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## Targeting FLT3 to treat leukemia

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### Abstract

**Introduction**—Approximately 23% of acute myeloid leukemia (AML) patients younger than 60 years of age carry a mutation in the transmembrane domain of the FMS-like tyrosine kinase-3 (FLT3) gene (FLT3/internal tandem duplications [ITD]). In normal karyotype AML, the presence of a FLT3/ITD mutation is associated with poor prognosis, as mirrored by a high risk of relapse even after allogeneic stem cell transplantation. The poor prognostic impact along with the observation that FLT3 is frequently overexpressed in the majority of AML cases has formed the platform for the development of FLT3-targeted strategies. To date, several FLT3 kinase inhibitors have been investigated in preclinical and clinical studies. However, as of yet, none of the studied FLT3 inhibitors has received FDA approval for routine clinical use in AML. This is in part due to the ‘off target’ effects observed with most inhibitors when administered at concentrations needed to achieve sustained levels of FLT3 inhibition, which are required to exhibit substantial cytotoxic effects against leukemic blasts. Furthermore, the development of resistance mutations has emerged as a clinical issue posing a threat to successful FLT3 inhibitor therapy.

**Areas covered**—In this review, the authors provide a brief summary of FLT3 inhibitors investigated thus far, and discuss current treatment approaches and strategies how to best incorporate FLT3 tyrosine kinase inhibitors (TKIs) into therapy.

**Expert opinion**—The combination of a FLT3 inhibitor with conventional chemotherapeutic regimens, epigenetic modifiers or inhibitors of FLT3 downstream and collateral effectors has emerged as a promising strategy to improve treatment outcome. The future of a tailored, molecular-based treatment approach for FLT3-mutated AML demands novel clinical trial concepts based on harmonized and aligned research goals between clinical and research centers and industry.

### Keywords

acute myeloid leukemia; drug resistance; FMS-like tyrosine kinase-3/internal tandem duplications mutation; hypomethylation

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#### Declaration of interest

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## 1. Introduction

Although the majority of acute myeloid leukemia (AML) cases present with a normal karyotype and therefore fall into the intermediate-risk category, recent advances in gene expression profiling has further honed risk stratification in this subgroup [1]. Herein, internal tandem duplications (ITD) of the juxtamembrane region of the FMS-like tyrosine kinase-3 (FLT3) receptor tyrosine kinase (Figure 1), present in a substantial proportion of AML patients of all age groups, have shown to be an important negative prognostic indicator of disease outcome [2,3], even in older patients [4]. Whereas point mutations in the activation loop of the kinase domain (FLT3/tyrosine kinase domain [TKD]) do not appear to have a significant prognostic impact [5], patients harboring an ITD have a dismal prognosis with a median survival of less than a year [2,3]. Interestingly, early reports could not establish an association between age and the incidence of FLT/ITD mutations. However, most of these studies were either conducted in small patient populations or focused on patients at the age of 60 or younger [3,6]. More recent data suggests that the incidence of FLT3/ITD mutations decreases with age, with an incidence of 16 – 20% in patients older than 60 years as compared to up to 35% in patients between 20 and 59 years [4,7,8].

ITD of the FLT3 receptor gene invariably lead to the constitutive activation of the receptor-tyrosine kinase (RTK) and its downstream signaling effectors, such as RAS/RAF/MEK/ERK kinases, STAT5 and PI3-kinase [9]. As a consequence, altered mechanisms of cellular proliferation and apoptosis promote cell survival thereby conferring a substantial growth advantage to leukemic stem and progenitor cells. From a clinical perspective, this frequently translates into a higher percentage of blood and bone marrow blasts, and a worse overall survival primarily due to a high relapse rate. Lines of evidence suggest that the allelic burden plays a pivotal role in predicting the level of ‘FLT3 addiction’ and clinical outcome [10–12]. For example, Pratz *et al.* showed that low allelic burden FLT3/ITD AML, which appears to more commonly present at the time of initial diagnosis, is less responsive to FLT3 inhibition compared with high allelic burden FLT3/ITD AML, a disease more frequently diagnosed at the time of relapse. Although numerous clinical trials were able to demonstrate that patients with FLT3/ITD AML frequently achieve remission rates similar to other AML patients, the disease usually relapses within a matter months in most cases. Relapsed FLT3/ITD AML represents a portentous clinical situation given the lack of treatment options available to these patients.

In a randomized trial of FLT3-mutated AML patients in first relapse, only 11% of patients with a first remission duration of < 6 months achieved a second remission, and only 29% of patients with a first remission duration of 6 months or longer achieved a second remission when treated with salvage chemotherapy (high-dose cytarabine or mitoxantrone/etoposide/cytarabine) [13] highlighting the urgent need for novel therapeutic strategies to improve the outcome in this patient group. To this end, the role of allogeneic stem cell transplantation (SCT) as a consolidation regimen for FLT3/ITD-mutated patients in first remission has been a topic of controversy among experts in the field [14]. Allogeneic SCT is a treatment modality that offers a potentially higher chance of cure for AML in general, and most studies suggest that allogeneic SCT reduces relapse and improves leukemia-free survival, indicating that this approach may be superior to standard induction and post-remission

chemotherapeutic protocols [15]. However, this approach is associated with a considerable morbidity and mortality rate, particularly in older patients. Based on the analyses of two clinical trial populations in the UK involving 1135 patients, Gale *et al.* did not find good evidence that FLT3 status should guide the decision to proceed with SCT [16]. On the other hand, investigators from a study of 872 cytogenetically normal adult AML patients younger than 60 years from four clinical trial populations in Germany, Austria and Belgium reported that the benefit of SCT is limited to FLT3/ITD<sup>+</sup> patients, as well as to patients harboring wild-type (wt) nucleophosmin 1 and CCAAT/enhancer-binding protein  $\alpha$  (CEBPA) in the absence of FLT3/ITD [17]. Herein, the authors reported a difference in event free but not overall survival with allogeneic SCT. In line with these findings, an analysis of 206 AML patients (n = 120 [FLT3/ITD<sup>+</sup>]; n = 86 [FLT3/ITD<sup>-</sup>]) in first CR (CR1) treated with either HLA-identical sibling or matched unrelated donor SCTs demonstrated an improvement in the 2-year relapse-free survival and leukemia-free survival in the FLT3-mutated group [18]. Differences in outcome with SCT between various trials might be a reflection of the difference in cohorts and subgroups studied. For example, while the study of Gale *et al.* focused on FLT3/ITD-mutated patients, the study by Schlenk *et al.* focused on the cohort of cytogenetically normal AML patients with unfavorable genotypes, including, but not limited to, the FLT3/ITD mutation. In the former study, the rate of allo-SCT during the first period of complete response (CR) was 63% in the setting of a treatment related mortality of 30%, whereas in the latter study these parameters were 82 and 21%, respectively. Moreover, AML patients in the study of Schlenk *et al.* were younger than 60 years, whereas in the study by Gale *et al.* patients older than 60 years were included in the study population. The combined evidence of multiple smaller studies suggest that FLT3/ITD-mutated AML patients in first remission may derive greater benefit from allogeneic SCT compared to patients treated with consolidation chemotherapy. However, this patient population remains at higher relapse risk than transplanted AML patients who are FLT3/ITD<sup>-</sup> [19–21].

