



Gilteritinib monotherapy for relapsed/refractory *FLT3* mutated acute myeloid leukemia: a real-world, multi-center, matched analysis

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Abstract

Patients with *FLT3*-mutated relapsed or refractory (R/R) acute myeloid leukemia (AML) have a dismal prognosis. Gilteritinib is a *FLT3* tyrosine kinase inhibitor (TKI) recently approved for patients with R/R AML. We aimed to characterize real-world data regarding gilteritinib treatment in *FLT3*-mutated R/R AML and to compare outcomes with matched *FLT3*-mutated R/R AML patients treated with chemotherapy-based salvage regimens. Twenty-five patients from six academic centers were treated with gilteritinib for *FLT3*-mutated R/R AML. Eighty percent were treated with a prior intensive induction regimen and 40% of them received prior TKI therapy. Twelve patients (48%) achieved complete response (CR) with gilteritinib. The estimated median overall survival (OS) of the entire cohort was eight (CI 95% 0–16.2) months and was significantly higher in patients who achieved CR compared to those who did not (16.3 months, CI 95% 0–36.2 vs. 2.6 months, CI 95% 1.47–3.7; p value = 0.046). In a multivariate cox regression analysis, achievement of CR was the only predictor for longer OS (HR 0.33 95% CI 0.11–0.97, p = 0.044). Prior TKI exposure did not affect OS but was associated with better event-free survival (HR 0.15 95% CI 0.03–0.71, p = 0.016). An age and ELN-risk matched comparison between patients treated with gilteritinib and intensive salvage revealed similar response rates (50% in both groups); median OS was 9.6 months (CI 95% 2.3–16.8) vs. 7 months (CI 95% 5.1–8.9) in gilteritinib and matched controls, respectively (p = 0.869). In conclusion, in the real-world setting, gilteritinib is effective, including in heavily pre-treated, TKI exposed patients.

Keywords Relapsed/refractory AML · *FLT3* · Gilteritinib

Introduction

Acute myeloid leukemia (AML) is the most common type of acute leukemia in adults [1]. Despite significant improvements in risk stratification and treatment, disease relapse occurs in up to 50% of patients [2] and is associated with dismal outcome, especially in patients who are considered

ineligible for allogeneic stem cell transplantation (alloSCT) [3, 4].

Mutations in the *Fms*-like tyrosine kinase receptor 3 (*FLT3*) are found in up to one-third of AML patients (abbreviated as *FLT3*-mutated AML), with 80% of these mutations being internal tandem duplications (ITD) [5], a trait

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associated with inferior prognosis due to higher relapse rate, especially in patients with high mutation allelic ratios [6–8].

Mutations in *FLT3* have been the target for several tyrosine kinase inhibitors (TKI)[9], both in newly diagnosed patients as well as in the relapse or refractory (R/R) disease setting [10–14].

Gilteritinib is a highly potent second-generation, type 1 FLT3 inhibitor that was found to be effective against *FLT3*-mutated AML [15]. Gilteritinib has recently been approved by the FDA and EMA as monotherapy for patients with R/R *FLT3*-mutated AML based on the results of the ADMIRAL phase 3 trial comparing gilteritinib to salvage chemotherapy [16]. Gilteritinib was associated with improved overall survival of 9.3 months vs. 5.5 months in patients in the Standard of care (SOC) group. However, while in the real-world setting most patients with *FLT3*-mutated AML receive midostaurin during induction based on the results of the RATIFY phase 3 trial [17], only 12% of patients in the ADMIRAL trial treated with *FLT3* inhibitor in the upfront setting. Thus, “real-world” analyses of patterns of therapy, response, and safety are warranted.

In this multicenter retrospective cohort study, we aimed to assess and characterize the real-world multi-center data regarding gilteritinib treatment in *FLT3*-mutated R/R AML and to compare outcomes with patients with R/R AML treated with SOC salvage regimens.

Methods

Study population

We conducted a multicenter nationwide retrospective cohort study in six academic centers in Israel. Gilteritinib treatment was identified by searching the electronic medical records systems of all participating centers and cross referencing these data with the departments’ AML database. We included all consecutive patients with R/R AML who had documented gilteritinib monotherapy treatment as salvage therapy between January 2019 and September 2021. Patient, disease, and treatment characteristics were collected using the electronic medical record system. The study was approved by the Institutional Review Board of each center.

