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Venetoclax Combined With FLAG-IDA Induction and Consolidation in Newly Diagnosed and **Relapsed or Refractory Acute Myeloid Leukemia**

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PURPOSE Sixty percent of newly diagnosed patients with acute myeloid leukemia (ND-AML) receiving frontline therapy attain a complete response (CR), yet 30%-40% of patients relapse. Relapsed or refractory AML (R/Rtra AML) remains a particularly adverse population necessitating improved therapeutic options. This phase Ib/II study evaluated the safety and efficacy of fludarabine, cytarabine, granulocyte colony-stimulating factor, and idarubicin combined with the B-cell lymphoma-2 inhibitor venetoclax in ND-AML and R/R-AML.

MATERIALS AND METHODS The phase IB portion (PIB) enrolled patients with R/R-AML using a 3 + 3 dose escalation and de-escalation algorithm for identification of maximum tolerated dose and dose-limiting toxicities. The phase II portion enrolled patients into two arms to evaluate response and time-to-event end points: phase IIA (PIIA): ND-AML and phase IIB (PIIB): R/R-AML.

RESULTS Sixty-eight patients have enrolled to date (PIB, 16; PIIA, 29; PIIB, 23). Median age was 46 years (range, 20-73). Grade 3 and 4 adverse events occurring in $\geq 10\%$ of patients included febrile neutropenia (50%), bacteremia (35%), pneumonia (28%), and sepsis (12%). The overall response rate for PIB, PIIA, and PIIB was 75%, 97%, and 70% with 75%, 90%, and 61%, respectively, achieving a composite CR. Measurable residual disease-negative composite CR was attained in 96% of ND-AML and 69% of R/R-AML patients. After a median follow-up of 12 months, median overall survival (OS) for both PII cohorts was not reached. Fifty-six percent of patients proceeded to allogeneic hematopoietic stem-cell transplantation (ND-AML, 69%; R/R-AML, 46%). In R/R-AML, allogeneic hematopoietic stem-cell transplantation resulted in a significant improvement in OS (median OS, NR; 1-year OS, 87%). One-year survival post-HSCT was 94% in ND-AML and 78% in R/R-AML.

CONCLUSION Fludarabine, cytarabine, granulocyte colony-stimulating factor, and idarubicin + venetoclax represents an effective intensive treatment regimen in ND-AML and R/R-AML patients, associated with deep remissions and a high rate of transition to successful transplantation.

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Protocol

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INTRODUCTION

Induction chemotherapy (IC) in acute myeloid leukemia (AML) typically combines an anthracycline (ie, daunorubicin or idarubicin) with the antimetabolite cytarabine (the often termed 7 + 3 regimen), resulting in complete remission (CR) rates of approximately 60%.¹ Anthracycline dose augmentation and development of synergistic multidrug regimens improves CR rates to 70%-80%²⁻⁶; however, 30%-40% of patients still ultimately relapse.^{7,8} Treatment regimens capable of producing long-term remissions are needed.

The multiagent regimen of fludarabine, cytarabine, granulocyte colony-stimulating factor (G-CSF), and idarubicin (FLAG-IDA) is an effective frontline treatment in fit patients with AML. Compared with alternate IC regimens, frontline FLAG-IDA induction results in composite CR (CRc) rates of 85%, a reduced cumulative incidence of relapse (38% v 55%), improved relapsefree survival, and median overall survival (OS) of approximately 5 years, albeit with increased myelosuppression.³ Patients with high-risk myelodysplastic syndrome or secondary AML (sAML) receiving FLAG-IDA achieved CR rates of 78%; 63% achieved a morphologic and cytogenetic CR after one or two treatment cycles.⁹

Patients with relapsed and/or refractory (R/R) AML experience inferior CR rates (20%-60%) with IC reinduction, and few patients obtain durable remissions with salvage therapy.^{10,11} Although no proven

CONTEXT

Key Objective

Is the addition of the B-cell lymphoma-2 inhibitor venetoclax (VEN) to standard fludarabine, cytarabine, granulocyte colonystimulating factor, and idarubicin (FLAG-IDA) induction and consolidation (FLAG-IDA + VEN) therapy safe and effective for patients with newly diagnosed acute myeloid leukemia (ND-AML) and/or relapsed or refractory AML (R/R-AML)?

Knowledge Generated

FLAG-IDA + VEN resulted in high rates of measurable residual disease–negative composite complete remission in patients with both ND-AML and R/R-AML, with a majority of patients able to effectively transition to allogeneic stem-cell transplantation. The combination was associated with an expected and manageable myelosuppression-related toxicity profile.