## 2. FLT3/ITD<sup>+</sup>AML – the evolution of the disease from diagnosis to relapse and treatment response at relapse

Whole genome sequencing studies of mutation profiles in AML cells indicate that AML is polyclonal at diagnosis, whereas a dominant clone tends to emerge at relapse. This is largely due to the fact that recurrent disease is frequently driven by distinct clonal evolution patterns consisting of either a founding clone that has acquired additional mutations over time and evolved into a relapsed clone, or a sub-clone of the founding clone that was not eliminated by initial therapy, acquired additional mutations and expanded, or both [22]. Growing evidence suggest that tumor cells remain dependent on the same oncogene signaling pathways as their malignant predecessors [23]. This is particularly evident in FLT3/ITD<sup>+</sup>AML, in which the cells at relapse are essentially addicted to signaling from the constitutively activated FLT3 RTK [10]. This addiction to FLT3 signaling is demonstrated by the fact that small molecule FLT3 inhibitors selectively kill AML cells harboring FLT3-activating mutations [24]. *In vitro* studies of these agents predicted that their use would overcome the differentiation block in this AML subtype [25,26]. This was in fact highlighted by several studies with different FLT3 inhibitors in relapsed FLT3/ITD<sup>+</sup> AML, which concordantly reported treatment responses that were associated with terminal myeloid

differentiation, some of them even with clinical manifestations of differentiation syndrome [27–29].

Primary refractory AML is rapidly fatal in most cases [30], whereas relapsed AML occasionally responds to salvage chemotherapy [31]. Recent data on prognostic factors in relapsed AML indicate that cytogenetics, FLT3/ITD status and duration of first remission are the most important predictors of survival in the relapsed setting [32]. FLT3/ITD<sup>+</sup> AML characteristically relapses after a shorter first remission than other AML subtypes [13]. From published data, relapsed/refractory FLT3/ITD<sup>+</sup> AML patients can be categorized into two groups: i) patients who are primary refractory or who have relapsed after a remission duration of 6 months or less and ii) patients who have relapsed after a remission duration of > 6 months. With currently approved therapies, the former group has an expected long-term survival of 0 – 3%, whereas the latter would be expected to achieve a second remission in 20% of the time and to have a 5 – 10% long-term survival [30,31,33]. FLT3 inhibition as a therapeutic modality offers the possibility of clinical benefit in these patient populations through the reduction of tumor volume in a relatively non-toxic fashion, resulting in improved survival either directly or by allowing for allogeneic transplant. The criteria for response to such compounds should be a reflection of the efficacy of this tumor reduction (Table 1).

### 3. FLT3 inhibitors investigated in clinical trials

*Midostaurin (PKC412; N-benzoylstaurosporin)*, a multi-targeted tyrosine kinase inhibitor (TKI) with activity against PKC- $\alpha$ , VEGFR2, KIT, PDGFR and FLT3 tyrosine kinases [34] demonstrated some encouraging anti-leukemic activity in multiple clinical trials. In a Phase IIB trial of 95 patients with either FLT3 wt or mutated AML or myelodysplastic syndrome (MDS), the administration of twice daily 50 mg midostaurin resulted in a greater than 50% reduction of peripheral blood or bone marrow blasts in 42% (wt) and 71% (FLT3 mutant) of patients [35]. When combined with conventional chemotherapy in younger, newly diagnosed AML patients, midostaurin induced high remission and survival rates in both FLT3-mutated and wt patients [36]. The combination of midostaurin with a hypomethylating agent, such as decitabine (DEC), furthermore, demonstrated synergistically enhanced antileukemic activity in a Phase I study in elderly AML patients [37]. Phase III clinical trials of midostaurin are currently ongoing. To this end, the CALGB is currently spearheading an international, randomized, double-blind, placebo-controlled Phase III trial (RATIFY) in treatment-naive FLT3-mutated AML patients younger than 60 years of age (ClinicalTrials.gov Identifier: NCT00651261). This trial encompasses induction chemotherapy (daunorubicin/cytarabine) and four postremission cycles of high-dose cytarabine combined with placebo or midostaurin, followed by midostaurin maintenance or placebo for 1 year. Accrual started in 2008 and completed in 2013 (R. Stone, pers. commun.). The indolocarbazole derivative *lestaurtinib (CEP-701)*, a small molecule inhibitor of JAK2, TrkA, TrkB, TrkC and FLT3, displayed cytotoxic, dose-responsive inhibition of FLT3 *in vitro* and *in vivo*. In addition, the administration of lestaurtinib was associated with an improvement in survival in mouse models of FLT3/ITD-mutated leukemia [38], which altogether provided the framework for further evaluation in clinical trials. Data from Phase I/II trials of single-agent lestaurtinib as salvage treatment for

partially heavily pretreated patients with refractory, relapsed or poor-risk AML carrying FLT3-activating mutations indicated clinical activity in the context of significant reductions of peripheral blood and bone marrow blasts [39]. These findings also held true in a Phase II trial of elderly, newly diagnosed and treatment-naive AML patients considered unfit for intensive chemotherapy [40]. However, combination studies of salvage chemotherapy followed by lestaurtinib in FLT3-mutated AML in first relapse did not show any significant difference in remission rates or benefit in survival. The authors reported that only a small proportion of patients achieved sustained FLT3 inhibition *in vivo* and therefore only limited conclusions regarding the efficacy of FLT3 inhibition combined with chemotherapy could be made [13]. *Sorafenib (Nexavar)*, a small molecule multi-targeted kinase inhibitor targeting VEGFR, PDGFR, Raf kinase and FLT3, has been approved by the FDA for the treatment of advanced renal cell cancer and hepatocellular carcinoma. Studies in Ba/F3 AML cell lines demonstrated that sorafenib effectively induces growth arrest and apoptosis in blasts harboring FLT3/ITD mutations. Furthermore, sorafenib exhibited marked anti-leukemic effects by decreasing the leukemic burden and prolonging median survival (36.5 days [sorafenib] vs 16 days [vehicle]) in a mouse leukemia xenograft model [41] indicating therapeutic efficacy in AML. In an open-labeled single-arm study, the majority of relapsed or refractory FLT/ITD<sup>+</sup> AML patients treated with sorafenib experienced clearance or near complete clearance of bone marrow blasts along with evidence of myeloid differentiation. However, the initial response was lost in most patients after 72 days [28]. When studying the effects of sorafenib combined with idarubicin and cytarabine in FLT3-mutated patients, Al-Kali *et al.* reported overall response rates of 100% (CR = 16/18, CR with incomplete platelet recovery = 2/18). However, after a median follow-up of 9 months more than half of the patients relapsed in the absence of any acquired FLT3 resistance mutations. The authors concluded that the combination of sorafenib with chemotherapy is feasible and effective in inducing remissions but that yet unknown mechanisms of resistance may develop over time [42]. In a Phase II study of relapsed or refractory, FLT3/ITD<sup>+</sup> AML, the combination of sorafenib and the hypomethylating agent azacitidine yielded response rates of 46%, consisting of CR, CR with incomplete count recovery (CRi) and partial response (PR) rates of 16, 27 and 3%, respectively [43], suggesting that the combination of the two drugs may represent a clinically valuable regimen for relapsed, FLT3/ITD<sup>+</sup> AML. *Quizartinib (AC220)*, a second-generation, small molecule inhibitor with activity against FLT3, PDGFR and KIT demonstrated unique selectivity and potency in cell culture assays and animal models [44]. Early phase trials of quizartinib revealed promising results as a single agent in patients with relapsed or refractory FLT3/ITD AML [45,46]. In contrast to other FLT3 inhibitors such as lestaurtinib and midostaurin which inhibit multiple other kinases besides FLT3, quizartinib is much more selective and associated with fewer side effects, less protein bound and therefore up to 50 times more potent *in vivo*. Furthermore, its half-life of greater than 24 h appears optimal given the fact that sustained FLT3 inhibition is essential to exert meaningful cytotoxicity against leukemic blasts [47,48]. In this context, treatment response is mainly represented by a rapid clearance of peripheral blasts through induction of apoptosis and terminal myeloid differentiation in the bone marrow compartment [27]. Preliminary data from several trials suggest that quizartinib, combined with conventional induction and consolidation chemotherapy in younger and older patients with newly diagnosed AML is feasible and safe based upon which multiple Phase III studies are currently being planned.

