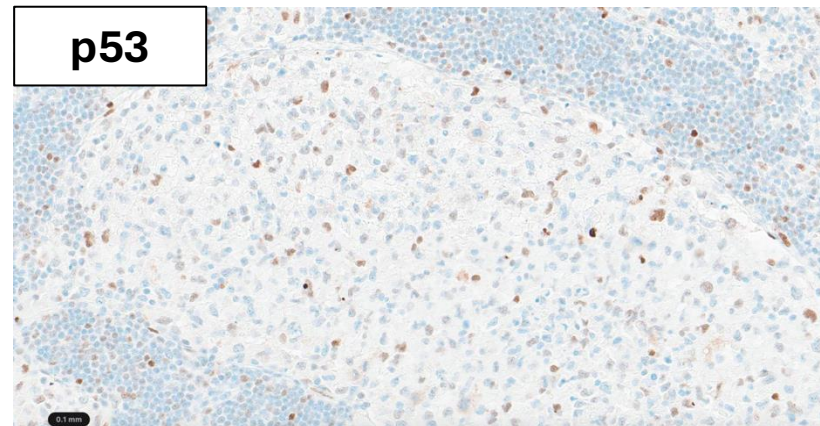
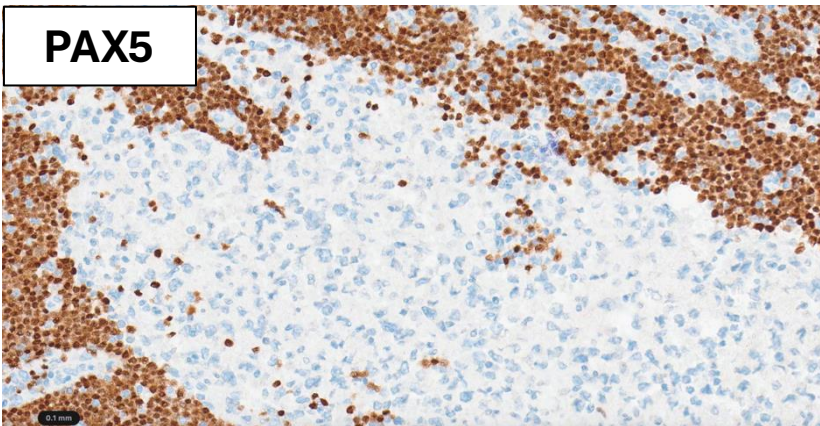
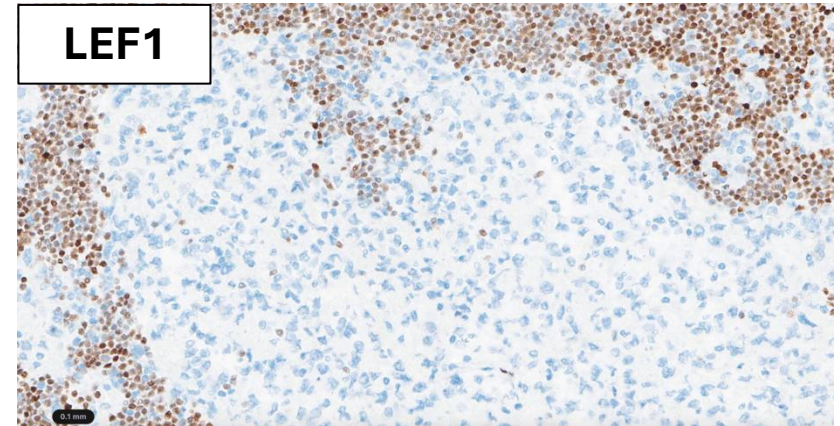
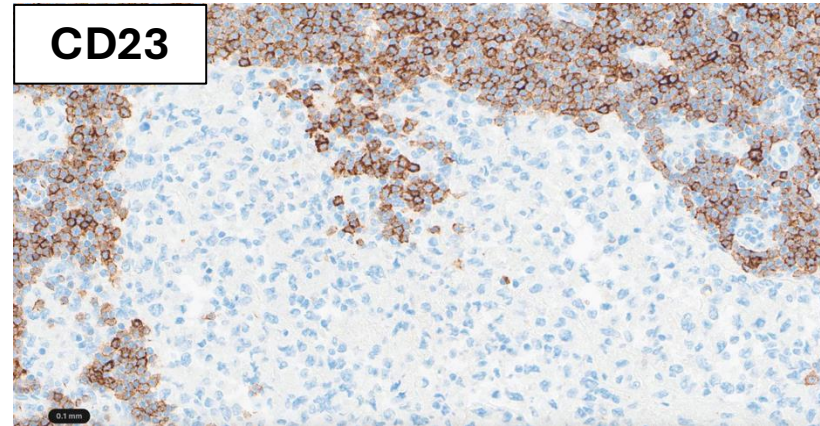
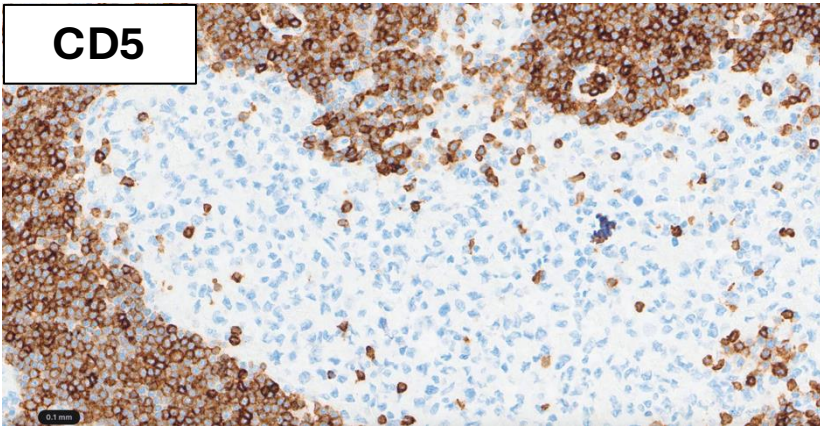