In addition, all patients who were treated upfront with intensive regimens were matched 1:1 with R/R AML patients who were treated with SOC salvage regimens from the surveillance database of the participating centers. Matching was performed according to age and European leukemia network (ELN) 2017 risk score criteria [18]. The selection of these parameters was based on their prognostic value [19, 20]. Patients who were treated in the upfront setting with hypomethylating agents (HMA) and venetoclax combination were seldom considered for salvage regimens other than

supportive therapy due to their dismal outlook [21]. Thus, these patients were excluded from the comparative analysis.

Outcomes

Efficacy outcomes included responses according to the recommendations of the European LeukemiaNet and the International working group for assessment and outcomes [18, 22]. Safety data included hematological and nonhematological adverse events (AE), classified according to the CTCAE criteria version 5.0.

Statistics

Categorical variables are presented as numbers and percentages. Continuous variables are presented as mean and standard deviation for normally distributed variables and as median and interquartile range 1–3 (IQR1–3) for non-normally distributed variables. Differences in continuous variables were estimated by *t* test or Mann–Whitney test, as applicable. Differences in categorical variables were estimated by the Fischer exact test. The probability of OS and EFS were estimated by the Kaplan–Meier method. The log-rank test was used to compare survival distributions.

Cox proportional hazards regression models were fitted to predict effect of covariates on OS and EFS in univariable models. Pre-defined covariates of age and type of induction therapy, as well as covariates with a *p* value ≤ 0.05 in the univariate model were retained in the cox regression multivariable model for OS and EFS. AlloSCT post gilteritinib use was calculated as time-dependent variable. All statistics were performed with IBM SPSS, version 28.0 (IBM Corp. Released 2021. *IBM SPSS Statistics for Windows*, Version 28.0. Armonk, NY: IBM Corp) and STATA software version 17.0 (StataCorp. 2021. *Stata Statistical Software: Release 17*. College Station, TX: StataCorp LLC).

Results

Population

Between January 2019 and September 2021, 25 patients were treated with gilteritinib for R/R AML. The median age at AML diagnosis was 61 years (IQR1–3 47–73). Twenty-two patients (88%) harbored *FLT3* mutations at initial diagnosis and 3 patients acquired *FLT3* mutations only during relapse—two with *FLT3-TKD* point mutations and one with *FLT-ITD* mutation. Induction with standard intensive chemotherapy (“7 + 3”) was given to 20 patients (80%) and 8 patients within this group (40%) were given *FLT3* inhibitors as part of their first line therapy (7 midostaurin, 1 quizartinib). Additional patients’ characteristics are presented in

Table 1. Overall, 18 patients (72%) achieved CR post induction therapy and of these, ten patients (56%) proceeded to alloSCT in CR1 (Table 1).

The median time from AML diagnosis to gilteritinib treatment was 12 (range 2–51) months and the median number of previous lines of therapy prior to gilteritinib was 2 (range 1–3), with 11 patients (44%) treated with AlloSCT prior to gilteritinib treatment. Gilteritinib was orally administered in 20 patients (80%) at the full dose of 120 mg, similar to the recommended dose in the ADMIRAL trial, and was reduced to 80 mg in 5 patients (20%) due to cytopenias. The length of each cycle was 28 days and the median number of cycles was 2 (range 1–34).

Response and toxicity

Twelve patients achieved CR (48%), three patients achieved CRi (12%), and ten patients (40%) had stable or progressive disease. After a median time of 7 months post gilteritinib initiation (range 1–34), 23 patients (92%) discontinued treatment. In most cases, this was due to progressive disease or relapse ($n=15$, 60%) followed by four patients (16%) due to adverse effects, three patients (12%) who proceeded to alloSCT and did not continue with gilteritinib and one patient (4%) due to completion of two years of treatment (Fig. 1).