#### 4. Resistance mutations

Although FLT3 inhibitors induce clinical remissions in the majority of patients, drug-resistant disease frequently develops within the first 12 months of treatment [28,49]. This in some ways parallels the clinical course of chronic myeloid leukemia (CML) treated with imatinib, where imatinib-resistant clones appear in a number of patients during therapy [50–52]. However, while drug resistance in CML is most commonly conferred by point mutations within the Abl kinase domain of the Bcr-Abl gene, the genetic basis for resistance to FLT3 inhibitors in AML blasts are less well understood. Based on studies on AML cell lines and primary cells, distinct patterns of resistance – FLT3-dependent and FLT3-independent – have been described, such as increased expression of the FLT3 receptor and its ligand, acquired mutations in the TKD of FLT3 or in other kinase genes, development of FLT3-independent pathways of signal transduction and increased expression of antiapoptotic proteins [53–57]. The acquisition of secondary FLT3 kinase domain (FLT3 KD) mutations have been recognized as one of the most common mechanisms of resistance to FLT3 inhibitors [49,58–60]. Moore *et al.* established FLT3 inhibitor-resistant cells by incubating MOLM-13 cells in the presence of increasing doses of MLN518, a selective FLT3 inhibitor. After an incubation period of ~ 7 weeks, a D835Y TKD mutation conferring high relative resistance to quizartinib and sorafenib was detected by FLT3 mutational analysis. In contrast, MOLM-13 cells cultured in parallel in the absence of MLN518 were negative for the mutation [61]. FLT3 KD mutations most often occur at residues D835 (F/V/Y) and Y842 (‘activation loop residue’) and F691L (‘gatekeeper residue’), leading to a gain of function in the context of constitutive tyrosine phosphorylation and activation of the RTK through stabilization of the activation loop of the open ATP-binding configuration [5,49]. Data from CML patients treated with imatinib have shown that resistance-conferring mutations in the Abl kinase domain occur prior to imatinib treatment in some patients [62]. Similarly, Man *et al.* demonstrated that FLT3/ITD/TKD clones are in some cases present prior to therapy with a FLT3 inhibitor. In their study of 13 relapsed or chemo-refractory FLT3/ITD-positive AML patients, the authors unveiled potential mechanisms contributing to the nonresponsiveness to sorafenib after an initial response. Utilizing polymerase chain reaction (PCR) assays followed by allele-specific *EcoRV* digestion Man *et al.* reported that a D835Y mutation that was previously not detectable in treatment-naive blasts became detectable upon engrafting in NOD/SCID mice in three out of six patients who have developed resistance to sorafenib, suggesting that the mutation was already present in the leukemia-initiating cell population. Furthermore, in one patient who did not show an ITD-D835 mutation in the leukemia-initiating clone in treatment-naive blasts, the double mutation subsequently emerged and became detectable when the engrafted mice were treated with sorafenib [28]. A novel mechanism of resistance to sorafenib, and potentially other FLT3 inhibitors, was recently reported by Man *et al.* Their work on FLT3/ITD-mutated cell lines demonstrated that tescalcin, a recently discovered protein which is frequently upregulated at leukemia progression, promotes an increase in intracellular pH (pH<sub>i</sub>) by direct interaction with the Na<sup>+</sup>/H<sup>+</sup> exchanger type 1. The authors postulated that lowering the pH<sub>i</sub> might abrogate FLT3 activity, induce apoptosis and foster cell cycle progression [63].

## 5. Patient categories

Emerging data from clinical trials suggest that the incorporation of FLT3 inhibitors into current AML therapy depends on several factors, such as the disease setting (newly diagnosed vs relapsed or refractory) and the intended treatment regimen (single agent vs combined therapy).

## 6. Newly diagnosed FLT3/ITD<sup>+</sup> AML – younger versus older patients

Because the disease consists of multiple leukemic clones at the time of diagnosis and therefore appears to be less FLT3 signaling ‘addicted’ [10], selective FLT3 inhibition by itself is unlikely to achieve clinically significant antileukemic effects. Although conventional induction regimens have yielded similar remission rates in FLT3/ITD<sup>+</sup> and FLT3/ITD<sup>-</sup> AML patients, several studies were conducted with the goal to make the best use of the FLT3 target by incorporating a FLT3 inhibitor into the induction regimen, thus improving treatment response. To this end, a Phase I/II study of sorafenib combined with idarubicin and cytarabine in young adults (age 18 to 65 years) with AML reported that FLT3-mutated patients were more likely to achieve a CR than FLT3 wt patients ( $p = 0.033$ ). In this study, 14 out of 15 patients (93%) with FLT3-mutated AML achieved a CR compared to 24 out of 36 patients (66%) with FLT3 wt disease [64]. In contrast, investigators of a more recent, randomized placebo-controlled trial reported that sorafenib combined with intensive chemotherapy is not beneficial for elderly patients. In this trial, treatment-related mortality and CR rates were indeed higher in the sorafenib arm when compared to the standard induction arm. However, the authors appreciated that the lack of benefit in the sorafenib arm may in part be due to upregulated FLT3 ligand (FL) levels post-chemotherapy and that different scheduling regimens of sorafenib, such as concurrent with chemotherapy, may result in a more favorable outcome [65]. Consequently, whenever a combined approach is used, the involved drugs should be sequenced in accordance to their cell cycle specificity and FL-inducing effects in order to avoid antagonism and optimize synergy [66]. As of yet, the most appropriate treatment regimen for newly diagnosed FLT3/ITD<sup>+</sup> AML remains undefined. Given the enormous molecular and genetic diversity of leukemic clones at presentation, the natural aggressiveness of the disease and the historically poor treatment outcome, intensive treatment protocols consisting of induction chemotherapy followed by SCT in CR1 represents a widely accepted treatment approach, if feasible. While the incorporation of FLT3 inhibitors into these regimens appears as an attractive option, additional clinical trials are warranted to expand our experience with these promising agents. Therefore, any newly diagnosed FLT3-mutated patient, irrespective of age, should be guided to a clinical trial whenever possible.