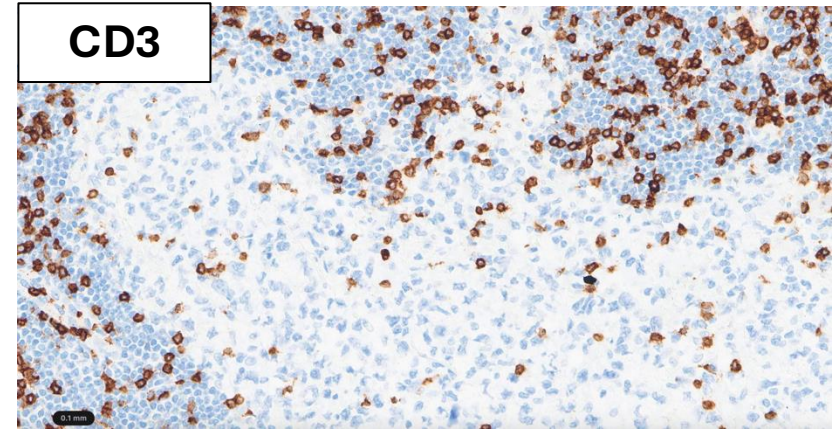
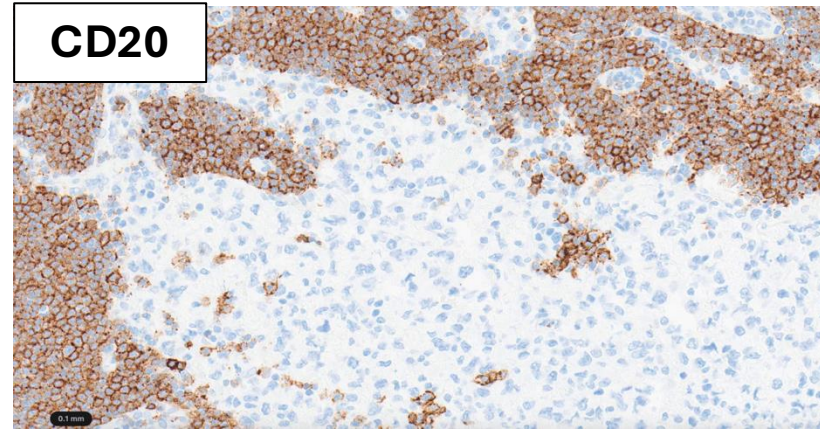
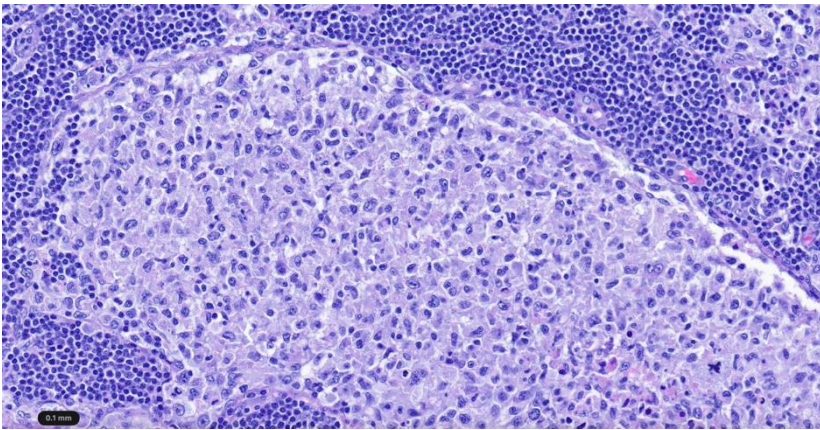
2. Splenomegaly without focal abnormal FDG uptake, consistent with CLL.



