Revumenib Monotherapy in Patients With Relapsed/Refractory KMT2Ar Acute Leukemia: Efficacy and Safety Results From the AUGMENT-101 Phase 1/2 Study

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INTRODUCTION

- Revumenib, a small-molecule inhibitor of the menin—histone-lysine N-methyltransferase 2A (KMT2A) interaction, is being investigated in adult and pediatric patients with relapsed/refractory (R/R) *KMT2A*-rearranged (*KMT2Ar*) and nucleophosmin 1–mutated (*NPM1m*) acute leukemias in the phase 1/2 AUGMENT-101 study (NCT04065399)¹
- In the phase 1 study, patients were assigned to 1 of 6 doseescalation cohorts designed to identify a recommended phase 2 dose (RP2D) for concomitant administration of a cytochrome P450 3A4 inhibitor (CYP3A4i; moderate or strong) or no CYP3A4i
- RP2D was determined based on review of pharmacokinetics, clinical activity, safety, and tolerability data
- An initial analysis of the phase 1 study has been reported¹
 Revumenib resulted in deep, durable responses in heavily pretreated R/R acute leukemias; 37.5% of responders proceeded to hematopoietic stem cell transplant (HSCT)
- Grade ≥3 treatment-related adverse events (TRAEs) were reported in 16.2% of patients; asymptomatic prolongation of the QT interval was the only dose-limiting toxicity identified
- Here we provide an update of >1-year additional experience of patients with R/R KMT2Ar acute leukemia in the phase 1 portion of the AUGMENT-101 study, which has completed enrollment, and report topline efficacy and safety results from the phase 2 portion of the study

METHODS

- The phase 1/2 study used a highly innovative trial design enabling an early phase pivotal study for both adult and pediatric, with KMT2Ar acute myeloid leukemia (AML), acute lymphoblastic leukemia (ALL), and mixed-phenotype acute leukemia (MPAL). The study also includes a separate NPM1m AML cohort
- Phase 1 methods have been previously published¹
- The pivotal phase 2 portion of the study was initiated after identification of an RP2D of 163 mg (or 95 mg/m² if body weight <40 kg) every 12 hours (q12h) with a strong CYP3A4i in 28-day cycles until unacceptable toxicity or lack of at least morphological leukemia-free state (MLFS) by end of cycle 4</p>
- Phase 2 study enrolled patients in 2 subgroups: those with R/R KMT2Ar acute leukemia (including AML, ALL MPAL) and patients with R/R NPM1m AML
- Phase 2 primary objectives were the assessment of the complete remission (CR)+CR with partial hematologic recovery (CRh) rate and evaluation of safety and tolerability of revumenib
- A planned interim analysis (IA) of adult and pediatric patients with KMT2Ar acute leukemia was conducted with data cutoff in July 2023, and topline data are reported here
- The efficacy population for the IA (n=57) included all phase 2 patients who had centrally confirmed KMT2Ar acute leukemia, had ≥5% blasts in bone marrow at baseline, had received ≥1 dose of study drug, and had started treatment at the same time as or before the 38th adult AML efficacy-evaluable patient
- The analysis was conducted when 57 patients (adult and pediatric) had completed 6 months of follow-up or discontinued therapy
- Enrollment of R/R NPM1m AML in cohort 2C is ongoing, and data for this cohort are not included in this analysis

