

Ivosidenib (AG-120) in mutant IDH1 relapsed/refractory acute myeloid leukemia: Results of a phase 1 study

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BACKGROUND

- Somatic mutations in the isocitrate dehydrogenase 1 (IDH1) gene occur in ~6-10% of patients with acute myeloid leukemia (AML).
- The mutant IDH1 (mIDH1) enzyme catalyzes the reduction of α-ketoglutarate to the oncometabolite D-2-hydroxyglutarate (2-HG),¹ and the resulting 2-HG accumulation leads to epigenetic dysregulation and impaired cellular differentiation.^{2,4}
- Ivosidenib (AG-120) is a first-in-class, oral, potent, targeted, small-molecule inhibitor of the mIDH1 enzyme.⁵
- Ivosidenib is under evaluation in an ongoing phase 1 dose escalation and expansion study of mIDH1 advanced hematologic malignancies, including relapsed/refractory acute myeloid leukemia (R/R AML).⁶
- On the basis of data from this study, ivosidenib received US FDA approval on July 20, 2018 for the treatment of adult patients with R/R AML with a susceptible IDH1 mutation, as detected by an FDA-approved test.
- The prognosis for patients with R/R AML is poor, with a median overall survival of 5.6 months,⁷ and there is no standard-of-care treatment.

OBJECTIVE

- To report updated efficacy, safety, mIDH1 variant allele frequency (VAF), and baseline co-mutation data from all patients with R/R AML receiving ivosidenib 500 mg once daily (QD) in the phase 1 study.

METHODS

- The ivosidenib phase 1, open-label, multicenter, dose escalation and expansion study includes the evaluation of safety, tolerability, maximum tolerated dose, pharmacokinetics and pharmacodynamics (including 2-HG levels), and clinical activity in patients with mIDH1 advanced hematologic malignancies (NCT02074839).⁶
- Single-agent ivosidenib is administered orally QD or twice daily (BID) in continuous 28-day cycles.
 - Doses in the escalation phase were 100 mg BID and 300, 500, 800, and 1200 mg QD.
 - 500 mg QD was selected for the expansion phase.
- The primary efficacy endpoint for R/R AML was the rate of complete remission plus complete remission with partial hematologic recovery (CR+CRh, Table 1).
- International working group (IWG) responses were reported by the investigator; CRh was derived by the sponsor.

Table 1. Definitions of CR and CRh

Response	Bone marrow blasts (%)	ANC/μL	Platelets/μL
CR (per modified IWG 2003 criteria) ⁸	<5	>1000	>100,000
CRh	<5	>500	>50,000

- Here we report data for all patients with R/R AML whose ivosidenib starting dose was 500 mg QD.
- The data cutoff date for this analysis was November 10, 2017.

RESULTS

- The baseline characteristics of 179 R/R AML patients who received ivosidenib 500 mg QD are shown in Table 2.
- 17 (9.5%) remained on treatment at data cutoff.
- 17 (9.5%) discontinued treatment to proceed to stem cell transplant.
- Median treatment duration was 3.9 months (range, 0.1–39.5).

- The majority of adverse events (AEs) were grade 1–2 (Table 3) and unrelated to treatment.
- AEs of interest (Table 4) were managed using standard-of-care treatments and ivosidenib dose modifications, as required.
- Ivosidenib induced durable responses (Table 5, Figures 1 and 2) and provided additional clinical benefits (Figure 3, Table 6).
- Transfusion independence was observed across all response categories in patients who were dependent at baseline.
- Ivosidenib induced IDH1 mutation clearance (IDH1-MC) in bone marrow mononuclear cells (BMMCs) from patients with a best overall response of CR or CRh (Table 7), and reduced mIDH1 VAF in BMMCs and neutrophils from patients with a best overall response of CR or CRh (Figure 4).
- 26% of patients with a best response of CR/CRh for whom molecular data were available had IDH1-MC in both BMMCs and neutrophils.
- Patients with IDH1-MC had improved durations of CR+CRh and overall survival versus patients with detectable mIDH1 (Figure 5).