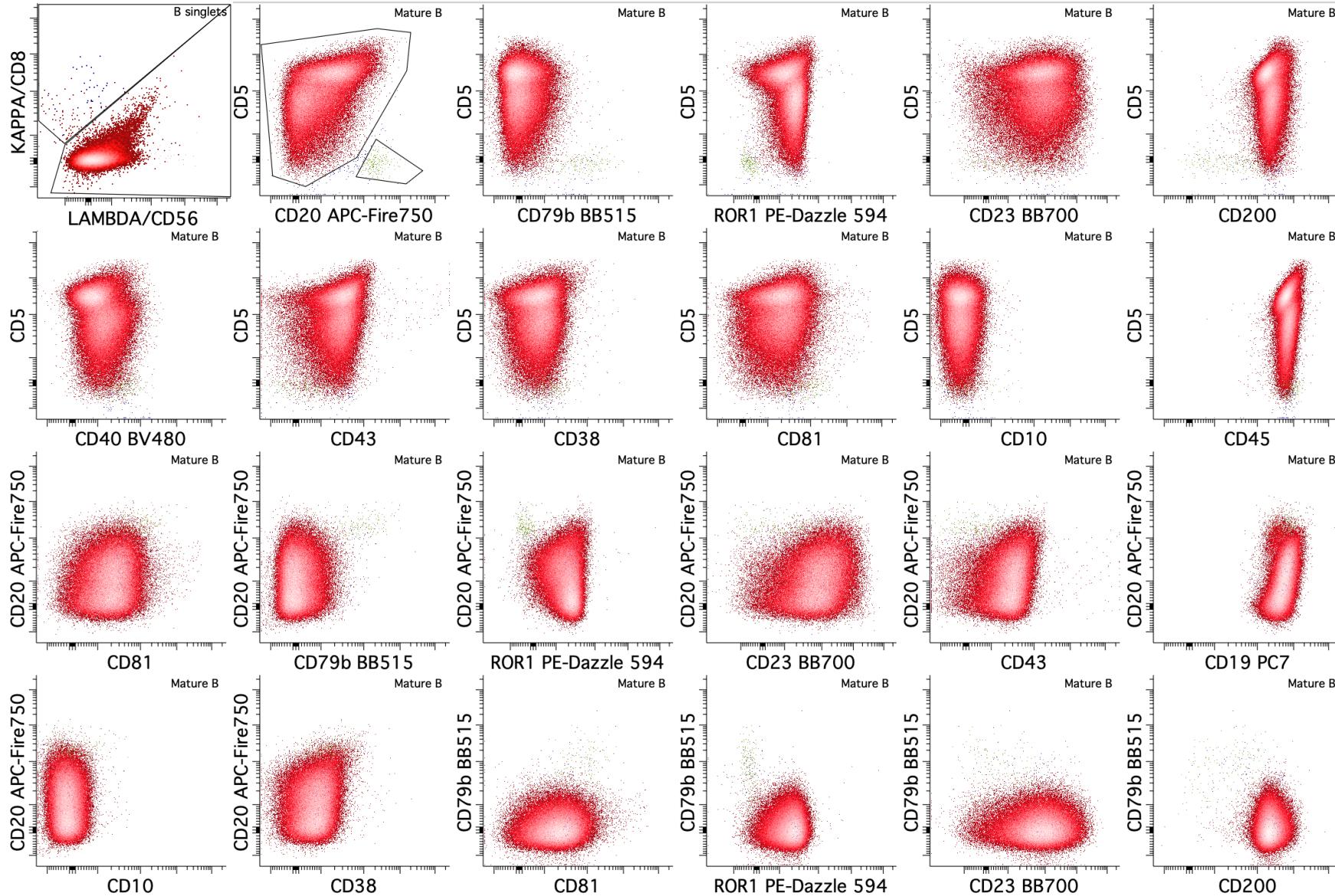








Flow Cytometry - Tibia



Abnormal B cell population detected (83.9% of WBC).

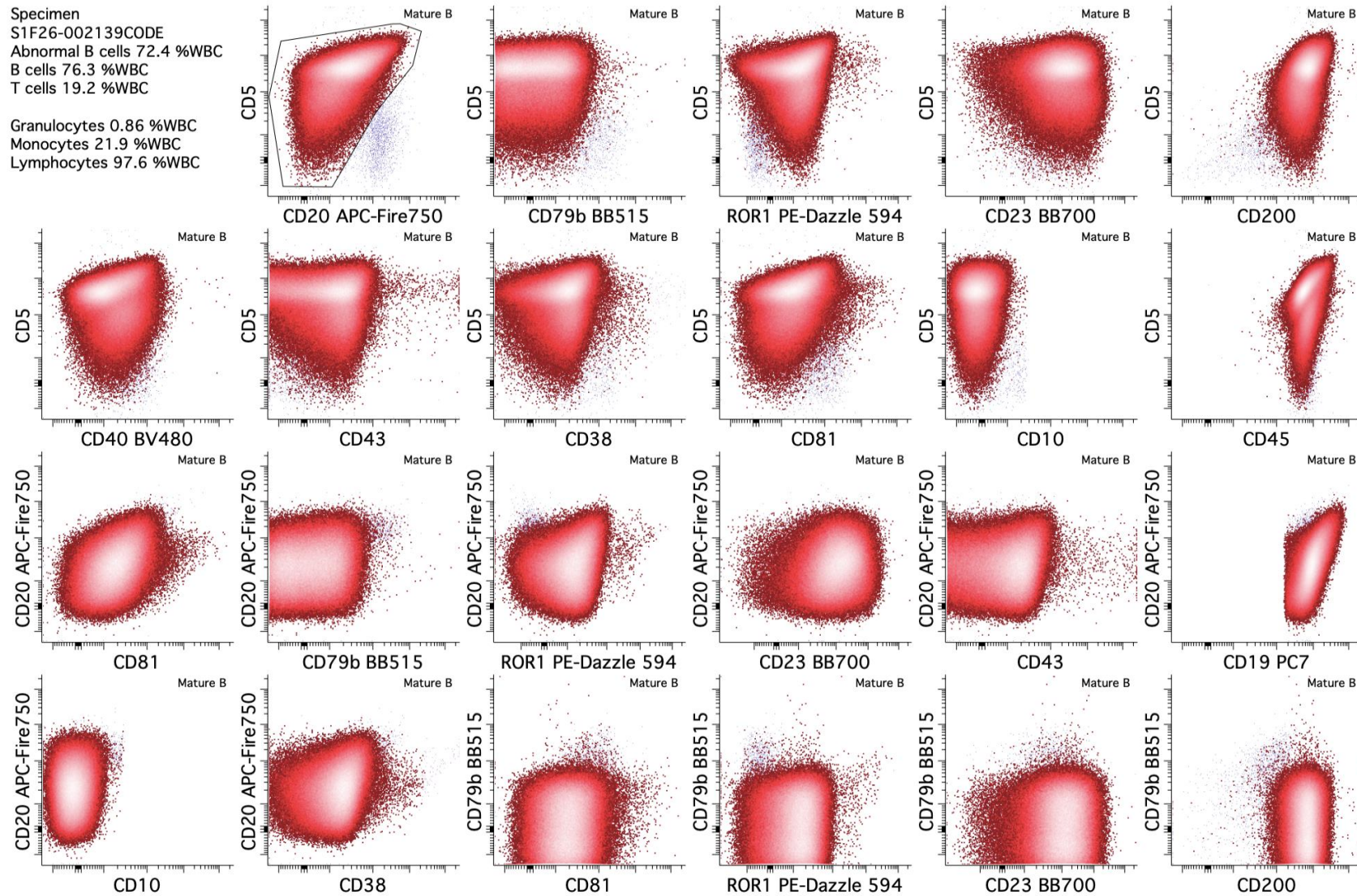
Immunophenotype:

- Abnormal: CD5 (positive), CD20 (dim), CD22 (negative to dim), CD23 (positive), CD43 (positive), CD79b (negative), CD81 (negative to dim), CD200 (bright), ROR1 (positive), surface Lambda (dim)
- Normal: CD19, CD40, CD45
- Negative: CD3, CD10, CD14, CD25, CD38, CD279 (PD1), surface Kappa

Flow Cytometry – Inguinal LN

Specimen
S1F26-002139CODE
Abnormal B cells 72.4 %WBC
B cells 76.3 %WBC
T cells 19.2 %WBC

Granulocytes 0.86 %WBC
Monocytes 21.9 %WBC
Lymphocytes 97.6 %WBC



Abnormal B cell population detected (59.9% of WBC).

