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Enhancing the Ability to Diagnose, Interpret and Apply Best Treatment Options for Cutaneous Lymphomas

CHALLENGING CASES OF CUTANEOUS LYMPHOMAS | #46

Molecularly-Annotated Case of CD30+ Gamma-Delta T-Cell Proliferation with Features of Primary Cutaneous Anaplastic Large Cell Lymphoma: a Wolf in Sheep's Clothing?

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Disclosures

I do not have any relevant financial relationships.

This presentation and/or comments will provide a balanced, non-promotional, and evidence-based approach to all diagnostic, therapeutic and/or research related content.





Clinical Presentation







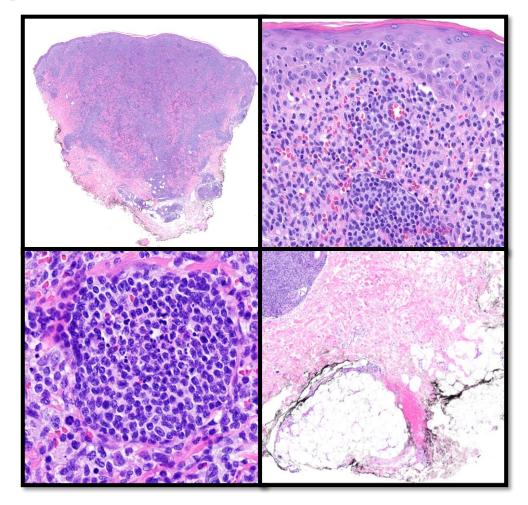
Clinical Presentation







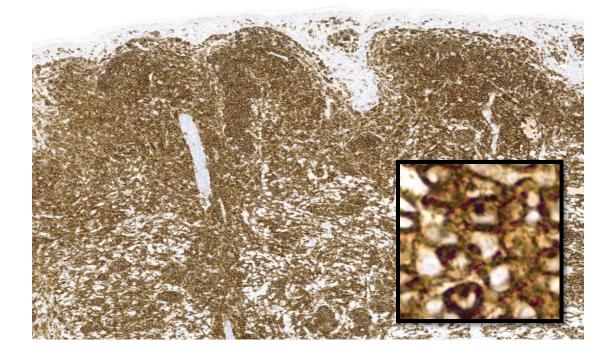
Biopsy Findings







Biopsy Findings







CD3

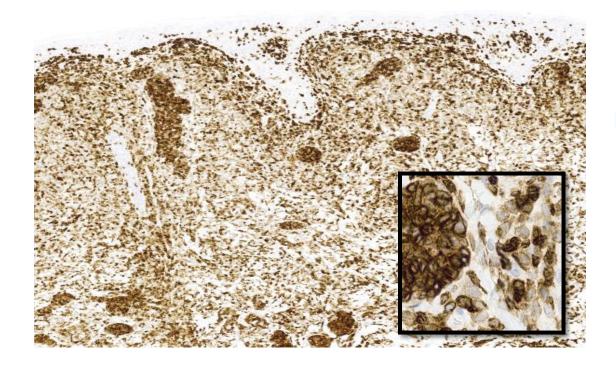


- ALK
- EBER
- CD56
- Granzyme
- **■** TIA-1
- CD4
- CD8
- CD2
- CD5
- PAX5





Biopsy Findings







CD3

■ TCR-Gamma



- ALK
- EBER
- CD56
- Granzyme
- TIA-1
- CD4
- CD8
- CD2
- CD5
- PAX5





Clinical Findings Favoring ALCL/LyP > pcGDTCL

- Indolent, non-ulcerating papules and nodules, spontaneous regression
- No mucosal involvement or lymphadenopathy by physical examination and PET CT
- No fever, sweats, or weight loss
- No peripheral blood lymphocytosis
- No cytopenias
- LDH normal





Histologic Findings Favoring ALCL/LyP > pcGDTCL

- No panniculitis-like fat involvement or angiodestructive-vasculitis-like features
- Small epidermotropic cells with larger cells in the dermis, more reminiscent of DUSP22-IRF4-rearranged CD30+LPD than epidermotropic pcGDTCL
- Widespread distention of dermal blood vessels
- No cytotoxic marker expression
- Loss of many T-cell markers
- Pan CD30 expression





Ancillary Test Findings

- FISH was negative for rearrangement involving *DUSP22-IRF4*
- Comprehensive genomic profiling revealed no reportable pathogenic mutations
- Comprehensive genomic profiling revealed no gene fusions
- Eight variants of unknown significance were detected
- Copy number profiling revealed aneuploidy





- Copy number profiling revealed aneuploidy
 - o Gain 1q

loss of 16q

loss of 19p13.3 (including *STK11*)

- Gain 1q is the most frequent alteration reported in pcGDTCL
- Copy number variations with a median of 4 arm-level events common in pcGDTCL
- STK11 (LKB1) is a tumor suppressor gene. Germline mutations seen in Peutz-Jeghers syndrome. Sporadic mutations present in malignancies including some renal, lung, and pancreatic cancers.





- Copy number profiling revealed aneuploidy
 - o Gain 1q

loss of 16q

loss of 19p13.3 (including *STK11*)

- Copy number variations are also seen in pcALCL, although often not arm-level
- Most frequently include gains involving 7q and 17, and losses involving 6q and 13





- Eight variants of unknown significance were detected
 - ADGRA2 EPHA7 ERBB2 LRP1B NOD1 RAF1 RICTOR WDR90
 - Many of the above genes have been found to be mutated in pcGDTCL
 - In general, however, pcGDTCL has been associated with reportable driver mutations involving MAPK, MYC, JAK/STAT and chromatin remodeling pathways, which often includes mutations in consensus cancer genes such as *CDKN2A*, *TP53*, or *ARID1A*
 - TERT gain of function mutations are also common





- Eight variants of unknown significance were detected
 - ADGRA2 EPHA7 ERBB2 LRP1B NOD1 RAF1 RICTOR WDR90
 - LyP and pcALCL have been associated with JAK/STAT pathway mutations
 - LRP1B (tumor suppressor) mutations have been identified in a subset of pcALCL, but it's among the most altered genes in cancer





- FISH was negative for rearrangement involving *DUSP22-IRF4*
- Comprehensive genomic profiling revealed no gene fusions
 - In addition to DUSP22-IRF4 rearrangements, fusions involving JAK/STAT pathway genes have been identified in CD30+LPD
 - Rearrangements involving TP63 and JAK2 have been described in pcGDTCL





ORIGINAL ARTICLE

ORIGINAL ARTICLE

TCR-γ Expression in Primary Cutaneous T-cell Lymphomas

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Lorenzo Cerroni, MD, PhD,||||| and Miguel Angel Piris, MD, PhD*§§

Abstract: Primary cutaneous νδ T-cell lymphomas (PCGD-TCLs) are considered a subgroup of aggressive cytotoxic T-cell lymphomas (CTCLs). We have taken advantage of a new, commercially available antibody that recognizes the T-cell receptor-y (TCR-y) subunit of the TCR in paraffin-embedded tissue. We have analyzed a series of 146 primary cutaneous T-cell lymphomas received for consultation or a second opinion in the CNIO Pathology Department. Cases were classified according to the World Health Organization 2008 classification as mycosis fungoides (MF; n = 96), PCGD-TCLs (n = 5), pagetoid reticulosis (n = 6), CD30+ primary cutaneous anaplastic large cell lymphomas (n = 5), primary cutaneous CD8+ aggressive epidermotropic CTCLs (n = 3), primary cutaneous CTCL. not otherwise specified (n = 4), and extranodal nasal-type NK/T-cell lymphomas primarily affecting the skin or subcutaneous tissue (n = 11). Sixteen cases of the newly named lym-

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phomatoid papulosis type D (LyP-D; n = 16) were also included. In those cases positive for TCR-7, a further panel of 13 antibodies was used for analysis, including TIA-1, granzyme B, and perforin. Clinical and follow-up data were recorded in all cases. Twelve cases (8.2%) were positive for TCR-y, including 5 PCGD-TCLs, 2 MFs, and 5 LvP-Ds, All 5 PCGD-TCL patients and 1 MF patient died of the disease, whereas the other MF patient and all those with LvP-D were alive. All cases expressed cytotoxic markers, were frequently CD3+/CD8+, and tended to lose CD5 and CD7 expressions. Eight of 12 and 5 of 11 cases were CD30+ and CD56+, respectively. Interestingly, 5/12 TCRγ-positive cases also expressed TCR-BF1. All cases analyzed were negative for Epstein-Barr virus-encoded RNA. In conclusion. TCR-v expression seems to be rare and is confined to cytotoxic primary cutaneous TCLs. Nevertheless, its expression is not exclusive to PCGD-TCLs, as TCR-γ protein can be found in other CTCLs. Moreover, its expression does not seem to be associated with bad prognosis by itself, as it can be found in cases with good and bad outcomes.

Key Words: TCR-γ, CTCL, prognosis (Am J Surg Pathol 2013;37:375–384)

P oripheral T-cell lymphomas are a heterogenous group of tumors comprising > 20 diseases. Although their molecular pathogenesis is not well understood, T-cell receptor (TCR) signaling is considered an essential component in lymphomagenesis. ¹⁻⁴

The $\dot{T}CR$ is a 'multimeric complex comprising 2 ligand-binding glycoproteins containing variable regions ($\alpha\beta$) or $\gamma\delta$ TCR heterodimers) that are expressed on the cell surface in association with 4 CD3 molecules. \dot{T} The $\alpha\beta$ TCR,CD3 complex is the most commonly represented among T cells and differs from that of the $\gamma\delta$ TCR/CD3 complex. $\gamma\delta$ T cells account for <55% of CD3 'cells in peripheral blood but constitute a more prevalent T-cell subset in certain tissues, including skin and intestine. Within lymphoid organs and tissues, they account for 15% of T cells in the spleen, 2% to 4% in lymph nodes, 1% in the thymus cortex, and 3% to 5% in the medulla. δ .