Relevance

VEN in combination with intensive induction and consolidation therapy in AML demonstrates the regimen is effective in both the ND-AML and R/R-AML setting. The high rates of observed measurable residual disease–negative responses suggest this regimen induces deep remissions and is effective for successfully bridging patients with ND-AML and R/R-AML to allogeneic transplantation.

optimal salvage regimen exists, FLAG-IDA is commonly used, resulting in a sobering CR or CR with incomplete hematologic recovery (CRi) rate of 21%, 3.5-month median OS, and 30-day mortality exceeding 10%.¹⁰⁻¹²

The B-cell lymphoma-2 (BCL-2) inhibitor venetoclax (VEN) combined with lower-intensity treatment regimens (azacitidine, decitabine, or low-dose cytarabine) is approved for unfit, older patients with AML^{13,14} and has rapidly emerged as a standard-of-care treatment option for this challenging patient population.¹³⁻¹⁶

Congruent with the synergy observed with VEN combinations incorporating lower-intensity treatments, VEN demonstrated preclinical synergy with standard chemotherapeutic agents, suggesting benefit beyond the older unfit AML population.¹⁷

Attenuated cytarabine and idarubicin (the 5 + 2 regimen) combined with VEN in older, adverse-risk patients with newly diagnosed acute myeloid leukemia (ND-AML) resulted in CRc (CR + CRi) rates of 72% and 30-day mortality of 6%.¹⁸ In R/R-AML, 69% of patients treated with FLA-IDA with VEN (days 1-7) achieved a CRc without significantly increased hematologic toxicity compared with a matched FLA-IDA cohort,¹⁹ suggesting optimization of VEN with IC may increase efficacy without untoward toxicity. Herein, we provide results of a phase Ib/II study of FLAG-IDA + VEN as frontline or salvage AML therapy.

MATERIALS AND METHODS

Inclusion Criteria

Patients age > 18 years with ND-AML (de novo AML, sAML, treated secondary AML [ts-AML], and therapy-related AML [t-AML]), high-risk myelodysplastic syndrome (defined by the presence of > OR = 10% blasts), or R/R-AML (defined as persistent leukemia without achievement of an International

Working Group–defined response following at least one cycle of induction chemotherapy or patients in first relapse or beyond) were eligible. Only R/R-AML patients were eligible for the phase Ib (PIB) portion. Patients with acute promyelocytic leukemia, significant cardiovascular comorbidities, known malabsorption syndromes, or who had received prior BCL-2 inhibitor therapy were excluded (full eligibility available within the Appendix, online only). All investigations were conducted under the approval of the institutional review committee and in accordance with the Declaration of Helsinki.

Cytogenetic and Molecular Analysis

Cytogenetic evaluation using standard metaphase karyotype analysis and molecular analysis via an 81-gene institutional next-generation sequencing platform was performed at study enrollment. Measurable residual disease (MRD) was assessed by 8-color multiparameter flow cytometry (FC) using leukemia-associated immunophenotype or different from normal assessment²⁰ with a minimum sensitivity of 10^{-3} to 10^{-4} (0.1%-0.01%).

Safety and Efficacy

The PIB (dose-escalation) portion applied a 3 + 3 dose escalation and de-escalation algorithm to determine the maximal tolerated dose (MTD) (details of PIB cohorts are provided in the Appendix) starting at the -1 dose level (VEN 200 mg) and escalating to dose level 0 (VEN 400 mg). Patients receiving at least one dose of VEN were included in the intention-to-treat safety and efficacy analysis. PIB patients receiving a minimum of 80% of planned VEN doses and at least 3 days of FLAG-IDA were evaluable for doselimiting toxicity (DLT). The phase II (dose-expansion) portion enrolled separate cohorts of ND-AML (PIIA) and R/R-AML (PIIB) patients at the recommended PII dose.

Treatment Administration

FLAG-IDA induction consisted of 28-day cycles of intravenous (IV) fludarabine (30 mg/m²) and cytarabine (1.5-2 g/m² IV) on days (D) 2-6, idarubicin (IV; ND-AML, 8 mg/m² D4-6; R/R-AML, 6 mg/m² D4-5), and filgrastim (5 mcg/kg D1-7). Consolidation used reduced durations of fludarabine and cytarabine (D2-4) and filgrastim (D1-5); idarubicin was permitted (D3-4) in up to two consolidation cycles at the discretion of the treating physician. At the recommended phase II dosing, VEN was administered on D1-14 during induction and D1-7 in consolidation. PEGylated filgrastim was permitted after D5 (induction) or D3 (consolidation) to replace remaining G-CSF doses. VEN dose adjustments for patients receiving CYP3A inhibitors such as azole antifungals followed US prescribing information recommendations (Appendix). Antimicrobial prophylaxis was recommended during periods of neutropenia.

Because of pronounced grade 3 and 4 neutropenia-related infectious complications and one DLT (typhlitis) in the original PIB dose -1 level (n = 8), the Protocol (online only) was amended to evaluate an alternate dose level -1, reducing the VEN induction duration to 14 days (from 21) and with attenuated cytarabine (1.5 g/m² from 2 g/m²). Venetoclax (D1-14) was then administered at 200 mg (alternate dose level -1, n = 5) and 400 mg (dose level 0, n = 3), and dose level 0 was confirmed as the recommended phase II dose for expansion. Additional details are provided in the Appendix. Following completion of induction or consolidation, continuous daily VEN maintenance was permitted on D1-28 of each 28-day cycle for up to 1 year in patients not proceeding to stem-cell transplantation.