Immunophenotype:

- Abnormal: CD5 (positive), CD20 (dim), CD22 (negative to dim), CD23 (positive), CD43 (positive), CD79b (negative), CD81 (negative to dim), CD200 (bright), ROR1 (positive), surface Lambda (dim)
- Normal: CD19, CD40, CD45
- Negative: CD3, CD10, CD14, CD25, CD38, CD279 (PD1), surface Kappa

MSK-IMPACT (Targeted NGS, DNA)

Tibia (lesion)

GENE	TYPE	ALTERATION	Allele Frequency
CHD2	Missense Mutation	D888Y (c.2662G>T)	3.49%
ATM	Missense Mutation	I2702R (c.8105T>G)	2.23%
KRAS	Missense Mutation	G12A (c.35G>C)	16.2%
DNMT3A	Splicing Mutation	X803_splice (c.2408+1G>A)	13.9%
DNMT3A	Missense Mutation	E733V (c.2198A>T)	12.12%
PIM1	Missense Mutation	H68N (c.202C>A)	13.31%
CIITA	Frameshift Insertion	A354Rfs*46 (c.1059dupC)	8.76%
TNFRSF14	Missense Mutation	G60S (c.178G>A)	29.54%
KLHL6	Missense Mutation	E568K (c.1702G>A)	11.58%
CD70	Missense Mutation	L167F (c.499C>T)	23.19%

Peripheral Blood

GENE	TYPE	ALTERATION	Allele Frequency
CHD2	Missense Mutation	D888Y (c.2662G>T)	47.77%
ATM	Missense Mutation	I2702R (c.8105T>G)	15.07%
ATM	Frameshift Deletion	F858Tfs*21 (c.2572_2575delTTTA)	1.4%

Molecular Studies – Peripheral Blood

SNP-ARRAY ANALYSIS

- Chromosome 11: low level loss in segment 11q21-q23.3, including the ATM gene
- Chromosome 13: loss in segment 13q14.2-q14.3, including the DLEU1, DLEU2, MIR16-1, and a portion of DLEU7 genes
- No evidence of 6q deletion by SNP-array analysis.
- No evidence of trisomy 12 by SNP-array analysis.
- No evidence of TP53 (17p13) gene deletion by SNP-array analysis.

NGS Clonality Testing – IGHV

- Clonal rearrangement involving the IG heavy chain gene is detected.
- IGHV mutation status is mutated

Clonal Sequence 1:

- IGHV Family: IGHV4-34
- IGHJ Family: IGHJ2
- Functionality: Productive
- Mutation Frequency: 6.48%
- Mutated

Clonal Sequence 2:

- IGHV Family: IGHV3-30
- IGHJ Family: IGHJ2
- Functionality: Unproductive
- Mutation Frequency: 11.82%
- Mutated

Final Diagnosis

- Atypical histiocytic proliferation most consistent with histiocytic sarcoma
- Involvement by chronic lymphocytic leukemia/small lymphocytic lymphoma

TABLE 1 Cases Submitted to Session 1: B-Cell Neoplasms to Histiocytic/Dendritic Cell Neoplasms

Case	Age, y	Sex	Submitter	Panel diagnosis	MAPK variations in HDCN	Clonal relatedness
23 ^a	62	M	Nichols	CLL/SLL with clonally related DLBCL variant Richter transformation and transdifferentiation to HS	<i>KRAS</i>	Shared trisomy 12 and <i>FGFR2</i> variations
168 ^a	48	F	Yabe	CLL/SLL with clonally related transdifferentiation to LCS	<i>BRAF</i>	Shared variations (<i>BRAF/NOTCH1</i>)
33	80	F	Lytle	CLL/SLL with clonally related transdifferentiation to HS	<i>MAP2K1</i>	Shared trisomy 12 and peaks of fragment analysis
52	54	M	Hollander	Low-grade B-cell lymphoma with clonally related transdifferentiation to HS	Not done	Shared peaks of fragment analysis
125	59	F	Wong	CLL/SLL with clonally related transdifferentiation to LCS	<i>BRAF</i>	Shared <i>ATM</i> deletion and peaks of fragment analysis
17	61	M	Ravindran	CLL/SLL with clonally related transdifferentiation to HS	<i>BRAF</i> and <i>KRAS</i>	Shared trisomy 12
106	61	F	LaValley	Composite CLL/SLL and HS with a history of DLBCL	<i>KRAS</i>	Shared trisomy 12 and other karyotypic abnormalities
28 ^a	56	F	Yu	MCL with clonally related transdifferentiation to HS	<i>BRAF</i> and <i>KRAS</i>	Shared <i>CCND1</i> rearrangements by FISH and peaks of fragment analysis
177 ^a	63	F	Grzywacz	Splenic MZL with clonally related transdifferentiation to HS	<i>KRAS</i>	Shared del7q by FISH and <i>MYD88</i> and <i>CREBBP</i> variations
247	53	F	Sakhdari	MZL with clonally related transdifferentiation to HS	Not detected	Shared peaks of fragment analysis, CDR3 sequence, and many variations
38 ^a	70	M	Rohr	FL, grade 1-2, with clonally related transdifferentiation into LCH, and later transformation to DLBCL	Not detected	Shared peaks of fragment analysis, <i>BCL-2</i> FISH, and many variations
54 ^a	70	F	Karrs	FL, grade 1-2, with clonally related transdifferentiation to HS	<i>KRAS</i>	Shared peaks of fragment analysis, <i>BCL-2</i> FISH, and many variations
34	31	F	Tashakori	FL, grade 1-2, with clonally related transdifferentiation to LCS	Not done	Shared <i>BCL-2</i> FISH
117	64	F	Merzianu	DLBCL with clonally related transdifferentiation to HS during CAR T-cell therapy	<i>KRAS</i>	Shared <i>BCL-2</i> FISH and many variations
239	65	F	Yuen	FL, grade 1-2, with clonally related transdifferentiation to LCS	<i>MAP2K1</i> and <i>PTEN</i>	Shared <i>BCL-2</i> FISH
45	58	F	Wotherspoon	FL, grade 1-2, with clonally related transdifferentiation to HS	Not detected	Shared <i>BCL-2</i> FISH and peaks of fragment analysis
74	60	F	Cook	FL, grade 1-2, with clonally related transdifferentiation to LCS and transformation to DLBCL	<i>MAP2K1</i>	Shared <i>BCL-2</i> FISH and many variations
86	50	M	Shet	Composite FL and HS	Not done	Unclear
178	57	F	Madrigal	FL, grade 1-2, with clonally related transdifferentiation to HS	<i>BRAF</i> and <i>PTEN</i>	Shared <i>BCL-2</i> FISH and peaks of fragment analysis
235	71	F	Hergott	FL, grade 1-2, with clonally related transdifferentiation to HS	Not detected	Shared <i>BCL-2</i> FISH
133 ^a	–	–	Zemmour	Clonally related LCH and BPDCN, with concurrent FL and HDCN, unclassifiable	Not detected	Shared variations between LCH and BPDCN but unclear on the nodal biopsy; many shared variations across tissue by whole-exome sequencing
139 ^a	56	F	Goodlad	FL, grade 1-2, with clonally related transdifferentiation to HDCN, unclassifiable	Not detected	Shared <i>BCL-2</i> by FISH
79	13	F	Aljudi	B-ALL to HS	Not done	Shared SNV array abnormalities
203	14	M	Fitzpatrick	B-ALL to HS	<i>NRAS</i>	Shared <i>DNMT3A</i> and <i>NRAS</i> variations