RESULTS

PHASE 1

EFFICACY

- As of July 24, 2023, 132 patients aged 0.8 to 82.0 years with R/R acute leukemia were enrolled in the phase 1 study and included in the overall population
- Among 77 patients with R/R KMT2Ar acute leukemia treated with revumenib in the 6 dose-escalation cohorts, CR+CRh rate was 31.2% (24/77) and overall response rate (ORR; ie, composite CR [CRc]+MLFS+partial remission) was 64.9% (50/77), with 38.0% (19/50) of responders proceeding to HSCT
- In a subgroup of adults with AML, CR+CRh rate was 37.3% (19/51) and ORR was 68.6% (35/51), with 40.0% (14/35) of responders proceeding to HSCT
- Of 132 patients in the overall population, 25.0% had a Grade ≥3 TRAE, with Grade 3 QTc prolongation in 8.3% of patients and Grade 3 differentiation syndrome in 2.3% of patients 10.6% of patients discontinued revumenib due to TRAEs
- Based on AUGMENT-101 phase 1 pharmacokinetics, clinical activity, and safety data, an RP2D of 276 mg (or 160 mg/m² if body weight <40 kg) q12h without a strong CYP3A4i was established
- For concomitant use with a strong CYP3A4i, an RP2D of 163 mg (or 95 mg/m² if body weight <40 kg) q12h was established

PHASE 2

BASELINE CHARACTERISTICS

- As of July 24, 2023, 94 patients had received ≥1 dose of study drug and were included in the KMT2Ar safety population (Table 1)
- At the time of the IA, 57 patients had the opportunity for sufficient follow-up to evaluate for efficacy
- 57.9% of patients were female and 22.8% of patients were pediatric
- Patients received a median of 2 prior lines of therapy (range, 1–11) and 45.6% had prior HSCT

Table 1. Phase 2 Patient Demographics and Baseline Characteristics^a

| Parameter | Efficacy population (n=57) | Safety population (n=94) ^b |
|--|----------------------------|---------------------------------------|
| Median age, y (range) | 34.0 (1.3–75.0) | 37.0 (1.3–75.0) |
| Sex, n (%) | | |
| Female | 33 (57.9) | 56 (59.6) |
| Race, n (%) | | |
| White | 43 (75.4) | 68 (72.3) |
| Non-White | 10 (17.5) | 14 (14.9) |
| Unknown | 4 (7.0) | 12 (12.8) |
| Leukemia type, n (%) | | |
| AML | 49 (86.0) | 78 (83.0) |
| ALL | 7 (12.3) | 14 (14.9) |
| MPAL/Other | 1 (1.8) | 2 (2.1) |
| Co-mutations, n (%) | | |
| FLT3 | 5 (8.8) | 7 (7.4) |
| RAS | 9 (15.8) | 12 (12.8) |
| p53 | 4 (7.0) | 5 (5.3) |
| Primary refractory, n (%) | 14 (24.6) | 18 (19.1) |
| Number of prior lines of therapy, median (range) | 2 (1–11) | 2 (1–11) |
| ≥3 prior lines of therapy, n (%) | 26 (45.6) | 41 (43.6) |
| Prior venetoclax, n (%) | 41 (71.9) | 61 (64.9) |
| Prior HSCT, n (%) | 26 (45.6) | 47 (50.0) |

ALL, acute lymphoblastic leukemia; AML, acute myeloid leukemia; *FLT3*, fms-related tyrosine kinase 3; HSCT, hematopoietic stem cell transplant; *KMT2Ar*, histone-lysine N-methyltransferase 2A rearrangements; MPAL, mixed-phenotype acute leukemia; *RAS*, rat sarcoma virus. ^aData cutoff: July 24, 2023. ^bDefined as patients with *KMT2Ar* acute leukemia having received at least 1 dose of revumenib.

EFFICACY

- With a median follow-up time of 6.1 months in the efficacy population, the ORR was 63.2% (36/57; 95% CI, 49.3–75.6); CRc was 43.9% (25/57; 95% CI, 30.7–57.6)
- The primary endpoint of the pivotal phase 2 study was met at the IA, with a CR+CRh rate of 22.8% (13/57; 95% CI, 12.7–35.8; 1-sided *P* value, 0.0036), with 70.0% (7/10) of those patients (with available data) achieving negative measurable residual disease (MRD) status (Table 2)
- Responses stratified by KMT2A rearrangements and major subgroups are shown in Table 3 and Figure 1, respectively
- CR+CRh rate was similar in adult (22.7% [10/44; 95% CI, 11.5–37.8]) and pediatric (23.1% [3/13; 95% CI, 5.0–53.8]) patients (Figure 1)
- However, the study was not powered to evaluate differences among subgroups as the number of patients per subgroup are small
- At the IA, median overall survival was 8.0 (95% CI, 4.1–10.9) months (Figure 2)
 CR+CRh responses were rapid (median, 1.9 [range, 0.9–4.6] months; Figure 3) and durable (6.4 [95% CI, 3.4–not reached] months)
- 38.9% (14/36) of patients proceeded to HSCT, with 7 patients resuming revumenib after HSCT
 3 additional patients remained eligible to initiate revumenib after HSCT at the time of data cutoff

