

# A Case of Systemic Mastocytosis Associated with Acute Myeloid Leukemia Terminating as Aleukemic Mast Cell Leukemia after Allogeneic Hematopoietic Stem Cell Transplantation

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In up to 40% of systemic mastocytosis (SM) cases, an associated clonal hematological non-mast cell lineage disease such as AML is diagnosed before, simultaneously with, or after the diagnosis of SM. A 40-yr-old man was diagnosed with AML with t(8;21)(q22;q22). Mast cells were not noted at diagnosis, but appeared as immature forms at relapse. After allogeneic hematopoietic stem cell transplantation (HSCT), leukemic myeloblasts were not observed; however, neoplastic metachromatic blasts strikingly proliferated during the state of bone marrow aplasia, and finally, aleukemic mast cell leukemia developed. As the disease progressed, we observed serial morphologic changes from immature mast cells with myeloblasts to only metachromatic blasts and atypical mast cells as mast cell leukemia; FISH analysis showed that the neoplastic mast cells originated from the same clone as the leukemic myeloblasts of AML.

**Key Words:** Systemic mastocytosis, Acute myeloid leukemia, Aleukemic mast cell leukemia, Allogeneic hematopoietic stem cell transplantation

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## INTRODUCTION

Mastocytosis is defined as a clonal, neoplastic proliferation of mast cells with accumulation in one or multiple organs [1]. The clinical presentation is heterogeneous, ranging from skin-limited disease (cutaneous mastocytosis) that spontaneously regresses to a highly aggressive variant with extracutaneous involvement (systemic mastocytosis, SM) associated with multiorgan failure and short survival [2]. SM with associated clonal hematological non-mast cell lineage disease, which is characterized by the presence of the *KIT* mutation, is the second most common type of SM, representing 40% of cases [2]. AML with *RUNX1-RUNX1T1*, frequently seen in cases of SM with AML, also occasionally has the *KIT*-activating mutational codon D816 [3, 4]. Both SM and *RUNX1-RUNX1T1*-positive AML with the *KIT* mu-

tation at codon D816 have been reported to be refractory to chemotherapy [3, 4]. However, the relationship between the 2 coexisting clonal disorders remains unknown. Here, we describe the clinical and laboratory features of neoplastic mast cell proliferation in a patient with *RUNX1-RUNX1T1* and *KIT*-D816 mutation-positive AML terminating as aleukemic mast cell leukemia after allogeneic hematopoietic stem cell transplantation (HSCT). To the best of our knowledge, no report has described the sequential morphologic alterations of neoplastic mast cells progressing to mast cell leukemia.