Table 2. Baseline characteristics

Characteristic	R/R AML 500 mg (n=179)
Women/men, n	89/90
Age, median (range), years	67.0 (18–87)
Age category, n (%)	
<60 years	47 (26.3)
60 to <75 years	92 (51.4)
≥75 years	40 (22.3)
ECOG Performance Status at baseline, n (%)	
1	36 (20.1)
2	99 (55.3)
3	42 (23.5)
De novo AML, n (%)	120 (67.0)
Secondary AML, n (%)	59 (33.0)
No. of prior therapies, median (range)	2.0 (1–6)
Prior AML therapy outcomes ^a , n (%)	
Relapsed after transplant	43 (24.0)
In 2nd or later relapse	26 (14.5)
Refractory to initial induction/reinduction therapy	106 (59.2)
In 1st relapse	17 (9.5)
In 2nd relapse	15 (8.4)
Other	5 (2.8)
Cytogenetic risk status by investigator, n (%)	
Intermediate	105 (58.7)
Poor	50 (27.9)
Unknown/missing	24 (13.4)
Most common baseline co-mutations ^b , %	
DNMT3A	34
mRNA splicing gene ^c	31
NPM1	25
RAS pathway ^d	24
ASXL1	19
RUNX1	18
PS3	14

^aNot mutually exclusive; patients may be in >1 category. ^bAssessed in 178 patients; mutations were identified using next-generation sequencing (FoundationOne[®] Home Panel) in the escalation phase and Rapid Home Panel in expansion. ^cIncludes SF3B1, SRSF2, U2AF1, U2AF2, and ZRSR2. ^dIncludes MAP3K4, NRAS, PTPN11, KRAS, NF1, BRAF, and KIT. ^eECOG = Eastern Cooperative Oncology Group.

Table 3. Most common AEs (≥20%) by preferred term, regardless of causality

R/R AML 500 mg (n=179)	Any grade, n (%)	Grade ≥3, n (%)
Any AE	179 (100)	148 (82.7)
Diarrhea	60 (33.5)	4 (2.2)
Leukocytosis	56 (31.3)	14 (7.8)
Nausea	56 (31.3)	1 (0.6)
Febrile neutropenia	52 (29.1)	52 (29.1)
Fatigue	51 (28.5)	3 (1.7)
ECG QT prolonged	46 (25.7)	18 (10.1)
Dyspnea	44 (24.6)	7 (3.9)
Edema peripheral	43 (24.0)	0 (0.0)
Pyrexia	41 (22.9)	2 (1.1)
Anemia	40 (22.3)	36 (20.1)
Cough	38 (21.2)	1 (0.6)

ECG = electrocardiogram

Table 4. Investigator-reported AEs of interest by preferred term

AEs of interest	n (%)	R/R AML 500 mg (n=179)
Grade ≥3 leukocytosis ^a	14 (8)	<ul style="list-style-type: none"> Managed with hydroxyurea None were fatal
Grade ≥3 ECG QT prolongation	18 (10)	<ul style="list-style-type: none"> Study drug was reduced in 2 patients and held in 13 patients (all grades) None were fatal QT-prolonging medications such as antifungals and fluoroquinolone anti-infectives were allowed on study with monitoring
IDH1-DS (all grades)	19 (10.6)	<ul style="list-style-type: none"> Resolved in 17 patients, ongoing in 2 patients at data cutoff Grade ≥3 IDH1-DS in 9 patients (5.0%) 7/19 patients with IDH1-DS had co-occurring leukocytosis Study drug held in 6 patients (3.4%) No instances of IDH1-DS led to dose reduction, permanent treatment discontinuation, or death Managed with corticosteroids and diuretics, and hydroxyurea if accompanied by leukocytosis Best response for the 19 patients with IDH1-DS: 5 CR, 3 CR/CRp, 2 MLFS, 8 SD, and 1 not evaluable

^aGrade 3: white blood cells >100,000/mm³; Grade 4: clinical manifestations of leukostasis, urgent intervention indicated. CRh = CR with incomplete hematologic recovery; CRp = CR with incomplete platelet recovery; DS = differentiation syndrome; MLFS = morphologic leukemia-free state; SD = stable disease.

Table 5. Response rates

	R/R AML 500 mg (n=179)
CR+CRh rate, n (%) [95% CI]	57 (31.8) [25.1, 39.2]
Time to CR/CRh, median (range), months	2.0 (0.9–5.6)
Duration of CR/CRh, median [95% CI], months	8.2 [5.6, 12.0]
CR rate, n (%) [95% CI]	43 (24.0) [18.0, 31.0]
Time to CR, median (range), months	2.8 (0.9–8.3)
Duration of CR, median [95% CI], months	10.1 [6.5, 22.2]
CRh rate, n (%)	14 (7.8)
Duration of CRh, median [95% CI], months	3.6 [1.0, 5.5]
Overall response rate, n (%) [95% CI]	75 (41.9) [34.6, 49.5]
Time to first response, median (range), months	1.9 (0.8–4.7)
Duration of response, median [95% CI], months	6.5 [5.5, 10.1]
Best response, n (%)	
CR	43 (24.0)
CR or CRp	21 (11.7)
MLFS	11 (6.1)
SD	68 (38.0)
PD	15 (8.4)
NA	21 (11.7)