Table 2. Phase 2 Response^a

| Parameter | Efficacy population (n=57) |
|---|---|
| ORR, n (%) | 36 (63.2) |
| Best response, n (%) | |
| CR | 10 (17.5) |
| CRh | 3 (5.3) |
| CRi | 1 (1.8) |
| CRp | 11 (19.3) |
| MLFS | 10 (17.5) |
| PR | 1 (1.8) |
| Other ^b | 21 (36.8) |
| CR+CRh rate, n (%) | 13 (22.8) |
| | 95% CI, 12.7–35.8; 1-sided <i>P</i> value, 0.0036 |
| CRc, n (%) | 25 (43.9) |
| Negative MRD status in CRc, n (%) ^c | 15/22 (68.2) |
| Negative MRD status in CR+CRh, n (%) ^c | 7/10 (70.0) |
| No. of responders who proceeded to HSCT, n (%) | 14/36 (38.9) |
| CR, complete remission; CRc, composite CR (CR+CRh+CRp+CRi); CRh, CR with partia | al hematologic recovery; CRi, CR with incomplete hematologic recovery; CRp, |

CR, complete remission; CRc, composite CR (CR+CRh+CRp+CRi); CRh, CR with partial hematologic recovery; CRi, CR with incomplete hematologic recovery; CRp, CR with incomplete platelet recovery; HSCT, hematopoietic stem cell transplant; MLFS, morphological leukemia-free state; MRD, measurable residual disease; ORR, overall response rate (CRc+MLFS+PR); PR, partial remission. ^aData cutoff: July 24, 2023. ^bIncludes no response, disease progression, and patients without postbaseline disease assessment. ^cMRD done locally; not all patients had MRD status reported.

Table 3. Phase 2 Responses Observed Across KMT2A Rearrangements

| | Summary of ORR | | Summary of CR+CRh rate | |
|-----------------------------------|------------------|------------------|------------------------|----------------------|
| KMT2A rearrangement/translocation | n/N ^b | ORR (95% CI) | n/N ^b | CR+CRh rate (95% CI) |
| 9;11 | 10/11 | 90.9 (58.7–99.8) | 2/11 | 18.2 (2.3–51.8) |
| 11;19 | 7/13 | 53.8 (25.1–80.8) | 2/13 | 15.4 (1.9–45.4) |
| 10;11 | 5/7 | 71.4 (29.0–96.3) | 2/7 | 28.6 (3.7–71.0) |
| 6;11 | 5/7 | 71.4 (29.0–96.3) | 2/7 | 28.6 (3.7–71.0) |
| 4;11 | 2/2 | 100 (15.8–100) | 0/2 | 9 (0.0–84.2) |
| 1;11 | 0/2 | 0 (0.0–84.2) | 0/2 | 9 (0.0–84.2) |
| 11;16 | 1/1 | 100 | 0/1 | 0 |
| 11;22 | 1/1 | 100 | 1/1 | 100 |
| Unknown KMT2A fusion partner | 5/13 | 38.5 (13.9–68.4) | 4/13 | 31.0 (9.1–61.4) |

CR, complete remission; CRc, composite CR (CR+CRh+CRp+CRi); CRh, CR with partial hematologic recovery; CRi, CR with incomplete hematologic recovery; CRp, CR with incomplete platelet recovery; *KMT2A*, histone-lysine N-methyltransferase 2A; MLFS, morphological leukemia-free state; ORR, overall response rate (CRc+MLFS+PR); PR, partial remission. ^aData cutoff: July 24, 2023. ^bN = total number of *KMT2A* rearrangements/translocations.

