Schnitzler syndrome associated with *MYD88* L265P mutation



Aaron M. Goodman, MD,^{a,b,c} Philip R. Cohen, MD,^d Alex Li, BA,^e Brian Hinds, MD,^f and Razelle Kurzrock, MD^{a,b} La Jolla and National City, California

Key words: Schnitzler; MYD88.

INTRODUCTION

Schnitzler syndrome (SS) is a rare autoinflammatory disease with unclear etiopathogenesis. The disease is associated with cytokine dysregulation involving interleukin (IL)-1 β and the inflammasome pathway and manifests with major clinical criteria, namely IgM paraprotein and urticarial rash, and minor criteria, including fever, lymphadenopathy, musculoskeletal pain, and thrombosis.¹

Anakinra is a recombinant version of human IL-1 receptor (IL-1R) antagonist and binds to IL-1R, inhibiting IL-1 α and IL-1 β signaling. SS typically responds dramatically to IL-1R blockade.² No recurrent genetic alterations have been identified in SS to date. Interestingly, a case series examining 21 SS patients sequenced 32 autoinflammatory genes without identifying a single genetic alteration in the cohorts.³

SS carries a 15% to 20% risk of an associated lymphoproliferative disorder, similar to that of other conditions with IgM paraproteinemia, including lymphoplasmacytic lymphoma (LPL)/Waldenström macroglobulinemia (WM), and systemic amyloidosis. The precise relationship amongst hematologic disorders is unknown.

WM is an IgM-secreting LPL with clinical features including cytopenias and lymphadenopathy and paraprotein-related manifestations. *MYD88* is a

gene involved in nuclear factor κ light chain enhancer of activated B cells (NF- κ B) signaling.
Mutations in MYD88 (eg, L265P) are the molecular
hallmark of WM and have been identified in more than 90% of WM cases. ⁴ This mutation was not
included in previous molecular panels investigating SS. 3
Here we present the first case, to our knowledge,

Abbreviations used:

interleukin

ESR:

IL-1R:

IRAK:

PD-L1:

LPL:

SS:

WM:

IL:

Here we present the first case, to our knowledge, of SS with *MYD88* L265P mutation and explore the proposed role of this alteration in the pathogenesis of SS. Consent was obtained in accordance with University of California San Diego Institutional Review Board guidelines.

erythrocyte sedimentation rate

lymphoplasmacytic lymphoma

Waldenström macroglobulinemia

Inerleukin-1 receptor-associated kinase

programmed death receptor-ligand 1

interleukin-1 receptor

Schnitzler syndrome

CASE

A 79-year-old man presented with a history of right inguinal lymphadenopathy for 7 years. Initial

From the Department of Medicine, Division of Hematology/Oncology^a and Division of Blood and Marrow Transplantation^c; the Department of Dermatology^f; the Center for Personalized Cancer Therapy^b; and the School of Medicine,^e University of California San Diego and San Diego Family Dermatology, National City.^d

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Neomed, and Actuate Therapeutics and speaker fees from Roche. Dr Kurzrock has stocks in IDbyDNA and has an ownership interest in Curematch Inc. The rest of the authors have no conflicts to disclose.

Correspondence to: Aaron M. Goodman, MD, UC San Diego Moores Cancer Center, 3855 Health Sciences Drive, La Jolla, CA, 92093-0658. E-mail: a1goodman@ucsd.edu.