Statistical Considerations

The dual primary objectives included safety and tolerability of FLAG-IDA + VEN, with identification of DLTs and determination of MTD (PIB), and assessment of overall activity (overall response rate [ORR]) per modified International Working Group criteria (PII).²¹ Secondary objectives included additional assessments of efficacy: CRc (CR + CRi + CR with partial hematologic recovery [CRh]), ORR (CR + CRh + CRi + morphologic leukemia-free state + partial response), OS (time from treatment initiation to death), event-free survival (EFS; time from treatment initiation until death or relapse, whichever occurred first: nonresponding patients were considered as progressing on cycle 1 day 1 for EFS), and duration of response (DOR; time from best response to relapse or death in responding patients only). Exploratory objectives included identification of biomarkers (ie cytogenetic and molecular mutations) predictive of VEN activity.

Futility and toxicity monitoring used a Bayesian method,²² applying monitoring rules to each arm separately (Appendix); 95% credible intervals were calculated for the primary objective for each PII arm and 95% exact CIs were computed for other response outcomes. Descriptive

statistics were assessed using the Fisher's exact test. Time-

RESULTS

Demographics

Sixty-eight patients (median age 46 years [range, 20-73]) have been enrolled (Table 1). Forty-one percent of ND-AML had sAML, ts-AML, or t-AML. European LeukemiaNet (ELN) risk across PIB, PIIA, and PIIB cohorts was favorable in 37.5%, 17%, and 26% of patients; intermediate in 12.5%, 45%, and 13%; and adverse in 50%, 38%, and 61%. Most R/R-AML patients (69%) were in salvage 1; 15% were in salvage 2 and 15% were in salvage 3 or greater. Forty-four percent PIB and 30% PIIB patients had received prior allogeneic hematopoietic stem-cell transplantation (alloHSCT).

Cytogenetic and Molecular Mutations at Study Enrollment

Diploid or other intermediate-risk cytogenetics (76%) were frequent in ND-AML, whereas adverse-risk or complex cytogenetics were common in R/R-AML (41%). Ten percent of patients harbored *KMT2A* rearrangements (*KMT2A*⁺). Epigenetic mutations (*DNMT3A, TET2, IDH1,* and *IDH2*) were more common in ND-AML compared with R/R-AML (41% v 15%; *P* value, .025), with enrichment of *IDH2* in ND-AML (24% v 5%; *P* value, .03). Conversely, mutations in *TP53* (18%) and *WT1* (13%) were frequent in R/R-AML (Appendix Fig A1, online only).

Treatment Characteristics

Patients received a median of 2 treatment cycles (range, 1-6; Appendix Table A1, online only). Sixty-nine percent (n = 47) received 1-2 cycles; 31% (n = 21) received ≥ 3 cycles. Four patients required reinduction (R/R-AML, 3; ND-AML, 1) with only the ND-AML patient responding (CRi). Median time to count recovery following induction (absolute neutrophil count \geq 500 and platelet count \geq 50,000) for PIB, PIIA, and PIIB patients was 37, 31, and 37 days (Appendix Figs A2A-F, online only) and was prolonged across all cohorts following cycle 2. Median time to count recovery for R/R-AML patients who underwent prior alloHSCT for cycles 1, 2, and 3 was 36, 41, and 37 days. Fifty-six percent of patients (n = 38: PIB: 38%, PIIA: 69%, and PIIB: 52%) transitioned to alloHSCT in remission after a median of 2 (range, 1-4) cycles. Sixtyseven percent (n = 20) of patients not undergoing alloHSCT received 1-2 cycles of therapy; 33% (n = 10) received \geq 3 cycles.

Cycle lengths extending \geq 40 days occurred in 19%, 59%, and 47% of patients completing cycles 1, 2, and 3. Myelosuppression was the leading cause of cycle delays, particularly following C2. Delayed count recovery requiring dose reductions in consolidation occurred in 24% (n = 11) of patients, and most frequently occurred in patients with

FLAG-IDA + VEN in Acute Myeloid Leukemia

TABLE 1. Patient Demographics

Parameter	Phase IIA ND-AML (n = 29)	Phase IB $R/R-AML (n = 16)$	Phase IIB R/R-AML (n = 23)	
Age, years	45 (20-65)	51 (20-73)	47 (22-66)	
Sex (male)	13	10	14	
VEN dose level				
Dose level -1 (VEN 200 mg, D1-21)	—	8	—	
Alternate dose level -1 (VEN 200 mg, D1-14)		5	_	
Dose level 0 (VEN 400 mg, D1-14)	29	3	23	
Median No. of prior therapies	_	2 (1-6)	1 (1-3)	
Prior HSCT	_	7	7	
Median duration of prior CR, months	_	15.1 (2.3-44)	12.6 (2.7-70)	
Salvage 1	—	8	19	
Salvage 2	—	3	3	
Salvage 3 or greater	_	5	1	
Median blast (%) at enrollment ^a	41 (4-85)	63 (6-94)	46 (1-89)	
Extramedullary leukemia	3	—	1	
AML type				
de novo AML	17	—	—	
sAML	5	—	—	
ts-AML	2	_	_	
t-AML	5	_	_	
R/R-AML	—	16	23	
ELN risk group				
Favorable	5	6	6	
Intermediate	13	2	3	
Adverse	11	8 14		
Cytogenetic group				
Favorable	_	4	2	
Diploid	13	2	8	
Other intermediate	8	2	3	
Adverse-risk or complex	4	4	9	
Insufficient mitoses	1	1	_	
KMT2A-rearranged	3	3	1	
Molecular mutations				
NPM1	3	2	3	
IDH1	3	1	_	
IDH2	7	1	1	
RUNX1	5	_	5	
ASXL1	2	_	3	
TP53	3	2	5	
Active signaling ^b	11	6	9	
Tumor suppressor ^c	5	5	8	