The major hematological toxicity was thrombocytopenia (grade \geq III $n=5$, 20% of patients) and the main non-hematological toxicity was grade II elevated liver enzymes ($n=6$, 24%, Table 2). No clinically meaningful QT interval prolongations were reported as were no events of differentiation syndrome.

Survival

At the time of last follow up, 7 patients (28%) were alive, with 30-day and 60-day mortality rates of 8% ($n=2$) and 28% ($n=7$), respectively. Out of 18 death events, 14 (78%) were attributable to leukemia, two (11%) were caused by infection without active disease, one related to acute heart failure post-transplant, and one to acute GVHD.

The estimated median OS of patients after gilteritinib initiation was 8 (95%CI 0–16.2) months (Fig. 2A). Median OS was higher in patients treated with intensive induction chemotherapy compared to those treated with induction of HMA-venetoclax (9.6 vs. 1.9 months, $p=0.03$). Among patients who achieved CR with gilteritinib, the estimated median OS was significantly higher than patients who did not achieve CR (16.3 (95%CI 0–36.2) months vs. 2.6 months, (95% CI 1.5–3.7); $p=0.046$; Fig. 2B). The estimated EFS in the cohort was 3.5 (95% CI 0–7.9) months, with higher median EFS among patients who achieved CR compared

to those who did not (7 (95%CI 0–14.2) vs. 1.86 (95%CI 0.77–3.0) months, $p=0.045$).

In univariate analysis, initial intensive induction therapy vs. low-intensity regimen predicted better OS with gilteritinib treatment (HR 0.33, CI 95% 0.11–0.98, $p=0.045$). Treatment with a FLT3 inhibitor during first induction and achievement of CR after gilteritinib treatment were associated with better OS, albeit with borderline significance (HR 0.29, 95% CI 0.08–1.01, $p=0.052$ and 0.37, 95% CI 0.14–1.03 $p=0.057$). Age, type of AML (de novo vs. secondary AML), ECOG score, ELN risk category, number of previous lines of treatment, time from first therapy to gilteritinib treatment, and alloSCT (prior to gilteritinib or following gilteritinib) did not significantly impact OS after gilteritinib use (Table 3). In a multivariate model that included CR post gilteritinib and FLT3-targeted therapy during induction, as well as pre-defined covariates—age, and type of induction therapy—only CR achievement post gilteritinib retained its predictive value (HR 0.33 95% CI 0.11–0.97, $p=0.044$).

In a univariate EFS analysis, treatment with FLT3 inhibitor during induction therapy was predictive of better EFS (HR 0.14, 95% CI 0.03–0.61, $p=0.009$). Intensive induction therapy and achieving CR post gilteritinib were associated with better EFS, albeit borderline significance (HR 0.35, 95% CI 0.12–1.0, $p=0.05$ and HR 0.39, 95% CI 0.15–1.02, $p=0.054$, respectively). In a multivariate model that included CR post gilteritinib and FLT3-targeted therapy during induction, as well as pre-defined covariates—age, and type of induction therapy—CR achievement as well as prior treatment with FLT3 inhibitor were associated with better EFS (HR 0.37 95% CI 0.12–0.94, $p=0.037$ and HR 0.15 95% CI 0.03–0.71, $p=0.016$).

Matched analysis of gilteritinib vs. intensive salvage regimens

In order to gain insight into comparative efficacy of gilteritinib in the real-world setting, we conducted a matched analysis that focused on the 20 R/R patients that were initially treated with intensive induction. Patients were matched by age and ELN risk score for the purposes of this analysis.

There were no statistical differences between gilteritinib and SOC groups in terms of baseline characteristics (supplementary S1Table 1) except for the number of previous lines of therapy and the duration from diagnosis to salvage therapy. All the patients in the SOC salvage therapy group (SOC group) were treated with one previous line of therapy, vs. median of two lines of therapy (range 1–3) in the gilteritinib group ($p=0.047$). As a result, the duration between diagnosis and salvage therapy was longer in the gilteritinib group as compared to the SOC group median (median 11 months, IQR1–3 4–37 vs. median 5 months, IQR1–3 1–9, respectively, $p=0.003$).

