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Case report: combined myeloid and lymphoid lineage disorders in patient with chronic eosinophilic leukemia and T-cell lymphoma

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Abstract

Aims: We present a rare clinical case of combined myeloid and lymphoid lineage disorders in patients with chronic eosinophilic leukemia and T-cell lymphoma.

Materials: The patient (female, 51) complained of general weakness, a permanent hyperthermia up to 38.5–39°C, constant painful dry cough, and also she expressed redness, peeling and itchy skin on her face. An external inspection showed a significant peripheral lymphadenopathy. A clinical blood analysis revealed the following: leukocytes – $21.9 \times 10^9/L$, segmented neutrophils – 47%, eosinophilia – 17% ($3.7 \times 10^9/L$), lymphocytes – 28% ($6.1 \times 10^9/L$), monocytes – 8%, hemoglobin – 96 g /L, and ESR – 32 mm /h. Up to 44% of eosinophils were found in the myelogram.

Results: The bulk of lymphoid bone marrow cells were represented by a homogeneous population of CD2+CD3+CD4+CD5+ T-cells. A significant portion of the T-cells had an early thymic phenotype CD4+CD8+CD3+. The eosinophils phenotype was represented by markers CD13+CD15+CD33+ with a lack of CD117 and CD23. Molecular biological studies identified chimerical gene transcripts FIP1L1/PDGRF α and monoclonal T-cell receptors. The level of mRNA expression of the interleukin – 5 gene was negative.

Summary: T-lymphocytes represented a clone of transformed cells confirmed by the molecular genetics data (receptor clonality) and flow cytometry (aberrant phenotype). The presence of mutations of the gene FIP1L1/PDGRF α showed the clonal nature of hypereosinophilia. A lack of increase in IL5 proved the lymphoproliferative variant of HES to be incorrect. Thus, using molecular genetics and immunological techniques we verified the presence of two diseases: chronic eosinophilic leukemia and T-cell lymphoma unspecified IVA.

Keywords: hypereosinophilic syndrome, T-cell lymphoma, case report, chronic eosinophilic leukemia