## CASE REPORT

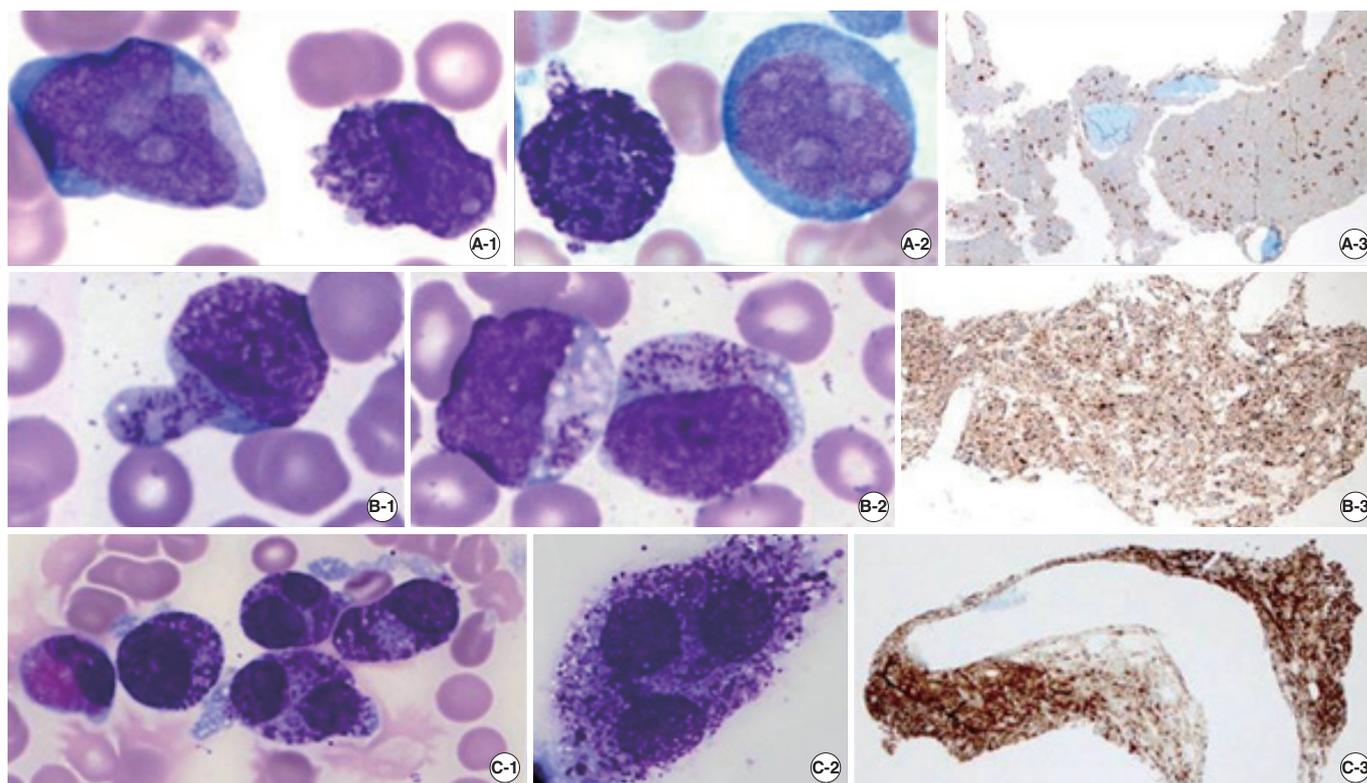
A 40-yr-old Korean man presented with dyspnea on exertion. In the peripheral blood (PB), the white blood cell (WBC) count was

$5 \times 10^9/L$  and consisted of 68% myeloblasts and no mast cells. The bone marrow (BM) was hypercellular and included 90.4% typical myeloblasts and no mast cells. The karyotype was 45,X,-Y,t(8;21)(q22;q22). He was diagnosed with AML with t(8;21)(q22;q22) according to the WHO criteria [1]. He received 1 cycle of combination chemotherapy with idarubicin and cytosine arabinoside and 2 cycles of high-dose chemotherapy with cytosine arabinoside within a period of 5 months. At the end of the first cycle of chemotherapy, complete remission of AML was achieved. The sequential high-dose chemotherapy cycle was complicated with intracranial hemorrhage and severe sepsis, but the patient recovered without any sequelae.

Eight months after the diagnosis, the patient relapsed with clonal evolution. The karyotype was 45,X,-Y,t(8;21)(q22;q22), del(17)(p11.2). In spite of salvage chemotherapy, hematological remission was not achieved. PB showed pancytopenia with WBC

count of  $1.9 \times 10^9/L$ , hemoglobin of 10.6 g/dL, and platelet count of  $28 \times 10^9/L$ . The differential count showed 16% myeloblasts and 1% circulating immature mast cells. BM aspirate smear showed 31.2% myeloblasts and 5.8% immature mast cells. The myeloblasts showed medium-sized, oval- to irregular-shaped nuclei, prominent nucleoli, and scant basophilic cytoplasm (Fig. 1A-1, A-2). Flow cytometric immunophenotyping showed that the leukemic myeloblasts expressed CD34, HLA-DR, CD13, CD33, CD117, CD65, CD15, and CD56, but lacked CD2 (Fig. 2A). BM trephine biopsy revealed persistent leukemic myeloblasts and interstitial infiltration of mast cells. Immunohistochemically, these mast cells were positive for tryptase and CD25 (Fig. 1A-3). Reverse transcriptase (RT)-PCR of the BM aspirate showed *RUNX1-RUNX1T1* fusion transcript and *KIT* mutation at codon 816. The patient was diagnosed with SM-AML with t(8;21)(q22;q22).