Table 1 Baseline patient, disease, and treatment characteristics

Patient Characteristics	All (n = 25)	Intensive induction (n = 20)	HMA-Venetoclax (n = 5)
Gender (n, % Male)	13 (52)	10 (50)	3 (60)
Age at diagnosis (median, IQR1-3)	58 (47–73)	56 (40–69)	77 (75–83.5)
Age at gilteritinib treatment (median, IQR1-3)	61 (47–73)	59 (42–72)	80 (77–86)
ECOG (n, %)			
0–1	20 (80)	16 (80)	4 (80)
2–3	5 (20)	4 (20)	1 (20)
Disease characteristics			
AML classification			
De novo AML	18 (72)	16 (80)	2 (40)
AML with AHN	6 (24)	3 (15)	3 (60)
Therapy-related AML	4 (16)	1 (5)	3 (60)
NPM1 mutated (n, %)	11 (44)	9 (45)	2 (40)
FLT3 mutated at diagnosis (n, %)	22 (88)	19 (95)	3 (60)
Type of FLT3 mutation (n, %)			
FLT3-ITD mutated	21 (84)	18 (90)	3 (60)
FLT3-TKD mutated	5 (20)	3 (15)	2 (40)
Karyotype at diagnosis (n, %)			
Normal karyotype	18 (72)	14 (70)	4 (80)
Complex karyotype	2 (8)	2 (10)	0 (0)
Others	2 (8)	1 (5)	1 (20)
Missing	3 (12)	3 (15)	0 (0)
ELN cytogenetic risk criteria (n, %)			
Favorable	6 (24)	4 (20)	2 (40)
Intermediate	10 (40)	9 (45)	1 (20)
Adverse	9 (36)	7 (35)	2 (40)
Treatment prior gilteritinib			
TKIs during 1s ¹ line of therapy (n, %)	8 (32)	8 (40)	0 (0)
Midostaurin	7 (28)	7 (35)	0 (0)
Quizartinib	1 (4)	1 (5)	0 (0)
Achieving CR1 (n, %)	18 (72)	14 (70)	4 (80)
AlloSCT prior to salvage therapy (n, %)	9 (36)	9 (45)	0 (0)
Lines of therapy prior to gilteritinib (n, %)			
1	12 (48)	8 (40)	4 (80)
2	8 (32)	7 (35)	1 (20)
3	5 (20)	5 (25)	0 (0)
Gilteritinib treatment			
Number of cycles (median, IQR1-3)	2 (1–7)	2 (1–7)	1 (1–6)
Maximal tolerated dose (n, %)			
80 mg/day	5 (20)	5 (25)	0 (0)
120 mg/day	20 (80)	15 (75)	5 (100)
AlloSCT post gilteritinib (n, %)	9 (36)	9 (45)	0 (0)
Time from diagnosis to gilteritinib treatment (months, median, IQR1-3)	12 (4–24)	11 (4–37)	14 (5–18)
Time from gilteritinib to last follow up (months, median, IQR1-3)	7 (2–14)	7 (2–14)	2 (1–10)

HMA hypomethylating agents, IQR1-3 interquartile range 25–75%, ECOG eastern cooperative oncology group, AML acute myeloid leukemia, AHN antecedent hematologic neoplasm, NPM1 nucleophosmine 1, FLT3 fms-like tyrosine kinase 3, ITD internal tandem duplication, TKD tyrosine kinase domain, ELN European leukemia network, TKI tyrosine kinase inhibitors, AlloSCT allogeneic hematopoietic cell transplantation

Fig. 1 Gilteritinib treatment by patient. Each bar represents one patient. CR complete response