Abbreviations: AML, acute myeloid leukemia; CR, complete response; ELN, European LeukemiaNet; HSCT, hematopoietic stem-cell transplantation; MLL, mixed-lineage leukemia; ND-AML, newly diagnosed acute myeloid leukemia; R/R-AML, relapsed or refractory acute myeloid leukemia; sAML, secondary AML; t-AML, treatment related AML; ts-AML, treated secondary AML; VEN, venetoclax.

^aIncludes patients with isolated extramedullary AML.

^bActive signaling: K/NRAS, FLT3-ITD/TKD, KIT, CBL, and PTPN11.

°Tumor suppressor: TP53, WT1, FBXW7, and PHF6.

s-AML, t-AML, ts-AML, or R/R-AML following C1 (89%) and C2 (63%), possibly reflecting poor marrow reserve in these populations. Sixty-one percent (n = 23) of patients transitioned to alloHSCT without full hematologic recovery (ie, absolute neutrophil count < 500 and/or platelet count < 50,000).

Adverse Events

Grade 3 and 4 adverse events (AEs) occurring in \geq 2 patients are displayed in Figure 1; all grade 3 and 4 AEs are provided in Appendix Table A2 (online only). Grade 3 and 4 AEs occurring in \geq 10% of patients included febrile neutropenia (50%), bacteremia (35%), pneumonia (28%), and sepsis (12%). Febrile neutropenia and pneumonia occurred at similar frequencies in R/R-AML and ND-AML. Bacteremia was more common in R/R-AML (46% v 21%; P = .04), particularly PIB patients (50%; Fig 1B). Typhlitis was only observed in the original PIB cohort. Grade 3 and 4 AEs in R/ R-AML patients who received prior alloHSCT were predominantly infectious (90%). Across all cohorts, 30- and 60day mortality was 0% and 4.4%. Deaths on study or within one week of discontinuation all occurred in R/R patients, including four nonresponding (sepsis, n = 2; pneumonia, n = 1; pulmonary hemorrhage, n = 1) and two responding patients (sepsis and hemophagocytic syndrome).

Efficacy

The ORR for PIB, PIIA, and PIIB cohorts was 75% (95% CI, 48 to 93), 97% (95% credible interval, 85 to 99), and 70% (95% credible interval, 47 to 83]) (Table 2) with CRc attained in 75% (48 to 93), 90% (73 to 98), and 61% (39 to 80; Fig 2); 67% of patients with R/R-AML (including 57% [n = 8]patients receiving prior alloHSCT) and 83% with sAML, t-AML, or ts-AML attained a CRc. No significant response difference was observed between patients with refractory versus relapsed AML (56% v70%; P, .53). Median time to best response was 30 days, with ongoing responses in 70% of patients. Twelve patients (R/R-AML, 11; ND-AML, 1) were refractory to FLAG-IDA-VEN induction. Eighty-three percent (95% CI, 70 to 92) of patients in CRc attained MRD negativity (MRD⁻) including 96% (95% CI, 80 to 99) and 69% of patients with ND-AML (de novo AML: 94%; sAML, t-AML, or ts-AML: 100%) and R/R-AML, respectively (PIB: 58% [95% CI, 28 to 85]; PIIB: 79% [95% CI, 49 to 95]).

After a median follow-up of 12 months, 12-month OS for the overall population, PIIA patients, and PIIB patients was 70% (95% CI, 58 to 83), 94% (95% CI, 84 to 100), and 68% (95% CI, 49 to 94). Median EFS and OS were 6 (95% CI, 3 to not estimated [NE]) and 9 (95% CI, 4.9 to NE) months in PIB versus 11 (95% CI, 2 to NE) months and not reached (NR; 95% CI, 6 to NE) in PIIB patients (Figs 3A and 3B). Median DOR was 6 months (95% CI, 3 to NE) in PIB and NR in PII patients. The study population was predominantly composed of younger patients (median age, 46 years); however; 73% (n = 8) of patients age \geq 60 years (including 100% age \geq 65 [n = 4]) attained a CRc.

No survival difference was observed between patients age ≤ 60 (n = 57) or age > 60 years (n = 11).

Median OS in R/R-AML was 13 months (95% CI, 7 to NR). Survival was not influenced on the basis of receipt of prior alloHSCT (median OS 13 months [95% CI, 7.64 to NE]) or prior CR duration, although survival in patients with prior CR durations < 12 months was similar to patients refractory at study enrollment (n = 9; 7 [6 to NE] v 8 [5 to NE] months; P = .94). Seventy-six percent of patients in salvage 1 or 2 and 17% in salvage 3 or greater achieved a CRc. Median OS in R/R-AML in salvage 1 (n = 27) or salvage 2 (n = 6) was significantly longer than that in patients in salvage 3 or greater (n = 6; median OS: 14 [10 to NE] v 4 [4 to NE] months; P = .003; Fig 4A).