He received HSCT from a HLA-haploidentical donor. The con-



**Fig. 1.** Serial findings in peripheral blood (PB), bone marrow (BM) aspirate (Wright stain,  $\times 1,000$ ), and a BM biopsy specimen (tryptase immunohistochemical stain,  $\times 40$ ) at relapse (A), day 29 post hematopoietic stem cell transplantation (HSCT) (B), and day 49 post-HSCT (C). (A-1) A leukemic myeloblast (left) and an immature mast cell with inconspicuous nucleoli (right) in PB. (A-2) An immature mast cell (left), and a leukemic myeloblast (right) in BM aspirate. (A-3) Hypercellular marrow with interstitial infiltration of mast cells in a BM biopsy specimen. (B-1) A metachromatic blast with a high nuclear-to-cytoplasmic (N/C) ratio, fine nuclear chromatin, inconspicuous nucleoli, hypogranular cytoplasm, and variable number of metachromatic granules in PB. (B-2) Metachromatic blasts in BM aspirate. (B-3) Hypercellular marrow with interstitial and focally aggregated infiltration of metachromatic blasts in a BM biopsy specimen. (C-1) and (C-2) Atypical mast cells with bilobed or polylobed nuclei, high or low N/C ratio, relatively dense nuclear chromatin, no nucleoli, pale cytoplasm, and abundant metachromatic granules in BM aspirate. (C-3) Hypercellular marrow with packed infiltration of atypical mast cells in a BM biopsy specimen.

ditioning regimen included busulfan and fludarabine. Twenty-nine days after HSCT, the patient's PB WBC count was  $0.8 \times 10^9/L$  with 17% metachromatic blasts. BM aspirates also contained metachromatic blasts (62.4%) showing a high nuclear-to-cytoplasmic (N/C) ratio, immature nuclear chromatin, inconspicuous nucleoli, and several metachromatic granules (Fig. 1B-1, B-2). Metachromatic blasts infiltrated as interstitial type and focally aggregated in a BM biopsy specimen (Fig. 1B-3).

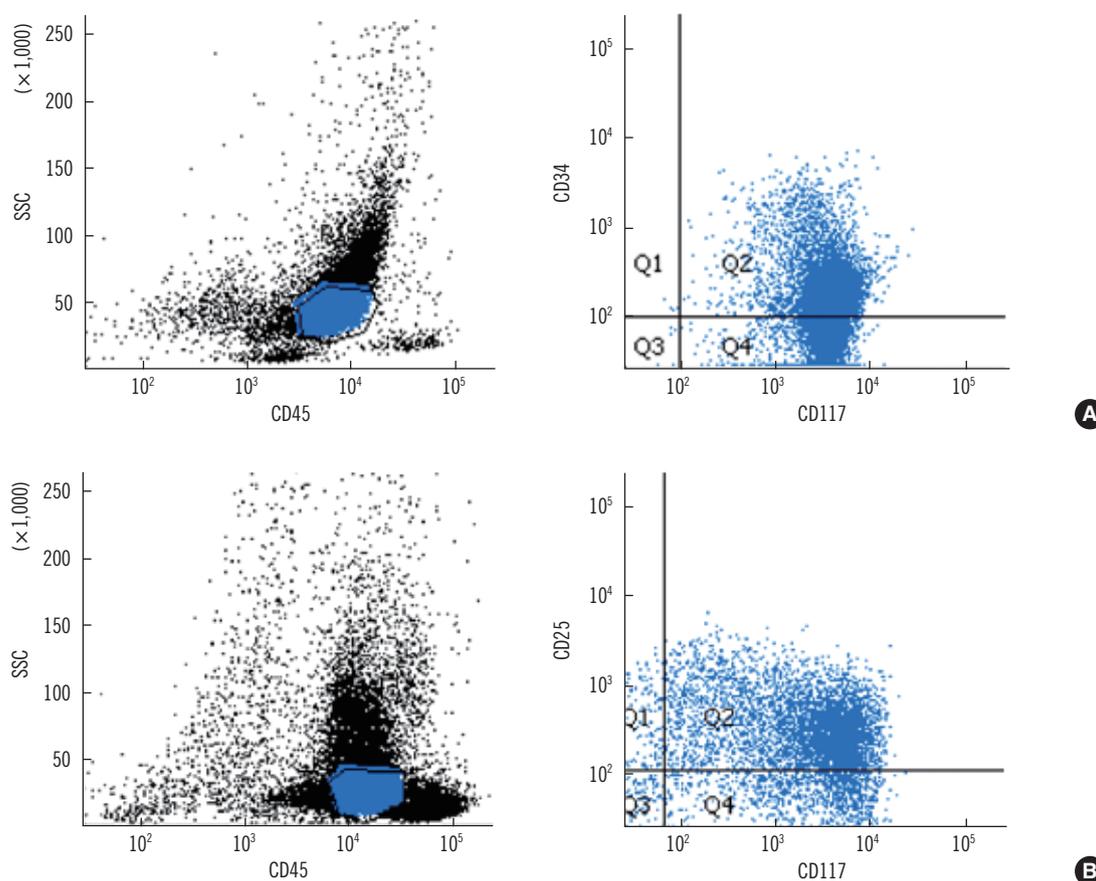
On day 49 after HSCT, pancytopenia worsened, with WBC count of  $0.2 \times 10^9/L$ , hemoglobin of 8.7 g/dL, and platelet count of  $11 \times 10^9/L$ ; however, myeloblasts or mast cells were not seen in the PB. Instead of them, atypical mast cells (type II) constituted 69.2% of all nucleated cells in a BM aspirate smear (Fig. 1C-1, C-2) [5] and were compactly infiltrated in a BM biopsy sample (Fig. 1C-3). Leukemic myeloblasts were not seen in the BM aspirate smear. On flow cytometric analysis, these atypical mast cells expressed CD13, CD33, CD117, CD25, and CD56, but lacked CD34 and HLA-DR, which were expressed strongly at re-

