

Ivosidenib (AG-120) in Mutant IDH1 AML and Advanced Hematologic Malignancies: Results of a Phase 1 Dose Escalation and Expansion Study

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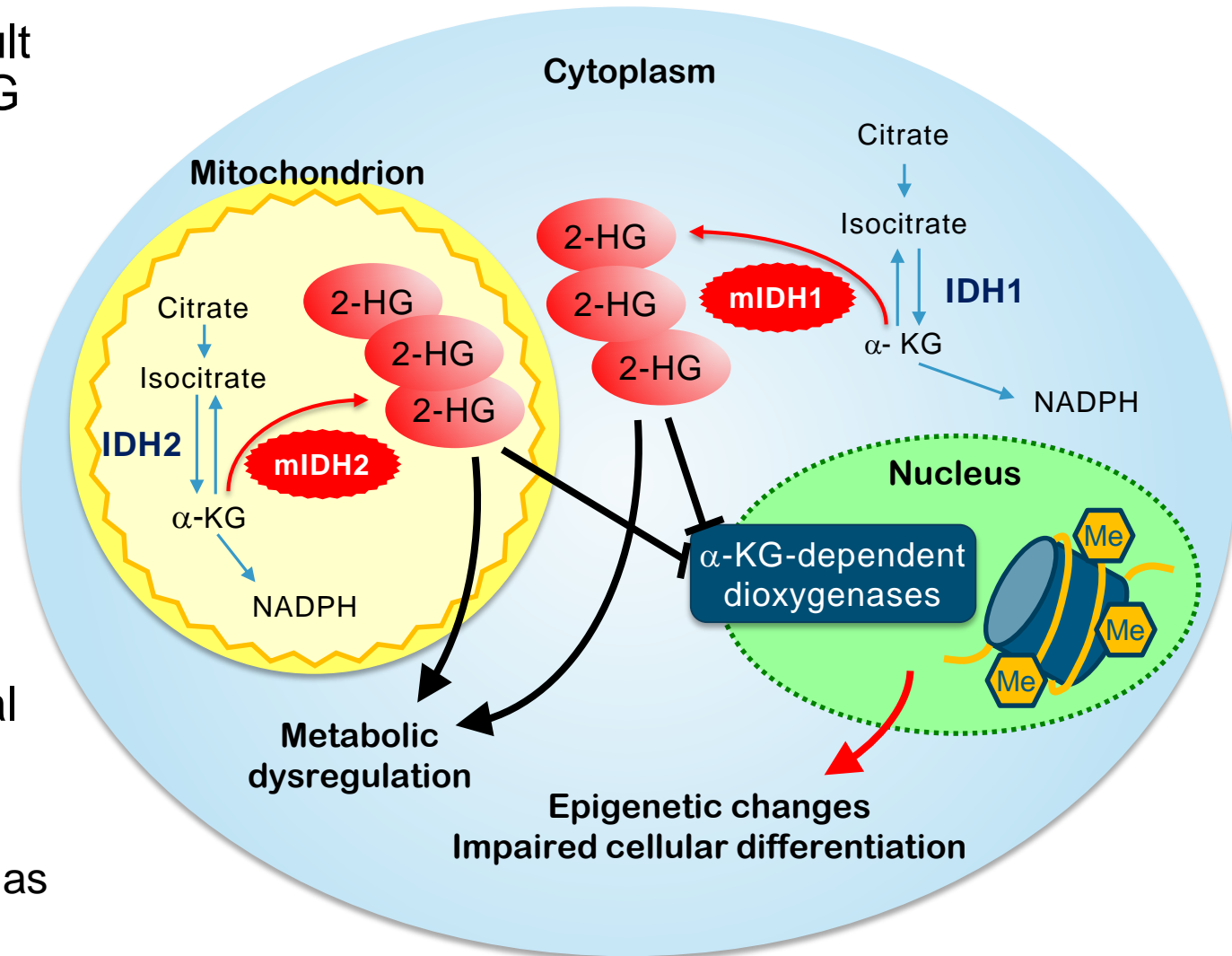
Isocitrate Dehydrogenase (IDH) Mutations as a Target in AML

- Somatic IDH1 and IDH2 mutations result in accumulation of oncometabolite 2-HG
 - epigenetic changes, impaired cellular differentiation

- mIDH identified in multiple solid and hematologic tumors

	mIDH1	mIDH2
% of AML patients	~6–10%	~9–13%

- **Ivosidenib (AG-120)**: an investigational first-in-class, oral, potent, reversible, targeted inhibitor of mIDH1 enzyme
 - under evaluation in multiple clinical trials as a single agent and in combinations



Study Design and Objectives

Single-arm, open-label, phase 1, multicenter trial

Dose escalation (n=78)