Cytogenetic and Molecular Subgroups

Overall, 100%, 85%, and 91% of ND-AML and 83%, 60%, and 59% of R/R-AML patients with ELN favorable-, intermediate-, and adverse-risk disease achieved a CRc. Four patients with extramedullary AML (ND-AML, 3; R/R-AML, 1) had durable responses, with three transitioning to alloHSCT. *KMT2A-rearranged* patients (n = 7; ND-AML, 3; R/R-AML, 4) attained a 100% CRc rate (80% MRD⁻ by reverse transcriptase polymerase chain reaction for *KMT2A*) with a resultant 12-month OS of 80% (95% CI, 52 to 100).

Biomarkers predictive of response were only discriminatory within R/R-AML, given the high response rate (97%) in ND-AML. R/R-AML with favorable-risk cytogenetics (n = 6) correlated with an unpredictably poor median EFS and OS of 4 (95% CI, 4 to NE) and 7.6 (95% CI, 4 to NE) months; median EFS and OS were 7 (95% CI, 3 to NE) and 11 (95% CI, 5 to NE) months in R/R-AML patients with complex or adverse-risk cytogenetics (n = 16).

Molecular subgroups (*NPM1*, *IDH1*, or *IDH2*; n = 7) conferring sensitivity to VEN-based therapy^{13,23,24} demonstrated a 100% CRc rate and a 12-month OS of 83% in R/R-AML. Conversely, tumor suppressor mutations (*TP53*, *WT1*, *FBXW7*, and *PHF6*; n = 13) associated with treatment resistance (CRc with mutation: 38% v without: 77%; P = .021) were enriched in nonresponders (73%) versus responders (18%; P = .002) and trended toward inferior OS (7 [5 to NE] v 16 [11 to NE] months; P = .054; Fig 4C). Signaling mutations (*RAS*, *FLT3*, *PTPN11*, *CBL*, and *KIT*; n = 15) similarly correlated with inferior survival in R/R-AML (median OS, 6 [4 to NE] v 16 [14 to NE] months; P = .0081; Fig 4D).

Ten patients (ND-AML, 3; R/R-AML, 7) had detectable *TP53* mutations (*TP53*⁺) at baseline. Sixty percent attained a CRc (ND-AML, 3/3; R/R-AML, 3/7) including 4 with MRD⁻ CRc by FC. Median DOR and OS in ND-AML were 3.4 (95% CI, 2 to NE) and 9 (95% CI, 9 to NE) months. In R/R-AML, median DOR and OS were 3.2 (95% CI, 2 to NE) and 7 (95% CI, 5 to NE) months. Of interest, *TP53*⁺ persisted in all four patients with MRD⁻ CRc and was identified in 64% (n = 7) of relapsed patients with available



FIG 1. Adverse events by (A) AML type and (B) cohort. AML, acute myeloid leukemia; ND-AML, newly diagnosed acute myeloid leukemia; R/R-AML, relapsed or refractory acute myeloid leukemia; SSTI, skin and soft tissue infection.

NGS, including three initially TP53 wild-type patients who developed *TP53*⁺ clones despite achieving a prior MRD⁻ CRc by FC.

Role of HSCT

Thirty-eight patients (ND-AML, 20 [69%]; R/R-AML, 18 [46%]) transitioned to alloHSCT, including 75% (n = 6) of responding R/R patients who had received a prior alloHSCT. A 3-month landmark analysis in R/R-AML demonstrated improved OS with consolidative alloHSCT in CRc versus without (median OS: NR [14 to NE] v7 [4 to NE] months; *P* value, .009; Fig 4B). Twelve-month OS was 87% in patients undergoing HSCT. After a median post-HSCT follow-up of 9 months, 12-month post-HSCT survival was 94% in ND-AML and 78% in R/R-AML. Thirty and 60-day post-HSCT mortality was 3%.

Three PIB patients transitioned to VEN maintenance, two patients who experienced significant infectious complications during induction precluding additional intensive chemotherapy and one following four cycles of FLAG-IDA + VEN induction or consolidation without plans for alloHSCT. Median EFS and OS with maintenance VEN were 8.8 (95% CI, 7 to NE) and 10.7 (95% CI, 10 to NE) months.

DISCUSSION

Despite AML induction regimens resulting in CR rates of 75%-85%,^{4,7,9} relapse remains the primary cause of

mortality.²⁵ Overall, 90% of ND-AML patients receiving FLAG-IDA + VEN achieved a CRc, including 81% with ongoing responses, with 12-month EFS of 85%. FLAG-IDA + VEN demonstrated a robust CRc rate of 83% in patients with sAML, ts-AML, or t-AML, an improvement compared with standard IC regimens in this higher-risk population.⁹

MRD-negative remissions translate into improved outcomes in ND-AML and R/R-AML and are increasingly considered an optimal IC end point, stratifying pre- and posttransplantation relapse risk.²⁶⁻²⁹ Development of potent frontline and salvage regimens capable of achieving this end point remain of particular interest. Ninety-six percent of ND-AML and 69% of R/R-AML receiving FLAG-IDA + VEN attained an MRD⁻ CRc, highlighting the regimen's capability of producing MRD-negative remissions. Additional follow-up is warranted to confirm the survival impact of MRD⁻ CRc within the ND-AML and R/R-AML cohorts.