B-cell lineage neoplasms transdifferentiating into histiocytic/dendritic cell neoplasms: diversity, differentiation lineage, genomic alterations, and therapy

Report from the 2021 SH/EAHP Workshop

Wenbin Xiao, MD, PhD,^{1,•} Catalina Amador, MD,² James R. Cook, MD, PhD,^{3,•} Magdalena Czader, MD, PhD,⁴ Sandeep Dave, MD, MS,⁵ Ahmet Dogan, MD, PhD,¹ Amy Duffield, MD, PhD,¹ John Goodlad, MD,^{6,•} Reza Nejati, MD,⁷ German Ott, MD,⁸ Mariusz Wasik, MD^{7,•}

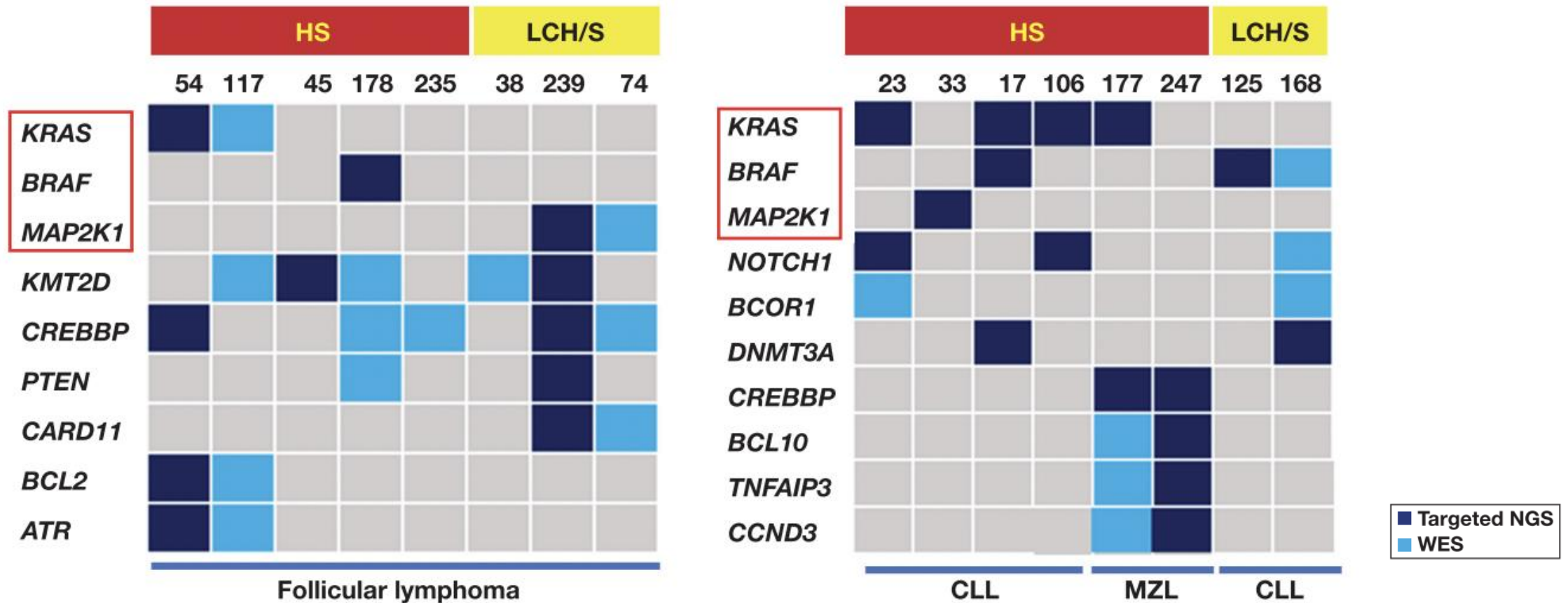
KEY POINTS

- Transdifferentiated histiocytic/dendritic cell neoplasms (HDCNs) show marked heterogeneity and overlapping immunophenotypic features.
- Variations in the MAPK pathway are enriched in HDCNs and may be therapeutic targets.
- Transdifferentiation into HDCNs occurs through both linear and divergent clonal evolution.