CRh includes 8 patients with investigator-assessed responses of CR/CRp and 5 with MLFS. Overall response rate includes CR, CR/CRh, MLFS, and PD. At the time of the database lock, among the 179 patients with R/R AML, 4 non-dose escalation and 1 from dose escalation were not positive for mIDH1 by the companion diagnostic test and none of these 5 patients achieved a CR or CRh. The patient from dose escalation was found to be positive for mIDH1 by the companion diagnostic test after database lock. CR+CRh rate was consistent across baseline age groups, including patients who were ≥65 years of age. NA = not assessed; PD = progressive disease.

Figure 1. Duration of treatment and best overall response in responders

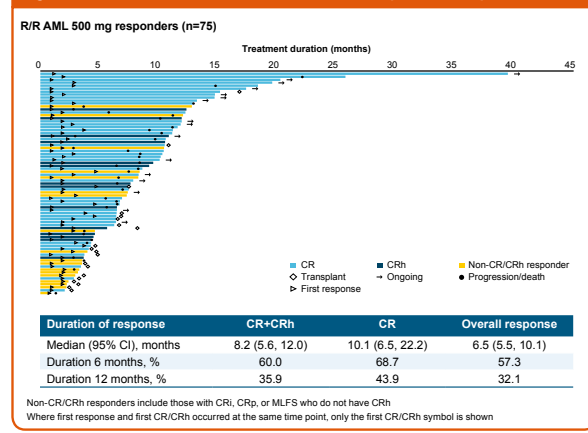
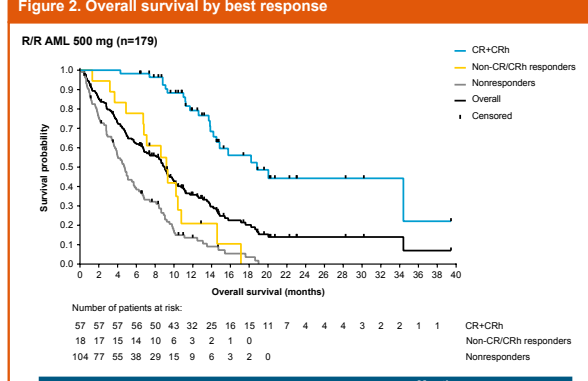


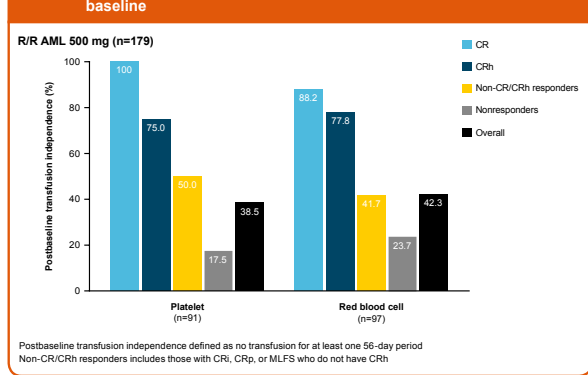
Figure 2. Overall survival by best response



Overall survival, median (95% CI)	Months
CR+CRh	18.8 (14.2, NE)
Non-CR/CRh responders	9.2 (6.7, 10.8)
Nonresponders	4.7 (3.7, 5.7)
All	9.0 (7.1, 10.0)
Overall follow-up, median (range)	15.3 (0.2–39.5)

Non-CR/CRh responders includes those with CR, CRp, or MLFS who do not have CRh; nonresponders includes all others, including those with best responses of SD, PD, or not evaluable. NE = not estimable.

Figure 3. Transfusion independence in patients who were dependent at baseline



Postbaseline transfusion independence defined as no transfusion for at least one 56-day period. Non-CR/CRh responders includes those with CR, CRp, or MLFS who do not have CRh.