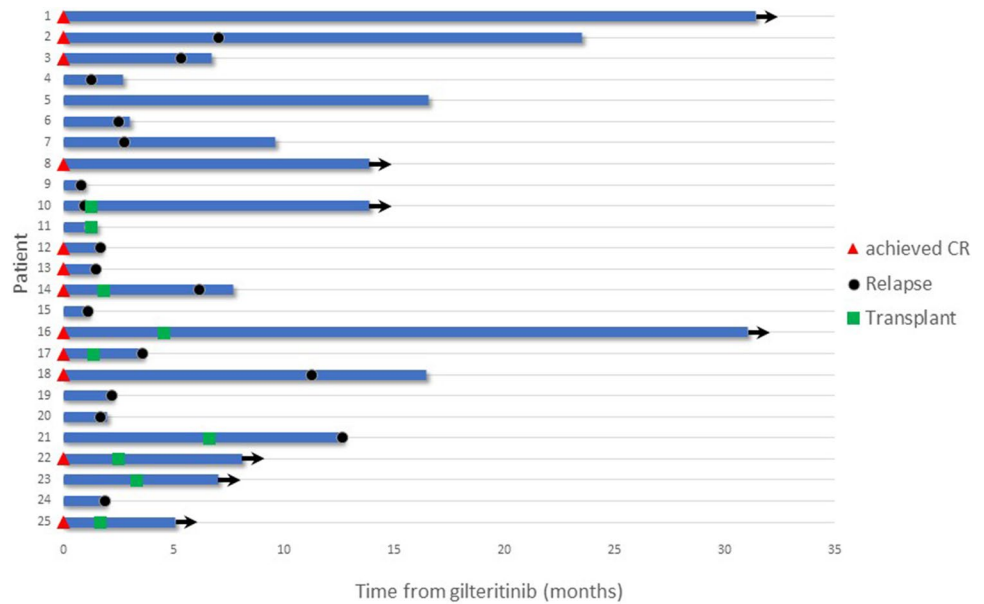


Table 2 Patient responses and toxicity

Responses (n, %)	All patients (n=25)	Intensive induction (n=20)	HMA + venetoclax induction (n=5)
CR	12 (48)	10/20 (50)	2 (40)
CRi/MLFS	3 (12)	2 (10)	1 (20)
PD	10 (40)	8 (40)	2 (40)
Drug discontinuation			
Relapse/progressive disease	15 (60)	12 (60)	3 (60)
Side effects	4 (16)	2 (10)	2 (40)
AlloSCT post gilteritinib	3 (12)	3 (15)	0 (0)
Completed 2 years of treatment	1 (4)	1 (5)	0 (0)
Toxicity			
Thrombocytopenia (grade ≥ III)	6 (24)	4 (20)	2 (40)
Anemia (grade ≥ III)	2 (8)	1 (5)	1 (20)
Infection	2 (8)	1 (5)	1 (20)
Elevated liver enzymes (all grade I–II)	6 (24)	4 (20)	2 (40)
Hyperbilirubinemia	2 (8)	2 (10)	0 (0)
Pancreatitis	1 (4)	1 (5)	0 (0)
Acute kidney injury	1 (4)	0 (0)	1 (20)
Confusion	1 (4)	1 (5)	0 (0)
stomatitis	1 (4)	1 (5)	0 (0)
QT prolongation	0 (0)	0 (0)	0 (0)
Differentiation syndrome	0 (0)	0 (0)	0 (0)

HMA hypomethylating agents, CR complete response, CRi complete response with incomplete recovery, MLFS morphologic leukemia-free state, PD progressive disease, AlloSCT allogeneic stem cell transplantation

Among the SOC group, 50% ($n=10$) were treated with FLT3 TKI during the induction treatment. Salvage consisted of high dose cytarabine and anthracycline in 16 patients (80%), HMA and venetoclax in two patients, one patient was treated with HMA only, and one patient was treated for

refractory AML with sequential alloSCT. Specific regimen details and other SOC group characteristics are shown in Table S1.

Response rates were similar in both groups with ten patients (50%) achieving CR in both groups ($p=1$).