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Lymphomatoid Papulosis With T-cell Receptor–Gamma Delta Expression

A Clinicopathologic Case-series of 26 Patients of an Underrecognized Immunophenotypic Variant of Lymphomatoid Papulosis

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Abstract: Lymphomatoid papulosis (LyP) has several histopathologic presentations. LyP featuring gamma-delta (y8) T-cell receptor expression may masquerade as and may be misdiagnosed as aggressive cutaneous T-cell lymphoma, particularly primary cutaneous y8 T-cell lymphoma (PCGDTL) or y8

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StemLine Therapeutics, Innate Pharma, and Dren Bio, He is also

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mycosis fungoides. We performed a clinicopathologic analysis of the largest series of LvP featuring γδ T-cell expression. We identified 26 patients with a diagnosis of LyP with γδ T cells from our institutions, as well as through a comprehensive review of the literature, and characterized these cases. Most cases were treated with topical steroids or not treated at all. The majority of cases showed a CD4 CD8+ phenotype and featured at least one cytotoxic marker. Histopathologic features included an intraepidermal or dermal infiltrate with large cells and frequent angiotropism. One case was initially misdiagnosed as PCGDTL requiring further therapy. Our case series, the largest international cohort of yo T cell predominant LyP cases, confirms marked clinicopathologic heterogeneity that may contribute to misdiagnosis, reasserting the need to identify classic clinical features, CD30+ T-cell components, and markers of cytotoxicity when dealing with this differential diagnosis. A limitation of this study includes somewhat limited follow-up, histologic, and im munophenotypic information for some cases.

Key Words: CD30* T-cell lymphoproliferative disorders, cutaneous T-cell lymphoma, dermatopathology, gamma-delta T-cells, immunohistochemistry, lymphomatoli papulosis, mycosis fungoides, primary cutaneous gamma-delta T-cell lym-

(Am J Surg Pathol 2024;00:000-000)

Lymphomatoid papulosis (LyP) is a subacute CD30* LT-cell lymphoproliferative disorder that manifests clinically as recurrent self-remitting papules or small nodules with diverse histopathologic correlates. This diagnosis is important to achieve, because LyP is clinically troublesome, because specific treatment protocols are available upon recognition of the diagnosis, and because, if incorrectly microscopically evaluated, the findings can

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Histopathology



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T-cell receptor- δ expression and $\gamma\delta$ + T-cell infiltrates in primary cutaneous $\gamma\delta$ T-cell lymphoma and other cutaneous T-cell lymphoproliferative disorders

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T-cell receptor- δ expression and $\gamma\delta$ + T-cell infiltrates in primary cutaneous $\gamma\delta$ T-cell lymphoma and other cutaneous T-cell lymphoproliferative disorders

Aims: The diagnosis of cutaneous $\gamma\delta$ T-cell lymphoma (GDTCL) requires the identification of $\gamma\delta$ chains of the T-cell receptor (TCR). Our aim in this study was, by using a new monoclonal antibody (mAb) against TCR δ , to evaluate TCR δ expression in formalin-fixed parafilin-embedded (FPPE) skin tissue from TCR γ + cutaneous T-cell lymphoma (CTCL), and to assess TCR δ expression within a spectrum of other cutaneous lymphoroliferative disorders (CLPDs).

Methods and results: Twelve cases (10 patients) with TCRγ+ CTCL and 132 additional CLPD cases (127 patients) were examined, including mycosis fungoides (MF) (n = 60), cutaneous GDTCL (n = 15), subcutaneous panniculist-like T-cell lymphona (SPTCL) (n = 11), and CD30* lymphoproliferative disorder (LPD) (n = 24). Clone H-41 against TCRδ was used on a Leica Bond-3 automated stainer to label FFPE slides. H-41 immunostaining was graded as percentage infiltrate:

high (50–100%), moderate (10–49%), and low (0–9%). In TCRγ+ tumours, 12 of 12 (100%) patients showed TCRô expression comparable to TCRγ expression. No (0%) TCRγ+ cases were negative for TCRô. In all CLPDs, TCRô expression was as follows: GDTCL, 16 of 20 cases (14 of 15 patients) high, two moderate, and two low: CD30* LPD, one of 24 cases high, live moderate, and 21 low: CD30* LPD, one of 24 cases high, two moderate, and was 10 km moderate, and was provided to the moderate of the moderate o

Conclusions: mAb H-41 against TCR\u00e3 matches TCR\u00e3 in immunostaining FFPE tissues from GDTCL, supporting H-41 as a replacement for mAb \u00b73.20. TCR\u00e3 expression in our study suggests that the true occurrence of \u00b7\u00e3+ non-GDTCL CTCL/CLPD may be lower than suggested by the recent literature.

Keywords: classification, immunohistochemistry, immunological techniques, lymphoproliferative disorders, T-cell lymphoma





Reports

DIGINIA

TCR-γ Expression in Prinary

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From the "CNIO Lymphoma Group: Departments of ### Thermatology, Fundación Jiménez Diaz. Department. Hospital (Frematology, 1898) Hospital Universitario L. Spathology Department, Hospital de Torrejon de Arj. Department Hospital de Torrejon de Arj. Department Hospital de Torrejon de Arj. Department Hospital Universitario Get Getafe, Getafe, †##Immunolistoche, Universitario de Getafe, Getafe, †##Immunolistoche, Universitario Getafe, Getafe, †##Immunolistoche, Universitario Hospital Universitario Jana Canalogo, Getafe, †##Immunolistoche, Universitario Manquela del Valencia Jana Getafe, †##Immunolistoche, Universitario Manquela del Valencia Jana Getafe, †##Immunolistoche, Universitario Manquela del Valencia, †*Postratogo Department, Inceptial Virgen del Rocio, Sevilla; Sglanhology Department, Iversitario Manquela del Valencia, Santander, "Poetr single Department, Iversitario Manquela del Valencia, "Poetr single Department, Iversitario Manquela del Valencia del Valenci

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Take-home messages from case series

- Most LyP subtypes can show gamma-delta expression
- Clinical course and outcomes as expected for LyP
- Papules, variable nodules, variable ulceration
- Variable expression of CD2, CD3, CD5, CD7, CD4, CD8
- Variable expression of cytotoxic markers
- Variable DUSP22-IRF4 rearrangement
- EBER-negative



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sign mAb H-41 against TCRδ matches TCRγ mostaining FFPE tissues from GDTCL. supply 41 as a replacement for mAb γ3.20. TCRδ in our study suggests that the true occur- δ+ non-GDTCL CTCL/CLPD may be lower at the study the recent literature.

te niques, lymphoproliferative disorders, T-cell

Cityof Hope



CLINICAL AND LABORATORY INVESTIGATIONS

British Journal of Dermatology

$\gamma\delta$ T-cell-rich variants of pityriasis lichenoides and lymphomatoid papulosis: benign cutaneous disorders to be distinguished from aggressive cutaneous $\gamma\delta$ T-cell lymphomas

M.E. Martinez-Escala, M. Sidiropoulos, J. Deonizio, P. Gerami, M.E. Kadin and J. Guitart 1,3

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chart review of a relatively small sample.

What does this study add?

What's already known about this topic?

are associated with aggressive cutaneous lymphomas

Background T cells with a $\gamma\delta$ phenotype have been associated with aggressive lym-

phomas. Yet, inflammatory skin disorders and low-grade lymphoproliferative dis-

Objectives To review our experience and determine the clinical relevance of the γδ

T-cell phenotype in lymphomatoid papulosis (LyP) and pityriasis lichenoides

Methods A retrospective dermatopathology file review looking for LyP and PL

characterized by a $\gamma\delta$ T-cell phenotype was performed. Clinical manifestations

Results Six of 16 cases of LyP and four of 23 cases diagnosed as PL during a

5-year period (2009-14) were identified. The median follow-up for the whole

group was 16 months (range 3-64), showing an indolent clinical course in all

Conclusions The detection of a predominantly γδ T-cell phenotype in papular

lymphoid-rich infiltrates in the absence of other lesions is not associated with a

clinically aggressive course. $\gamma\delta$ T-cell-rich variants of LyP and PL may reflect a

spectrum of related conditions. This is a single academic centre retrospective

Most lymphoid-rich inflammatory skin conditions are composed of T cells express

ing $\alpha\beta$ T-cell receptors (TCRs) and, in general, TCR $\gamma\delta$ T-cell infiltrates of the skir

Regardless of the presence or absence of lymphoid atypia, a subset of γδ T-cell-rich

cutaneous lymphoid infiltrates are self-limited and indolent. Pityriasis lichenoid

and lymphomatoid papulosis may present as a predominant $\gamma\delta$ T-cell infiltrate.

orders have rarely been described with a predominant $\gamma\delta$ T-cell infiltrate.

and course, histological features and molecular data were analyzed.

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Summarv

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Cutaneous Pathology

Atypical cutaneous $\gamma \delta$ T cell proliferation with morphologic features of lymphoma but with clinical features and course of PLEVA or lymphomatoid papulosis

Reactive lymphoid infiltrates of the skin composed predominantly of gamma-delta (γδ) T cells are not well described in the literature. Herein we report a case of an otherwise healthy 4-year-old male who presented with a waxing and waning papular rash characterized by small, discrete crusted papules spread across his trunk, face and extremities. Clinical evaluation revealed no evidence of systemic disease. Microscopic examination revealed a dermal, perivascular infiltrate of highly atvoical lymphocytes with a γδ T cell phenotype, worrisome for primary cutaneous γδ T cell lymphoma. The clinical course, however, was that of a reactive condition and prompted consideration of a diagnosis of pityriasis lichenoides et varioliformis acuta (PLEVA) and lymphomatoid papulosis (LyP). In many ways, this case defies current classification schemes and seems to expand the spectrum of reactive γδ T cell infiltrates of the skin.

Keywords: T lymphocytes, atypical features, CUTANEOUS LYMPHOMAS, hematopathology,

King RL, Yan AC, Sekiguchi DR, Choi JK. Atypical cutaneous γδ T cell proliferation with morphologic features of lymphoma but with clinical features and course of PLEVA or lymphomatoid

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Journal of **Cutaneous Pathology**

Indolent course of cutaneous gamma-delta T-cell lymphoma

Cutaneous gamma-delta T-cell lymphoma (y&TCL) is a rare malignancy that typically displays an aggressive clinical course. We present an unusual case of a 57-year-old woman with a 3-year history of lower extremity nodules. Histopathologic, immunophenotypic and molecular genetic studies revealed a clonal, predominantly pannicular gamma-delta T-cell infiltrate, leading to a diagnosis of cutaneous γδTCL. The clinical course was characterized by rapid improvement within months of starting systemic corticosteroids, with relapse in ulcerations but no new lesions more than 3 years after onset of disease Our case and seven previously reported patients with indolent and relatively localized cutaneous γδTCL provide evidence that not all cases of this entity carry a poor prognosis. This indolent subset adds complexity to treatment of cutaneous γδTCL.