## 7. Relapsed/refractory FLT3/ITD<sup>+</sup> AML – younger versus older patients

Relapsed or refractory FLT3/ITD<sup>+</sup> AML constitutes a clinical dilemma, irrespective of the patient’s age group. In theory, being derived from an initially declining, heterogenous leukemic clone, the leukemic blasts presenting at relapse are likely more homogenous in their molecular and genetic architecture, more ‘addicted’ to distinct, survival promoting signaling pathways and therefore expected to be more ‘drugable’. However, virtually, all

patients with relapsed/refractory FLT3/ITD<sup>+</sup> AML succumb to their disease within a few months, despite intensive treatment regimens. Lines of evidence suggest that the dismal prognosis at relapse may be correlated with a larger initial tumor burden [67], which is consistent with the highly proliferative nature of FLT3/ITD<sup>+</sup> AML. In a recent study of quizartinib in patients with relapsed/refractory AML who failed second-line chemotherapy or hematopoietic SCT, selective FLT3 inhibition with quizartinib yielded high response rates. In one cohort of FLT3/ITD<sup>+</sup> (n = 99) and FLT3/ITD<sup>-</sup> (n = 38) AML patients older than 18 years with relapsed/refractory disease, single-agent quizartinib induced composite CR rates (CRc, CRc = CR with incomplete platelet recovery [CRp] + CR with incomplete hematologic recovery [CRi]) of 44% with a median duration of 11.3 weeks and median OS of 23.1 weeks in the FLT3/ITD<sup>+</sup> group. The CRc rate in FLT3/ITD<sup>-</sup> patients was 34% with a median response duration of 5.0 weeks and a median OS of 25.6 weeks. In another cohort of elderly patients (60 years and older) with either FLT3/ITD<sup>+</sup> (n = 92) or FLT3/ITD<sup>-</sup> (n = 41) AML, CRc rates of 54 and 32% were observed, respectively. The median CRc duration was 12.7 weeks with a median OS of 25.3 weeks in the FLT3/ITD<sup>+</sup>, and 22.1 weeks with a median OS of 19.0 weeks in the FLT3/ITD<sup>-</sup> group. Combining the cohorts, 79% of FLT3/ITD<sup>+</sup> AML patients who did not respond to prior therapy achieved at least a PR. Moreover, a total of 37% of patients were successfully bridged to an allogeneic hematopoietic SCT and achieved a median OS of 33.3 weeks, whereas patients not bridged to transplant showed a median OS of 17.7 weeks [47,68]. Based on these observations, treatment of relapsed/refractory FLT3/ITD<sup>+</sup> AML patients of all age groups with a selective FLT3 inhibitor such as quizartinib, followed by hematopoietic SCT, if feasible, has emerged as a promising strategy to improve the outcome of a patient population that is known to be very difficult to treat. The clinical value of FLT3 inhibitors as maintenance therapy in the post-transplant setting, as well as for patients unable to undergo SCT, remains to be further evaluated on a wider scale.

## 8. Relapsed/refractory FLT3/ITD<sup>+</sup> with TKI-resistance mutation, *de novo* D835 mutation

In the relapsed or refractory setting where previous treatment may have fostered the development of FLT3-mutated clones, selective FLT3 targeting has come forth as a beneficial strategy. However, resistance-conferring mutations in the FLT3 TKD, commonly at residue D835, are increasingly seen following FLT3 inhibitor therapy. Likewise, FLT3/D835 mutations have frequently been described *de novo* in the absence of ITD mutations, which has furthered the development of second-generation compounds. To this end, ponatinib (AP24534), a multikinase inhibitor initially designed to be effective against TKI-resistant, mutated CML, also demonstrated activity against FLT3/ITD<sup>+</sup> AML in preclinical studies. Herein, Zirm *et al.* reported that ponatinib exerted pro-apoptotic effects against FLT3/ITD and FLT3/TKD (N676D, F691I or G697R) mutated cell lines, which was further mirrored in the potent inhibition of the FLT3/ITD protein as well as its downstream effectors STAT5, AKT and ERK1/2 in western blot assays [69]. Similarly, upon evaluating the *in vitro* effects of ponatinib against TKI-resistance associated, FLT3/ITD mutant isoforms, another group showed that ponatinib conferred antileukemic activity against F691L-mutated blasts but not against blasts carrying D835 mutations, which exhibited a



high level of resistance, suggesting that other treatment strategies are needed for these patients [70].

Crenolanib, a benzamidine quinolone derivative and potent, selective FLT3 inhibitor, has shown promising activity in preclinical and clinical studies [71–73]. In cell culture assays, crenolanib demonstrated substantially higher binding affinity for the dual FLT3/ITD/D835V mutant than quizartinib and conferred pro-apoptotic and anti-proliferative effects in FLT3/ITD<sup>+</sup> cell lines in the nanomolar range. In addition, treatment with crenolanib was associated with improved survival in a FLT3/ITD<sup>+</sup> leukemia murine bone marrow transplant model [73]. When comparing the antileukemic effects of crenolanib to sorafenib and quizartinib in primary AML samples harboring different FLT3 mutations, including the D835 mutation, only crenolanib demonstrated activity, with an IC<sub>50</sub> in FLT3 dephosphorylation of 2 nM in culture medium. Notably, crenolanib is 33-fold less active against c-Kit than against FLT3 and may therefore confer less myelosuppressive effects than quizartinib. This hypothesis was corroborated in colony-forming assays, where 200 nM of crenolanib yielded markedly less inhibition of erythroid colonies than quizartinib at the same dose (abundance of erythroid colonies: 75% [crenolanib] vs 23% [quizartinib]) [72]. Clinical data on crenolanib in 19 heavily pretreated patients with relapsed or refractory, mutated AML (FLT3/ITD, FLT3/D835, FLT3/ITD/D835) were recently reported [74]. In this trial, nanomolar concentrations of crenolanib (median concentration: 473 nM) were well tolerated and associated with a rapid molecular and clinical CR with full count recovery in one patient, a CR with incomplete count recovery in two patients and a PR in four patients. Four patients were bridged to transplant. In sum, crenolanib represents a promising treatment option for FLT3/D835-mutated AML patients. A number of clinical trials to further evaluate the effects of crenolanib in TKI-naive and TKI pretreated FLT3-mutated AML patients are currently under way (NCT01657682, NCT01522469).

## 9. Preclinical and clinical rationales for combined treatment approaches

AML is a disease of the elderly patient. Unfortunately, this patient population has a generally dismal prognosis secondary to a disease biology that frequently arises out of a preexisting disorder, such as MDS. Furthermore, elderly patients are typically unable to tolerate the intensive induction and consolidation regimens used in younger patients. Although younger patients with FLT3/ITD<sup>+</sup> AML seem to benefit from allogeneic transplant, this is not an option for many older patients. An effective treatment regimen for the elderly patient with FLT3/ITD<sup>+</sup> AML, therefore, represents a significant unmet need in this field. Many hematologists and oncologists frequently choose not to aggressively treat elderly patients or to guide them to a clinical trial, due to concerns for reduced tolerance as indicated by a report from the Swedish Leukemia Registry [75]. However, taking into consideration that AML patients who achieve a remission often report improved quality of life and require less hospitalizations and supportive care, one may conclude that remission induction may be beneficial not only from a clinical but also from an economical point of view.

## 10. Hypomethylating agents have proven beneficial in AML, including in elderly patients

Methylation refers to the biochemical process of controlling gene expression without changing the DNA sequence. Deviant methylation of gene promoter CpG islands represent commonly observed epigenetic alterations in cancer cells [76] and may play a critical role in leukemogenesis [77]. Indeed, hypermethylation of tumor suppressor gene promoters, such as ER, P15 and/or P16, have been described in a wide range of human hematopoietic tumors, including AML [78–80]. As a consequence, hypomethylating agents are currently under comprehensive evaluation for their antileukemic potential in AML, either as single agents or in combination with other compounds.