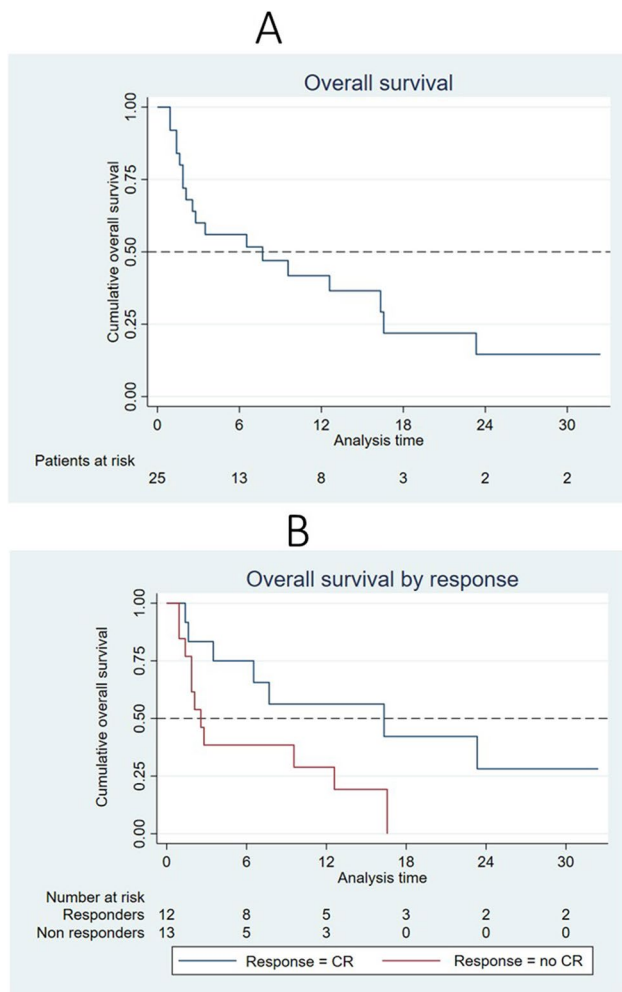


Fig. 2 Kaplan–Meier curve: Overall survival of patients treated with Gilteritinib. **A** Entire group. **B** Comparison of patients who achieved complete remission following Gilteritinib treatment with patient who did not. CR complete response

Likewise, 9 patients (45%) in both the gilteritinib group and in the SOC group proceeded to alloSCT.

The OS among the gilteritinib group was 9.6 (CI 95% 2.3–16.8) months as compared to 7 months (CI 95% 5.1–8.9) in the matched controls (log rank $p=0.869$, Fig. 3A). Similarly, the EFS was 5.1 months (CI 95% 0–10.3) and 3.3 months (CI 95% 0–7.3, Fig. 3B) in the gilteritinib and the SOC group, respectively (log rank $p=0.607$).

Discussion

Herein, we report the results of real-world data regarding treatment with gilteritinib for R/R AML. The median OS in our cohort was 8 months (CI 95% 0–16.2) and is largely comparable to the results demonstrated in the ADMIRAL trial. Achieving CR after gilteritinib and initial treatment

with TKI during induction therapy predicted better EFS following gilteritinib treatment, whereas only the former was predictive of OS as well. In a matched analysis of gilteritinib vs. SOC intensive regimens, the median OS was slightly longer with gilteritinib but did not reach statistical significance.

The outcome of patients with R/R *FLT3-mutated* AML is extremely dismal. Analysis of a French registry demonstrated a median survival of 7 to 8 months for R/R patients with *FLT3-ITD* in the era before second-generation *FLT3* inhibitors were utilized for this indication [23, 24]. The ADMIRAL trial demonstrated that monotherapy with gilteritinib was associated with more than doubling of the remission rates for patients with R/R *FLT3-mutated* AML as compared to salvage chemotherapy (54.3% vs. 21.8%, respectively). This was translated into significantly improved survival in patients treated with gilteritinib (median overall survival of 9.3 vs. 5.6 months, respectively; HR 0.64 (95% CI, 0.49–0.83)) and was the basis for the approval of gilteritinib for this indication by the FDA and EMA in 2018 and 2019, respectively. The results of a similar phase 3, randomized open-label study in Asia (the COMMODORE trial) were recently reported and showed similar superiority for gilteritinib over SOC salvage chemotherapy [25].