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Specimen	Morphology	Flow cytometry	Cytogenetics
Right inguinal	Reactive follicular hyperplasia, clusters of	κ -preponderant (κ :λ ratio of 3:1) B cells	3 copies of
node	plasma cells with mature morphology	(43%); T cells exhibit increased CD4:CD8 ratio at 9:1	MALT1
Bone marrow	No morphologic evidence of lymphoma	Monotypic κ-restricted B-cell population identified	N/A
		CD19 ⁺ , CD20 ⁺ , CD22 ⁺ , CD23 ⁺ , CD25 ⁺ , FMC7 ⁺	
		CD5 ⁻ , CD10 ⁻ , CD11c ⁻ , CD103 ⁻	
*Left inguinal node	Follicular hyperplasia with marked polyclonal plasmacytosis	Monotypic κ-restricted B cells, CD5 ⁺ , CD19 ⁺ , CD20 ⁺ , CD22 ⁺ , partial CD23 ⁺ , partial FMC7 ⁺ , CD10 ⁻ , CD38 ⁻ , EBER ⁻ , ISH ⁻ , HHV8 ⁻ , Ki67 high	N/A
Bone marrow	IgM(⁺) plasma cell neoplasm involving hypercellular marrow with trilineage hematopoiesis. No B-lineage cells present.	No definitive B cells and plasma cells; 8.4% small mature T cells with CD4:CD8 ratio of 2.8:1; 1.1% NK cells.	Normal male karyotype
Skin	Interstitial neutrophilic infiltrate with	N/A	N/A

Table I. Pathology specimens (lymph nodes, bone marrow, skin), morphology, flow cytometry, cytogenetics, and genomic profiling/biomarker analysis

Comprehensive genomic profiling and biomarker analysis

Test	Findings
Tissue NGS of lymph node ^{*†}	MAP2K1 P124S
	MYD88 L265P
	ARID1A D1810fs* [†]
	DNMT3A R736C
Blood cfDNA [‡]	TP53 L145P (0.2% cfDNA)
	TERT promoter SNV (7.7% cfDNA)
TMB [†]	1 mutation/mb
PD-1/L1 IHC of lymph node tumor cells [§]	PD-1 expression: low positive
	PD-L1 expression: Negative

cfDNA, cell free DNA; *EBER*, Epstein Barr virus—encoded small RNAs; *HHV8*, human herpes virus 8; *IHC*, immunohistochemistry; *ISH*, in-situ hybridization; *NGS*, next-generation sequencing; *NK*, natural killer; *PD*, programmed death receptor; *TMB*, tumor mutational burden. *Tissue NGS done on left inguinal lymph node as noted.

[†]Performed by Foundation One Heme panel (406 gene panel).

epitheliotropism. No vasculitis.

[‡]Performed by Guardant 360 (73 gene panel).

 $^{\$}$ Performed by Pathline emerge using clone NAT105 for PD-1 staining and clone SP142 for PD-L1 staining.

past workup found an IgM paraprotein, without malignancy, associated with κ light chain restriction in the bone marrow (CD5⁻/CD10⁻ B cells) (Table I).

Over the next 7 years, he suffered from arthralgias, bone pain, and an episodic urticarial eruption. He had a persistent leukocytosis, anemia, and elevated erythrocyte sedimentation rate (ESR). A lymph node biopsy was performed, which found monotypic CD5⁻/CD10⁻ B cells again without malignancy (Table I). Positron emission tomography—computed tomography found mild fluorodeoxyglucose-avid bilateral axillary and inguinal lymph nodes, so he was empirically treated with rituximab monotherapy without improvement.

The patient eventually presented to our clinic with persistent fatigue, chronic anemia, arthralgias, and urticarial rash (Fig 1). Repeat bone marrow biopsy found an IgM⁺ κ -restricted plasmacytosis with hypercellular marrow. No B cells were identified, which excluded LPL/WM. Cytogenetic tests found a normal male karyotype. *MYD88* L265P mutation was identified on next-generation sequencing (Table I). A skin biopsy found interstitial neutrophilic infiltrates compatible with neutrophilic urticarial

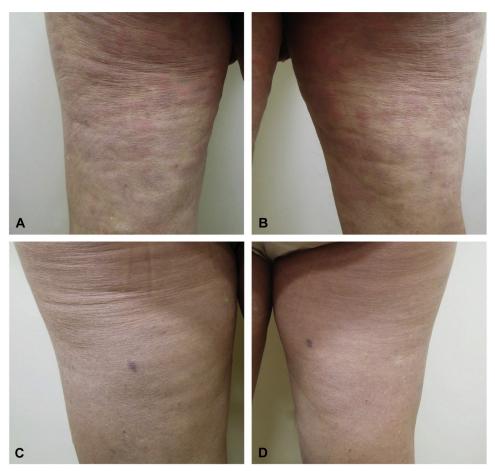


Fig 1. Erythematous urticarial plaques most prominent on the left (**A**) and right (**B**) posterior thighs. Follow-up examination on day 9 after daily subcutaneous anakinra, 100 mg, shows complete resolution of skin lesions. Focal scar and dyspigmentation on the left (**C**) and right (**D**) thighs represent biopsy sites.