Responses in R/R-AML vary greatly (CR rate, 20%-60%) by treatment selection and line of salvage therapy and often lack durability.^{10,12,30} FLAG-IDA salvage in older (median age, 62 years) patients results in CR rates of approximately 20% and OS of 3.5 months¹⁰ Here, FLAG-IDA + VEN in a largely younger R/R-AML population improves upon historical outcomes with a CRc rate of 67% and median OS of

TABLE 2. Patient Outcomes

Outcome	AII (N = 68)	Phase IIA ND-AML (n = 29)	Phase IB R/R-AML (n = 16)	Phase IIB $R/R-AML (n = 23)$
ORR, No. (% [CI])	56 (82 [71 to 91])	28 (97 [85 to 99])ª	12 (75 [48 to 93])	16 (70 [47 to 83]) ^a
CRc (CR + CRi + CRh), No. (% [95% CI])	52 (76 [65 to 86])	26 (90 [73 to 98])	12 (75 [48 to 93])	14 (61 [39 to 80])
CR, No. (%)	37 (53)	20 (69)	6 (38)	11 (48)
CRh, No. (%)	10 (15)	5 (17)	2 (13)	3 (13)
CRi, No. (%)	5 (7)	1 (3)	4 (25)	—
MRD ⁻ CR (flow cytometry), No. (% [95% CI])	43 (83 [70 to 92])	25 (96 [80 to 99])	7 (58 [28 to 85])	11 (79 [49 to 95])
MLFS	4	2	—	2
No response	12	1	4	7
DOR (median, months)	NR	NR	6 (3 to NE)	NR
EFS				
Median, months (95% CI)	18 (10.1 to NE)	NR	6 (3 to NE)	11 (2 to NE)
6-month, % (95% CI)	70 (59 to 81)	89 (78 to 100)	50 (31 to 82)	59 (41 to 84)
12-month, % (95% CI)	56 (44 to 71)	85 (72 to 100)	31 (15 to 65)	41 (21 to 77)
OS				
Median, months (95% CI)	NR	NR	9 (4.9 to NE)	NR (6 to NE)
6-month, % (95% CI)	81 (71 to 91)	100	63 (43 to 91)	68 (49 to 94)
12-month, % (95% CI)	70 (58 to 83)	94 (84 to 100)	38 (20 to 71)	68 (49 to 94)

Abbreviations: AML, acute myeloid leukemia; CR, complete response; CRc, composite CR; CRh, CR with partial hematologic recovery; CRi, CR with incomplete hematologic recovery; DOR, duration of response; EFS, event-free survival; HSCT, hematopoietic stem-cell transplantation; MLFS, morphologic leukemia-free state; MRD, measurable residual disease; ND-AML, newly diagnosed acute myeloid leukemia; NE, not estimated; NR, not reached; ORR, overall response rate; OS, overall survival; R/R-AML, relapsed or refractory acute myeloid leukemia.

^a95% credible intervals per Protocol-defined primary efficacy outcome (95% credible interval estimation assumed ORR follows a prior distribution of beta [1.4, 0.6] and beta [0.6, 1.4] in PIIA and PIIB, respectively). All other reported intervals represent exact 95% Cls.

13 months. Patients receiving salvage 1 or 2 attained a 76% CRc rate, with a median OS of 14 months. Patients receiving FLAG + IDA + VEN as salvage 3 or greater experienced reduced CRc rates (17%) and median OS (4 months), reflecting the unmet therapeutic need in this population.¹²

Across ELN risk groups, FLAG-IDA + VEN resulted in high CRc rates. Diploid and intermediate-risk cytogenetics predicted favorable outcomes. Inferior survival was observed in patients with adverse-risk or complex cytogenetics, although FLAG-IDA + VEN improved outcomes compared with contemporary analyses of IC in this cytogenetic subgroup.^{31,32} Favorable-risk cytogenetics, implicated in upregulation of alternative BCL-2 proteins and a monocytic phenotype, may contribute to VEN resistance, partially accounting for the poor outcomes observed within this favorable but multiply relapsed subgroup.^{33,34}

NPM1-, IDH1-, or *IDH2-*mutated AML had favorable responses to FLAG-IDA + VEN, whereas tumor suppressor mutations, in particular *TP53,* resulted in primary and secondary resistance and similar to signaling mutations predicted inferior survival in R/R-AML. Acknowledging the small sample sizes and exploratory nature of included

molecular subgroup analyses, cautious interpretation and confirmation within larger study populations are warranted.

FLAG-IDA + VEN permitted transition to alloHSCT in 69% of ND-AML and 46% of R/R-AML patients, improving OS in R/R-AML compared with those not undergoing alloHSCT in CRc. Comparisons are limited by confounding reasons for patients not receiving alloHSCT; however, the high pre-transplant MRD⁻ CRc rate and favorable posttransplant survival (1-year post-HSCT OS, 78%) suggest FLAG-IDA + VEN is effective to bridge R/R-AML patients to alloHSCT.

