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# Clinical outcome of patients with acute promyelocytic leukemia and FLT3 mutations

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# Abstract

Acute promyelocytic leukemia (APL) represents a subset of AML with t(15;17), responsiveness to ATRA, and a favorable prognosis, yet the impact of FLT3ITD mutations over wild-type (wt)FLT3 remains unclear. We retrospectively analyzed the outcome of 26 APL patients treated at our center according to their FLT3 receptor mutation status. We show that APL patients with an ITD mutation (n = 9) have a lower fibrinogen at presentation (103.5 vs. 235 mg/dl, p = 0.04) and a worse disease free survival (DFS) (p = 0.0114) but similar overall survival (OS) compared to patients with a wt FLT3 (n = 13). Our data suggests that APL with FLT3ITD represents a subset of APL patients who have a higher risk of relapse and should be treated with aggressive therapies upfront, possibly by including targeted therapy in the form of FLT3 inhibitors.

Acute promyelocytic leukemia (APL) is a distinct subtype of acute myeloid leukemia characterized by promyelocytes in the blood and bone marrow, coagulopathy, and characteristic translocation between chromosomes 15q22 and 17q21 [1]. The t(15;17) results in the formation of a fusion protein between the promyelocytic leukemia gene (PML) and retinoic acid receptor alpha (RARA). The PML-RARA fusion protein results in a block in myeloid differentiation and is capable of inducing APL in murine models [2,3].

APL patients have been classified into different prognostic categories according to their white blood cell (WBC) and platelet count at presentation [4]. Unlike other AML subtypes where the cytogenetics and the presence of molecular mutations form one of the most important prognostic factors, the value of additional karyotypic abnormalities and/or molecular mutations is unclear in these patients [5,6]. Internal tandem duplication mutations within the Fms-like tyrosine kinase receptor (FLT3ITD) are found in ~30% of patients with AML. Survival data for FLT3ITD-APL is limited to a few studies with small numbers and is conflicting, with some studies showing a worse DFS and OS with others not showing a

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correlation between the FLT3ITD and prognosis [7–11]. A recent meta-analysis suggests the FLT3ITD is adversely associated with DFS and OS in patients with APL [12].

We analyzed the clinical outcome of patients with APL to determine the relationship between FLT3 receptor status and prognosis.

There were a total of 26 patients with APL for whom FLT3 mutation status was known. There were 13 patients with wt FLT3, 9 with FLT3ITD mutation (1 also had TKD), and the remaining 4 had an isolated TKD mutation. Of the 13 patients with wt FLT3-APL, four had prior chemotherapy and/or radiation (The first patient had received radiation for early stage breast cancer, the second patient received adjuvant chemotherapy and radiation therapy for breast cancer, the third patient received mitoxantrone for multiple sclerosis, and the forth patient received CHOP chemotherapy regimen for diffuse large B cell lymphoma). No patients in FLT3ITD-APL or FLT3TKD-APL groups had prior chemotherapy or radiation.

Additional cytogenetic abnormalities (ACAs) were seen in 6 patients in the wt FLT3 group (two of these patients had a secondary malignancy), 4 patients in the FLT3ITD group and 2 in the FLT3TKD group. Trisomy 8 was the most common ACA and was found in 3 patients.

Patient characteristics such as age at diagnosis, gender, time to CR, hemoglobin (HgB), platelet count, WBC count at presentation, and coagulation parameters are shown and compared between groups in Table I. One patient in the FLT3ITD and one in the TKD group had microgranular variant APL. Because only a few patients had an isolated FLT3TKD mutation, we restricted our analysis to patients who had either wt FLT3 or FLT3ITD.

The CR rates for patients with wt FLT3 and FLT3ITD were 92 (12/13) and 100% (9/9), respectively. Patients with the FLT3ITD had a lower fibrinogen at presentation than those with wt FLT3 (103.5 vs. 235 mg/dl, P = 0.04). The patients with FLT3ITD appeared to have a higher WBC but it was not statistically significant (P = 0.1). There were no significant differences among the other characteristics described between the two patient groups.

Of the wt FLT3-APL patients, 9/13 were enrolled and treated on CALGB 9710 [13]. Of the remaining 4 patients, two were treated according to the PETHEMA regimen [14], one with a history of breast cancer and significant anthracycline exposure underwent induction with ATRA and AsO<sub>3</sub> [15], and the last patient died before therapy could be initiated. Of the 9 patients with FLT3ITD-APL, 5 were treated on CALGB 9710, 2 were treated according to 9710 but off protocol, and 2 with the PETHEMA regimen [13,14].