Figure 1. Forest plot illustrating the estimated rate of CR+CRh for prespecified subgroups in the phase 2 efficacy population of AUGMENT-101.^a

| | CR+CRh rate (95% CI) | | No. of participants |
|---|--|---------------|-------------------------------------|
| Overall | 22.8 (12.7–35.8) | ├── | 57 (100.0) |
| Age group <18 years ≥18 years | 23.1 (5.0–53.8) 22.7 (11.5–37.8) | ├ | 13 (22.8) 44 (77.2) |
| Sex Female Male | 21.2 (9.0–38.9) 25.0 (9.8–46.7) | | 33 (57.9) 24 (42.1) |
| Race White Non-White Unknown | 20.9 (10.0–36.0) 30.0 (6.7–65.2) 25.0 (0.6–80.6) | | 43 (75.4) 10 (17.5) 4 (7.0) |
| Histology AML ALL Acute leukemia of ambiguous lineage | 24.5 (13.3–38.9) 14.3 (0.4–57.9) 0.0 (0.0–97.5) | | 49 (86.0) 7 (12.3) — 1 (1.8) |
| Prior HSCT Yes No | 23.1 (9.0–43.6) 22.6 (9.6–41.1) | | 26 (45.6) 31 (54.4) |
| Primary refractory Yes No | 35.7 (12.8–64.9) 18.6 (8.4–33.4) | | 14 (24.6) 43 (75.4) |
| No. of prior lines of therapy 1 2 ≥3 | 47.1 (23.0–72.2) 14.3 (1.8–42.8) 11.5 (2.4–30.2) | | 17 (29.8) 14 (24.6) 26 (45.6) |
| | | 0 20 40 60 80 | 100 |

ALL, acute lymphoblastic leukemia; AML, acute myeloid leukemia; CR, complete remission; CRh, CR with partial hematologic recovery; HSCT, hematopoietic stem cell transplant. ^aData cutoff: July 24, 2023.

Figure 2. Kaplan-Meier estimate of overall survival in the phase 2 efficacy population of AUGMENT-101.^a