Keywords: connective tissue disease panniculitis, gamma-delta T-cell lymphoma, panniculitic lymphoma

Endly DC, Weenig RH, Peters MS, Viswanatha DS, Comfere NI. Indolent course of cutaneous gamma-delta T-cell lymphoma. J Cutan Pathol 2013; 40: 896-902. © 2013 John Wiley & Sons A/S. Published by John Wiley & Sons Ltd

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Cutaneous gamma-delta T-cell lymphoma (γδTCL) represents less than 1% of primary cutaneous lymphomas and currently is a provisional entity in the subset of peripheral T-cell lymphomas (TCLs), unspecified in the current World Health Organization-European Organization for Research and Treatment of Cancer (WHO-EORTC) classification of lymphoid neoplasms.1 Composed of a clonal population of mature gamma-delta T cells with a cytolytic phenotype, γδTCL was previously designated as a subcutaneous panniculitis-like T-cell lymphoma (SPTCL) subset. The new distinction was made after γδTCL was recognized to have a distinctively adverse prognosis with 11% 5-year survival rate compared with 82% for SPTCL 2 In contrast with the lymphocytes of SPTCL that express CD8 but not CD56 or CD4, νδTCL is characterized by expression of CD56 but not CD8 and CD4. Cytolytic protein expression is present in both variants of subcutaneous TCL.

Lupus erythematosus (LE) panniculitis continues to represent a clinicopathologic challenge with

respect to its distinction from the two subcutaneous T-cell lymphoma variants. 3,4 Shared presentations of LE panniculitis and subcutaneous T-cell lymphomas include erythematous to violaceous dermal or subcutaneous nodules that may ulcerate, and histopathologic features of interface changes, hyaline fat necrosis and lobular pannicular infiltration by cytolytic lymphocytes that exhibit adipocyte rimming, karyorrhexis and atypia.5 We present a case of cutaneous y&TCL that followed an indolent clinical course and review the literature regarding similar cases.

A 57-year-old woman was referred to our dermatology department with a 3-year history of tender, ulcerating nodules of the legs associated with swelling of the ankles. Past medical history included chronic lower extremity ulcers of unknown etiology dating back to 1997, which healed after skin grafting; persistent patches of 'parapsoriasis'





CLINICAL AND LABORATORY INVESTIGATIONS

BJD British Journal of Dermatology

$\gamma\delta$ T-cell-rich variants of pityriasis lichenoides and lymphomatoid papulosis: benign cutaneous disorders to be distinguished from aggressive cutaneous $\gamma\delta$ T-cell lymphomas

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Summary

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Conflicts of interest

DOI 10.1111/bjd.13364

Background T cells with a $\gamma\delta$ phenotype have been associated with aggressive lymphomas. Yet, inflammatory skin disorders and low-grade lymphoproliferative disorders have rarely been described with a predominant $\gamma\delta$ T-cell infiltrate. Objectives To review our experience and determine the clinical relevance of the $\gamma\delta$ T-cell phenotype in lymphomatoid papulosis (LyP) and pityriasis lichenoides

Methods A retrospective dermatopathology file review looking for LyP and PL characterized by a $\gamma\delta$ T-cell phenotype was performed. Clinical manifestations and course, histological features and molecular data were analyzed.

Results Six of 16 cases of LyP and four of 23 cases diagnosed as PL during a 5-year period (2009–14) were identified. The median follow-up for the whole group was 16 months (range 3–64), showing an indolent clinical course in all cases.

Conclusions The detection of a predominantly $\gamma\delta$ T-cell phenotype in papular lymphoid-rich infiltrates in the absence of other lesions is not associated with a clinically aggressive course, $\gamma\delta$ T-cell-rich variants of LyP and PL may reflect a spectrum of related conditions. This is a single academic centre retrospective chart review of a relatively small sample.

What's already known about this topic?

 Most lymphoid-rich inflammatory skin conditions are composed of T cells expressing αβ T-cell receptors (TCRs) and, in general, TCR γδ T-cell infiltrates of the skin are associated with aggressive cutaneous lymphomas.

What does this study add?

 Regardless of the presence or absence of lymphoid atypia, a subset of y\u00f3 T-cell-rich cutaneous lymphoid infiltrates are self-limited and indolent. Pityriasis lichenoides and lymphomatoid papulosis may present as a predominant y\u00f3 T-cell infiltrate. J Cutan Pathol 2015: 42: 1012–1017 doi: 10.1111/cup. 12601 John Wiley & Sons. Printed in Singapore © 2015 John Wiley & Sons A/S.
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J Cutan Pathol 2013: 40: 896–902 dai: 10.1111/cup.12091 John Wiley & Sons. Printed in Singapore

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Cutaneous Pathology

Patient	Sex/age						Length of follow-up	
no.	(years)	Ethnicity	Clinical presentation	Distribution	Treatment	Status	(months)	Diagnosis
1	M/26	U	Crusted papules	Generalized	U	U	U	PL
2	F/3	W	Crusted papules	Generalized	Oral ATB	AWD	14-1	PL
3	F/33	W	Crusted and eroded papules	Generalized	Topical steroids	AWOD	2.3	PL
4	M/13	W	Crusted papules hypo- and hyperpigmentation	Trunk, upper and lower extremities	Oral ATB, NB-UVB	AWD	29·1	PL
5	F/53	W	Papules, overlap with eczematous patches	Trunk and lower extremities	NB-UVB	AWOD	62-6	LyP
6	M/57	W	Crops of papules	Upper and lower extremities	None	AWD	33.0	LyP
7	M/48	W	Papules	Upper and lower extremities	Topical steroids	AWOD	13.6	LyP
8	M/71	U	Single lesion	Left lower extremity	Surgical excision	AWOD	22.0	LyP
9	F/30	W	Papules, some eroded	Trunk, upper and lower extremities	Oral ATB	AWOD	14.9	LyP
10	F/44	U	Solitary papule	Neck	Surgical excision	AWOD	1.0	LyP

M, male; F, female; U, unknown; W, white; ATB, antibiotics; NB-UVB, narrowband ultraviolet B; AWD, alive with disease; AWOD, alive without disease; PL, pityriasis lichenoides; LyP, lymphomatoid papulosis.





Am J Dermatopathol • Volume 44, Number 10, October 2022

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The First Case of Gamma-Delta Primary **Cutaneous Anaplastic** Large Cell Lymphoma?

We have read with much interest the report by Meawad et al1 on a case of a primary cutaneous gamma-delta T-cell lymphoma (PCGD-TCL) mimicking ALK-1 negative anaplastic large cell lymphoma (ALCL).

It caught our attention that the tumor cells showed many features which are characteristic of ALCL such as hallmark cells, anaplasia, pleomorphism, and CD30 positivity. Lack of ALK expression is also typical of primary cutaneous ALCL.2 Nevertheless, the authors made the diagnosis of PCGD-TCL based on the panniculitic pattern, the rimming, and mainly on the immunostaining for T-cell receptor (TCR)-delta.

We would like to remark that prominent involvement of the subcutaneous fat is identified in some cases of primary cutaneous ALCL.3 We are also aware that, to the best of our knowledge, no cases of gamma-delta ALCL have been published so far, although Pulitzer et al4 reported one case of a "CD30 lymphoproliferative disorder

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addition, there are cases of gamma-delta lymphomatoid papulosis, which informed. is another CD30+ lymphoproliferative disorder closely related to primary cutaneous ALCL.5

The current case was negative for CD4 and CD8. This is not a handicap for She man pignicition a claim keliasus. Am

CD4 and CD8. This is not a handicap for
Buonaccorsi JN, Prieto VG, Torres-Cabal
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C, et al. Diagnostic utility and comparative
AL CLI because only one-third cases. ALCL, because only one-third of cases are positive for CD4, and negativity for CD4 and CD8 is seen in almost half of the cases.3 In addition, most of PCGD-TCL are CD30 negative, and when positive, they usually show low-medium expression of the marker.6

Primary cutaneous ALCI, usually presents with eosinophils.3 which are presents with eosinophils, which are rarely seen in PCGD-TCL. Although it Biomedical Research of A Coruña (INIBIC), is not sharply appreciated, Figure 2b seems to show some eosinophils accompanying the tumoral cells.

Regarding the clinical presentation, lesions of PCGD-TCL often are ulcerated, involvement of mucosae is frequent, B symptoms occur in most cases, and lactate dehydrogenase is normally elevated. In addition, involvement of soft tissue and lymph nodes by the PCGD-TCL are usually present at the time of diagnosis.7 None of these features were reported in the case report of the authors.

Multiple lesions are not uncommon in CD30 positive lymphomas: Although most frequent in lymphomatoid papulosis, such presentation has also been described in some cases of primary cutaneous ALCL,3 so much so that the latter can sometimes clinically mimic the former

Cases of PCGD-TCL have been described after treatment with etanercept, but this drug has also been associated with primary cutaneous ALCL.8

We believe that the current case would better fit into the diagnosis of TCR delta positive primary cutaneous ALCL. Therefore, we agree that this case is extraordinary, as in our opinion, it represents the first well-documented case of gamma-delta primary cutaneous

The good response to treatment is 8. De Angelis F, Di Rocco A, Minotti C, et al also more in consonance with primary cutaneous ALCL. However, long-term Ethics approval: Approved.