## 11. FLT3 inhibition confers pro-apoptotic effects on peripheral leukemic blasts but induces differentiation in bone marrow blasts

It has recently been shown that potent and selective inhibition of FLT3 results in rapid clearance of peripheral blasts by induction of apoptotic cell death in most cases of FLT3/ITD-mutated AML. However, rather than inducing apoptosis in bone marrow blasts, selective FLT3 inhibition led to terminal myeloid differentiation occurring during the 4 weeks of treatment followed by a surge of leukemia-derived neutrophils in the peripheral blood during the next 4 weeks, as reported by studies conducted by Sexauer *et al.* Although these neutrophils resembled normal neutrophils in morphology and lacked FLT3 protein expression, they still carried the FLT3/ITD mutation which indicated that they were derived from the malignant clone. In line with a differentiation process, flow cytometric analysis revealed loss of expression of CD34 and CD117, along with gained expression of CD15 [27]. From a clinical point of view, the reported neutrophil surge into the peripheral blood was often accompanied by a steroid-responsive differentiation syndrome similar to the differentiation syndrome observed in acute promyelocytic leukemia patients treated with all *trans*-retinoic acid. A recently published case series by Fathi *et al.* further substantiated this finding by reporting that dermatopathologic evaluation of skin nodules presenting after initiation of FLT3 inhibition for FLT3/ITD-mutated AML revealed neutrophilic infiltrates that were uniformly positive for the FLT3/ITD transcript on PCR analysis [29].

## 12. FL surge post-chemotherapy impedes effects of FLT3 inhibitors

FL constitutes an early hematopoietic growth factor and ligand for the FLT3 receptor. FL is predominantly synthesized and released by bone marrow microvascular endothelial cells [81], T cells [82] and by leukemic cells in an autocrine fashion [83]. FL stimulates the proliferation and colony formation of human hematopoietic progenitor cells and acts synergistically with other cytokines such as G-CSF, M-CSF, IL-3 and SCF. As such, highly elevated serum levels of FL have been described in stem-cell-deficient conditions in the bone marrow [84,85] as an attempt to restore a depleted hematopoietic stem cell compartment. A rapid and massive increase in FL levels that is frequently sustained for several weeks has also been described in response to induction and consolidation chemotherapy for AML. This poses a major problem for FLT3-targeted therapy, because FL

acts directly on the mutated receptor, elicits a proliferative response in leukemic blasts [86] and therefore impedes the effects of the most potent FLT3 inhibitors as demonstrated in several clinical trials. For example, in a randomized trial of lestaurtinib administered in sequence with chemotherapy (mitoxantrone, etoposide and cytarabine) for FLT3-mutated AML patients in first relapse (Cephalon 204 trial), the FLT3 inhibitor failed to improve the clinical outcome. In this trial, *in vivo* FLT3 inhibition was observed in only 58% of patients which correlated with the remission rate. Quantification of FL baseline levels in the plasma of the lestaurtinibarm patients were consistently < 20 pg/ml but markedly rose to levels greater than 1000 pg/ml in many cases, and partially remained elevated at the outcome assessment [13]. A similar increase in FL levels was observed in the MRC AML 15 trial, a randomized trial of lestaurtinib administered in sequence with chemotherapy (cytarabine, daunorubicin and etoposide) for newly diagnosed AML patients with FLT3-activating mutations. Here, mean FL levels rose higher with each subsequent course of chemotherapy [57]. Results from both trials indicated that FL levels may remain elevated for several weeks after initiation therapy. *In vitro* studies on Molm14 cells, which harbor the FLT3/ITD mutation, further confirmed the FL induced, obviating effects on FLT3 inhibitors. When incubating Molm14 cells in the presence and absence of exogenous FL, the presence of FL, even at low concentrations (1 ng/ml), was associated with significant autophosphorylation of the FLT3 receptor that cannot be overcome by the most potent FLT3 inhibitors lestaurtinib, midostaurin, sorafenib, quizartinib and KW-2449. Of note, although being constitutively activated in FLT3/ITD, the FLT3 receptor remains highly sensitive to stimulation by FL [87]. It appears therefore likely that successive rounds of chemotherapy promote survival of FLT3/ITD-positive leukemic blasts through upregulation of FL. In line with this hypothesis, many newly diagnosed patients with FLT3/ITD AML achieve remission in response to conventional chemotherapy but eventually relapse during consolidation.

### 13. DNMTis are less intense and do not induce a surge of FL

A number of clinical trials have clearly demonstrated that hypomethylating agents have single-agent activity, consisting of remission induction and improvement of survival and quality of life, in elderly patients with AML and MDS [88–91]. Based on their potential clinical benefits and favorable toxicity profile, hypomethylating agents have become attractive compounds in the treatment of AML, either as monotherapy or combined with other agents. Furthermore, the finding that these agents do not appear to induce large magnitude increases in FL levels [43] translates into a clear rationale of combining DNA methyltransferase inhibitors (DNMTis), as well as other inhibitors from the same drug class, with a potent FLT3 inhibitor. As the degree of FLT3 inhibition closely correlates with the degree of induced cytotoxicity, a combination of a FLT3 inhibitor with an agent that leaves FL levels unaffected offers the theoretic potential to achieve a higher degree of antileukemic activity.

### 14. DNMTis and FLT3 TKIs confer apoptotic and differentiation-inducing effects

Several studies on cancer cells, including AML cell lines and primary AML cells, indicate that hypomethylating agents, at clinically relevant doses, confer pro-apoptotic effects. In one

study, treatment of Kasumi cells with DEC resulted in apoptotic cell death, expression of TNF-related apoptosis-inducing ligand (TRAIL) and demethylation of the CpG-A element [92]. These findings are in line with the report by Lund *et al.*, who found that DEC induces apoptosis in HRAS (G12V) transformed cells by upregulation of endogenous TRAIL in concert with favorable regulation of co-factors acting in TRAIL-mediated apoptosis [93]. Similarly, stimulation of the MDS/AML derived SKM-1S cells with vidaza resulted in upregulation of the pro-apoptotic protein FOXO3A as well as its targets BIM and PUMA [94]. In a different study, treatment of KG-1a cells with either DEC or 5-azacytidine for 48 h resulted in an increased fraction of cells undergoing apoptosis as detected by flow cytometry. Interestingly, DEC-treated cells demonstrated a more potent upregulation of pro-apoptotic markers, whereas a higher degree of cell kill was observed with 5-azacytidine indicating that 5-azacytidine-induced cell death may be mediated by mechanisms other than apoptosis alone [95].

The regulation of apoptosis in FLT-3-mutated leukemic cells is largely regulated by the constitutively activated FLT3 receptor and its downstream effectors STAT5, PI3K-Akt and ERK which jointly interact with the pro- and anti-apoptotic proteins BAD, and Bcl2, Bcl-X<sub>L</sub>, and Mcl-1, respectively. As leukemic transformation is mainly mediated by accelerated proliferation and suppressed apoptosis [96,97], inhibition of FLT3 signaling may theoretically result in increased induction of apoptosis and thereby eliminate leukemic cells. Indeed, FLT3 inhibition by *bis*(1H-indol-2-yl) methanones effectively induced apoptosis in FLT3/ITD transfected murine myeloid cells and in primary FLT3/ITD-positive blasts as indicated by PARP cleavage. Correlative studies demonstrated dose-dependent de-phosphorylation of FLT3/ITD as well as its downstream effectors STAT5, PI3K-Akt and ERK [98]. In a different study, FI-700, a selective FLT3 inhibitor, potently induced G1 arrest and apoptosis in the FLT3/ITD-expressing AML cell lines Molm13 and MV4-11R cells after a 48 h incubation period. On a molecular level, these effects were mediated by rapid and marked downregulation of the anti-apoptotic protein Mcl-1 at 2 h, whereas other Bcl-2 protein members were largely unaffected [99]. Similarly, 24-h treatment of MV4-11 and M14 cells with quizartinib, to date the most effective and selective FLT3 inhibitor, resulted in induction of apoptosis in a dose-dependent manner along with a rapid decrease in FLT3, STAT5 and Erk1/2 phosphorylation [100].