lapse (Fig. 2B). The karyotype was 45,X,-Y,t(8;21)(q22;q22),del(17)(p11.2)[5]/46,XX[15]. FISH using a Vysis RUNX1/RUNX1T1 Dual Color, Dual Fusion Translocation Probe (Abbott Laboratories, Abbott Park, IL, USA) revealed *RUNX1-RUNX1T1* rearrangement as nuc ish(RUNX1T1, RUNX1)x3(RUNX1T1 con RUNX1x2)[58/100] (Fig. 3). The atypical mast cells also showed a *RUNX1-RUNX1T1* fusion transcript and a *KIT* 816 mutation on RT-PCR. At this time, the patient was diagnosed with aleukemic mast cell leukemia.

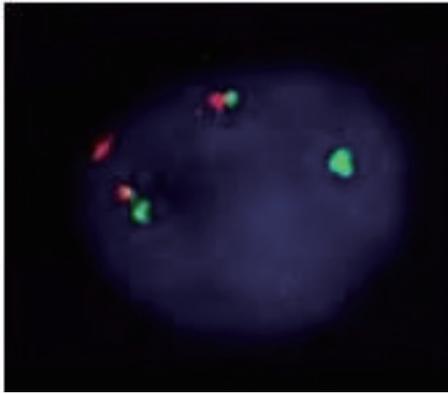
He died 4 months after HSCT because of a new intracranial hemorrhage.

## DISCUSSION

This case was a diagnostic challenge because it appeared to be a case of AML with t(8;21)(q22;q22) without evidence of mastocytosis at the initial diagnosis but terminated as mast cell leukemia after HSCT. In cases of SM-AML with t(8;21)(q22;q22) in



**Fig. 2.** Immunophenotypic findings of bone marrow aspirate at relapse (A) and on day 49 post-HSCT (B). (A) Leukemic myeloblasts with CD45 intermediate expression show the positivity of immature myeloid markers (CD34 and CD117) at relapse. (B) Atypical mast cells with CD45 intermediate to bright expression show the positivity of neoplastic mast cell markers (CD117 and CD25) on day 49 post-HSCT. Abbreviation: SSC, side scatter.