At the phase II dosing regimen, FLAG + IDA + VEN was tolerable, with no early mortality observed. Grade 3 and 4 AEs occurring with FLAG-IDA + VEN were primarily infectious. Febrile neutropenia, bacteremia, and pneumonia accounted for the majority of AEs with observed rates similar to induction regimens across varying AML types, particularly sAML or R/R-AML.^{3,4,10,35} Despite G-CSF utilization, delayed count recovery following C2 was common, with cycle lengths exceeding 40 days in 59% of patients, similar to prior investigations of FLAG-IDA.³ Approximately one quarter of patients required dose reductions in consolidation, and 61% transitioned to HSCT without complete count recovery. Myelosuppression was most



FIG 2. (A) Swimmers plot of all study participants and (B) response across study cohorts. Horizontal arrows indicate patients remaining on study. CR, complete response; CRc, composite CR; CRh, CR with partial hematologic recovery; CRi, CR with incomplete count recovery; HSCT, hematopoietic stemcell transplantation; MLFS, morphologic leukemia-free state; MRD, measurable residual disease; ND-AML, newly diagnosed acute myeloid leukemia; NR, not reached; PD, progressive disease; R/R-AML, relapsed or refractory acute myeloid leukemia.

pronounced in patients with sAML or t-AML, or R/R-AML. G-CSF administration, frequent monitoring of peripheral blood counts, antimicrobial prophylaxis, and surveillance for infectious complications are essential for optimal patient support.

FLAG-IDA + VEN represents an effective intensive induction regimen for ND-AML and R/R-AML, with particular utility as a bridge to alloHSCT in the R/R-AML population. Confirmation of these results in ongoing dose-expansion



FIG 3. (A) EFS and (B) OS by cohort. EFS, event-free survival; ND-AML, newly diagnosed acute myeloid leukemia; OS, overall survival; R/R-AML, relapsed or refractory acute myeloid leukemia.



FIG 4. Outcomes in R/R-AML on the basis of (A) salvage number and (B) 3-month landmark analysis of HSCT in patients attaining CRc. Outcomes in R/R-AML with versus without mutations in (C) tumor suppressor and (D) active signaling genes. alloHSCT, allogeneic hematopoietic stem-cell transplantation; HSCT, hematopoietic stem-cell transplantation; ND-AML, newly diagnosed acute myeloid leukemia; OS, overall survival; R/R-AML, relapsed or refractory acute myeloid leukemia.

cohorts and through randomized comparison with standardof-care induction regimens is necessary to confirm the safety and effectiveness of FLAG-IDA + VEN both as a

frontline induction regimen in newly diagnosed AML and an optimal salvage regimen in fit patients with relapsed or refractory AML.

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AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST

Venetoclax Combined With FLAG-IDA Induction and Consolidation in Newly Diagnosed and Relapsed or Refractory Acute Myeloid Leukemia

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APPENDIX.

SUPPLEMENTARY METHODS

PIB Dosing Cohorts

The PIB (dose-escalation) arm applied a 3 + 3 dose escalation and deescalation algorithm to determine the MTD starting at the -1 dose level (venetoclax [VEN] 200 mg) and escalating to dose level 0 (VEN 400 mg). Fludarabine, cytarabine, granulocyte colony-stimulating factor, and idarubicin (FLAG-IDA) induction consisted of 28-day cycles of intravenous (IV) fludarabine (30 mg/m²) and cytarabine (1.5-2 g/m² IV) on days (D) 2-6, idarubicin (IV; newly diagnosed acute myeloid leukemia [ND-AML]: 8 mg/m² D4-6; relapsed or refractory acute myeloid leukemia [R/R-AML]: 6 mg/m² D4-5), and filgrastim (5 mcg/kg D1-7). Consolidation used reduced durations of fludarabine and cytarabine (D2-4) and filgrastim (D1-5); idarubicin was permitted (D3-4) in up to two consolidation cycles at the discretion of the treating physician. PEGylated filgrastim was permitted after D5 (induction) or D3 (consolidation) to replace remaining granulocyte colonystimulating factor doses. VEN dose adjustments for patients receiving azole antifungals followed FDA recommendations.

Because of increased rates of grade 3 and 4 infectious complications in the original PIB patients treated at dose level -1 (n = 8) including one dose-limiting toxicity (DLT) of typhlitis, the Protocol was amended to use a reduced duration of venetoclax during induction and attenuated cytarabine (1.5 g/m² in induction and consolidation). At this alternate dose level -1, no DLTs occurred and the dose was escalated to dose level 0, evaluating 400 mg of venetoclax. Dose level 0 was also determined to be safe and was selected for the phase II doseexpansion arms.

Note that in the original study design, the definition of DLT evaluable required the receipt of two cycles of therapy. Because of efficacy observed at even the dose -1 level, many patients in remission transitioned to allogeneic hematopoietic stem-cell transplantation after one induction cycle, rendering them unevaluable for DLT and requiring replacement. A Protocol amendment, which updated the DLT evaluation period to one cycle of therapy, was approved by all applicable regulatory bodies in August 2018 and dose level 0 opened for patient enrollment in September 2018.