It has been postulated that reduced expression or loss of function of the CEBPA, a major regulator of hematopoiesis, plays a key role in leukemogenesis by inhibiting myeloid differentiation [101,102]. In this context, functional deprivation of C/EBP- $\alpha$  can occur either through aberrant methylation in the upstream promoter [103] or through ERK1/2-mediated phosphorylation [26], or both. *In vitro* studies in Kasumi-1 and CD34<sup>+</sup> RUNX-ETO expressing cord blood cells demonstrated that clinically achievable concentrations of DEC induced morphologic signs of terminal differentiation, while maintaining normal hematopoietic self-renewal capacity. In this study, DEC-induced hypomethylation was greatest at promoter CpGs that are hypomethylated with myeloid maturation and accompanied by cellular differentiation [104]. An *in vitro* comparison of the antileukemic effects of 5-azacytidine and DEC in AML cell lines indicated that both drugs share mechanistic, DNA-specific effects but display different effects on protein synthesis, cell

cycle regulation and gene expression. Gene expression analysis in KG-1a cells using microarrays demonstrated that 5-azacytidine regulated > 1000 genes and was more potent than DEC at two different time points. Moreover, pathway analysis revealed that 5-azacytidine predominantly downregulated genes involving cell cycle, cell division and apoptosis, whereas DEC, in contrast to 5-azacytidine, more potently upregulated genes involved in cell differentiation ( $p = 0.00024$  [vidaza],  $p = 4.4E-12$  [DEC]) [95]. *In vitro* studies on FLT3/ITD-expressing 32D cells demonstrated that inhibition of FLT3 by lestaurtinib induced granulocytic differentiation [105]. This release of the differentiation block appears to represent an ‘on-target’ effect of FLT3 inhibition because *in vitro* differentiation has successfully been induced with several other FLT3 inhibitors [26,27].

To summarize, by virtue of their differentiating effects, their lack of induction of an FL surge, and their relative tolerability, hypomethylating agents represent an intriguing class of drug to combine with FLT3 inhibitors. A number of such trials are actively accruing patients at his time.

## 15. Conclusions

ITD and point mutations in the FLT3 gene constitutively activate the cytokine receptor and lead to uncontrolled proliferation of leukemic blasts by upregulated anti-apoptotic and growth signaling pathways. While remissions, although short lived, can in most cases be achieved with initial induction therapy they are much less likely to occur in the relapsed setting which provides a strong rationale for the current approach to consider consolidation therapy with allogeneic SCT in CR1. Consistent with a complex, adaptive system seen in many aggressive malignancies [106], FLT3-mutated AML evolves from a polyclonal state at diagnosis to a dominant, ‘FLT3 addicted’ clone at the time of relapse. To tackle this issue, several FLT3 inhibitors have been developed and evaluated in clinical trials (Table 2). While a rapid clearance of leukemic blasts in the peripheral blood is frequently seen, most FLT3 inhibitors only confer limited effects on bone marrow blasts. Recent reports on more potent, second-generation FLT3 inhibitors, such as quizartinib, suggest that high levels of sustained FLT3 inhibition induces cell cycle arrest and terminal myeloid differentiation rather than apoptotic cell death. The protective effect of the bone marrow microenvironment on leukemic blasts, partially mediated through persistent activation of survival signaling pathways and expression of FL, represents a significant obstacle in successfully eradicating the disease and founded the framework for combined treatment approaches with other agents. To this end, preclinical and clinical studies identified hypomethylating compounds and MEK inhibitors as promising agents [43,107], particularly for elderly patients who are frequently not able to tolerate intensive treatment regimens. Although the identification of a FLT3 mutation has not yet changed remission-induction and consolidation strategies using approved protocols, the incorporation of a FLT3 inhibitor into these treatment paradigms appear inevitable in the near future. Another treatment paradigm for the use of a FLT3 TKI may be derived from the treatment of Philadelphia chromosome-positive acute lymphocytic leukemia (Ph<sup>+</sup> ALL) and high-risk CML where TKIs are sometimes offered as maintenance therapy after SCT. Herein, several trials demonstrated that the patients receiving a prophylactic TKI after transplant were more likely to remain minimal residual disease (MRD) negative [108–110]. However, prior to the use of a TKI in the post-transplant setting

for remission maintenance therapy several issues need to be addressed, such as the timing of the TKI (early after the transplant vs at the time of MRD detection), the duration of therapy, and the willingness of the patient to undergo frequent MRD monitoring. Several advanced phase, pivotal clinical trials will need to be completed to optimize the management of FLT3/ITD-mutated patients in the newly diagnosed, relapsed/refractory and post-transplant setting.

## 16. Expert opinion

The most effective management of a patient with FLT3/ITD-mutated AML requires an understanding of the disease biology, including the evolution from diagnosis to relapse and the prognostic impact of the mutations in the context of other genetic lesions. Several compounds that target specific signaling pathways have brought major advances in leukemia therapy over the past decade. However, when administered as single agents development of resistance through upregulation of compensatory, collateral pathways is frequently encountered. In addition, bone marrow stromal cells further leukemic cell survival and mediate drug resistance mainly through activation of critical survival pathways such as Akt, Erk and Stat3 [111,112]. In part due to their undeviating response to FL, FLT3/ITD<sup>+</sup> leukemia stem and progenitor cells, furthermore, appear to be better protected by bone marrow stromal cells than their wt counterparts [113]. Lines of evidence suggest that the development of resistance might be delayed or prevented if a combination of different targeted agents is used, irrespective of each agent's activity in the monotherapeutic setting. Hence, combining a FLT3 inhibitor with conventional chemotherapeutic regimens, epigenetic modifiers or inhibitors of FLT3 downstream and collateral effectors has emerged as a potential strategy to overcome *de novo* and acquired resistance. Illustrating the potential benefits of a combined, targeted treatment approach, Weisberg *et al.* reported that the combination of BAG956, a dual PI3K/PDK-1 inhibitor, and midostaurin yielded additive to synergistic antiproliferative effects when tested against FLT3-mutated cell lines and AML primary cells [114]. More recently, Yang *et al.* showed that persistent activation of downstream ERK signaling mediates resistance to FLT3 inhibitor-induced apoptosis in leukemic blasts co-cultured with FL and stroma. In their study, combined inhibition of FLT3 and MEK signaling resulted in synergistic cytotoxicity in relapsed/refractory, and additive cytotoxicity in newly diagnosed FLT3/ITD<sup>+</sup>, primary AML samples [107], thereby providing a strong rationale for a potential benefit of combining FLT3 and MEK inhibitors. The hope is that a significant improvement in prognosis for all FLT3/ITD-mutated patients can be achieved by FLT3 TKI-based treatment protocols. Ongoing clinical trials might not only further our understanding of the molecular events leading to FLT3-mutated AML but also deliver more detailed information about mechanisms of resistance, which altogether will hopefully result in the development of novel targeted agents. The future of a molecular-based treatment approach for FLT3-mutated AML faces the challenge to effectively study novel compounds within small subgroups and molecular entities. When compared to solid malignancies such as breast, prostate, lung or colon cancer, AML presents with a relatively low incidence rate. Hence, large randomized trials conducted by a limited number of centers appear impracticable. In order to identify, recruit and enroll sufficient patient numbers, novel concepts of cooperative research must be employed, which demands closer cooperation between clinical and research centers, and industry. Novel clinical trial designs

based on harmonized and aligned research goals need to supplant traditional clinical trial concepts, where the enrollment of large patient numbers compensated for the potential risks of neglecting individual molecular patient characteristics, and, ultimately, for obtaining inconclusive data. In the future, clinical trials need to aim for small, well-defined and carefully selected patient collectives.

