

Sézary Syndrome (and mycosis fungoides)

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Mycosis Fungoides and Sézary syndrome

- Morphologically indistinguishable skin, blood and lymph node involvement.
- The immunophenotype of SS and most MFs is essentially identical.
- MF can progress to an erythrodermic phase identical to SS.
- MF and SS share the same staging system.
- Both entities are commonly accepted in the same clinical trials.







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Blood Staging in Sézary Syndrome and Mycosis Fungoides



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- Currently, no widely utilized phenotypic, genetic or molecular biomarkers for prognosis.
- Prognosis in CTCL relies largely on tumor
- Quantitative assessment of peripheral blood tumor burden is essential for staging and prognosis.

Α

Probability of Survival (%)

40

IIIB IVA1

Time Since Diagnosis (years)

20 20



ISCL/EORTC staging system for CTCL Prognostic implications

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Basic immunophenotypic features of Sézary cells

- Partially or completely negative in 50-80% of cases. Frequent variable loss in reactive CD4 CD7 T-cell subsets.
- CD26 Partially or completely negative in 80-100% of cases. Frequent variable loss in reactive CD4 T-cell subsets.



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Basic immunophenotypic features of Sézary cells

Positive. Slight dim expression in 40-70% of cases. Occasional partial or complete CD2 negativity.



Positive. Slight dim expression in 40-80% of cases. Rare partial or complete negativity. CD3



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Basic immunophenotypic features of Sézary cells



Basic immunophenotypic features of Sézary cells

- CD4 Positive. Slight dim expression in 30-50% of cases. Rare partial or complete negativity.
- Slightly dim expression inconsistently reported in 10-30% of cases. Rare partial or CD5 complete negativity



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Other useful immunophenotypic features of Sézary cells

Positive on 20% to 80% of cases, might depend on the antibody utilized. CD158k (KIR3DL2)



CD164 Variably overexpressed in most cases.



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Other useful immunophenotypic features of Sézary cells CCR4 and PD-1



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Identification and quantitation of Sézary cells by flow cytometry



Gate on Sézary cells: Phenotypically homogenous cluster of T-cells with features different than

> zary cell (with CBC: mate, in

- % abno x ALC. al cells (of lymphocytes)
- mal cells (of leukocytes) % abno x WBC

Immunophenotypic features of Sézary cells



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Retrospective study on 28 patients with erythrodermic CTCL



Absolute Sézary cell counts by FC can be integtrated into the ISCL/EORTC staging scheme.





Sézary cells often show two phenotypically distinct subsets



Idrees A, Sokol L, Horna P. Poster at: International C Cytometry Society Meeting 2013; Ft Lauderdale, FL

Sézary cells with two phenotypically distinct subsets





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Sézary cells with two phenotypically distinct subsets





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The immunophenotype of Sézary cells can vary depending on the anatomic location

C.

Right thigh

- Amph

В.

Chest

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Naïve/memory phenotype of peripheral blood Sézary cells



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Moins-Teisserenc H, et al. J Invest Dermatol 2015;135:247

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Naïve/memory T-cell phenotype of Sézary cells





Roelens M, et al. Blood 2017;130:1468

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Naïve/memory phenotyope of Sézary cells Is highly variable.

- No difference between MF and SS.
- Can vary with time and depending on anatomic site.
- Likely indicative of a functional/activation state, rather than a "cell or origin".







Feng B, et al. Mod Pathol. 2010;23:284-295.

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- Technically demanding and time consuming.
 Set up of 8 additional tubes after initial T-cell analysis.
- · Not a simple analysis, requires expertise
- · Significant increase in costs.
- 24 additional antibodies
- · Limited ability to analyze phenotypically distinct T-cell subsets.
- 30% of the TCR-Vβ repertoire not covered by the analysis.

2 TCR-Cβ genes: TRBC1 and TRBC2. germline DNA α recombination Their selection is random and mutually exclusive. rearranged 00000 α DNA Similar to kappa and lambda. transcripti splicing protein (T-cell recept translation splicing $V_{\beta 1}D_{\beta 1}J_{\beta}$ CBI 8 = arranged DNA A recombination 7035 Ca

Assessment of T-cell clonality by TCR Cβ restriction

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Assessment of T-cell clonality by TCR Cβ restriction Examples of Sézary syndrome

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Assessment of T-cell clonality by TCR Cβ restriction

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MATO	Shi M, et al. Cytometry B Clin Cytom. 2019 [Epub ahead of print]
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Assessment of T-cell clonality by TCR C β restriction CD4+ T-cells subsets from patients with no T-cell malignancy



CLINIC C	Shi M, et al. Cytometry B Clin Cytom. 2019 [Epub ahead of print]
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Quantitation of clonal CD4 T-cells by TCR C β restriction in patients with CTCL 33 peripheral bloods from 24 CTCL patients



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