

# Ivosidenib (AG-120) in IDH1-mutant newly diagnosed acute myeloid leukemia: Updated results from a phase 1 study

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### BACKGROUND

Somatic mutations in isocitrate dehydrogenase (IDH) 1 and 2 result in accumulation of the oncometabolite D-2-hydroxyglutarate (2-HG),<sup>1,2</sup> which leads to epigenetic changes and impaired cellular differentiation.<sup>3,4</sup> Mutant IDH (mIDH) has been identified in multiple solid and hematologic tumors. In acute myeloid leukemia (AML), mIDH1 is found in ~6–10% of patients,<sup>5,6</sup> and is associated with an adverse prognosis.<sup>7,8</sup> Ivosidenib (AG-120) is a first-in-class, oral, potent, reversible, targeted inhibitor of the mIDH1 enzyme,<sup>9</sup> and is under evaluation in multiple clinical trials as a single agent and in combinations. Ivosidenib has been approved by the US FDA for the treatment of adult patients (with a susceptible *IDH1* mutation as detected by an FDA-approved test) with relapsed or refractory (R/R) AML, or with newly diagnosed (ND) untreated AML who are ≥75 years of age or have comorbidities precluding chemotherapy. The approval was on the basis of data from the ongoing phase 1 study of mIDH1 advanced hematologic malignancies (Figure 1).<sup>11</sup>

### OBJECTIVE

To report updated data on the safety and efficacy of ivosidenib in patients with ND AML not eligible for standard therapy, who are enrolled in the first-in-human, phase 1, dose escalation and expansion study of patients with mIDH1 advanced hematologic malignancies.

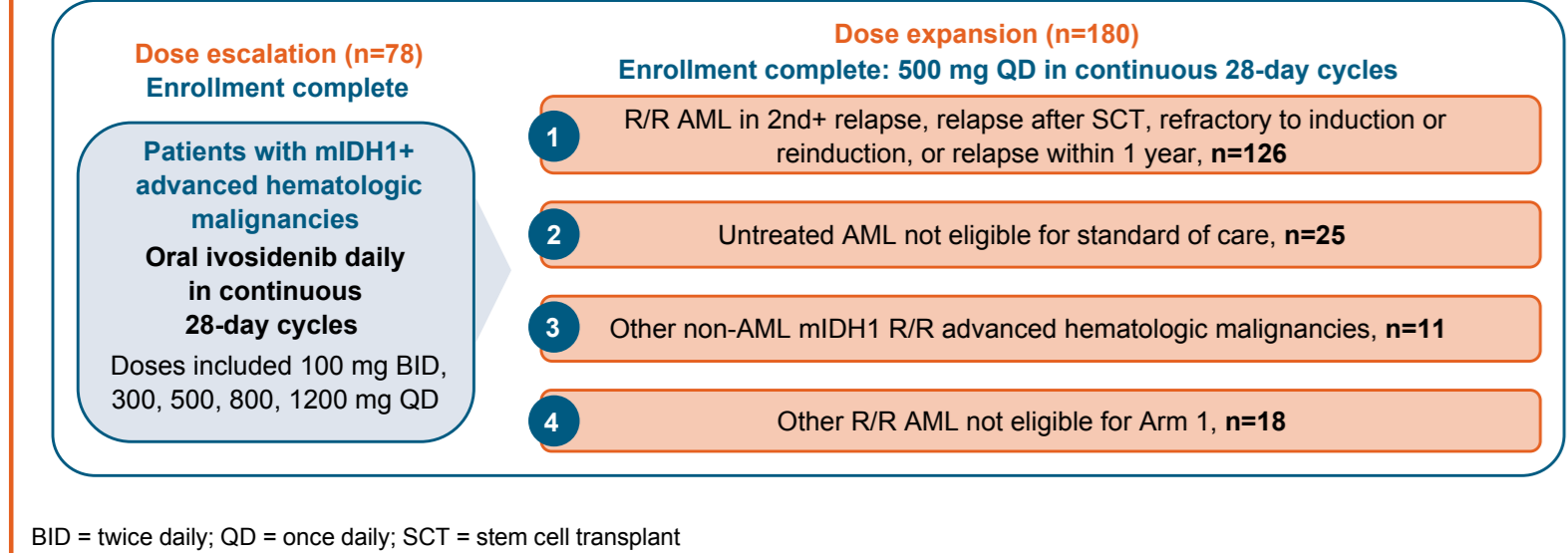
### METHODS

This study is an open-label, multicenter trial (ClinicalTrials.gov NCT02074839, Figure 1).