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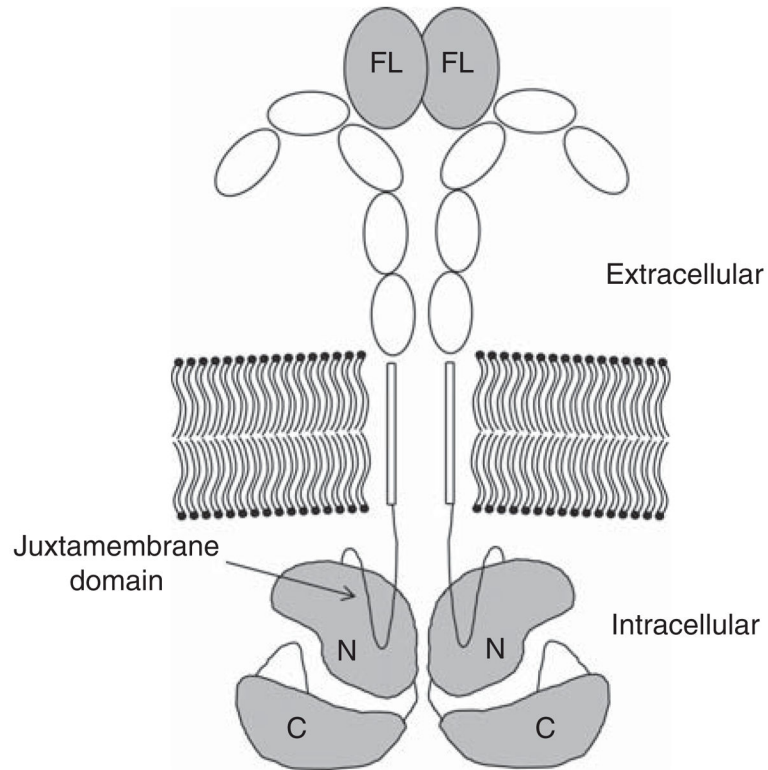
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**Article highlights**

- FMS-like tyrosine kinase-3 (FLT3)/internal tandem duplications mutated AML evolves from diagnosis to relapse.
- FLT3 inhibition induces apoptosis of peripheral blasts but induces differentiation of blasts in the bone marrow.
- Although FLT3 inhibitors induce clinical remissions in most patients, drug-resistant disease develops rapidly.
- The incorporation of a FLT3 inhibitor into current treatment concepts bears the potential to improve outcome.
- DNA methyltransferase inhibitors (DNMTis) do not induce a surge of FLT3 ligand, and they confer apoptotic and differentiation inducing effects and therefore represent an intriguing class of drugs to combine with FLT3 inhibitors.
- Owing to the complexity of AML pathogenesis, future clinical trials need to focus on small, well-defined and carefully selected patient collectives demanding closer cooperation between research centers.

This box summarizes key points contained in the article.



**Figure 1. The FLT3 receptor**

FLT3 ligand (FL) binds in dimeric form to induce receptor dimerization. FLT3/ITD mutations are inserted into the juxtamembrane domain.

C: C-lobe of kinase domain; FLT3: FMS-like tyrosine kinase-3; ITD: Internal tandem duplications; N: N-lobe of kinase domain.

**Table 1**

Proposed classification scheme for response to FLT3 inhibitors used as single agents in patients with relapsed/refractory FLT3/ITD acute myeloid leukemia (AML), based on what is known about the clinical characteristics of the study population and mechanism of action of FLT3 inhibitors.

No response	Persistent blasts in the bone marrow aspirate (Wright–Giemsa stain) and/or core sample, with characterization of leukemia-associated phenotype by multi-parameter flow cytometry, comprising 20% of total cellularity, corresponding to the 2008 WHO classification of acute myeloid leukemia
Minor response	Absence of circulating blasts in the peripheral blood Persistent marrow blasts 5% but < 20% of total cellularity FLT3-mutant allele detectable by polymerase chain reaction (PCR) from a bone marrow aspirate sample
Major response	Absence of circulating blasts in the peripheral blood Minimal residual disease, blasts < 5% of total marrow cellularity FLT3-mutant allele detectable by PCR from a bone marrow aspirate sample
Major molecular response	Absence of circulating blasts in the peripheral blood No morphologic evidence of leukemia by morphology or flow cytometry FLT3-mutant allele not detectable by PCR from a bone marrow aspirate sample

This classification scheme is independent of any platelet or red cell transfusion requirements. The categories of No Response and Minor Response essentially amount to the persistence of disease that is readily discernible by conventional light microscopic techniques. A Major Response is the reduction of the tumor burden to a minimal residual disease (MRD) state, in which the presence of the FLT3/ITD clone is detectable only with multi-parameter flow cytometry and PCR. A Major Molecular Response indicates the absence of any detectable evidence of the FLT3/ITD clone by any conventionally available methodologies. The latter two categories would be expected to be associated with clinical benefit, a hypothesis which could be tested in well-designed randomized trials using overall survival as an end point.

FLT3: FMS-like tyrosine kinase-3; ITD: Internal tandem duplications.



Table 2

FLT3 inhibitors under active clinical investigation either as single agent or in combination.