KEY WORDS

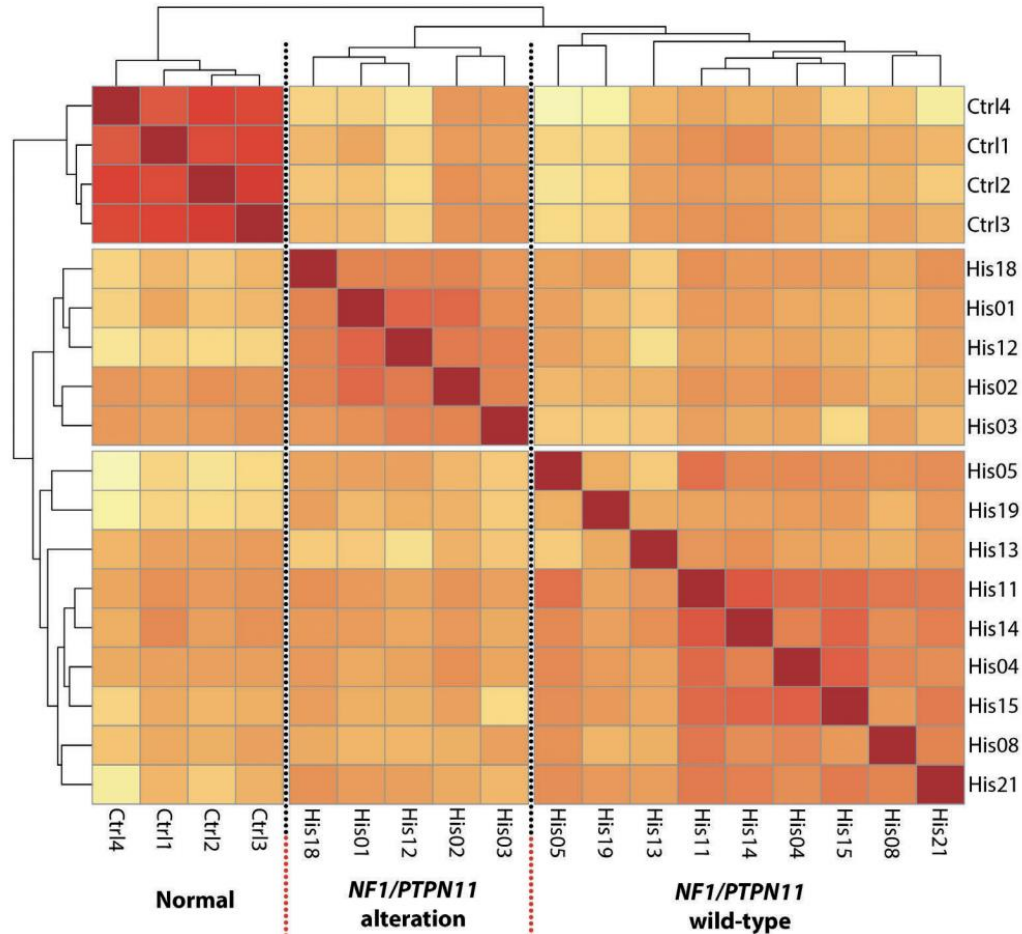
B-cell lymphoma; histiocytic/dendritic cell neoplasm; transdifferentiation

Mutational profile of histiocytic/dendritic cell neoplasms associated with B-cell lymphoma

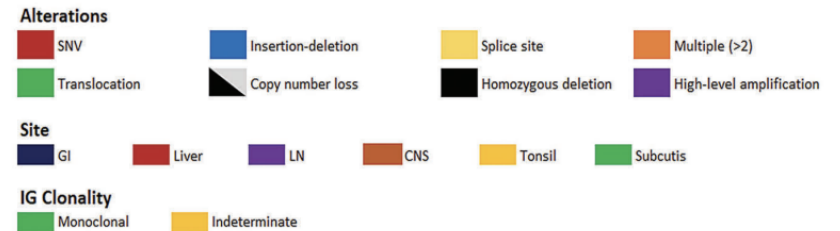
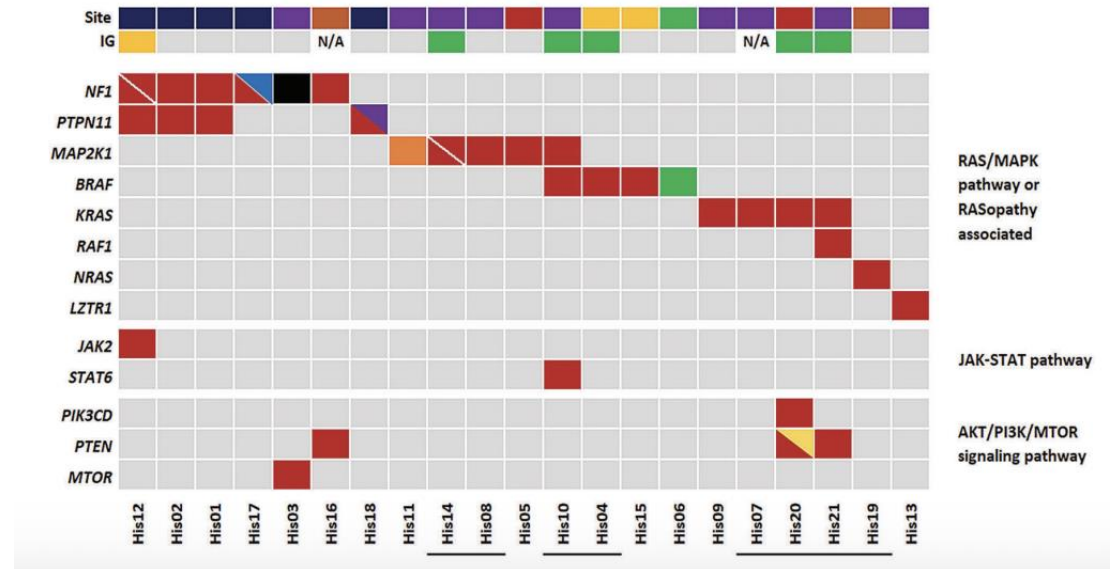


- B-cell neoplasms (FL > CLL > MZL > ALL) can give rise to HDCN.
- Clonal relatedness confirmed by shared mutations, IG rearrangements, FISH.
- MAPK pathway activation is the critical switch.