Patients in our analysis differed from those in the ADMIRAL trial. One-third of patients analyzed in the current study (and 42% of intensively treated patients) received midostaurin during induction as compared to only 12% of patients in the ADMIRAL trial. Furthermore, the median number of previous lines of therapy prior to gilteritinib was 2 (range 1–3) in our analysis, over one-third of patients had ELN adverse risk disease (as compared to only 10% in the ADMIRAL), and 44% of patients in our analysis received gilteritinib for post-alloSCT relapse as compared to 20% in the registration trial. In this heavily pre-treated patient population, the survival was at a median of 8 months. Survival was estimated at 9.6 months for patients that were initially deemed eligible for intensive therapy and was higher (though not statistically different) from the 7 months reported in matched controls. Of note, this match analysis focus on intensively treated patients with the gilteritinib group having more previous lines of therapy that may account for the blunted difference in outcome between the groups as compared to the ADMIRAL trial.

Patients who responded and achieved remission with gilteritinib derived the most benefit from therapy with an estimated median OS of 16.3 months and predictors of response for this patient population may aid in guiding therapy assignment.

The relative efficacy of gilteritinib in patients previously exposed to *FLT3* inhibitors in our analysis is reassuring and is in line with a recent retrospective analysis [26]. In

Table 3 COX Regression analysis for overall survival and event-free survival in patients treated with gilteritinib as salvage therapy

Overall survival		
<i>Univariate analysis (Hazard ratio, CI 95%, p value)</i>		
Age at gilteritinib use	1.02 (0.99–1.05)	0.25
de novo AML vs. secondary AML	0.76 (0.28–2.04)	0.56
ELN 2017 risk stratification (adverse vs. favorable and intermediate risk groups)	1.70 (0.63–4.58)	0.29
ECOG score (0–1 vs. 2–3)	0.90 (0.29–2.75)	0.85
First line of treatment (HMA + venetoclax compared to “7 + 3”)	3.01 (1.03–8.82)	0.045
TKI use in first line therapy	0.29 (0.08–1.01)	0.05
AlloSCT prior to gilteritinib treatment	0.92 (0.34–2.46)	0.87
AlloSCT following gilteritinib therapy	0.44 (0.14–1.34)	0.15
Number of previous lines (compared to 1)	1.37 (0.45–4.13)	0.46
2 lines of therapy	1.68 (0.53–5.29)	0.58
3 lines of therapy		0.38
Achieving CR after gilteritinib therapy	2.68 (0.97–7.39)	0.057
<i>Multivariate analysis (Hazard ratio, CI 95%)</i>		
Age at gilteritinib use	1.01 (0.98–1.04)	0.627
First line of treatment (HMA + venetoclax compared to “7 + 3”)	2.07 (0.55–7.85)	0.29
TKI use in first line therapy	0.36 (0.10–1.34)	0.13
Achieving CR after gilteritinib therapy	0.33 (0.11–0.97)	0.044
Event free survival		
<i>Univariate analysis (Hazard ratio, CI 95%, p value)</i>		
Age at gilteritinib use	1.01 (0.98–1.04)	0.74
de novo AML vs. AHM-AML or therapy related AML	0.58 (0.21–1.57)	0.28
ELN 2017 risk stratification (adverse vs. favorable and intermediate risk groups)	1.40 (0.55–3.59)	0.48
ECOG score (0–1 vs. 2–3)	0.99 (0.32–3.08)	0.99
First line of treatment (HMA + venetoclax compared to “7 + 3”)	2.89 (1.0–8.40)	0.051
TKI use in first line therapy	0.14 (0.032–0.61)	0.009
AlloSCT prior to gilteritinib treatment	0.77 (0.29–2.08)	0.61
AlloSCT following to gilteritinib therapy s	0.54 (0.19–1.54)	0.25
Time from first line to gilteritinib therapy	1.0 (0.99–1.01)	0.19
Number of previous (compared to 1)	1.44 (0.48–4.33)	0.61
2 lines of therapy	1.77 (0.56–5.60)	0.51
3 lines of therapy		0.33
<i>Multivariate analysis (Hazard ratio, CI 95%)</i>		
Age at gilteritinib use	0.97 (0.93–1.01)	0.135
First line of treatment (HMA + venetoclax compared to “7 + 3”)	3.80 (0.86–16.72)	0.08
TKI use in first line therapy	0.15 (0.03–0.71)	0.016
Achieving CR after gilteritinib therapy	0.34 (0.12–0.94)	0.037

ELN European leukemia network, ECOG eastern corporation oncology group, HMA hypomethylating agent, TKI tyrosine kinase inhibitor, AlloSCT allogeneic stem cell transplantation, CR complete remission

addition, A previous combined analysis of the CHRYSALIS and ADMIRAL clinical trials demonstrate that patients previously exposed to FLT3 inhibitors derived benefit from gilteritinib albeit to a lesser extent from of those not previously exposed to FLT3 inhibitors [27].