TKI	Trial phase	Patient population	Single agent/ combination	Ref./Identifier	Status
Sunitinib	I	Patients with all AML FAB classification types, except AML M3 (n = 29)	Single agent	[115]	Completed
	I	Patients with primary or secondary AML; relapsed after at least 1 cycle of conventional chemotherapy; refractory to at least 1 cycle of induction chemotherapy; or not candidates for induction chemotherapy; any AML FAB subtype; at least 18 years of age (n = 16)	Single agent	[116]	Completed
	I/II	Patients aged 60 years or higher with AML with activating FLT3 mutations (FLT3/ITD, FLT3/TKD); fit enough for intensive chemotherapy (n = 22)	Combination with standard induction and consolidation therapy	[117]	Completed
Sorafenib	I	Patients with refractory or relapsed AML; median age 61.5 years; median of 2 prior therapies (n = 16)	Single agent	[41]	Completed
	I	Relapsed, chemotherapy-refractory or frail FLT3/ITD AML patients ineligible for alternative treatments (n = 65)	Single agent	[118]	Completed
	I/II	AML patients younger than 65 years (n = 61)	Combination with conventional cytotoxic chemotherapy (AraC, idarubicin)	[64]	Completed
	II	Elderly AML patients (n = 197)	Combination with AraC/daunorubicin induction followed by consolidation and sorafenib maintenance	[119]	Completed
	I/II	AML by FAB criteria or high-risk MDS defined as IPSS category of intermediate-2 or greater; patients 60 years and older	Combination with low-dose cytarabine	NCT00516828	Completed
	I	FLT3/ITD AML patients who have undergone allogeneic HSCT; age 18 to 75 years	Single agent	NCT01398501	Recruiting
	I	FLT3/ITD AML patients in CR or PR who plan to undergo bone marrow transplantation; age 18 years or older	Single agent	NCT01578109	Recruiting
	I	Relapsed/refractory FLT3/ITD AML; age 18 years and older	Combination with G-CSF and plerixafor/mozobil	NCT00943943	Ongoing, not recruiting
	I	AML patients age 60 years and older and not candidates/refuses standard induction treatment or who have one of the following: poor-risk cytogenetics, AML following antecedent hematologic disorders, or therapy-related AML; patients with relapsed or refractory AML age 18 years are also eligible	Combination with several regimens, including bortezomib and DEC	NCT01861314	Recruiting
	I/II	Patients with newly diagnosed, refractory or relapsed AML, poor-risk or complex cytogenetics, del5, del7 or FLT3/ITD; age 18 years or older	Combination with vorinostat and bortezomib	NCT01534260	Recruiting
	I	Patients with AML or MDS; patients with APL if refractory to ATRA and arsenic trioxide; age 18 years and older	Combination with vorinostat	NCT00875745	Ongoing, not recruiting
	I	Patients with MDS, CMML or AML who have failed prior therapy	Combination with 5-azacitidine	NCT01254890	Ongoing, but not recruiting
	III	Patients with newly diagnosed AML; age up to 29 years	Combination with bortezomib	NCT01371981	Recruiting

TKI	Trial phase	Patient population	Single agent/ combination	Ref./Identifier	Status
Midostaurin	I	Previously untreated AML patients; age 18 – 60 years (n = 69)	Combination with conventional cytotoxic chemotherapy (daunoru-bicin/cytarabine induction, Ara-C consolidation)	[36]	Completed
	II	Patients with FLT3 mutated, relapsed/refractory AML or high-grade MDS; not candidates for chemotherapy; median age of 62 years (n = 20)	Single agent	[120]	Completed
	I	Patients with relapsed or refractory AML; Patients with secondary AML; age 18 years and older	Combination with bortezomib and conventional chemotherapy	NCT01174888	Recruiting
	II	Patients with suspected diagnosis of AML or related precursor neoplasm, or acute leukemia of ambiguous lineage; age 18 – 70 years	Combination with conventional chemotherapy	NCT01477606	Ongoing, not recruiting
	II	Patients with FLT3/ITD-mutated AML who have undergone allogeneic HSCT in CR1; age 18–60 years	Combination with conventional chemotherapy	NCT01883362	Recruiting
	II	Patients with c-KIT or FLT3/ITD mutated t(8;21) AML; age 18 – 65 years	Combination with conventional chemotherapy	NCT01830361	Recruiting
	II	Patients with newly diagnosed AML; except t(15;17); age 60 years and older	Combination with DEC	NCT01846624	Recruiting
	II	Patients with AML, MDS (RAEB-1, -2) or CMML who are either relapsed or refractory to standard therapy, or are considered inappropriate candidates for standard therapy; age 18 years and older	Combination with RAD001	NCT00819546	Ongoing, not recruiting
	I/II	Patients with MDS, CMML, AML or biphenotypic or bilineage leukemia who have failed prior therapy; patients must have evidence of FLT3-activating mutations; age 18 years and older	Combination with 5-azacitidine	NCT01202877	Ongoing, not recruiting
	I/II	Elderly AML patients (APL excluded); age 60 years and older	Combination with 5-azacitidine	NCT01093573	Recruiting
	III	AML patients with activating FLT3 mutations; patients with antecedent MDS allowed, provided they received no prior cytotoxic therapy (including azacitidine or DEC); age 18–59 years	Combination with standard cytotoxic chemotherapy (daunoru-bicin/cytarabine induction with cytarabine consolidation)	NCT00651261	Ongoing, not recruiting
Lestaurtinib	I/II	Patients with refractory, relapsed or poor-risk AML; median age 61 years (n = 14)	Single agent	[39]	Completed
	III	Patients with AML and activating FLT3 mutation (FLT3/ITD or FLT3/D835); age 18 years and older (n = 224)	Combination with standard cytotoxic chemotherapy	[13]	Completed
	III	British MRC trial, AML patients < 60 years considered suitable for intensive chemotherapy (n = 2800)	Combination with standard induction and consolidation regimens	ISRCTN55675535	Completed
Crenolanib	II	Patients with relapsed/refractory primary AML or AML secondary to antecedent hematologic disorder; age 18 years and older	Single agent	NCT01522469	Active, not recruiting
	I/II	Patients with primary AML relapsed or refractory after prior therapy; AML secondary to antecedent chemotherapy or radiation therapy, or AML due to prior MDS/MPN with presence of either FLT3/ITD and/or other FLT3 activating mutation; age 18 years and older	Single agent	NCT01657682	Recruiting
PLX3397	I/II	Patients with primary AML, prior-chemotherapy-related AML or AML secondary to an antecedent hematologic disorder; in at least first relapse or refractory AML; positive for FLT3/ITD; age 18 years and older	Single agent	NCT01349049	Recruiting
Quizartinib	I	Patients with primary or secondary AML (excluding APL), refractory to standard chemotherapy, relapsed after one or more cycles of induction	Single agent	[45]	Completed

TKI	Trial phase	Patient population	Single agent/ combination	Ref./Identifier	Status
		chemotherapy or not candidate for standard chemotherapy; age 18 year or older chemotherapy or not candidate for standard chemotherapy; age 18 year or older			
	II (cohort 1)	Elderly AML patients with relapsed/refractory disease; age 60 years and older (n = 134)	Single agent	[68]	Accrual completed
	II (cohort 2)	AML patients with relapsed disease or refractory to second-line salvage chemotherapy or relapsed after HSCT; age 18 years and older (n = 137)	Single agent	[47]	Accrual completed
	II	Patients with primary AML or AML secondary to MDS; relapsed or refractory after one second-line salvage regimen or after HSCT; FLT/ITD positive; age 18 years and older	Single agent	NCT01565668	Ongoing, not recruiting
	I	Patients with newly diagnosed AML; age 18–60 years	Combination with standard 7+3 induction, high-dose cytarabine consolidation, and quizartinib maintenance	NCT01390337	Ongoing, not recruiting
	I	AML patients who have received a high dose or a reduced intensity conditioning allogeneic HSCT during first or second remission; age 18 years and older	Single agent	NCT01468467	Ongoing, not recruiting
	I/II	Patients with AML (excluding APL and CML in blast crisis) or high-risk MDS; age 60 years and older	Combination with standard daunorubicin, ara-C and etoposide	NCT01236144	Completed
	I/II	AML, MDS or CMML patients with relapsed/refractory disease; age 18 years and older	Combination with azacitidine or cytarabine	NCT01892371	Recruiting

AML: Acute myeloid leukemia; APL: Acute promyelocytic leukemia; ATRA: All *trans*-retinoic acid; CML: Chronic myeloid leukemia; CR: Complete response; DEC: Decitabine; FLT3: FMS-like tyrosine kinase-3; ITD: Internal tandem duplications; MDS: Myelodysplastic syndrome; PR: Partial response; TKI: Tyrosine kinase inhibitor.