Copyright © 2022 Wolters Kluwer Health, Inc. All follow-up will probably give us more reliable information on the prognosis of

8. de Carvalho N, Fametani F, Ciardo S, et al. borderline" which showed an intense this lymphoma, and we believe that the deflectance confocal microscopy correlates of and diffuse TCR delta expression. In entire scientific community would be

Ana Taibo, MD* David Cassarino, MD, PhD+ Angel Fernandez-Flores, MD. PhD±'§"

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University of A Coruña (UDC), A Coruña

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EXTRAORDINARY CASE REPORT

Primary Cutaneous Gamma-Delta T-Cell Lymphoma Mimicking anaplastic lymphoma kinase-1-Negative Anaplastic Large Cell Lymphoma: A Case Report

Hanv Meawad, MD, MSc.* Joo Y, Song, MD.* Matthew L, Ulrickson, MD.† and Dennis D. Weisenburger, MD*

Abstract: Primary cutaneous gamma-delta T-cell lymphoma is a rare and aggressive neoplasm, representing less than 1% of all cutaneous T-cell lymphomas. In this article, we report the case of a 49-year-old woman who presented with a history of generalized skin rash and a recent mass on the left upper extremity, as well as right inguinal soft tissue swelling and splenomegaly. Histologic examination of the mass revealed a diffuse subcutaneous infiltrate of large anaplastic and CD30-positive lymphoid cells with rimming of the adinocytes. This case demonstrates unusual cytologic features in primary cutaneous gamma-delta T-cell lymphoma that mimic the features of anaplastic lymphoma kinase-1-negative anaplastic large-

Key Words: primary cutaneous gamma-delta T-cell lymphoma, ana-

(Am J Dermatopathol 2022;44:62-65)

INTRODUCTION

Primary cutaneous gamma-delta T-cell lymphoma (PCGD-TCL) is a rare and aggressive neoplasm that was identified as a distinct entity in the 2008 World Health Organization classification.1 This entity primarily affects the skin and is composed of clonal and activated cytotoxic gamma-delta T cells. The entity includes cases that were previously designated as "subcutaneous panniculitis-like T-cell lymphoma" but was separated from cases with the alpha-beta phenotype because PCGD-TCL has a worse survival, a higher incidence of necrosis and ulceration, and a higher frequency of hemophagocytic syndrome.2 Patients with PCGD-TCL usually present with plaques which ulcerate and form necrotic nodules that rapidly increase in size and occur most frequently on the extremities. Development of mucosal lesions and involvement of other extranodal sites are frequently observed. However, the bone marrow, lymph nodes, and spleen are not typically involved at the time of initial presentation.2 Two distinct molecular subtypes of PCGD-TCL

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have been identified. The Vdelta-1 subtype classically involves the epidermis and dermis, whereas the Vdelta-2 sub type is characterized by panniculitis and a more aggressive clinical course.3 Noticeable subcutaneous involvement can also occur in other T-cell neoplasms with a predilection for the skin, such as subcutaneous panniculitis-like T-cell lymphoma (SPTCL) and extranodal natural killer (NK)/T-cell lymphoma.4,5

Although epidermal involvement with skin ulceration and concomitant hemophagocytic syndrome are commonly seen in PCGD-TCL, these findings are not considered pathognomonic of the disease and accurate diagnosis depends on careful morphologic and immunophenotypic characteriza tion that includes T-cell receptor (TCR)-delta expression by immunohistochemistry or flow cytometry.2,6

CASE REPORT

A 49-year-old White woman, with a history of rheumatoid arthritis treated with etanercept and methotrexate, presented with mild neutropenia. Bone marrow biopsy at the time was hypercellular at 75% with no significant dysplasia or evidence of lymphoma. She was noted to have mild splenomegaly at 12 cm. A clinical diagno of Felty syndrome was made, and she was managed with oral methotrexate and intermittent rituximab with stable blood counts.

Three years later, she developed the onset of multiple

taneous nodules, initially in the left upper extremity and ther few nodules overlying the right lower quadrant of the abdomen and right lower extremity as seen in Figure 1A. These lesions were tender, mildly erythematous, and 2-3 cm in size. Ultrasound of the lesions suggested fat necrosis. Her spleen remained 11 cm in size, and computed tomography scan of the abdomen and pelvis did not reveal any significant adenopathy. Repeat bone marrow biopsy was negative for evidence of lymphoma.

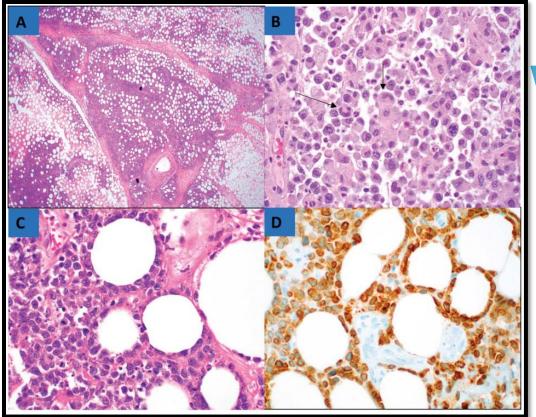
She underwent excision of these subcutaneous nodules and splenectomy. Positron emission tomography scan staging shortly after these procedures identified multiple fluorodeoxyglucose-avid cutaneous nodules in the extremities and lower abdomen without significant adenorathy, as seen in Figures 1B-D.

Histologic examination of biopsies of the mass lesions showed infiltration of the subcutaneous fat by sheets of large chromatin and prominent nucleoli. Many "hallmark cells" were also present, along with frequent mitotic figures. Rimming and necrosis present, along with request mixture injuries, relating and necroiss of fat cells were also observed. No significant epidermotropism was noted. Immunohistochemistry performed on paraffin sections showed strong expression of CD3 on the large cells without expression of CD4, CD8, or CD5. There was also expression of

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Nevertheless, the authors made the diagnosis of PCGD-TCL based on the panniculitic pattern, the rimming, and mainly on the immunostaining for T-cell receptor (TCR)-delta.

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Enhancing the Ability to Diagnose, Interpre left posterior medial thigh, and

with a γδ T-cell phenotype

EXTRAORDINARY CASE REPORT

Primary Cutaneous Gamma-Delta T-Cell Lymphoma Mimicking anaplastic lymphoma kinase-1-Negative Anaplastic Large Cell Lymphoma: A Case Report

Hanv Meawad, MD, MSc.* Joo Y, Song, MD.* Matthew L, Ulrickson, MD.† and Dennis D. Weisenburger, MD*

Abstract: Primary cutaneous gamma-delta T-cell lymphoma is a rare and aggressive neoplasm, representing less than 1% of all cutaneous T-cell lymphomas. In this article, we report the case of a 49-year-old woman who presented with a history of generalized skin rash and a recent mass on the left upper extremity, as well as right

been identified. The Vdelta-1 subtype classically involves the epidermis and dermis, whereas the Vdelta-2 subtype is characterized by panniculitis and a more aggressive clinical course.3 Noticeable subcutaneous involvement can also occur in other T-cell neoplasms with a predilection for the skin, such as subcutaneous panniculitis-like T-cell lym



FIGURE 1. A, Skin lesion on the right lower extremity. B-D, Positron emission tomography scan staging at diagnosis reveals multiple fluorodeoxyglucose-avid subcutaneous lesions at the left upper extremity, lower abdomen.

EXTRAORDINARY CASE REPORT

DUSP22-IRF4 Rearranged CD30-Positive Primary Cutaneous Lymphoproliferative Disorder With Gamma/ Delta Phenotype

Yasmin H. Fattah, MD,* David Crasto, DO,* Shuo S. Liu, MD, PhD,* Yuliya Linhares, MD,† Franz Kerdel, DO,‡ Andrew Hanly, MD,§ and Laszlo J. Karai, MD, PhD§

Abstract: CD30-positive primary cutaneous lymphoproliferative disorders (CD30+PCLPD) are a heterogeneous group of cutaneous T-cell lymphoma (CTCL) that includes lymphomatoid papulosis (LvP) and primary cutaneous anaplastic large cell lymphoma. They exist as a clinical and pathological spectrum, which display significant overlap and variability. The diagnosis is made based on correlation between clinical and histopathologic findings. LyP with 6p25.3 rearrangement subtype represents <5% of LyP cases and is defined by DUSP22-IRF4 rearrangement on 6p25.3 locus. The renorted cases express the alpha/beta T-cell recentor and follow an indolent clinical behavior typical of LyP. The same rearrangement is detected in 28% of anaplastic large cell lymphoma. We hereby present an extraordinary case of CD30+PCLPD with DUSP22-IRF4 rearrangement and novel expression of gamma/delta T-cell immunophenotype in a young patient. Although the gamma/delta T-cell immunophenotype has been described in many other T-cell lymphomas, this is the first reported association with CD30+PCLPD with

Key Words: CD30+ primary cutaneous lymphoproliferative disorders, C-ALCL, LyP with DUSP22-IRF4 rearrangement, γδ T cells (Am J Dermatopathol 2023:45:831-834)

lymphoproliferative disorders (CD30+PCLPDs) are the second most common group of cutaneous lymphomas, after mycosis fungoides, accounting for 30% of cases. 1,2 They include lymphomatoid papulosis (LvP) and primary cutaneous anaplastic large cell lymphoma (C-ALCL), which display a spectrum of morphologic features.3,4 C-ALCL is characterized by the appearance of necrotic and/or ulcerated papulonodular eruption, which may be solitary or grouped, favoring the head/neck region, along with the upper extremity. Median age of onset is the sixth decade of life. Most cases are anaplastic lymphoma kinase-1 negative.5,6 LyP is a monoclonal disorder that arises from a single transformed T cell and is characterized by

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diameter, one on the flexural aspect of the left forearm, and the other on thigh, sharply demarcated from the surrounding uninvolved skin (Fig. 1A). The remainder of her physical examination was unremarkable, including axillary and inguinal lymph node examination. An initial punch biopsy of the left arm lesion was performed followed by evaluation. The pathologic findings were remarkable for dermal based, atypical lymphocytic infiltrate, with Grenz zone and no epidermotropism (Figs. 1B, C). The atypical lymphocytes are medium to large in size, with no involvement or destruction of the follicular or adnexal structures, permeating through the collagen bundles with infiltration of the subcutaneous tissue (Figs. 1D-F). A polymorphic reactive inflammatory milieu was not identified. The lymphocytes were atypical T cells with mostly retained CD3 (Fig. 2A), strongly

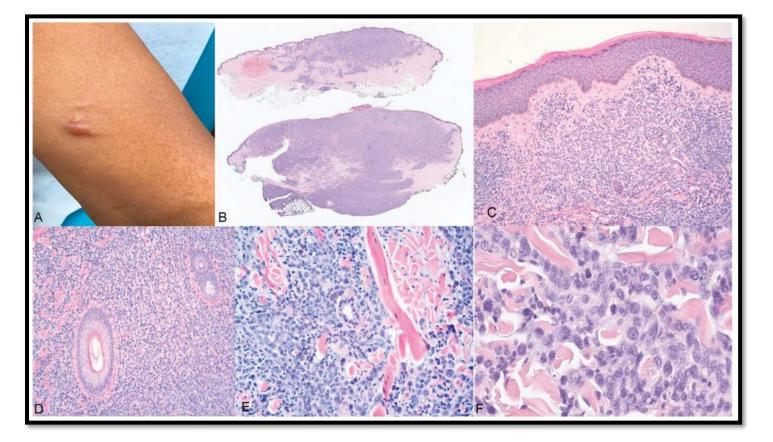
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multifocal crops of recurrent papulonodular lesions that favor the trunk and extremities, with a median age of onset of 45 years.7 LyP histological spectrum is classified by the World Health Organization into (A through E) subtypes based on morphologic features and stratified with either CD4positive or CD8-positive T cells of alpha/beta (αβ) or gamma/delta (γδ) immunophenotype. In 2013, the new subtype of LvP with 6p25.3 rearrangement was described8 and subsequently recognized in 2018 by the World Health Organization classification of LyP as a sixth subtype of LyP.9 We hereby present a novel case of CD30+PCLPD with DUSP22-IRF4 rearrangement demonstrating γδ phenotype in

a young patient that had an unusual clinical course. We illuminate the complexity of classifying such a case under the current spectrum of CD30+PCLPDs, which suggests that this might be an outlier and possibly a new subtype of CD30+PCLPD with unique histology and prognosis.