Baseline co-occurring mutations were assessed using a targeted next-generation sequencing (NGS) panel that detects common variants in hematologic malignancies.

*mIDH1* variant allele frequency (VAF) in bone marrow mononuclear cells was detected using BEAMING Digital PCR (Sysmex Inostics; lower limit of detection for *mIDH1*, 0.02–0.04%).

Figure 1. Study design



The primary efficacy endpoint was complete remission (CR) plus CR with partial hematologic recovery (CRh, Table 1).

Additional endpoints were overall response rate (ORR), CR, duration of response, and time to response. ORR comprised CR, CR with incomplete hematologic or platelet recovery (CRi/CRp), partial remission (PR), and morphologic leukemia-free state (MLFS, Table 1).

Table 1. Response definitions\*

Table with 4 columns: Response, Bone marrow blasts, %, Absolute neutrophil count, /μL, Platelets, /μL. Rows include CR, CRh, CRi, CRp, MLFS, and PR.

IWG responses, including CR, reported by investigator; CRh derived by sponsor \*Modified IWG 2003 criteria<sup>12</sup> <sup>†</sup>≥50% reduction IWG = International Working Group

Enrollment completed on May 8, 2017. The data cutoff date for this analysis was November 2, 2018. The analysis set comprised all patients with ND AML whose ivosidenib starting dose was 500 mg QD, (n=34; n=9 from dose escalation and n=25 from dose expansion, Arm 2). Efficacy endpoints were assessed in patients with ND AML confirmed positive for *mIDH1* by the companion diagnostic test (n=33). Disposition, treatment duration, baseline characteristics, and baseline co-occurring mutations are shown in Tables 2 and 3, and Figure 2.

### RESULTS

Table 2. Disposition and treatment duration

Table with 2 columns: Patients with ND AML receiving 500 mg ivosidenib n=34, and various disposition and treatment duration metrics.

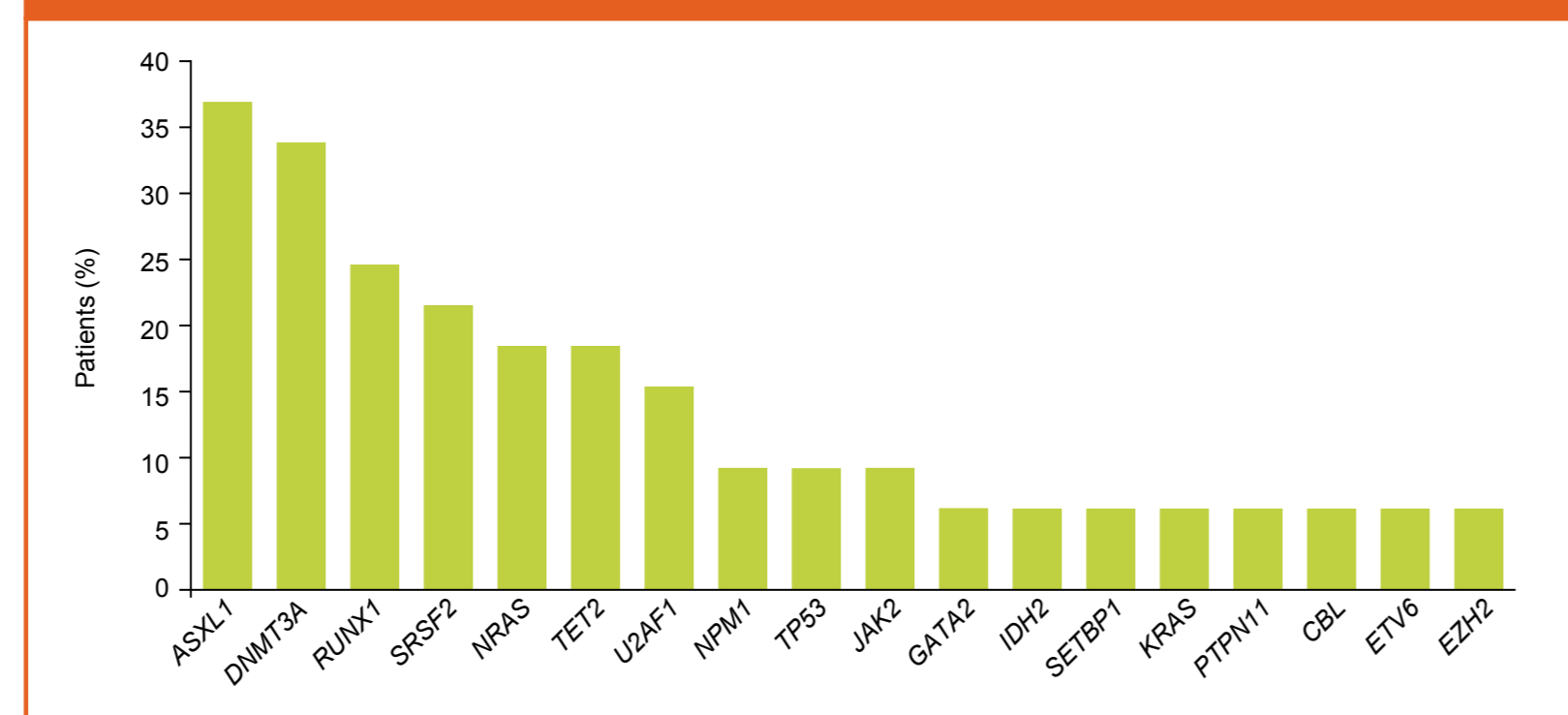
Data cutoff: November 2, 2018 AE = adverse event; PD = progressive disease