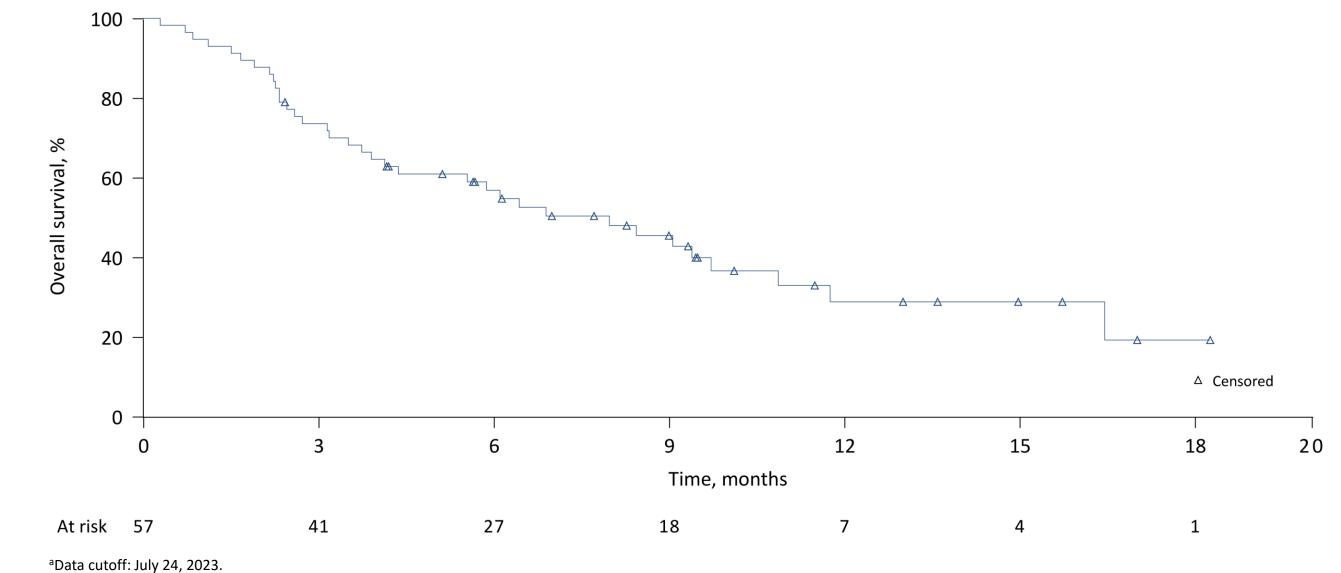
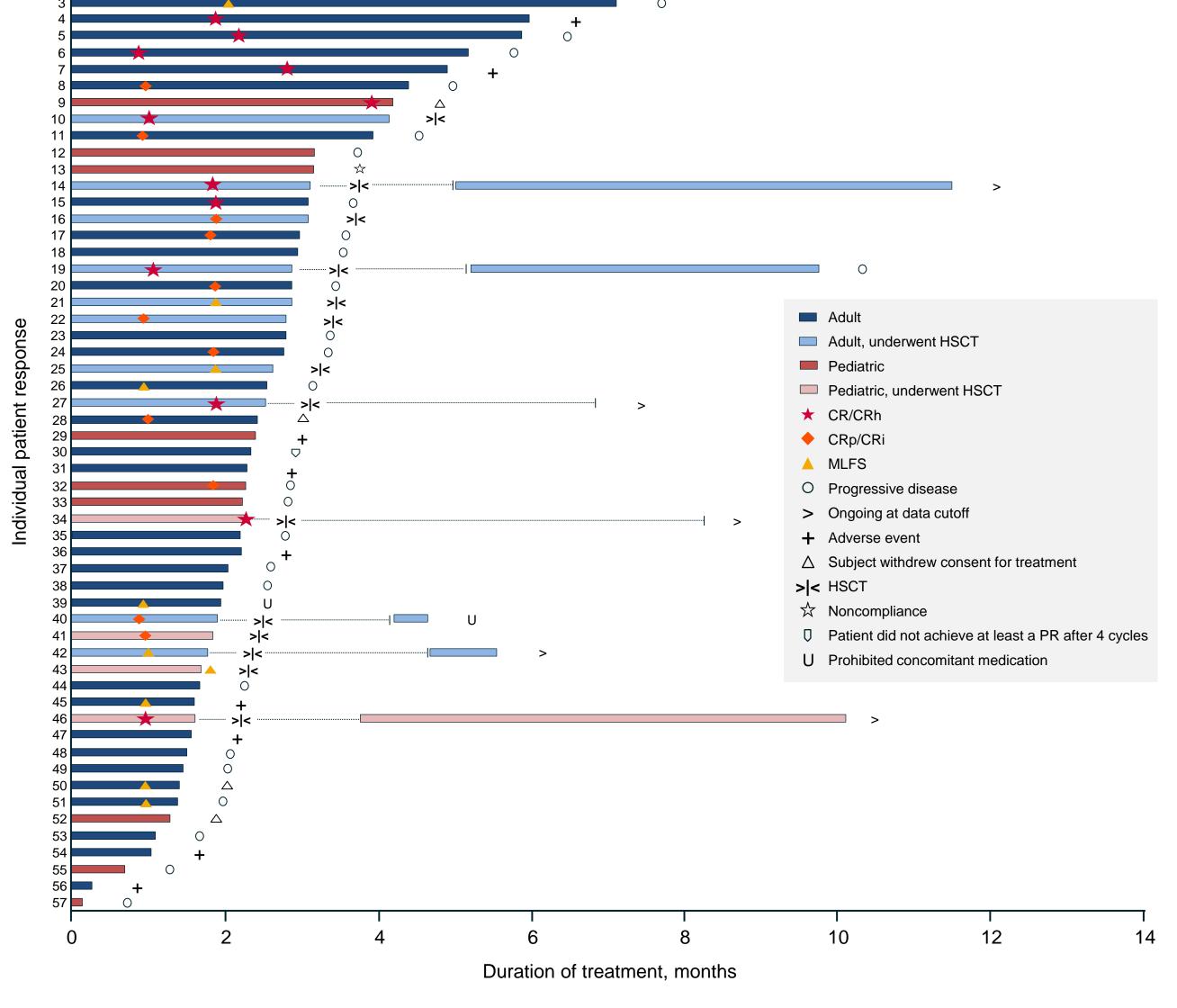