dermatosis related to SS (Fig 2). Anakinra, 100-mg subcutaneous daily injections, were started, which within days led to near-complete resolution of rash and arthralgias, and hemoglobin plus ancillary inflammatory markers (thrombocytosis, ESR, and C-reactive protein) normalized (Fig 3). He remains symptom free in clinical remission on anakinra, 100 mg/d, 1 year into treatment.

DISCUSSION

We show a common alteration in LPL/WM that also represents a novel finding in SS. *MYD88* plays an important role in IL-1 signaling, mediating the association between the IL-1R and its IL-1 receptor—associated kinase (IRAK). *MYD88* L265P gain-of-function mutation promotes cell survival by forming a protein complex of IRAK1 and IRAK4, contributing to increased expression of pro-inflammatory pathways.⁵ Given the apparent specificity of the L265P mutation for LPL and role in promoting survival, SS and LPL appear connected and likely represent a disease continuum. Although not previously reported, this mutation may exist in a subset of SS patients who may be at the highest risk of lymphoproliferative disorder development. Therapeutically, Bruton tyrosine kinase inhibitor ibrutinib has shown significant activity in MYD88-mutated LPL; this agent also deserves evaluation in SS.⁶ Further study of the genomics of SS is required with assessment of the prevalence of L265P mutation in a larger cohort.

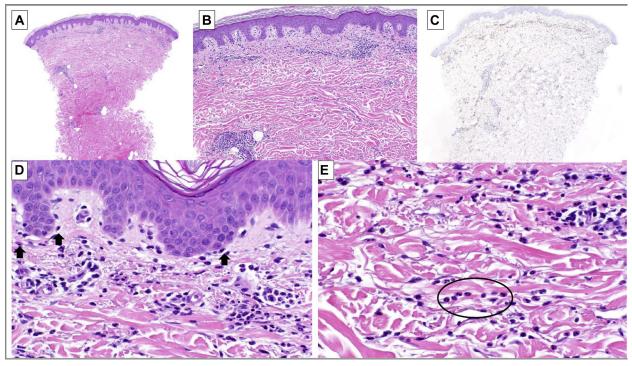


Fig 2. Histopathologic examination found sparse mononuclear cells around the papillary and mid-dermal vascular plexus (**A**). The mononuclear cells exhibit a slight interstitial and perijunctional pattern (**B**). Myeloperoxidase labels the infiltrate confirming myeloid lineage (**C**). Neutrophils are positioned along the dermoepidermal junction. Neutrophilic epitheliotropism (*arrows*) and superficial vessels are patent without fibrinoid necrosis (**D**). Some neutrophils extend in single-file configuration between collagen bundles (*circled*) (**E**). (**A**, **B**, **D**, and **E**, Hematoxylin-eosin stain; original magnifications: **A**, ×40; **B**, ×100; **D** and **E**, ×200; **C**, myeloperoxidase immunoperoxidase; original magnification: ×40).

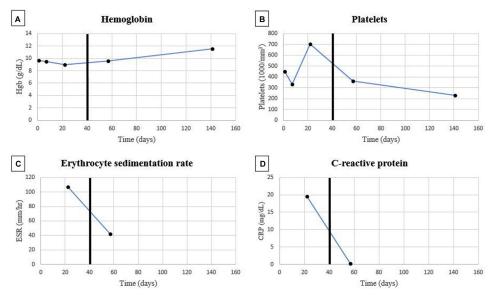


Fig 3. The trends in hemoglobin (**A**), platelets (**B**), ESR (**C**), and CRP (**D**) are shown before and after treatment with 100-mg daily subcutaneous anakinra. The dark black lines represent the start date of anakinra.

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