Table 3. Baseline characteristics

Table with 2 columns: Characteristic and Patients with ND AML receiving 500 mg ivosidenib n=34. Lists demographic and clinical characteristics.

Data cutoff: November 2, 2018 \*One patient enrolled in dose escalation tested positive for the IDH1-D54N mutation with local testing and did not test positive for the IDH1-R132 mutation with the companion diagnostic test

Figure 2. Baseline co-mutation rates: Patients with ND AML receiving 500 mg ivosidenib (n=33)\*



Data cutoff: November 2, 2018 Assessed using NGS panel for hematologic malignancies; mutations occurring in ≥2 patients shown \*One patient enrolled in dose escalation tested positive for the IDH1-D54N mutation with local testing and did not test positive for the IDH1-R132 mutation with the companion diagnostic test

Common AEs of any causality are shown in Table 4. Most AEs were grades 1–2 and unrelated to treatment. AEs of interest were managed using standard-of-care treatments and ivosidenib dose modifications as required (Figure 3).

Table 4. AEs regardless of causality: Patients with ND AML receiving 500 mg ivosidenib (n=34)

Table with 3 columns: Most common AEs (≥20% of patients), Any grade, n (%), Grade ≥3, n (%). Lists various adverse events.

Table 5. Responses in patients receiving 500 mg ivosidenib

Table with 2 columns: Patients with ND AML receiving 500 mg ivosidenib n=33, and various response metrics like CR+CRh rate, ORR, etc.

Data cutoff: November 2, 2018 \*ORR includes CR, CRi/CRp, MLFS, and PR mIWG = modified IWG; NE = not estimable; SD = stable disease; NA = not assessed

Figure 3. AEs of interest: Patients with ND AML receiving 500 mg ivosidenib (n=34)

Text boxes describing AEs of interest: Leukocytosis, Differentiation syndrome (DS), and ECG QT prolongation, including incidence and management details.

CR+CRh and CR rates were 42.4% and 30.3%, respectively (Table 5). CR+CRh rate was 26.7% (4/15) in patients who had previously been treated with HMAs, and 55.6% (10/18) in those who had not. Ivosidenib induced durable responses (Table 5 and Figure 4). Of 21 patients who were transfusion dependent at baseline, 43% became transfusion independent for ≥56 consecutive days while on treatment (Figure 5). In seven of 12 patients (58%) who were transfusion independent at baseline, independence was maintained for ≥56 days while on treatment. Ivosidenib induced IDH1 mutation clearance in bone marrow mononuclear cells (Table 6). The most frequent co-occurring mutations and mutational burden by clinical response are shown in Figure 6.

Table 6. IDH1 mutation clearance: Patients with ND AML receiving 500 mg ivosidenib (n=30)

Table with 2 columns: n, IDH1 mutation clearance n (%). Lists response categories and their corresponding mutation clearance rates.