The encouraging response rates in the intensively treated cohort are in stark contrast to the very poor survival in patients that were sequenced gilteritinib after failure of HMA and venetoclax combination (median OS 1.9 months). These data are in line with the poor survival reported for the

subgroup of patients treated with low-intensity approaches in the ADMIRAL trial and with previous outcomes reported for patients with HMA-venetoclax failure [21]. Based on our data, it seems that gilteritinib does not effectively salvage patient in this clinical setting. Combining gilteritinib to venetoclax with or without HMA's seems to have synergistic effects that result in high response rates even in patients previously exposed to FLT3 inhibitors. Such combinations may represent an attractive approach for low intensity treated FLT3-mutated patients [28–30].

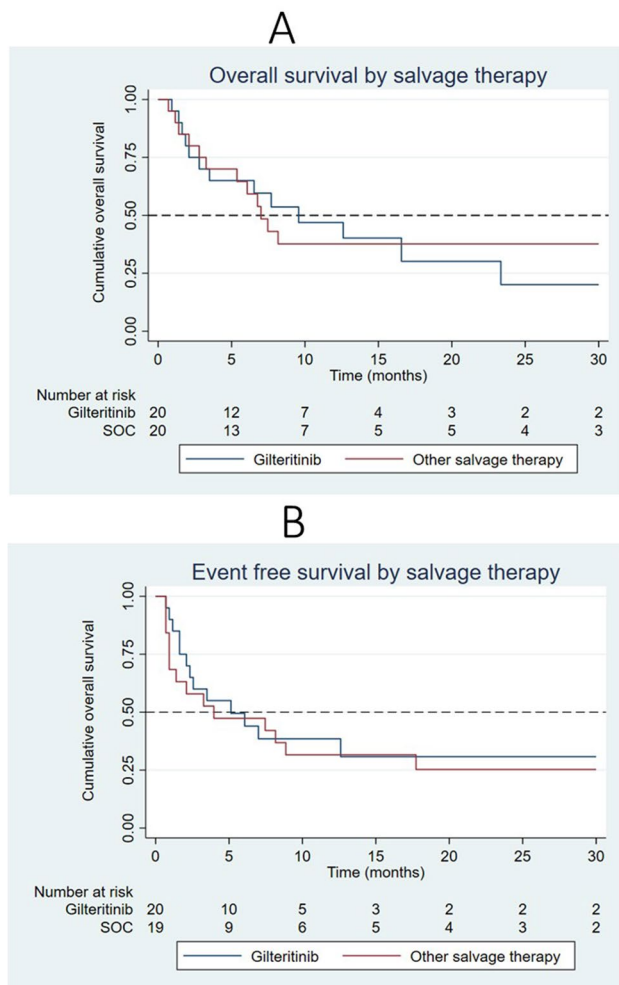


Fig. 3 Kaplan–Meier curve: Survival comparison between patients treated with gilteritinib to those treated with intensive salvage regimens. **A** Overall survival. **B** Event-free survival. SOC standard of care

This study is limited by its retrospective design and the relatively small number of patients. Nonetheless, it represents the first matched analysis in a real-world setting to our knowledge and provides insights as to treatment and response patterns with current therapeutic approaches for patients with *FLT3*-mutated AML.

In summary, gilteritinib is effective and well-tolerated in our patient cohort. In our analysis, patients seemed to receive gilteritinib at advanced time points in the course of therapy as compared to the clinical trials and earlier sequencing of this agent in the *FLT3*-mutated R/R setting may potentially optimize outcomes for this agent. Combination strategies for gilteritinib with venetoclax and other targeted approaches may represent another way to further improve responses and should be sought within clinical trials.

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Declarations

Ethics approval All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

Conflict of interest The authors declare no conflict of interest.

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