Table 6. Exposure-adjusted incidence of febrile neutropenia and grade ≥3 infections

	R/R AML 500 mg				Overall (n=179)
	CR (n=43)	CRh (n=14)	Non-CR/CRh responders (n=18)	Nonresponders (n=104)	
All grade febrile neutropenia ^a	2.0 (1.0, 3.8)	3.7 (1.4, 9.8)	6.1 (2.7, 13.5)	12.1 (8.8, 16.5)	5.9 (4.5, 7.6)
Grade ≥3 infections ^b	2.6 (1.5, 4.6)	6.4 (3.1, 13.5)	13.1 (7.6, 22.6)	21.3 (16.8, 27.0)	10.2 (8.4, 12.4)

Incidence rate reported as 100 patients/month (95% CI), calculated as total number of specific AEs per total person exposure time in months × 100 for all patients with the same best overall response. ^aPreferred term, including febrile bone marrow aplasia preferred term. ^bBased on MedDRA V20.0 System Organ Class of infections and interstitions.

Figure 4. Longitudinal mean mIDH1 VAF by best overall response

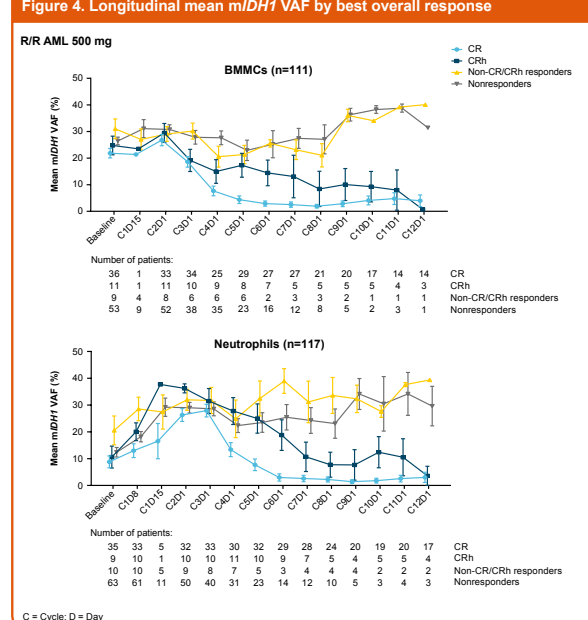


Figure 5. Duration of response and overall survival in patients with IDH1-MC

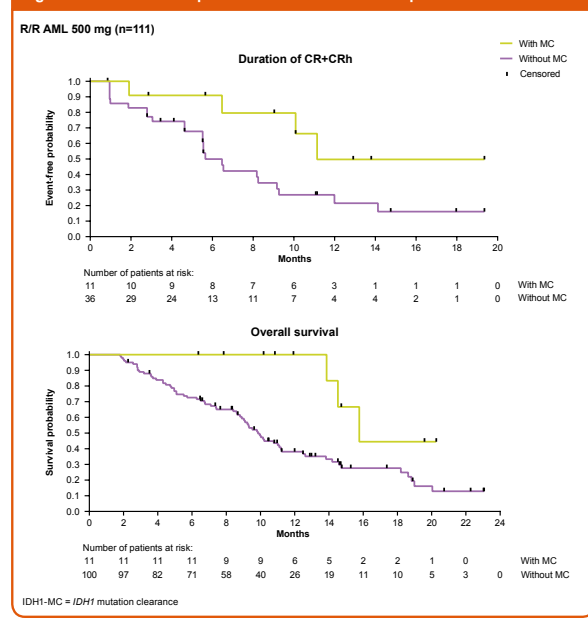


Table 7. IDH1 mutation clearance in BMMCs

Response	n	IDH1 mutation clearance, ^a n (%)	Detectable IDH1 mutation, n (%)
CR+CRh	47	11 (23)	36 (77)
CR	36	10 (28)	26 (72)
CRh	11	1 (9)	10 (91)
Others	64	0	64 (100)
Non-CR+CRh responders	9	0	9 (100)
Nonresponders	55	0	55 (100)

p-value^b <0.001

^aDefined as reduction in mIDH1 VAF to below the limit of detection of 0.02–0.04% (2–4×10⁻⁴) by digital PCR for at least one on-study time point. ^bp-value based on Fisher's exact test comparing IDH1 mutation clearance in patients who had a best overall response of CR+CRh with patients who had other responses (non-CR/CRh responders and nonresponders).

CR = complete remission; CRh = complete remission with partial hematologic recovery; CRp = complete remission with partial platelet recovery; DS = differentiation syndrome; MLFS = morphologic leukemia-free state; SD = stable disease.

NE = not estimable.

NA = not assessed.

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