We analyzed the relationship between FLT3 mutational status and DFS and OS. We determined that patients with FLT3ITD-APL have an inferior DFS compared to wt FLT3-APL (Fig. 1) (P= 0.0114). However there was no significant difference in OS between the two groups (Fig. 2) (P= 0.3857). One patient of the 13 with wt FLT3-APL relapsed; this patient was treated with AsO<sub>3</sub> reinduction followed by autologous SCT and remains in remission 14 months later. In contrast, 5/9 patients with FLT3ITD relapsed. Of these, 3 received an autologous SCT, one received an allogeneic SCT in CR2, and the fifth patient died within one month of relapse before treatment could be initiated. Two of the 3 patients who received autologous SCT relapsed again and underwent allogeneic SCT in CR3 for one and CR4 for the other.

The ACAs are often observed in APL, although they are not associated with worse prognosis with respect to DFS and OS [4,5,16]. The FLT3ITD mutation is present in 12 to 38% of APL cases [6,9,17], and is associated with the hypogranular variant (M3v) of APL, a

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The exact impact of the FLT3ITD mutation on the prognosis of APL is unclear. In a study of 203 APL patients, FLT3ITD was associated with a higher WBC and a higher incidence of induction-related deaths. However, the remission frequencies, DFS, and OS were similar for ITD and wt patients [7]. Other studies have observed similar CR rates and induction-related deaths regardless of FLT3 mutation status [6,7,17–19]. We observed a similar WBC between ITD and wt patients in our study and the presence of FLT3ITD mutation did not affect CR rates and induction death frequencies in our patient cohort.

Some studies have shown FLT3ITD to be associated with a poor OS in APL patients [6,8,20], however FLT3ITD did not emerge as an independent prognostic factor in multivariable analysis in these studies. A recent metaanalysis by Beitinjaneh et al. showed that FLT3ITD is associated with a poor DFS and OS in patients with APL [12]. Our data confirms the observation that APL patients with a FLT3ITD have a higher relapse risk but differs in that we observe a similar OS given their encouraging outcome following subsequent salvage therapies.

Our findings should be interpreted with caution as our study is limited by small patient numbers, the retrospective nature of our analysis. Notwithstanding these limitations, our data raise the possibility that APL therapy may be improved for patients with FLT3ITD, possibly by incorporating FLT3 inhibitors.

# Methods

#### Patients

We conducted a retrospective analysis of all APL patients who presented to the Dana-Farber Cancer Institute/Brigham and Women's Hospital and Massachusetts General Hospital between 2002 and 2008 who underwent testing for FLT3 mutations under an IRB-approved protocol. We analyzed outcomes for all adult APL patients who were newly diagnosed and were <60 years of age and had been tested for FLT3 mutations. Patient characteristics, CR rates, DFS, and OS were assessed by medical record review under an IRB-approved protocol.

#### **Mutation analysis**

FLT3 mutations, ITD length, sequence and allelic ratio, were determined as previously described [21].

#### End points and statistics

CR was defined by the presence of a normocellular bone marrow containing less than 5% blasts and showing trilineage maturation with an absolute neutrophil count of more than 1000/ul and a platelet count of more than 100,000/ul [22]. DFS is defined as the duration from the date patient achieved CR to the date of relapse or date of death if death occurred prior to relapse. Patients who did not relapse were censored on the last known alive date. OS is defined as the duration from date of diagnosis to date of death or were censored on the last known date alive if patients were still alive as the time analysis. Kaplan-Meier method was used to estimate median DFS and OS and log-rank p-values are presented. We assessed the associations of clinical characteristics and the types of mutation patients had using Kruskal-Wallis test. A p-value <0.05 was interpreted as statistically significant.