CASE REPORT

A 41-year-old African American woman presented to the dermatology clinic with a recent development of lesions on her extrem ities. She reported that the individual lesions had increased in size since their appearance, rendering significant worry and ultimately prompting her to seek medical consultation. Otherwise, she was without any active medical problems and denied any additional signs or symptoms suggestive of systemic disease. Physical examination demonstrated 2 skin-colored papules, approximately 8-10 mm in a subsequent diagnostic excision for more thorough and accurate positive CD30 (Fig. 2B), completely lost CD5, high Ki-67 prolifer-ative index, and partial loss of CD2 (Fig. 2C). Interestingly, the malignant T cells had a double-negative CD4 (Fig. 2D), and CD8 immunophenotype (Fig. 2E) and were positive for gamma T-cel receptor (TCR) (Fig. 2F). Cytotoxic T-cell markers TIA-1 and







Summary

- Our patient's clinical presentation and histopathologic findings over 3 years of follow-up support the possibility of gamma-delta-positive pcALCL with extensive LyP-like lesions
- While two arm-level copy number variations including gain of 1q raise alternative concern for pcGDTCL, the absence of consensus cancer gene mutations in MAPK, MYC, and JAK/STAT pathways would be unusual for pcGDTCL
- No rearrangements or fusions were identified
- The diagnosis is not unambiguous; therefore, our patient will be managed conservatively and followed closely





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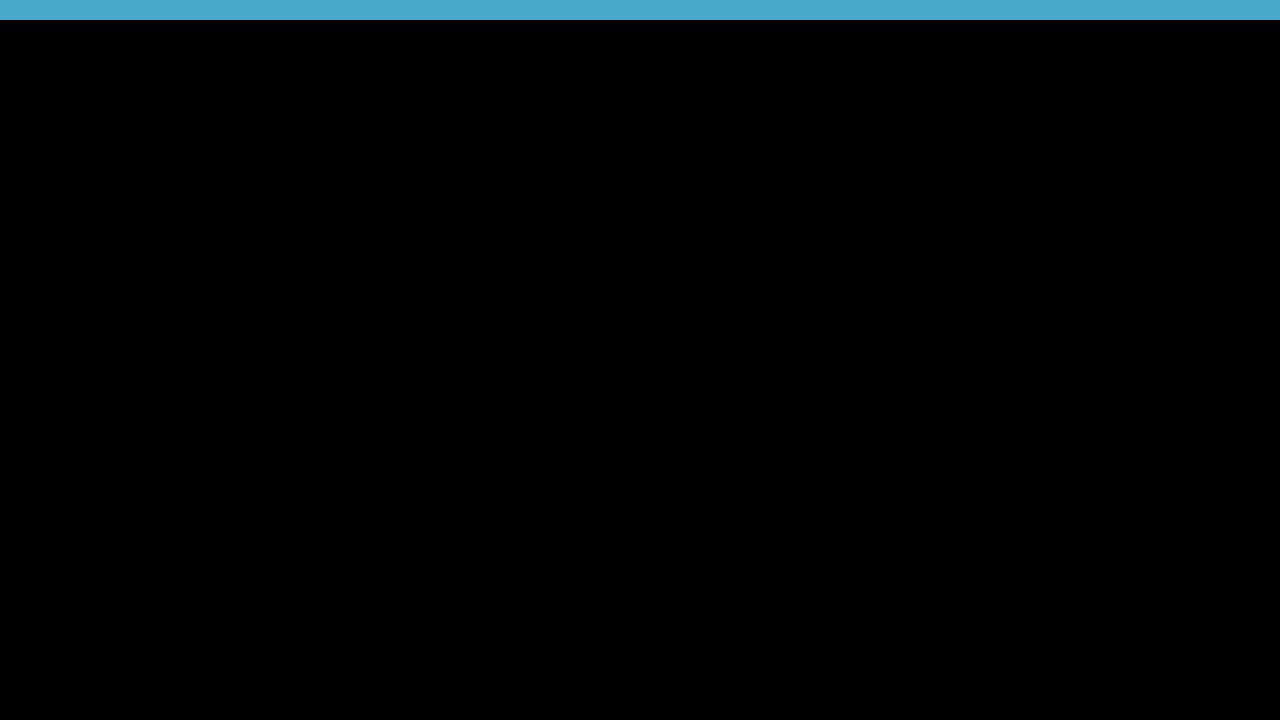


Thank you!

Thank you!







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	AL MALIGNANCY 6, INSERTION/DE				THE DETECTION	N OF BASE		
	•	ADGRA2 (GPR124)		AKT2		ALK	AMER1 (FAM123B	or WTX)
			ARAF	ARFRP1	ARHGAP26 (GRAF)		ARID1A	ARID2
			ATR	ATRX	AURKA			AXL
	BAP1		BCL10		BCL2		BCL6	BCL7A
			BLM	BRAF		BRCA2	BRD4	BRIP1
			BTLA				CBFB	CBL
			CCND3			CD22	CD274 (PD-L1)	CD36
CD58	CD70	CD79A	CD79B	CDC73		CDK12	CDK4	CDK6
CDK8	CDKN1B	CDKN2A	CDKN2B	CDKN2C	CEBPA	CHD2	CHEK1	CHEK2
	CIITA	CKS1B	CPS1	CREBBP	CRKL	CRLF2	CSF1R	CSF3R
	CTNNA1		CUX1				DDX3X	DNM2
			DUSP2				EED	EGFR
			EPHA3				ERBB2	ERBB3
			ETS1	ETV6			FAF1	FANCA
	FANCD2		FANCF	FANCG	FANCL		FBXO11	FBXO31
	FGF10		FGF19		FGF3		FGF6	FGFR1
			FHIT				FLT4	FLYWCH1
	FOXO1		FOXP1	FRS2	GADD45B		GATA2	GATA3
	GNA11		GNA13	GNAQ	GNAS	GRIN2A	GSK3B	GTSE1
	HDAC4		HGF	H1-2 (HIST1H1C)		H1-3 (HIST1H1D)		
H1-4 (HIST1H1E)		H2AC6 (HIST1H2AC)		H2AC11 (HIST1H2A		H2AC16 (HIST1H2A		
H2AC17 (HIST1H2AM)		H2BC4 (HIST1H2BC)		H2BC11 (HIST1H2BJ)		H2BC12 (HIST1H2BK)		
H2BC17 (HIST1H2B	-,	H3C2 (HIST1H3B)		HNF1A	HRAS	HSP90AA1		ID3
	IDH2		IKBKE				IL7R	INHBA
	INPP5D (SHIP)		IRF4	IRF8			JAK2	JAK3
			KDM2B		KDM5A	KDM5C	KDM6A	KDR
			KMT2A (MLL)		KMT2D (MLL2)		LEF1	LRP1B
			MAGED1					MAP3K1
	MAP3K6		MAPK1					MEF2B
			MIB1				MPL	MRE11 (MRE11A)
	MSH3		MTOR		MYC			MYD88
	NCOR2		NF1	NF2	NFE2L2		NKX2-1	NOD1
	NOTCH2		NRAS	NSD2 (WHSC1 or N			NTRK1	NTRK2
			P2RY8	PAG1	PAK3		PASK	PAX5
			PCLO DIV2CA	PDCD1		PDCD1LG2 (PD-L2)		PDGFRA
	PDK1		PIK3CA DDKAD1A		PIK3R1	PIK3R2	PIM1 DTEN	PLCG2 DTDN11
	PPP2R1A		PRKAR1A	PRKDC PADSO	PRSS8	PTCH1	PTEN	PTPN11 DASGEEIA
			RAD21	RAD50 RICTOR		RAF1 ROS1	RARA	RASGEF1A DLINY1
	RELN		RHOA SDHC				RPTOR SETD2	RUNX1 SF3B1
								SF3B1 SMO
	SMAD2		SMARCA1 SOX10		SMARCB1 SPEN	SMC1A SPOP	SMC3 SRC	SRSF2
			STAT5A	SOX2 STAT5B			SKC SUFU	SRSF2 SUZ12
			TCL1A (TCL1) TNFRSF11A	TENT5C (FAM46C) TNFRSF14			TLL2 TP53	TMEM30A TP63
TMSB4XP8 (TMSL3								
	TRAF3		TSC1				TYK2	U2AF1 7NE217
	VHL		WT1	XBP1	XPO1	YY1AP1	ZMYM3	ZNF217
ZNF24 (ZSCAN3)	ZNF/03	ZRSR2						