\*Defined as a reduction in mIDH1 VAF to levels below the limit of detection of 0.02–0.04% (2–4 × 10<sup>-5</sup>) by digital PCR for at least one on-study time point <sup>†</sup>p-value based on Fisher's exact test comparing IDH1 mutation clearance in patients with a best overall response of CR+CRh with patients with other responses (non-CR+CRh responders and nonresponders)

Data cutoff: November 2, 2018 \*ORR includes CR, CRi/CRp, MLFS, and PR mIWG = modified IWG; NE = not estimable; SD = stable disease; NA = not assessed

Figure 4. Duration of treatment and best overall response: Patients with ND AML receiving 500 mg ivosidenib (n=33)

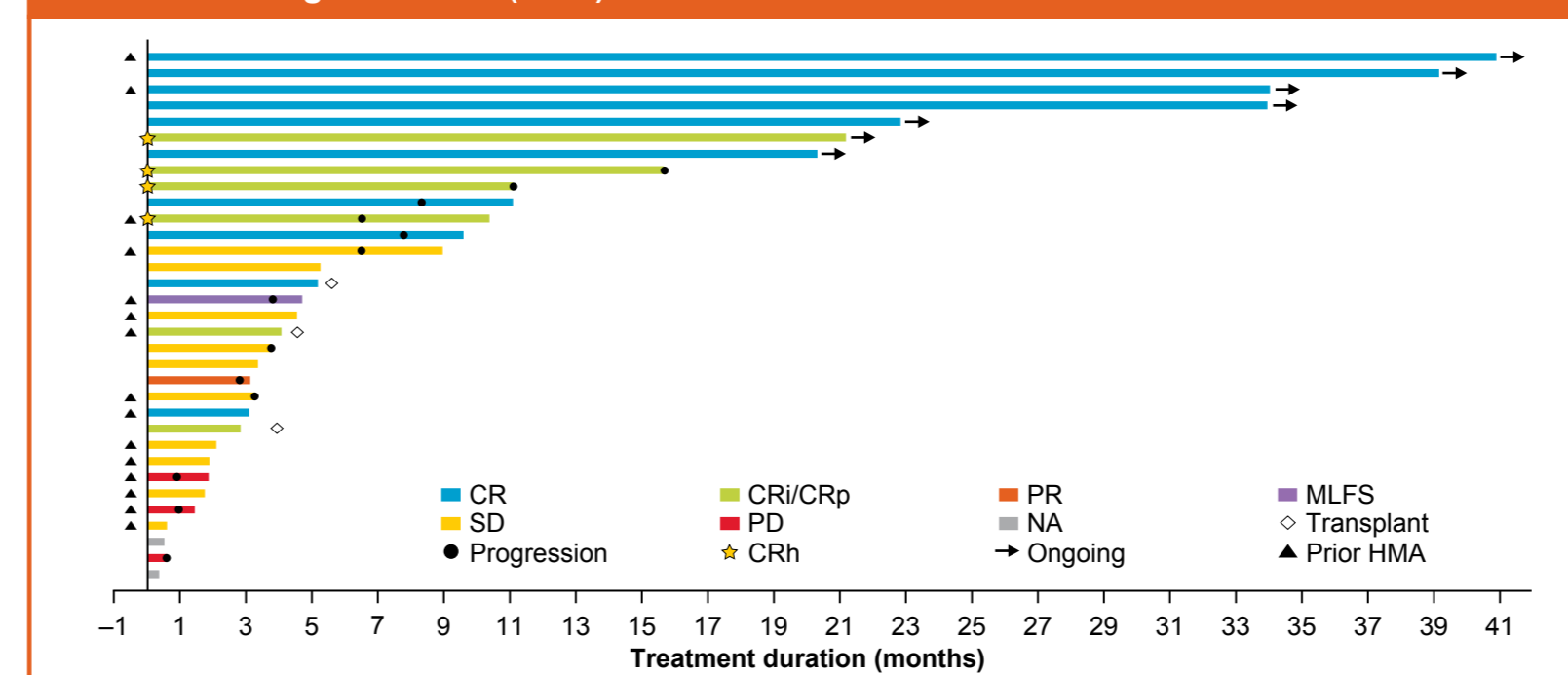
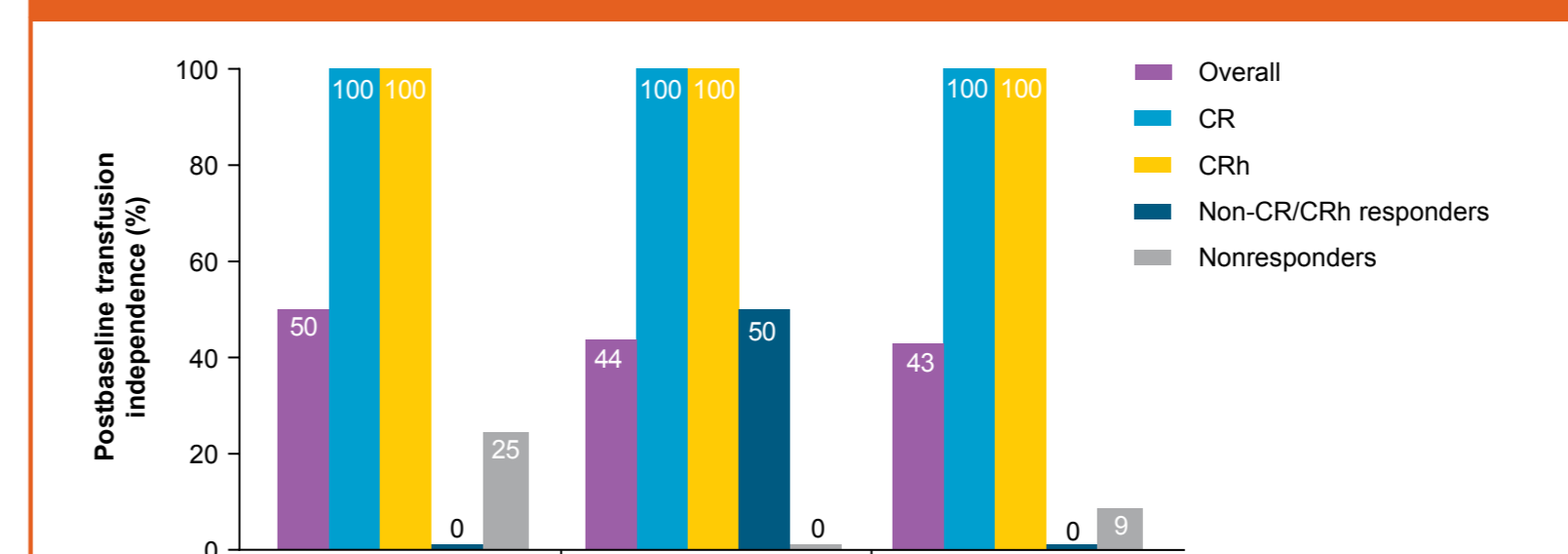