PIB Treatment Administration

Induction dosing schema after Protocol amendment

Dose Level	VEN Duration	VEN Dose, mg	Cytarabine Dose, g/m²
Original dose level -1	D1-21	200	2
Alternate dose level -1	D1-14	200	1.5
Dose level 0	D1-14	400	1.5

Consolidation dosing schema

Dose Level	VEN Duration	VEN Dose, mg	Cytarabine Dose, g/m²
Original dose level –1	D1-14	200	2
Alternate dose level -1	D1-7	200	1.5
Dose level 0	D1-7	400	1.5

ADDITIONAL STATISTICAL CONSIDERATIONS

Futility and Toxicity Monitoring for Dose-Expansion Cohorts (PIIA and PIIB)

Futility and toxicity monitoring used a Bayesian method,²² applying monitoring rules to each arm separately. The ND-AML cohort will be stopped early if a > 98% probability exists that the overall response rate (ORR) with FLAG-IDA + VEN was less than the overall response rate under standard-of-care treatment (SOC) plus 15% (ie, it is less likely that the study treatment will improve ORR by 15% over SOC) or if a > 88% probability existed that the DLT rate was > 30%. Similarly, the R/R-AML cohort will be stopped early if there is a > 99% probability that the ORR with FLAG-IDA + VEN was less than the ORR under SOC plus 10% or if > 90% probability existed that the DLT rate was > 30%.



FIG A1. Genomic landscape of AML cohorts. AML, acute myeloid leukemia; CR, complete response; CRh, CR with partial hematologic recovery; CRi, CR with incomplete count recovery; MLFS, morphologic leukemia-free state; NR, not reached.



FIG A2. Median time to count recovery (ANC \geq 500; platelet count \geq 50,000 cells/ μ L) by (A-C) cycle and (D-F) study cohort. ANC, absolute neutrophil count.

TABLE A1. Treatment Characteristics

Parameter	All (N = 68)	Phase IIA: ND-AML ($n = 29$)	Phase IB: R/R -AML ($n = 16$)	Phase IIB: R/R -AML ($n = 23$)
Median No. of cycles	2 (1-6)	2 (1-5)	2 (1-6)	2 (1-4)
Median cycle length (days) ^a				
Cycle 1 (No. [range, days])	33 (48 [23-59])	31 (26 [27-59])	36 (9 [31-55])	35 (13 [23-47])
Cycle 2 (No. [range, days])	41 (27 [26-91])	42 (14 [27-60])	47 (4 [26-91])	37 (9 [26-90])
Cycle 3 (No. [range, days])	39 (15 [21-69])	39 (10 [25-69])	40 (2 [34-45])	39 (3 [21-40])
Transitioned to maintenance	3	—	3	—
Transitioned to HSCT	38 (56%)	20 (69%)	6 (38%)	12 (52%)
Median time to best response, days	30	29	35	27
Median duration of response, months	NR (14.5-NE)	NR (17-NE)	6 (3-NE)	NR (7-NE)
30-day mortality				

Abbreviations: HSCT, hematopoietic stem-cell transplantation; ND-AML, newly diagnosed patients with acute myeloid leukemia; NR, not reached; R/R-AML, relapsed or refractory acute myeloid leukemia.

^aFor patients proceeding with a subsequent cycle of treatment.

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	All Patients (N = 68), No. (%)	Phase IIA: ND-AML $(n = 29)$		Phase IB: $R/R-AML$ (n = 16)		Phase IIB: R/R -AML (n = 23)	
Grade 3/4 AEs		Grade 3	Grade 4	Grade 3	Grade 4	Grade 3	Grade 4
Febrile neutropenia	34 (50)	14		8		12	
Bacteremia	24 (35)	6	_	7	1		10
Pneumonia	19 (28)	8	_	3	1	7	_
Sepsis	8 (12)	_	3	_	4	_	1
Abdominal pain	5 (7)	2		1	_	2	_
Skin and soft tissue infection	4 (6)	3	_	_	_	1	_
Colitis or typhlitis ^a	3 (4)	1	_	1	1	_	_
GI disorders	2 (3)	_	_	_	_	2	_
GI hemorrhage	2 (3)	1	1	_	_	_	_
Hyperglycemia	2 (3)	1	_	_	1	_	_
Hypotension	2 (3)		_	_	1	1	_
Intracranial hemorrhage	2 (3)	_	_	_	_	1	1
Urinary tract infection	2 (3)	1	_	_	_	_	1
Cardiac disorder	1 (1.5)	1	_	_	_	_	_
Chest pain	1 (1.5)	1	_	_	_	_	_
Conjunctivitis	1 (1.5)	_	_	1	_	_	_
Epistaxis	1 (1.5)		_	_	_	1	_
Hypertension	1 (1.5)	_	_	_	1	_	_
Infusion reaction	1 (1.5)	1	_	_	_	_	_
Leukocytosis	1 (1.5)	1	_	_	_	_	
Periorbital cellulitis	1 (1.5)	1	_	_	_	—	—
Rectal pain	1 (1.5)	_	_	1	_	_	_

Abbreviations: AE, adverse event; DLT, dose-limiting toxicity; ND-AML, newly diagnosed acute myeloid leukemia; R/R-AML, relapsed or refractory acute myeloid leukemia.

1

^aColitis occurred in one PIIA patient. Typhlitis (G3 and G4) occurred in one PIB patient and was considered a DLT.

1 (1.5)

Upper respiratory tract infection