ALK BCL2 BCL6 BCR BBAF CCND1 CRIF2 EGFR EPOR ETV1 ETV4 ETV5 EWSR1 FGFR2 IGH IGK IGL JAK1 JAK2 KMTZA (MILL) MYC NTRKI PDGFRB RAFI RAFI RET ROS1 TIMPRSS2 TRG PDGFRB PDGFRB RAFI RAFI HEMATOLOGIC-LA MALIGNANCY RNA GENE LIST: FOR THE DETECTION OF SELECT REARRAGEMENTS* ART ARX ABI1 ABL ARL ACS16 AFDN (MILT® or #6) AFT AFF4 ALX ARHGAP26 (GRA)- ARHGEFT2 ARDIA ARRT ASX11 ATT AFG5 ATTC BCL0 BL(18 BCL2 BCL3 BCL6 BCL7A BCL9 BCOR BCR BIRC3 BRAF BCL1B BCL2 BCL3 BCL6 BCL7A BCL7A BCL9 BCOR CCND1 CCND2 CCND3 CD274 (PD-11) CDK6 CDX2 CEP4	HEMATOLOGICAL MALIGNANCY DNA GENE LIST: FOR THE DETECTION OF SELECT REARRANGEMENTS								
JAK1 JAK2 KMTAA (MILD) MYC NTRKI PDGFRA PDGFRA RAFI RARA RET ROST TMPRSS2 TRG *** PDGFRA PDGFRB RAFI RARI ABIT ABL2 ACSL6 AFDN (MILT) or FS AFFT AFF4 ALK ABIT ABL2 ACSL6 AFDN (MILT) or FS AFFT AFF4 ALK BC10 BCL11A BCL1B BCL2 BCL3 BCL6 BCL7A BCL9 BCOR BCR BIRG3 BRAF BTGI CAMTAI CARS1 (CARS) CBFA2T3 CBFB CBL CCND1 CCND2 CCXD3 CD274 (P9-L1) CDK6 CDX2 CEPA3 (FGRFRIOP) CHC2 CHC1 CHN1 CREBBP CRIF2 CSF1 CTTONIBI DDT3 DDX10 DDX6 DEK DUSP22 CEPA3 (FGRFRIOP) CHC2 CHN1 CRB312 CRB312 CRB312 CRB312 CRB312 CRB312 CRB312 CRB312 <t< td=""><td>ALK</td><td>BCL2</td><td>BCL6</td><td>BCR</td><td>BRAF</td><td>CCND1</td><td>CRLF2</td><td>EGFR</td><td>EPOR</td></t<>	ALK	BCL2	BCL6	BCR	BRAF	CCND1	CRLF2	EGFR	EPOR
RET ROS1 TMPRSS2 TRG HEMATOLOGISC MALIGNANCY RNA GENE LIST: FOR THE DETCTION OF SELECT REARRANGEMENTS* ABII ABI2 ACSL6 AFDM (MILT4 or AFG) AFT ALIC ARHGAP26 (GRAF*) ARHGETI2 ARIDTA ASXL1 ATT ATG5 ATIC BCL10 BCL11B BCL2 ARIDTA ASXL1 ATT ATG5 ATIC BCR BCL17B BCL17B ARIDTA ASXL1 CARSTA CBB BCL2 CBL3 BCL3 BCL17A BCP3 CBL3 BCR CITTA CLTC CLTCL CLTCL CLTCL CLTCL CLTCL CLTCL	ETV1	ETV4	ETV5	ETV6	EWSR1	FGFR2	IGH	IGK	IGL
ABII ABL1	JAK1	JAK2	KMT2A (MLL)	MYC	NTRK1	PDGFRA	PDGFRB	RAF1	RARA
ABI1 ABL1 ABL2 ACSL6 AFDN (MLT4 or Jet) AFF1 AFF4 ALK ARHGAP26 (GRAF) ARHGAPE1 ARIDINA ARNT ASXI ATF1 ATG5 ATC BCL10 BCL118 BCL19 BCL2 BCL3 BCL6 CEVA BCL7A BCL9 BCCR BCR BIRC3 BRAF BTG1 CAMTA1 CARS1 (CARS) CEFA2T3 CBFB CBL CCND1 CCND2 CCND3 CCD274 (PD-L1) CDCA CAX2 CEFA2T3 CBFB CBL CCKC CILTA CLP1 CLTC CLTCL1 CARTS1 (CARS) CDL31 CREB11 CEB12 CKEBPP CRLF2 CST9 CLTNB1 DD173 DDX10 DDX6 DEX CHB312 CEB32 EGFR CRLF2 CST9 CLTCL1 CTCL11 CNTR(CPCB) DDX6 DEX DDX76 CPD15 CER312 CER312 CER312 CER312 CER312 CER312 CER312 CER312	RET	ROS1	TMPRSS2	TRG					
ABI1 ABL1 ABL2 ACSL6 AFDN (MLT4 or Jet) AFF1 AFF4 ALK ARHGAP26 (GRAF) ARHGAPE1 ARIDINA ARNT ASXI ATF1 ATG5 ATC BCL10 BCL118 BCL19 BCL2 BCL3 BCL6 CEVA BCL7A BCL9 BCCR BCR BIRC3 BRAF BTG1 CAMTA1 CARS1 (CARS) CEFA2T3 CBFB CBL CCND1 CCND2 CCND3 CCD274 (PD-L1) CDCA CAX2 CEFA2T3 CBFB CBL CCKC CILTA CLP1 CLTC CLTCL1 CARTS1 (CARS) CDL31 CREB11 CEB12 CKEBPP CRLF2 CST9 CLTNB1 DD173 DDX10 DDX6 DEX CHB312 CEB32 EGFR CRLF2 CST9 CLTCL1 CTCL11 CNTR(CPCB) DDX6 DEX DDX76 CPD15 CER312 CER312 CER312 CER312 CER312 CER312 CER312 CER312									
ARHGAP26 (GRAF) ARHGEF12 ARID1A ARNT ASXL1 ATF1 ATG5 ATIC BC110 BC111A BC11B BC12 BC3 BC16 BC17A BC19 BCOR BCR BIRC3 BRAF BTG1 CAMTA1 CARS1 (CARS) CBF3 CBFB CBL CCND1 CCND2 CCP03 CD274 (PD-L1) CDK6 CDX2 CEP43 (FGFR10P) CHC2 CHN1 CKC CITA CLP1 CLTC CTC11 CNTRL (CEP110) COL1A CREB1 CREB12 CHN1 CREBBP CRIF2 CSF1 CTNNB1 DDT3 DDX10 DDX6 DEX CDE922 CEP63 CEP43 (FGFR10P) CHC2 CHN1 CREB12 CREB12 CEP43 (FGFR10P) CD10 CREB12 CHN1 CREB12 CHN1 CEP43 (FGFR10P) CD10 CPH2 CHN1 CPE812 CHN1 CPF1 CN1 CPF8 CPF1 CPF1 CPF1 CPF1 CN2 CPF1 CPF1 CPF1 </td <td colspan="9">HEMATOLOGICAL MALIGNANCY RNA GENE LIST: FOR THE DETECTION OF SELECT REARRANGEMENTS*</td>	HEMATOLOGICAL MALIGNANCY RNA GENE LIST: FOR THE DETECTION OF SELECT REARRANGEMENTS*								
BCL10 BCL11A BCL11B BCL2 BCL3 BCL6 BCL7A BCL9 BCOR BCR BIRC3 BRAF BTG1 CAMTA1 CARSI (CARS) CBFA2TG CBFB CBL CCND1 CCND2 CCVD3 CD274 (PD-L1) CDK6 CP43 (FGFRIOP) CHIC2 CHIA CIC CITC1 CITC1 CNTRL (CEPTIO) COL1A1 CREB3L1 CREB3L2 CREBBP CRLF2 CSF1 CITNB1 DDIT3 DDX10 DDX6 DEK DUSP22 EGFR EIF4A2 ELF4 ELL ELN EML4 EP900 EPOR EPS15 EGRB2 ERG ET51 ETV1 ETV4 ETV5 ETV6 EWSR1 FCR2B2 FCRL4 FEV FGFR1 FGFR2 FGFR3 FLI FNBP1 FNX01 HAC9 (HISTIH4) HARPUDI HEY1 HIP1 HIF4 HIM6A1 HIM6A1 HNX11 HAC9 (HISTIH4) HACXP1 HOXC11 HOXC	ABI1	ABL1	ABL2	ACSL6	AFDN (MLLT4 or)	4 <i>F6)</i>	AFF1	AFF4	ALK
BCR BIRG3 BRAF BTG1 CAMTA1 CARS1 (CARS) CBFA2T3 CBFB CBL CCND1 CCND2; CCND3 CD274 (FD-L1) CDK6 CDX2 CEP43 (FGFRIOP) CHIC2 CHINI CIC CITTA CLIP CLIC CLICT CNTRL (CEP110) COLIA CREB3L1 CREB3L2 CRBBP CRLF2 CSF1 CTNNB1 DDIT3 DDX10 DDX6 DEK DUSP22 EGFR EIF4A2 ELF4 ELL ELN EML4 EP300 EPOR EPS15 ERBB2 ERG ETS1 ETV1 ETV4 ETV6 EVSR1 FCGR2B FCRL4 FEV FGFR1 FGFR2 FGFR2 FGFR3 FLI HNB4 POX01 FOX01 FOX03 FOX01 FOX04 FOXP1 FST13 FUS GAS7 GLI1 MB75 GPHN H4C9 (HISTIH4I) HAC9 (HISTIH4I) HAC9 (HISTIH4I) HAC9 (HISTIH4I) HAC9 (HISTIH4I) HAC9 (HISTIH4I) HAX3 HAX1<	ARHGAP26 (GRAF)		ARHGEF12	ARID1A	ARNT	ASXL1	ATF1	ATG5	ATIC
CCND1 CCND2: CCND3 CD274 (PD-L1) CDK6 CDX2 CEP43 (FGFRIOP) CHC2 CHN1 CIC CITA CLP1 CLTC CLTCL1 CNTRL (CEP110) COL1A1 CREB311 CREB322 CREBBP CRIF2 CSF1 CTNNB1 DBX DDX10 DDX6 DEK DBX922 EGFR EIF442 ELF4 ELL ELN EML4 EP300 EP0R EPS15 ERBB2 ERG ETS1 ETV1 ETV4 ETV5 ETV6 EWSR1 FCG22B FCXL4 FEV FGFR1 FGFR2 FGFR3 FLI FNBP1 FOX01 FOX03 FOX03 FOX04 FOXP1 FST13 FUS GSFR3 FLI FNBP1 FOX04 HAC9 (HST)HH19 HLF HMG61 HMG62 HOXA11 HOXA13 HAC9 (HST)H19H19 HERPUD HEV1 HIP1 HLF HMGA1 HMG61 HMG2 HOXA11 HOXA13 HAC9 (HST)H19H19 HERPUD	BCL10	BCL11A	BCL11B	BCL2	BCL3	BCL6	BCL7A	BCL9	BCOR
CIC CIITA CLP1 CLTC CLTCL1 CNTRL (CEP110) COLIA1 CREB3L1 CREB3L2 CREBBP CRLP2 CSF1 CTNNB1 DDIT3 DDX10 DDX6 DEK DUSP22 EGFR EIF4A2 ELF4 ELL EIN EML4 EP900 EPOR EPS15 ERBB2 ERG ETS1 ETV1 ETV4 ETV5 ETV6 EWSR1 FCGR2B FCRL4 FEV FGFR1 FGFR2 FGFR3 FLI1 FNBP1 FOXO1 FOXO3 FOXO4 FOXP1 FSTL3 FUS GAS7 GLI1 GMP5 GPNN H4C9 (HISTIH4I) HERPUDI HEY1 HIP1 HLF HMGA1 HMGA2 HOXA11 HOXA13 HOXA3 HOXA4 HMGA2 HOXA11 HOXA13 HXA3 HXA3 HXFA ITIL HMGA2 HXA31 HXA3 HXA3	BCR	BIRC3	BRAF	BTG1	CAMTA1	CARS1 (CARS)	CBFA2T3	CBFB	CBL
CREBBP CRIF2 CSF1 CTNNB1 DDIT3 DDX10 DDX6 DEK DUSP22 EGFR EIF4A2 ELF4 ELL ELN EML4 EP300 EPOR EPS15 ERBB2 ERG ETS1 ETV1 ETV4 ETV5 ETV6 EWSR1 FCG2B FCRL4 FEV FGFR1 FGFR2 FGFR3 FLI1 FMBP1 FOXO1 FOXO3 FOXO4 FOXP1 FSTL3 FUS GAS7 GLI1 GMPS GPHN H4C9 (HISTIH4I) HERPUDI HEY1 HIP1 HLF HMGA1 HMGA2 HOXA11 HOXA3 HOXA3 HOXC13 HOXC13 HOXD11 HOXD3 HSP90AA1 HSP90AB1 IGH IGK IGL IKZF1 IL21 HAXA IKX JAK1 JAK2 JAK3 JAZF1 KAT6A (MYST3) KOSR KIFSB KMT2A (MLI) LASP1 LCP1 LMO1 MLD1 LPP LYL1	CCND1	CCND2	CCND3	CD274 (PD-L1)	CDK6	CDX2	CEP43 (FGFR10P)	CHIC2	CHN1