Table with 4 columns: Duration of response, Duration of CR+CRh, CR, Overall response. Lists median durations and Kaplan-Meier survival rates.

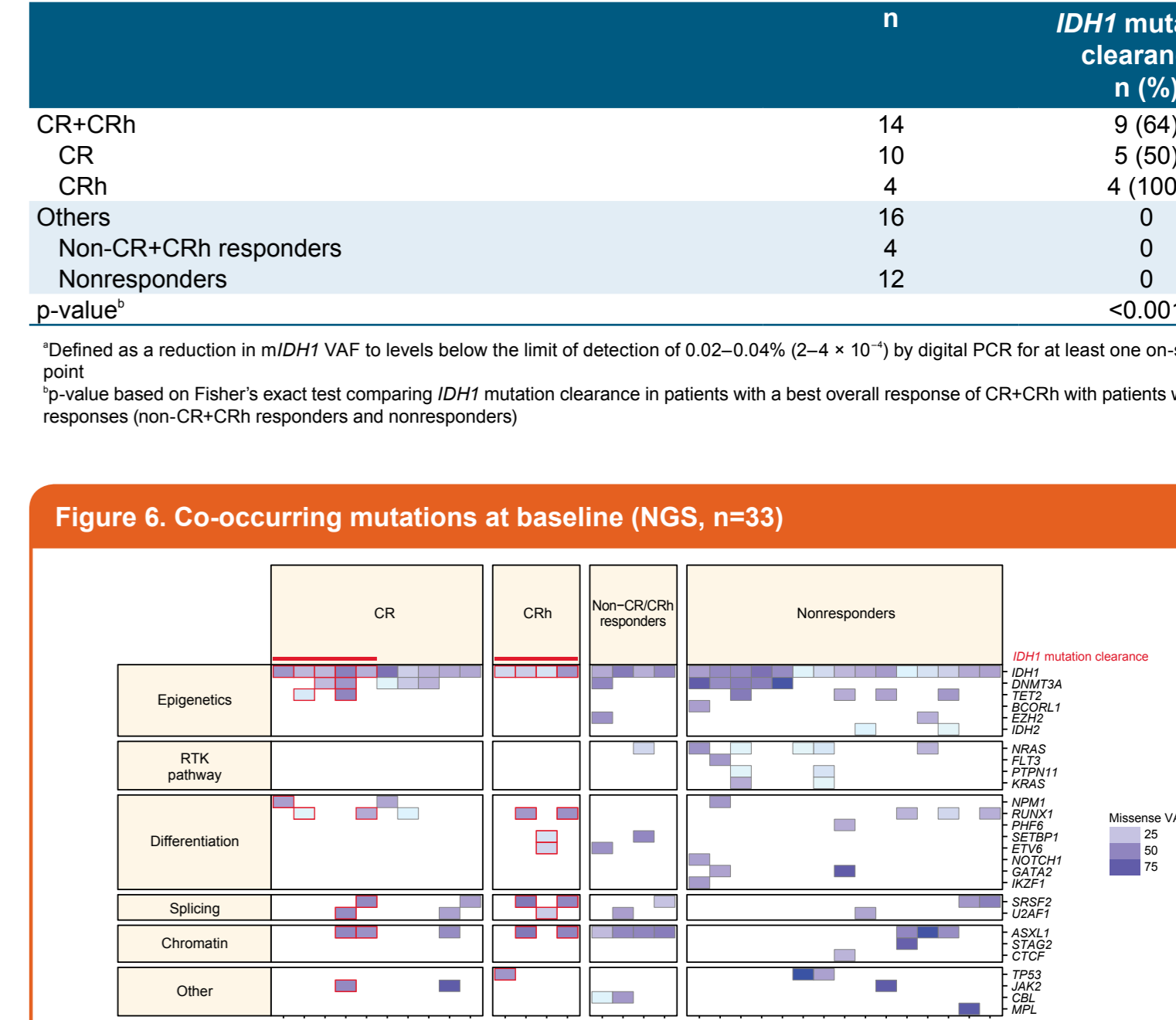
Data cutoff: November 2, 2018 HMA = hypomethylating agent

Figure 5. Transfusion independence across response categories in patients who were dependent at baseline (n=21): Patients with ND AML receiving 500 mg ivosidenib (n=33)



Postbaseline transfusion independence defined as no transfusion for at least one 56-day period. Data cutoff: November 2, 2018. Non-CR/CRh responders include patients who achieved CRi, CRp, or MLFS but not CRh

Figure 6. Co-occurring mutations at baseline (NGS, n=33)



In this heatmap, each column corresponds to a single patient, arranged by best overall response to ivosidenib. Detected known or likely oncogenic mutations are denoted by boxes and shaded by VAF category RTK = receptor tyrosine kinase

### CONCLUSIONS

- This cohort of patients with mIDH1 ND AML not eligible for standard therapy represent a molecularly defined elderly population with a poor prognosis. Ivosidenib induced durable responses, including in patients who had previously been treated with HMAs: Overall CR+CRh rate of 42.4%, median duration not estimable, lower bound of 95% CI 4.6 months. ORR of 54.5%, median duration not estimable. Ivosidenib was well tolerated. AEs of interest were managed with standard-of-care treatments and brief ivosidenib dose interruptions as required. There was a low rate of grade ≥3 AEs, including febrile neutropenia. Transfusion independence was observed across response categories. Ivosidenib induced IDH1 mutation clearance in five of 10 patients with CR and in all four patients with CRh. These results support the ongoing studies of ivosidenib in patients with mIDH1 AML: Phase 1 ivosidenib + azacitidine in patients with ND AML not eligible for intensive chemotherapy (see Poster 7011, DiNardo CD et al). AGILE: global, phase 3, ivosidenib + azacitidine versus placebo + azacitidine in patients with ND AML not eligible for intensive chemotherapy.<sup>13</sup> Phase 1 ivosidenib in combination with standard AML induction and consolidation therapy in patients with ND AML eligible for intensive chemotherapy.<sup>14</sup> HOVON 150 AML/AML-SG 29-18: phase 3, multicenter, randomized, placebo-controlled ivosidenib in combination with induction, consolidation, and maintenance therapy in patients with ND AML or MDS with excess blasts-2 eligible for intensive chemotherapy (ClinicalTrials.gov NCT03839771).

Acknowledgments, Disclosures, References, and QR code for more information.