Figure 3. Swimmer plot illustrating the duration of treatment and key study events of patients in the phase 2 efficacy population of AUGMENT-101.^a



CR, complete remission; CRh, CR with partial hematologic recovery; CRi, CR with incomplete hematologic recovery; CRp, CR with incomplete platelet recovery; HSCT, hematopoietic stem cell transplant; MLFS, morphological leukemia-free state; PR, partial remission. aData cutoff: July 24, 2023.

FETY

- Overall, a manageable safety profile was observed among the 94 patients in the safety population (Table 4)
- 6.4% and 12.8% of patients discontinued revumenib due to TRAEs and treatment-emergent adverse events, respectively
- Most common TRAEs (≥20%) were nausea (27.7%), differentiation syndrome (26.6%), and QTc prolongation (23.4%)
- There were no treatment discontinuations due to differentiation syndrome, QTc prolongation, or cytopenias

Table 4. Phase 2 Safety Profile⁶

| All terms | Safety population (n=94) ^b 93 (98.9) | |
|---|---|--|
| Any grade TEAE, n (%) | | |
| Any grade TEAEs that occurred in ≥30% patients, n (%) | | |
| Nausea | 42 (44.7) | |
| Febrile neutropenia | 36 (38.3) | |
| Diarrhea | 33 (35.1) | |
| Vomiting | 29 (30.9) | |
| Grade ≥3 TEAE, n (%) | 86 (91.5) | |
| Grade ≥3 TEAE that occurred in ≥10% patients, n (%) | | |
| Febrile neutropenia | 35 (37.2) | |
| Anemia | 17 (18.1) | |
| Decreased neutrophil count | 15 (16.0) | |
| Decreased white blood cell count | 15 (16.0) | |
| Differentiation syndrome | 15 (16.0) | |
| Decreased platelet count | 14 (14.9) | |
| QTc prolongation | 13 (13.8) | |
| Sepsis | 11 (11.7) | |
| Hypokalemia | 10 (10.6) | |
| Serious AE, n (%) | 72 (76.6) | |
| TEAE leading to dose reduction, n (%) | 9 (9.6) | |
| TEAE leading to discontinuation, n (%) | 12 (12.8) | |
| TEAE leading to death, n (%) | 14 (14.9) | |

AE, adverse event; TEAE, treatment-emergent AE. ^aData cutoff: July 24, 2023. ^bDefined as patients with *KMT2Ar* acute leukemia having received at least 1 dose of revumenib.

CONCLUSIONS

- The phase 1 study demonstrated promising efficacy and safety in heavily pretreated patients as previously reported¹
- At the phase 2 IA, this pivotal study of revumenib in an independent cohort of patients with KMT2Ar acute leukemia met its primary endpoint with a CR+CRh rate of 22.8% (13/57; 95% CI, 12.7–35.8) and a 1-sided P value of 0.0036
- Durable MRD-negative remissions and high rates of transplants were observed among responders
- Overall, a manageable safety profile was observed; discontinuations and dose reductions due to adverse events were low

Enrollment in the KMT2Ar cohorts was stopped early for efficacy after meeting

- the primary efficacy endpoint at the phase 2 IA
 A New Drug Application for *KMT2Ar* leukemia has been initiated under the
- Enrollment in the independent NPM1m AML cohort is ongoing

FDA Real-Time Oncology Review program based on these data

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REFERENCE: **1.** Issa GC, Aldoss I, DiPersio J, et al. *Nature*. 2023;615(7954):920–924.