EGFRIF4A2ELF4ELLELNEML4EP300EPOREP515ERBB2ERGETS1ETV1ETV4ETV5ETV6EWSR1FCGR2BFCRL4FEVFGFRIFGFRIFGFR3FLIFNBP1FOXO1FOXO3FOXO4FOXP1FST13FUSGAS7GLI1GMPSGPHN $HCCO1$ HCCO1HERPUDIHEV1HIP1HLFHMGA1HMGA2HOXA11HOXA13HOXA3HOXA9HOXCI1HOXCI3HOXD11HOXD13HSP90A1HSP90A81IGHIGKIGLIKZF1IL21RIL3IRF4ITKJAK1JAK2JAK3JAZF1KAT6A (MYST3)KDSRKIF5BKMT2A (MLL)LASP1LCP1LMO1LMO2LPPLY1MAFMAFBMAHTMDS2MECOMMLF1MLIT1(ENL)MLLTIO (AF10)MLT3MLT6MN1MNX1MRTFA (MKL1)MS12MSNMUC1MYBMYCMYH11MYH9NACANBEAPI (BCL8)NCOA2NDRG1NF1NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1+ MSET)NSD3 (WHSC1L)NTRK1NTRX2NTRX3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAHIB2PAX3PAX5PAX7PEX1PCM1PCKTPICH1PTCH1PTCH1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15	CIC	CIITA	CLP1	CLTC	CLTCL1	CNTRL (CEP110)	COL1A1	CREB3L1	CREB3L2
ERBB2ERGETSIETV1ETV4ETV5ETV6EWSR1FCGR2BFCRL4FEVFGFR1FGFR2FGFR3FLI1FNBP1FOXO1FOXO3FOXO4FOXP1FSTL3FUSGAS7GLI1GMPSGPHNH4C9 (HISTIH4I)HERPUD1HEY1HIP1HLFHMGA1HMGA2HOXA11HOXA13HOXA3HOXA9HOXC11HOXC13HOXD11HOXD13HSP90AB1IGHIGKIGLIKZF1IL21RIL3IRF4ITKJAK1JAK2JAK3JAZF1KAT6A (MYST3)KDSRKIF5BKMT2A (MLL)LASP1LCP1LM01LM02LPPLVL1MAFMAFBMALT1MDS2MECOMMLF1MLIT1 (ENL)MLLT10 (AF10)MLIT3MILT6MN1MNX1MRTFA (MKL1)MS12MSNMUC1MYBMYCMYHI1MYH9NACANBEAPI (BCL8)NCOA2NDRG1NF1NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1 or MMSET)NSD3 (WHSC1L1)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAHIB2PAX3PAX5PAX7PBX1PCM1PCSK7PCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFBAPDGFRBPER1PHF1PICALMPIM1PLAG1RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1T1 (ETO)RUNX2SEC31A <t< td=""><td>CREBBP</td><td>CRLF2</td><td>CSF1</td><td>CTNNB1</td><td>DDIT3</td><td>DDX10</td><td>DDX6</td><td>DEK</td><td>DUSP22</td></t<>	CREBBP	CRLF2	CSF1	CTNNB1	DDIT3	DDX10	DDX6	DEK	DUSP22
FCRL4 FEV FGFR1 FGFR2 FGFR3 FLI1 FNBP1 FOXO1 FOXO3 FOXO4 FOXP1 FSTL3 FUS GA57 GLI1 GMPS GPHN H4C9 (HISTIH4I) HERPUD1 HEY1 HIP1 HLF HMGA1 HMGA2 HOXA11 HOXA3 HOXA3 HOXA9 HOXC11 HOXC13 HOXD11 HOXD13 HSP90A11 HSP90AB1 IGH IGK IGL IKZF1 IL21R HJ3 IRF4 ITK JAK1 JAK2 JAK3 JAZF1 KAT6A (MYST3) KDSR KIF5B KMT2A (MLL) LASP1 LCP1 LMO1 LMO2 LPP LYL1 MAF MAFB MALT1 MDS2 MECOM MLF1 MLIT1 (ENL) MLLT0 (AF10) MLLT3 MLLT6 MN1 MXN1 MRTFA (MKL1) MS12 MSN MUC1 MYB MYC MYH11 MYH9 NACA NBEAP1 (BCL8) NCOA2 NDRG1 NF1 <td>EGFR</td> <td>EIF4A2</td> <td>ELF4</td> <td>ELL</td> <td>ELN</td> <td>EML4</td> <td>EP300</td> <td>EPOR</td> <td>EPS15</td>	EGFR	EIF4A2	ELF4	ELL	ELN	EML4	EP300	EPOR	EPS15
FOXO4FOXP1FSTL3FUSGAS7GLI1GMPSGPHNH4C9 (HIST1H4I)HERPUD1HEY1HIP1HLFHMGA1HMGA2HOXA11HOXA13HOXA3HOXA9HOXC11HOXC13HOXD11HOXD13HSP90AB1IGHIGKIGLIKZF1IL2IRIL3IRF4ITKJAK1JAK2JAK3JAZF1KAT6A (MYST3)KDSRKIF5BKMT2A (MLL)LASP1LCP1LM01LM02LPPLYL1MAFMAFBMALT1MDS2MECOMMLF1MLLT1 (ENL)MLT10 (AF10)MLLT3MLLT6MN1MNX1MRTFA (MKL1)MS12MSNMUC1MYBMYCMYH11MYH9NACANBEAPI (BCL8)NCOA2NDRG1NF1NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1 or MSET)NSD3 (WHSC1L1)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAHIB2PAX3PAX5PAX7PBX1PCM1PCSK7PDCDILG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOUZAF1PPPICBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETHOHRHOHRNF213SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2 </td <td>ERBB2</td> <td>ERG</td> <td>ETS1</td> <td>ETV1</td> <td>ETV4</td> <td>ETV5</td> <td>ETV6</td> <td>EWSR1</td> <td>FCGR2B</td>	ERBB2	ERG	ETS1	ETV1	ETV4	ETV5	ETV6	EWSR1	FCGR2B
HERPUD1 HEY1 HIP1 HLF HMGA1 HMGA2 HOXA11 HOXA13 HOXA3 HOXA9 HOXC11 HOXC13 HOXD11 HOXD13 HSP90AB1 IGH IGK IGL IKZF1 IL2IR IL3 IRF4 ITK JAK1 JAK2 JAK3 JAZF1 KAT6A (MYST3) KDSR KIF5B KMT2A (MLL) LASP1 LCP1 LM01 LMO2 LPP LYL1 MAF MAFB MALT1 MDS2 MECOM MLF1 MLLT1 (ENL) MLLT10 (AF10) MLLT3 MLLT6 MN1 MNX1 MRTFA (MKL1) MS12 MSN MUC1 MYB MYC MYH11 MYH9 NACA NBEAPI (BCL8) NCOA2 NDRG1 NF1 NF2 NIN NOTCH1 NPM1 NR4A3 NSD1 NSD2 (WHSCHT-WMSET) NSD3 (WHSCIL) NTRK1 NTRK2 NTRK3 NUMA1 NUP214 NUP98 NUTM2A OMD P2RY8 PAFAH1B2	FCRL4	FEV	FGFR1	FGFR2	FGFR3	FLI1	FNBP1	FOXO1	FOXO3
HOXA9HOXC11HOXC13HOXD11HOXD13HSP90A1HSP90AB1IGHIGKIGLIKZF1IL21RIL3IRF4ITKJAK1JAK2JAK3JAZF1KAT6A (MYST3)KDSRKIF5BKMT2A (MLL)LASP1LCP1LM01LM02LPPLYL1MAFMAFBMALT1MDS2MECOMMLF1MLLT1 (ENL)MLLT10 (AF10)MLLT3MLLT6MN1MNX1MRTFA (MKL1)MS12MSNMUC1MYBMYCMYH11MYH9NACANBEAP1 (BCL8)NCOA2NDRG1NF1NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1+WHSCT)NSD3 (WHSC1L)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAHIB2PAX3PAX5PAX7PBX1PCM1PCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PICALMPIM1PIK7PMLPOU2AF1PPP1CBPRDM1PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213SEPTIN5 (SEPTS)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUND-2SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBLIX1TLX3TMPRSS2TNFRSF11ATOP1TP63 <td>FOXO4</td> <td>FOXP1</td> <td>FSTL3</td> <td>FUS</td> <td>GAS7</td> <td>GLI1</td> <td>GMPS</td> <td>GPHN</td> <td>H4C9 (HIST1H4I)</td>	FOXO4	FOXP1	FSTL3	FUS	GAS7	GLI1	GMPS	GPHN	H4C9 (HIST1H4I)
IGLIKZF1IL21RIL3IRF4ITKJAK1JAK2JAK3JAZF1KAT6A (MYST3)KDSRKIF5BKMT2A (MLL)LASP1LCP1LM01LM02LPPLYL1MAFMAFBMALT1MDS2MECOMMLF1MLLT1 (ENL)MLLT10 (AF10)MLLT3MLLT6MN1MNX1MRTFA (MKL1)MS12MSNMUC1MYBMYCMYH11MYH9NACANBEAPI (BCL8)NCOA2NDRG1NF1NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1 or MMSET)NSD3 (WHSCIL)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAH1B2PAX3PAX5PAX7PBX1PCM1PCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOU2AF1PPPICBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RP122RPN1RUNX1RUNX1TI (ETO)RUNX2SEC31ASEPTINS (SEPT5)SEPTING (SEPT6)SEPTING (SEPT9)SETSH3GL1SLC1A2SNX29 (RUND-2A*)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3 </td <td>HERPUD1</td> <td>HEY1</td> <td>HIP1</td> <td>HLF</td> <td>HMGA1</td> <td>HMGA2</td> <td>HOXA11</td> <td>HOXA13</td> <td>HOXA3</td>	HERPUD1	HEY1	HIP1	HLF	HMGA1	HMGA2	HOXA11	HOXA13	HOXA3
JAZF1KAT6A (MYST3)KDSRKIF5BKMT2A (MLL)LASP1LCP1LMO1LMO2LPPLYL1MAFMAFBMALT1MDS2MECOMMLF1MLLT1 (ENL)MLLT10 (AF10)MLLT3MLLT6MN1MNX1MRTFA (MKL1)MS12MSNMUC1MYBMYCMYH11MYH9NACANBEAPI (BCL8)NCOA2NDRG1NF1NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1 or MSET)NSD3 (WHSC1L1)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAH1B2PAX3PAX5PAX7PBX1PCM1PCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOU2AF1PPP1CBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1T1 (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2A)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	HOXA9	HOXC11	HOXC13	HOXD11	HOXD13	HSP90AA1	HSP90AB1	IGH	IGK