Two Molecular Subgroups of HS

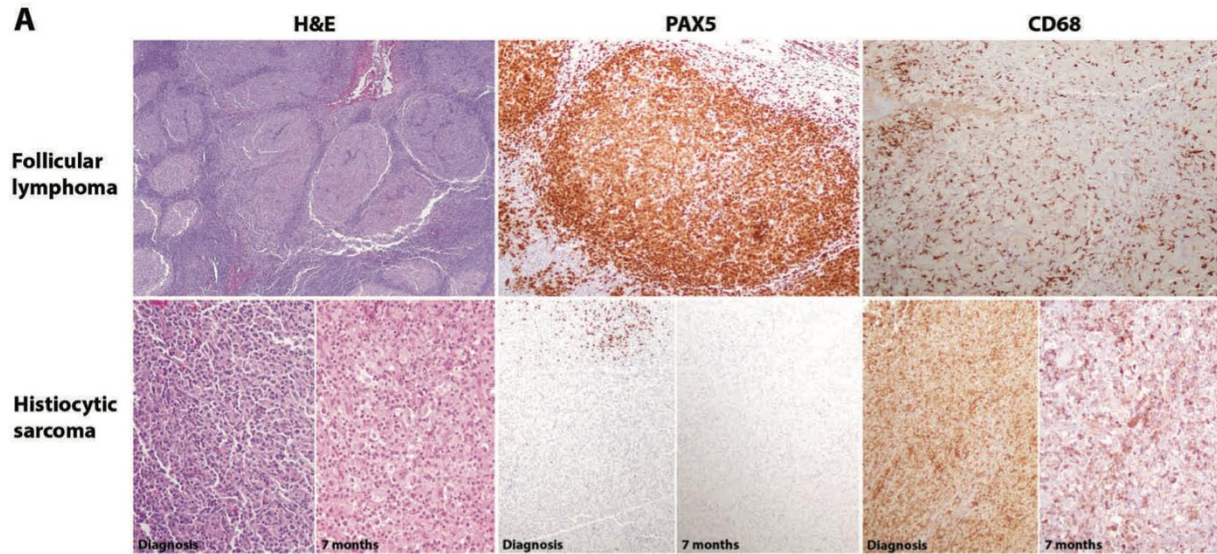


Selected molecular alterations in primary histiocytic sarcoma (pHS) cases

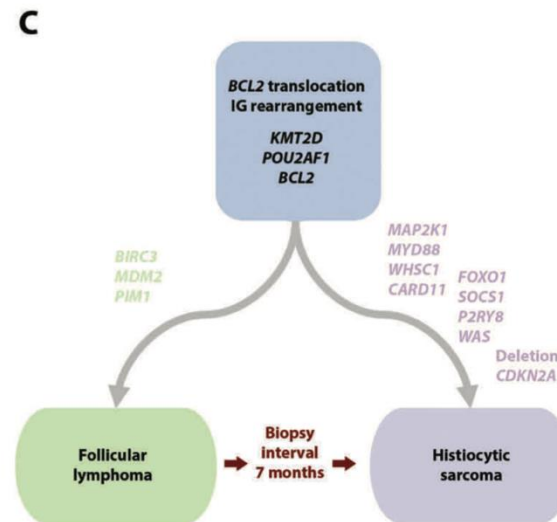
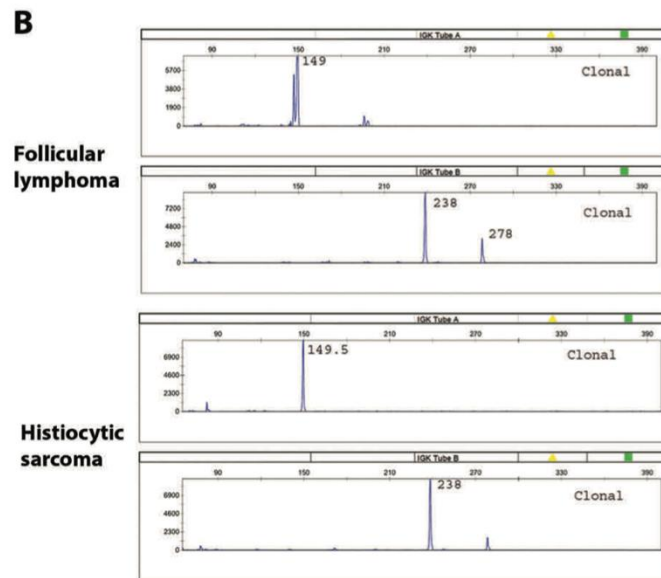


- NF1/PTPN11 (GI-tropic) vs. WT (B-cell-like).
- WT group often has clonal IG and B-lymphoma mutationseven in "primary" HS.

