Letter to the Editor

Diagnostic Genetics



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PCM1-JAK2 Fusion in a Patient With Acute Myeloid Leukemia

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Dear Editor,

Over 34 myeloid neoplasm cases with a *PCM1-JAK2* fusion have been reported worldwide since 1999, including one case reported in Korea [1, 2]. These cases share common features, such as splenomegaly, eosinophilia, myelofibrosis, and male predominance. Most cases have been diagnosed as myelopro-liferative neoplasm or myelodysplastic/myeloproliferative neoplasm, in particular, as chronic eosinophilic leukemia and atypical chronic myeloid leukemia. However, acute myeloid/lymphoid leukemia cases have also been reported [1]. Therefore, the 2016 WHO revision recognized the *PCM1-JAK2* fusion gene as a provisional entity [3], joining the existing category of "myeloid and lymphoid neoplasms with eosinophilia and abnormalities of platelet-derived growth factor receptor α (*PDGFRB*), or fibroblast growth factor receptor 1 (*FGFR1*)."

A *PCM1-JAK2* diagnosis can be made with or without eosinophilia, if the presence of the genetic rearrangement is proven. However, identification of the genomic breakpoint in the *PCM1-JAK2* fusion is quite difficult because it varies by case; as many as 14 different fusion transcripts from 15 patients have been reported [2, 4-6]. Moreover, the genetic lesions involved in the re-

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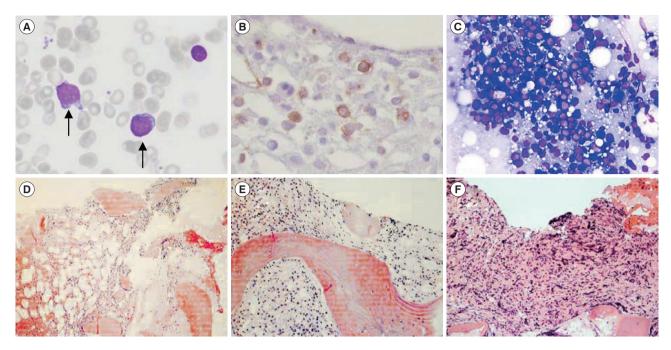


Fig. 1. Bone marrow (BM) findings. (A) Myeloblasts showing less cytoplasm and distinct prominent nucleoli at initial diagnosis (April 2011) (Wright stain, ×400); (B) CD34-positive myeloblasts in the BM biopsy (November 2011) (immunohistochemistry stain, ×400); (C) increased immature cells in touch preparation (March 2012) (Wright stain, ×400); (D–F) serial BM biopsies showing fibrosis progression (April 2011, November 2011, and March 2012, respectively) (hematoxylin & eosin stain ×200).

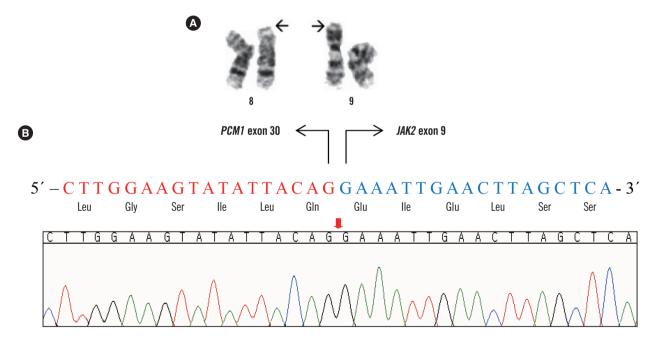


Fig. 2. The t(8;9)(p21;p24) translocations and *PCM1-JAK2* fusion gene. (A) Karyogram of BM showing 46,XX,t(8;9)(p21;p24), (B) Sequence of the chimeric *PCM1-JAK2* gene showing in-frame fusion between exon 30 of *PCM1* and exon 9 of *JAK2*.

classification [7]. A cytogenetic examination could not be performed successfully because of inadequate specimens. The patient experienced complete remission following induction chemotherapy with an idarubicin and cytosine arabinoside regimen and subsequently received three courses of high dose consolidation therapy with cytosine arabinoside. In April 2011, the patient re-developed pancytopenia, which lasted for three weeks; she was transferred to Seoul St. Mary's Hospital, and the BM biopsy confirmed increased myeloblasts (13%; Fig. 1A). The patient received re-induction chemotherapy (FLANG regimen; 30 mg/m²/day fludarabine, 1 g/m²/day cytosine arabinoside, 10 mg/m²/day mitoxantrone, and 300 µg/day G-CSF for five days) twice because of persistent myeloblasts in the BM. After seven months, in November 2011, the BM revealed increased myeloblasts (8%; Fig. 1B). The patient received an allogeneic peripheral blood stem cell transplant from an HLA-matched unrelated donor. Unfortunately, her BM study revealed a relapse of acute myeloid leukemia at the 3-month follow-up post transplantation (myeloblasts 60%, Fig. 1C). In this case, well-known phenotypes of the PCM1-JAK2 fusion, such as splenomegaly and eosinophilia, were not observed; however, myelofibrosis, a morphological feature that matches myeloid neoplasms with PCM1-JAK2 was detected (Fig. 1D, 1E, and 1F). Chromosomal analyses of the specimen in November 2011, demonstrated abnormalities in the short arm of chromosome 8 and 9 (Fig. 2A): 46,XX,t(1;14)(p36.1;q11.2),t(2;6)(q35;p21.1),t(8;9)(p21;p24) [10]/46,XX[3]. We amplified the PCM1-JAK2 fusion transcript from the BM specimen with the detected chromosomal rearrangement by reverse-transcription PCR with primers designed using PRIMER 3 (available from: http://primer3.sourceforge. net): PCM1 exon 28 forward (5'-GAGCGTATGAAGACTG-3') and JAK2 exon 9 reverse (5'-GGCCATGACAGTTGCTTTGT-3'). Sanger sequencing also confirmed an in-frame fusion between PCM1 exon 30 and JAK2 exon 9 (Fig. 2B). Targeted next-generation sequencing, including 46 myeloid neoplasm-associated genes, identified a DNMT3A c.2644C>T, p.Arg882Cys mutation.

The *PCM1-JAK2* fusion, together with the *JAK2* V617F, *MPL* mutations, causes overexpression of the *JAK2* pathway [5]. Therefore, a *JAK2* inhibitor, ruxolitinib, could be used as a therapeutic agent; in fact, several cases with positive treatment efficacy have been reported [8, 9]. Because eosinophilia and myelofibrosis accompany 50–70% of myeloid neoplasms with *PCM1-JAK2* [10], careful genetic evaluation is necessary for adequate diagnosis and the application of targeted therapy, even when a patient does not manifest well-known characteristics.

Authors' Disclosures of Potential Conflicts of Interest

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

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