LPPLYL1MAFMAFBMALT1MDS2MECOMMLF1MLLT1 (ENL)MLLT10 (AF10)MLLT3MLLT6MN1MNX1MRTFA (MKL1)MS12MSNMUC1MYBMYCMYH11MYH9NACANBEAPI (BCL8)NCOA2NDRG1NF1NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1 or MMSET)NSD3 (WHSC1L1)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAH1B2PAX3PAX5PAX7PBX1PCM1PCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOU2AF1PPP1CBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1T1 (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	IGL	IKZF1	IL21R	IL3	IRF4	ITK	JAK1	JAK2	JAK3
MLLT10 (AF10)MLLT3MLLT6MNIMNX1MRTFA (MKL1)MS12MSNMUC1MYBMYCMYH11MYH9NACANBEAP1 (BCL8)NCOA2NDRG1NF1NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1 or MMSET)NSD3 (WHSCIL1)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAHIB2PAX3PAX5PAX7PBX1PCM1PCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOU2AF1PPP1CBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1T1 (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TCF3 (E2A)TCL1A (TCL1)TECTET1TFE3TFGTFPTTFRCTLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	JAZF1	KAT6A (MYST3)	KDSR	KIF5B	KMT2A (MLL)	LASP1	LCP1	LMO1	LMO2
MYBMYCMYHIIIMYH9NACANBEAPI (BCL8)NCOA2NDRGINF1NF2NFKB2NINNOTCHINPMINR4A3NSD1NSD2 (WHSC1 or MMSET)NSD3 (WHSC1LI)NTRK1NTRK2NTRK3NUMAINUP214NUP98NUTM2AOMDP2RY8PAFAHIB2PAX3PAX5PAX7PBX1PCMIPCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPERIPHF1PICALMPIMIPLAGIPMLPOU2AF1PPP1CBPRDMIPRDMI6PRRXIPSIPIPTCHIPTK7RABEPIRAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPNIRUNX1RUNX1TI (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2X)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TCF3 (E2A)TCL1A (TCL1)TECTET1TFE3TFGTFPTTFRCTLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	LPP	LYL1	MAF	MAFB	MALT1	MDS2	MECOM	MLF1	MLLT1 (ENL)
NF2NFKB2NINNOTCH1NPM1NR4A3NSD1NSD2 (WHSC1 or MMSET) of MMSET)NSD3 (WHSC1L1)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAH1B2PAX3PAX5PAX7PBX1PCM1PCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOU2AF1PPP1CBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1T1 (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2A)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TCF3 (E2A)TCL1A (TCL1)TECTET1TFE3TFGTFPTTFRCTLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	MLLT10 (AF10)	MLLT3	MLLT6	MN1	MNX1	MRTFA (MKL1)	MSI2	MSN	MUC1
NSD3 (WHSC1L1)NTRK1NTRK2NTRK3NUMA1NUP214NUP98NUTM2AOMDP2RY8PAFAHIB2PAX3PAX5PAX7PBX1PCM1PCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOU2AF1PPP1CBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1T1 (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2A)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TCF3 (E2A)TCL1A (TCL1)TECTET1TFE3TFGTFPTTFRCTLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	MYB	MYC	MYH11	МҮН9	NACA	NBEAP1 (BCL8)	NCOA2	NDRG1	NF1
P2RY8PAFAHIB2PAX3PAX5PAX7PBX1PCM1PCSK7PDCD1LG2 (PD-L2)PDE4DIPPDGFBPDGFRAPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOU2AF1PPP1CBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1TI (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2A)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TCF3 (E2A)TCL1A (TCL1)TECTET1TFE3TFGTFPTTFRCTLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	NF2	NFKB2	NIN	NOTCH1	NPM1	NR4A3	NSD1	NSD2 (WHSC1 or MMSET)	
PDE4DIPPDGFBPDGFRBPER1PHF1PICALMPIM1PLAG1PMLPOU2AF1PPP1CBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1TI (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2A)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TCF3 (E2A)TCL1A (TCL1)TECTET1TFE3TFGTFPTTFRCTLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	NSD3 (WHSC1L1)	NTRK1	NTRK2	NTRK3	NUMA1	NUP214	NUP98	NUTM2A	OMD
PMLPOU2AF1PPP1CBPRDM1PRDM16PRRX1PSIP1PTCH1PTK7RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1TI (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2A)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TCF3 (E2A)TCL1A (TCL1)TECTET1TFE3TFGTFPTTFRCTLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	P2RY8	PAFAH1B2	PAX3	PAX5	PAX7	PBX1	PCM1	PCSK7	PDCD1LG2 (PD-L2)
RABEP1RAF1RALGDSRAPIGDS1RARARBM15RETRHOHRNF213RNF217-AS1 (STL)ROS1RPL22RPN1RUNX1RUNX1TI (ETO)RUNX2SEC31ASEPTIN5 (SEPT5)SEPTIN6 (SEPT6)SEPTIN9 (SEPT9)SETSH3GL1SLC1A2SNX29 (RUNDC2A)SRSF3SS18SSX1SSX2SSX4STAT6SYKTAF15TAL1TAL2TBL1XR1TCF3 (E2A)TCL1A (TCL1)TECTET1TFE3TFGTFPTTFRCTLX1TLX3TMPRSS2TNFRSF11ATOP1TP63TPM3TPM4TRIM24	PDE4DIP	PDGFB	PDGFRA	PDGFRB	PER1	PHF1	PICALM	PIM1	PLAG1
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	PML	POU2AF1	PPP1CB	PRDM1	PRDM16	PRRX1	PSIP1	PTCH1	PTK7
SEPTIN5 (SEPT5) SEPTIN6 (SEPT6) SEPTIN9 (SEPT9) SET SH3GL1 SLC1A2 SNX29 (RUNDC2A) SRSF3 SS18 SSX1 SSX2 SSX4 STAT6 SYK TAF15 TAL1 TAL2 TBL1XR1 TCF3 (E2A) TCL1A (TCL1) TEC TET1 TFE3 TFG TFPT TFRC TLX1 TLX3 TMPRSS2 TNFRSF11A TOP1 TP63 TPM3 TPM4 TRIM24	RABEP1	RAF1	RALGDS	RAP1GDS1	RA RA	RBM15	RET	RHOH	RNF213
SS18 SSX1 SSX2 SSX4 STAT6 SYK TAF15 TAL1 TAL2 TBL1XR1 TCF3 (E2A) TCL1A (TCL1) TEC TET1 TFE3 TFG TFPT TFRC TLX1 TLX3 TMPRSS2 TNFRSF11A TOP1 TP63 TPM3 TPM4 TRIM24	RNF217-AS1 (STL)		ROS1	RPL22	RPN1	RUNX1	RUNX1T1 (ETO)	RUNX2	SEC31A
TBL1XR1 TCF3 (E2A) TCL1A (TCL1) TEC TET1 TFE3 TFG TFPT TFRC TLX1 TLX3 TMPRSS2 TNFRSF11A TOP1 TP63 TPM3 TPM4 TRIM24	SEPTIN5 (SEPT5)	SEPTIN6 (SEPT6)	SEPTIN9 (SEPT9)	SET	SH3GL1	SLC1A2	SNX29 (RUNDC2A)	SRSF3
TLX1 TLX3 TMPRSS2 TNFRSF11A TOP1 TP63 TPM3 TPM4 TRIM24	SS18	SSX1	SSX2	SSX4	STAT6	SYK	TAF15	TAL1	TAL2
	TBL1XR1	TCF3 (E2A)	TCL1A (TCL1)	TEC	TET1	TFE3	TFG	TFPT	TFRC
TRIP11 TTL TYK2 USP6 YPEL5 ZBTB16 ZMYM2 ZNF384 ZNF521	TLX1	TLX3	TMPRSS2	TNFRSF11A	TOP1	TP63	ТРМ3	TPM4	TRIM24
	TRIP11	TTL	TYK2	USP6	YPEL5	ZBTB16	ZMYM2	ZNF384	ZNF521

ADGRA2 (GPR124)

NM_032777.9: c.3010_3011delinsAA (p.G1004K) chr8:37698866-37698867 EPHA7

NM_004440.3: c.2024A>C (p.Q675P) chr6:93967903

ERBB2

NM_004448.2: c.3484C>T (p.P1162S) chr17:37884013

LRP1B

NM_018557.2: c.9415C>A (p.P3139T) chr2:141242922

NOD1

NM_006092.2: c.434T>C (p.L145P) chr7:30492599

RAF1

NM_002880.3: c.94A>G (p.I32V) chr3:12660127 RICTOR

NM_152756.3: c.2719C>T (p.R907C) chr5:38953634

WDR90

NM_145294.4: c.2417G>A (p.G806D) chr16:707165