#### Acknowledgments

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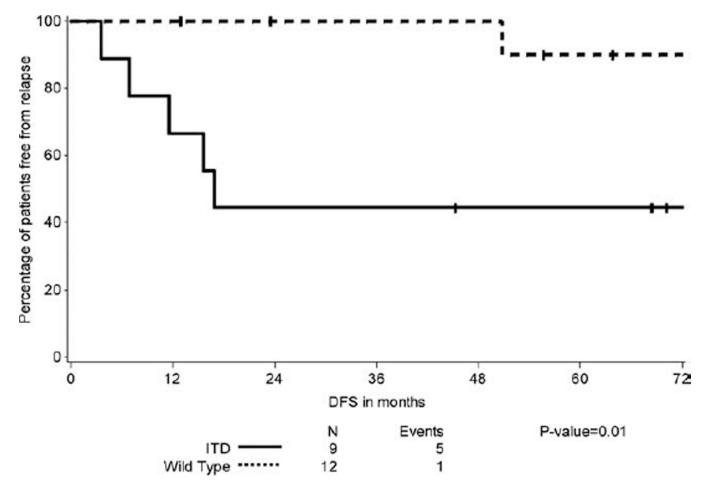
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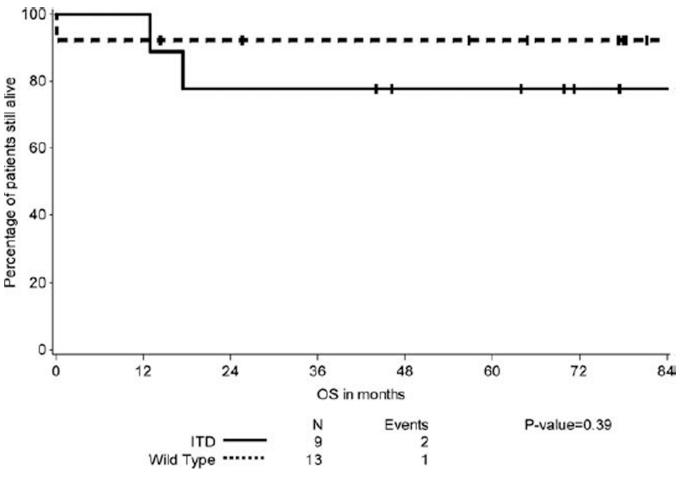
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**Figure 2.** Overall survival of patients with FLT3ITD-APL compared with wt FLT3-APL.

**TABLE I** 

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Comparison of clinical characteristics between wt FLT3 and FLT3ITD patients.