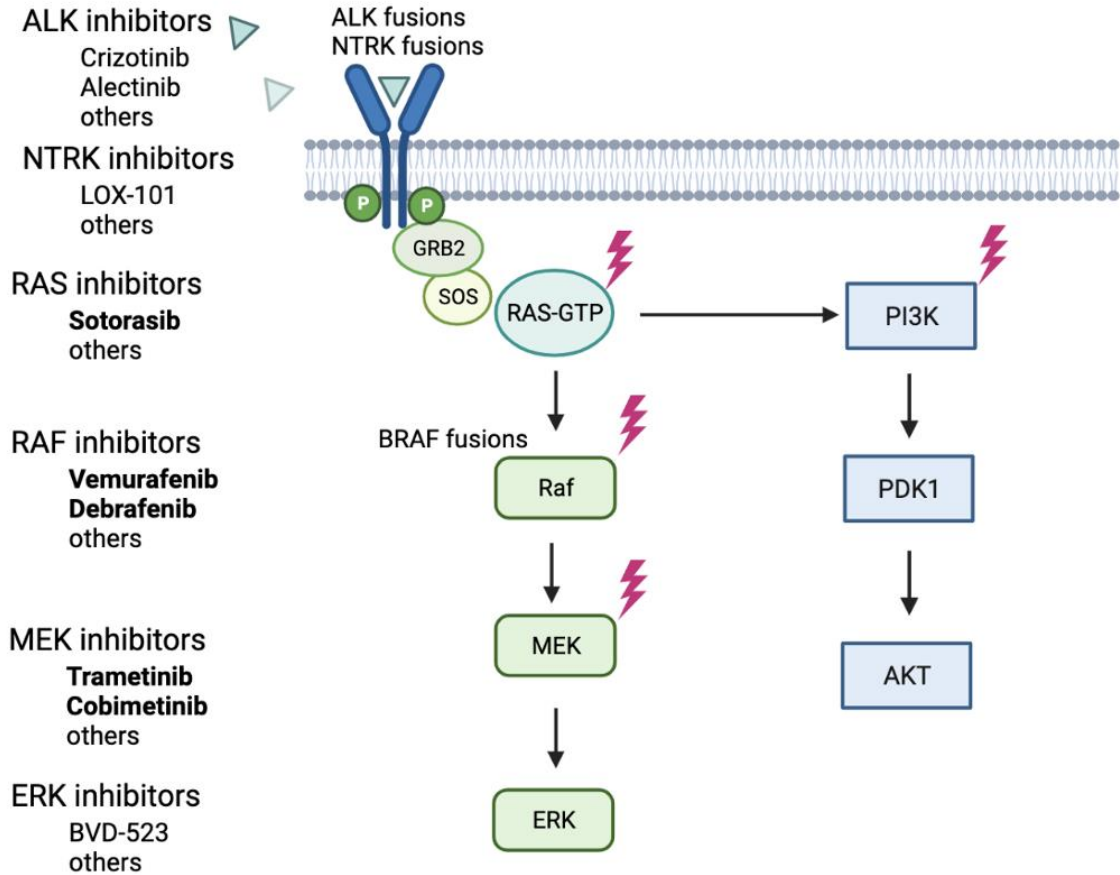
Secondary HS — Clonal Architecture



- Shared mutations = common precursor.
- RAS/MAPK acquired = transformation step.
- Clonal divergence in all matched pairs.



Treatment Implications



- MEK inhibitor cobimetinib → 89% ORR.
- BRAF inhibitor vemurafenib → 61.5% ORR.

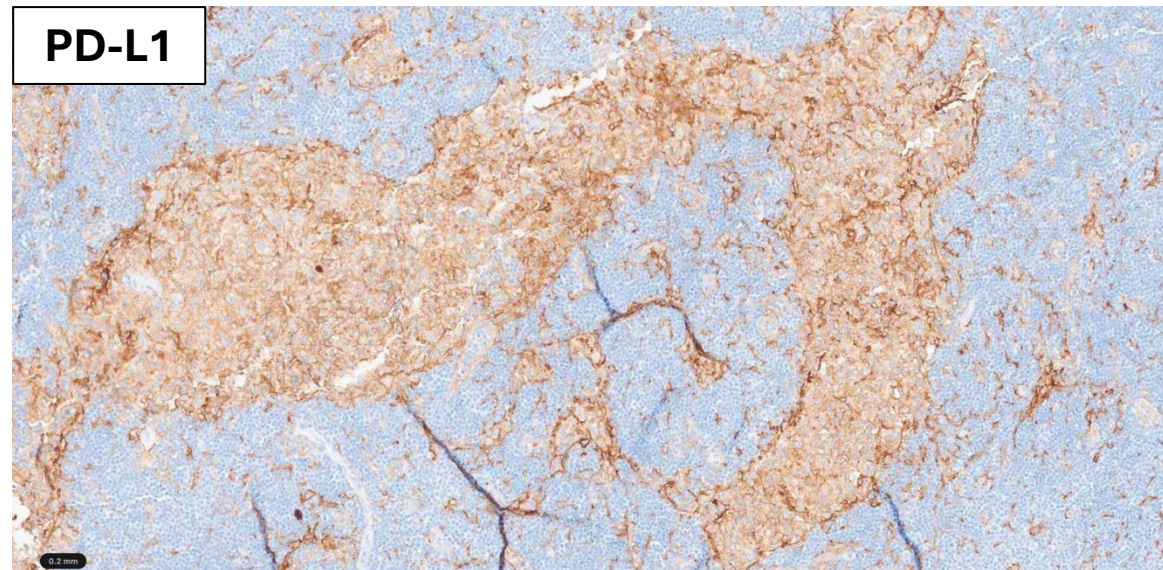
Table S6. Therapy and outcome of HDCNs after transdifferentiation

Case #	Diagnosis	treatment of HS/DC	live or death	follow up (months)
23	CLL/HS	Palliative Gemcitabine	died	<2
17	CLL/HS	Salvage chemo	died	<1
106	CLL/HS	RICE XRT	alive	?
177	MZL/HS	RCHOP for DLBCL	alive	12
38	FL/LCH	Mustard based chemo	died	4
117	DLBCL/HS	None	died	<1
239	FL/LCS	REPOCH	alive but progressed	?
45	FL/HS	RCHOP	alive	1
74	FL/LCS	XRT	died	24
139	FL/HS-DC	Local XRT	alive	?
178	FL/HS	R-ICE then Vemurafenib, Dabrafenib and trametinib	alive	24

Abbreviations: RICE, rituximab, fractionated ifosfamide, carboplatin, etoposide; REPOCH, rituximab, etoposide phosphate, prednisone, vincristine sulfate, cyclophosphamide, and doxorubicin hydrochloride; RCHOP, rituximab, cyclophosphamide, hydroxydaunorubicin hydrochloride, vincristine and prednisone; XRT, radiotherapy.

Treatment plan

- Cobimetinib, 20 mg/day (MEK inhibitor)
 - 21 days on, 7 days off
- Pembrolizumab
- Radiation tx



Acknowledgements

Memorial Sloan Kettering Cancer Center, Hematopathology Service

- Ahmet Dogan
- Megan So-Young Lim
- Oscar Lin
- Yanming Zhang
- Mikhail Roshal
- Filiz Sen
- Sergej Konoplev
- Scott Ely
- Michael Kluk
- Wenbin Xiao
- Mariko Yabe
- Pallavi Kanwar Galera
- Alexander Chan
- Kseniya Petrova-Drus
- Menglei Zhu
- Ying Liu
- Shenon Sethi
- Leonardo Boiocchi
- Kimon Argyropoulos
- Nicholas Barasch
- Ting Zhou
- Priyadarshini Kumar
- Qi Gao
- Jessica R. Chapman
- Ahmed Ayad
- Nicholas Haslett
- Chen Lossos

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