

Bilineage involvement of the fusion gene in a patient with FIP1L1-PDGFRA positive chronic eosinophilic leukemia and T-cell lymphoblastic lymphoma: A CASE REPORT



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INTRODUCTION

There are successful attempts treating FIP1L1-PDGFRA positive chronic eosinophilic leukemia (CEL) with low dose imatinib mesylate. However, FIP1L1-PDGFRA fusion gene can induce both proliferation and differentiation of eosinophils, neutrophils, monocytes, lymphocytes and mast cells. A recent report noted three cases with association of FIP1L1-PDGFRA positive CEL and T-cell lymphoblastic lymphoma (T-LBL).

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CASE:

A 42 years-old man addmitted to the hospiltal.

Clinical presentation:

- General fatigue
- Hepatosplenomegaly
- Cervical and axilar lymphadenopathy.

Peripheral blood:

- Leucocytosis 51 x 10⁹/l
- Marked eosinophilia 22 x 10⁹/l
- Evated promyelocytes and myelocites
- No blasts.

Hystological findings:

• Bone marrow biopsy exhibited diffuse eosinophilic hyperplasia with iron deposition, II° reticulin fibrosis and secondary/reactive mastocytosis and minor population of CD3+ TdT+ precursors (<5%): histological features compatible with diagnosis of chronic eosinophilic leukaemia (CEL) with possible minimal T-LBL. The flow cytometry was recommended. (*Figure 2*).

Cervical lymph node biopsy showed T-LBL.

•Immunophenotype: TdT+; CD34-; CD1a+; CD117-; CD3+; CD2+ ; CD4+; CD5+; CD7+; CD10+/-; CD79a-/+; MPO-. (*Figure 1*).







Figure 1: A. HE, 10x. T-LBL: the diffuse infiltration of blastic cells with dense neovascularisation and poorly visible interstitial macrophages ("starry sky" pattern).

B. HE, 40x. T-LBL: the medium sized blastic cells with the scant cytoplasm, ovoid and slightly irregular and lobated nuclei and multiple mitoses.

C. CD3, 20x. The diffuse and strong cytoplasmic positivity of the tumor cells.



Figure 3: RT-PCR analysis of *FIP1L1-PDGFRA* fusion transcripts in lymph node and bone marrow samples. Obtained PCR products (Lanes 2 and 3) correspond to a fusion of *FIP1L1* exon 10 and *PDGFRA* exon 12. *ABL* was used as an endogenous control (Lanes 5 and 6). The sizes of DNA Ladder are indicated.

Treatment:

6 cycles of ACVBP chemotherapy was administered resulting

complete remission of T-LBL

 decrease of eosinophil counts to upper normal range.

However, eosinophilia progressed again to more than 51 x 10⁹/l in a couple of months after completion of chemotherapy.

Molecular diagnostics:

• G-band karyotyping showed no abnormalities and BCR-ABL was absent in the peripheral blood.

• FIP1L1-PDGFRA transcripts (*Figure 4*) by RT-PCR were identified in

• Peripheral blood and bone marrow (*Figure 3*).

• Lymph nodes affected by T-LBL (Figure 3).

Final diagnosis:

FIP1L1-PDGFRA positive chronic eosinophilic and Tlymphoblastic lymphoma. D. TdT, 20x. The diffuse and strong nuclear positivity of the tumor cells.
E. Ki67, 20x. High proliferative activity in the tumor cells (~90%).
F. CD1a, 20x. The diffuse and weak-moderate cytoplasmic positivity of the tumor cells.







Figure 4. Sequencing of PCR product. Sequence analysis of PCR product of rearranged *FIP1L1-PDGFRA* confirmed an in-frame fusion between *FIP1L1* exon 10 and 5'-end truncated *PDGFRA* exon 12 with intervening 6

nucleotides.

Follow up:

• Blood and bone marrow counts normalized after the initiation of imatinib mesylate 100 mg/d.

• 2 years after the final diagnosis the patient is alive in complete clinical, hematological and molecular remission (*Figure 5*) on imatinib 100 mg/d.

CONCLUSION

• FIP1L1-PDGFRA positive CEL in conjunction with T-cell LBL suggests a bilineage cell involvement probably arising from an early hematopoietic progenitor.

• Identification of FIP1L1-PDGFRA in cases of T-LBL with concurrent eosinophilia is essential in selecting the targeted tyrosine kinase therapy.

Figure 2. A. HE, 20x. CEL: the hypercellular bone marrow with the prominent hyperplasia of the eosinophilic lineage.
B. HE, 40x. CEL: diffuse hyperplasia of the eosinophilic lineage with scattered erythroid islands and single megakariocytes.
C. TdT, 20x. The TdT+ (CD3+) precursor population in the bone marrow was elevated, but do not exceed 5% limit in the course of disease.



Figure 5. MRD monitoring of *FIP1L1-PDGFRA* fusion transcripts in PB and BM. Quantification of *FIP1L1-PDGFRA* transcripts was performed as described elsewhere (Jovanovic, J. V. et al. Low-dose imatinib mesylate leads to rapid induction of major molecular responses and achievement of complete molecular remission in FIP1L1-PDGFRA-positive chronic eosinophilic leukemia, Blood 2007, Vol 109, No 11, p4635-4640). The rate of *FIP1L1-PDGFRA* fusion transcripts was normalized to the *ABL* gene. The relative quantities of *FIP1L1-PDGFRA* transcripts at different sampling intervals are indicated.