steteNMedian, RangeMedian, Rangegnosis9 $44(19, 60)$ 12 $41(20, 60)$ $n$ le/female)9 $5/4$ 13 $6/7$ $n$ le/female)9 $5/4$ 13 $6/7$ $n$ le/female)9 $1.22(0.89, 1.74)$ 13 $1.27(0.92, 2.5)$ $n$ le10.7(8.6,12.5)13 $10.7(5.2, 12.3)$ $n$ 8 $10.7(8.6,12.5)$ 13 $10.7(5.2, 12.3)$ $n$ 8 $16.4(0.5, 65.7)$ 13 $1.27(0.92, 2.5)$ $n$ 8 $16.4(0.5, 65.7)$ 13 $1.27(0.92, 2.5)$ $n$ 8 $17.8(10.8, 21.4)$ 13 $14.3(11.2, 21.4)$ $n$ 8 $17.8(10.8, 21.4)$ 13 $14.3(11.2, 21.4)$ $n$ 8 $103.5(60, 251)$ 12 $235(62, 381)$ $n$ 9 $0.2578(0.04, 0.85)$ $121.660$			FLT3ITD		wtFLT3	
L Diagnosis944(19, 60)1241(20, 60)r (male/female)9 $5/4$ 13 $6/7$ lo CR1 (months)9 $1.22$ (0.99, $1.74$ )13 $1.27$ (0.92, $2.5$ )b8 $10.7$ (8.6,12.5)13 $10.7$ (5.2, 12.3)t8 $16.4$ (0.5, $65.7$ )13 $1.5$ (0.5, $7.5$ )et8 $38.5$ (9, 123)13 $42$ (8, 204)ec)8 $17.8$ (10.8, $21.4$ )13 $14.3$ (11.2, $21.4$ ) $f(sec)$ 8 $28.3$ (22.1, $42.2$ )13 $27.2$ (21.3, $49.9$ )ogen8 $103.5$ (60, $251$ )12 $235$ (62, $381$ )ultic ratio9 $0.2578$ (0.04, $0.85$ ) $1260$ $1260$	Patient characteristic	Z	Median, Range	Z	Median, Range	P value
tr (male/female)       9 $5/4$ 13 $6/7$ to CR1 (months)       9       1.22 (0.89, 1.74)       13       1.27 (0.92, 2.5) $8$ 10.7.(8.6,12.5)       13       10.7 (5.2, 12.3) $6$ 8       16.4 (0.5, 65.7)       13       1.5 (0.5, 7.5) $6$ 8       16.4 (0.5, 65.7)       13       1.5 (0.5, 7.5) $6$ 8       38.5 (9, 123)       13       42 (8, 204) $ec$ )       8       17.8 (10.8, 21.4)       13       14.3 (11.2, 21.4) $f(sec)$ 8       28.3 (22.1, 42.2)       13       27.2 (21.3, 49.9) $\sigma$ 0       0.35.5 (60, 251)       12       235 (62, 381) $\sigma$ 0       0.2578 (0.04, 0.85)       12       236 (62, 381)	Age at Diagnosis	6	44(19, 60)	12	41(20, 60)	0.88
io CR1 (months)       9       1.22 (0.89, 1.74)       13       1.27 (0.92, 2.5) $8$ 10.7.(8.6,12.5)       13       10.7 (5.2, 12.3) $6$ 8       16.4 (0.5, 65.7)       13       1.5 (0.5, 7.5) $41$ 8       38.5 (9, 123)       13       1.5 (0.5, 7.5) $42$ 8       38.5 (9, 123)       13       42 (8, 204) $ecc$ 8       17.8 (10.8, 21.4)       13       14.3 (11.2, 21.4) $d(sec)$ 8       28.3 (22.1, 42.2)       13       27.2 (21.3, 49.9) $ogen$ 8       103.5 (60, 251)       12       235 (62, 381) $dlicratio$ 9       0.2578 (0.04, 0.85)       10.4       0.85	Gender (male/female)	6	5/4	13	6/7	1.0
b       10.7.(8.6,12.5)       13       10.7 (5.2, 12.3)         b       8       16.4 (0.5, 65.7)       13       1.5 (0.5, 7.5)         st       8       38.5 (9, 123)       13       42 (8, 204)         ec)       8       17.8 (10.8, 21.4)       13       42 (8, 204) $d_{cc}$ 8       17.8 (10.8, 21.4)       13       14.3 (11.2, 21.4) $d_{(sc)}$ 8       28.3 (22.1, 42.2)       13       27.2 (21.3, 49.9)         ogen       8       103.5 (60, 251)       12       235 (62, 381)         olden       9 $0.2578 (0.04, 0.85)$ 12       235 (62, 381)	Time to CR1 (months)	6	1.22 (0.89, 1.74)	13	1.27 (0.92, 2.5)	0.89
8       16.4 (0.5, 65.7)       13       1.5 (0.5, 7.5)         8       38.5 (9, 123)       13       42 (8, 204)         c)       8       17.8 (10.8, 21.4)       13       14.3 (11.2, 21.4)         c)       8       28.3 (22.1, 42.2)       13       27.2 (21.3, 49.9)         gen       8       103.5 (60, 251)       12       235 (62, 381)         elic ratio       9       0.2578 (0.04, 0.85)       233 (66, 231)	Hgb <sup>a</sup>	8	10.7.(8.6,12.5)	13	10.7 (5.2, 12.3)	0.49
8         38.5 (9, 123)         13         42 (8, 204)           8         17.8 (10.8, 21.4)         13         14.3 (11.2, 21.4)           8         28.3 (22.1, 42.2)         13         27.2 (21.3, 49.9)           8         103.5 (60, 251)         12         235 (62, 381)           9         0.2578 (0.04, 0.85)         30 (21.66)	$\mathrm{WBC}^b$	8	16.4 (0.5, 65.7)	13	1.5 (0.5, 7.5)	0.1
8       17.8 (10.8, 21.4)       13       14.3 (11.2, 21.4)         8       28.3 (22.1, 42.2)       13       27.2 (21.3, 49.9)         8       103.5 (60, 251)       12       235 (62, 381)         9       0.2578 (0.04, 0.85)         9       30 (21.66)	Platelet	8	38.5 (9, 123)	13	42 (8, 204)	0.64
8         28.3 (22.1, 42.2)         13         27.2 (21.3, 49.9)           8         103.5 (60, 251)         12         235 (62, 381)           9         0.2578 (0.04, 0.85)         30 (21.66)	PT <sup>C</sup> (sec)	8	17.8 (10.8, 21.4)	13	14.3 (11.2, 21.4)	0.16
8 103.5 (60, 251) 12 9 0.2578 (0.04, 0.85) 9 30 (21, 66)	$aPTT^d$ (sec)	8	28.3 (22.1, 42.2)	13	27.2 (21.3, 49.9)	0.29
6 6	Fibrinogen	8	103.5 (60, 251)	12	235 (62, 381)	0.04
6	ITD Allelic ratio	6	$0.2578\ (0.04,0.85)$			
	ITD length	6	30 (21, 66)			
	$^{b}$ White blood count;					
$b_{m{b}}$ White blood count;	$^{\mathcal{C}}$ Prothrombin time;					
<pre>bWhite blood count; CProthrombin time;</pre>	d <sub>Activated</sub> partial thromb	oplas	tin time.			
$b_{ m W}$ hite blood count; $c_{ m Frothrombin time;}$ d/ctivated partial thromboplastin time.						