**Fig. 3.** Interphase FISH using Vysis *RUNX1/RUNX1T1* Dual Color, Dual Fusion Translocation Probe (Abbott) on day 49 post-HSCT. The 1 green (*RUNX1* probe on 21q22), 1 orange (*RUNX1T1* probe on 8q22), and 2 fusion signals were consistent with *RUNX1/RUNX1T1* rearrangement.

the literature, diagnoses of SM were made either before or at the same time as the diagnosis of AML or after chemotherapy [6-10]. Our patient was diagnosed with SM-AML at the time of relapse after chemotherapy; however, we cannot exclude the possibility that SM was present at the initial presentation but was masked by excessive blast accumulation.

In its morphologic aspect, the neoplastic mast cell has been classified into 3 subtypes: metachromatic blast and atypical mast cell types I and II [5]. Metachromatic blasts demonstrate blast-like nuclear morphology with several metachromatic granules. Atypical mast cells type I show elongated cytoplasmic projections, oval eccentric nuclei, and hypogranulation, whereas atypical mast cells type II typically have bilobed or polylobed nuclei. In aggressive SM cases and in mast cell leukemia, many neoplastic mast cells exhibit more prominent cellular atypia such as metachromatic blasts or atypical mast cells type II. In our case, we found extremely enhanced proliferation of neoplastic mast cells and morphologically metachromatic blasts without leukemic myeloblasts on day 29 after HSCT. These metachromatic blasts were medium-sized cells with fine nuclear chromatin and pale basophilic cytoplasm containing a variable number of metachromatic granules. On day 49 after HSCT, these cells had transformed to atypical mast cells type II with bilobed nuclei or multiple nuclei, more condensed nuclear chromatin, pale cytoplasm, and coarse metachromatic granules.

*KIT* mutations are associated with enhanced proliferation of mast cells and are frequently detectable in a group of patients with core-factor binding leukemias, such as AML with *inv(16)* or *t(8;21)(q22;q22)*. In such cases, D816V *KIT* is the most common mutation found [11]. We could clearly detect the *KIT*-D816 mu-

tation, not only from the BM aspirate containing 31.2% of leukemic myeloblasts and 5.8% of immature mast cells but also from BM aspirates with 62.4% of metachromatic blasts (day 29 post-HSCT), 69.2% of atypical mast cells (day 49 post-HSCT), and no leukemic myeloblasts.

Published data regarding SM-AML have shown different aspects of the clonal origins of mast cells. The presence of the *KIT* mutation at codon D816 or the *RUNX1-RUNX1T1* rearrangement in mast cells but not in leukemic myeloblasts in the same patient shows that mast cells and leukemic myeloblasts are possibly of different clonal origins or 2 separately developed subclones [6, 7]. However, in other cases, a common clonal origin of mast cells and leukemic myeloblasts has been suggested according to FISH analysis with a probe specific for *RUNX1-RUNX1T1* on sorted mast cells [9, 10, 12]. We proved a D816 *KIT* mutation by using RT-PCR and *RUNX1-RUNX1T1* expression by using FISH in both cell lineages. Thus, we provide evidence that neoplastic mast cells and myeloid leukemic blasts are likely to develop from common hematopoietic progenitors. On the other hand, an article regarding the origin of mast cells after allogeneic HSCT reported that the proliferated mast cells had originated from the donor [13], but in our case, a male patient received a transplant from a female donor, and so we could easily exclude that possibility by cytogenetic analysis.

Clinically significant bleeding can be caused by a high level of heparin after mast cell degranulation [14]. Thus, we conclude that a bleeding diathesis resulting from excessive atypical mast cells is strongly associated with intracranial hemorrhage, the main cause of this patient's death. Several studies about SM-AML reported that after successful HSCT, the neoplastic mast cells gradually decline with time [8-10], probably because of a graft-versus-mast cell effect or gradual apoptosis [9]. Because continued mastocytosis can elevate the risk of bleeding, it may be necessary to consider aggressive interventions like donor lymphocyte infusion [15]. Thus, in cases of SM-AML, whether HSCT is successful or not, a careful morphologic examination of the mast cells is necessary.

In conclusion, we report a case of SM-AML that terminated as aleukemic mast cell leukemia after HSCT, and describe the serial morphologic changes in neoplastic mast cells.

### Authors' Disclosures of Potential Conflicts of Interest

No potential conflicts of interest relevant to this article were reported.

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