Enrollment complete

Patients with mIDH1+ advanced hematologic malignancies

Oral ivosidenib daily in continuous 28-day cycles

Doses included 100 mg BID, 300, 500, 800, 1200 mg QD

Dose expansion (n=180)

Enrollment complete: 500 mg QD in continuous 28-day cycles

- 1 R/R AML in 2nd+ relapse, relapse after SCT, refractory to induction or reinduction, or relapse within 1 year, n=126
- 2 Untreated AML not eligible for SOC, n=25
- 3 Other non-AML mIDH1 R/R advanced hematologic malignancies, n=11
- 4 Other R/R AML not eligible for Arm 1, n=18

Study objectives

- Primary** Safety and tolerability, MTD and/or RP2D, clinical activity in mIDH1 R/R AML enrolled in expansion Arm 1
- Secondary** DLTs, pharmacokinetics and pharmacodynamics (including 2-HG), preliminary clinical activity in advanced hematologic malignancies
- Exploratory** Determination of comutations and mIDH1 variant allele frequency (VAF)

Primary Efficacy Endpoint and Analysis Sets

Key analysis sets

- Safety Analysis Set (N=258): All treated patients
- Primary R/R AML Analysis Set (n=125):
 - The first 125 treated patients from Arm 1 of expansion (n=92) and eligible dose escalation patients (n=33) treated at 500 mg QD who were enrolled \geq 6 months prior to the primary analysis cutoff date of 12 May 2017

Primary efficacy endpoint for R/R AML: CR+CRh rate

Response	Bone marrow blasts (%)	ANC	Platelets
CR ¹	< 5	> 1000	> 100,000
CRh	< 5	> 500	> 50,000

IWG responses, including CR, reported by Investigator. CRh derived by Sponsor

Disposition and Treatment Duration

	All treated patients (N=258)	Primary R/R AML Set (n=125)
Ongoing treatment, n (%)	62 (24.0)	12 (9.6)
Discontinued treatment, n (%)	196 (76.0)	113 (90.4)
Progressive disease	104 (40.3)	66 (52.8)
Adverse event	33 (12.8)	17 (13.6)
Bone marrow transplant	22 (8.5)	12 (9.6)
Death	16 (6.2)	8 (6.4)
Withdrawal of consent	12 (4.7)	4 (3.2)
Investigator decision	7 (2.7)	5 (4.0)
Other	2 (0.8)	1 (0.8)
Discontinued study, n (%)	159 (61.6)	92 (73.6)
In post-transplant follow-up	8 (3.1)	4 (3.2)
In survival follow-up	29 (11.2)	17 (13.6)

- Median treatment duration: Primary R/R AML Set, 3.9 months (range 0.1–25.8)

Baseline Characteristics (n=125)

Characteristic	Primary R/R AML Set (n=125)
Women / men, n	60 / 65
Age in years, median (range)	67.0 (18–87)
ECOG PS at screening, n (%)	
0	27 (21.6)
1	64 (51.2)
De novo AML, n (%)	83 (66.4)
Secondary AML, n (%)	42 (33.6)
History of MDS	18 (14.4)
Therapy-related AML	14 (11.2)
No. of prior therapies, median (range)	2.0 (1–6)
Prior AML therapy outcomes ^a , n (%)	
Relapsed after transplant	36 (28.8)
In 2nd or later relapse	20 (16.0)
Refractory to initial induction/reinduction therapy	86 (68.8)
Relapsed ≤ 1 year of initial therapy	13 (10.4)

Characteristic	Primary R/R AML Set (n=125)
Cytogenetic risk status by investigator, n (%)	
Favorable	0
Intermediate	66 (52.8)
Poor	38 (30.4)
Unknown	4 (3.2)
Missing	17 (13.6)
Co-mutation rates ^b , n (%)	
FLT3	10 (8.1)
ITD	3 (2.4)
TKD	7 (5.6)
NPM1	24 (19.4)
CEBPA	3 (2.4)

^aNot mutually exclusive, patients may be in more than one category; ^bAssessed in 124 patients

Most Common AEs Regardless of Causality ($\geq 15\%$) (N=258)

All treated patients, N=258	Any grade, n (%)	Grade ≥ 3 , n (%)
Any AE	255 (98.8)	200 (77.5)
Diarrhea	86 (33.3)	6 (2.3)
Leukocytosis	78 (30.2)	17 (6.6)
Nausea	76 (29.5)	3 (1.2)
Fatigue	74 (28.7)	8 (3.1)
Febrile neutropenia	65 (25.2)	64 (24.8)
Dyspnea	61 (23.6)	9 (3.5)
Anemia	60 (23.3)	49 (19.0)
Electrocardiogram QT prolonged	58 (22.5)	23 (8.9)
Edema peripheral	56 (21.7)	0 (0.0)
Pyrexia	53 (20.5)	4 (1.6)
Decreased appetite	51 (19.8)	4 (1.6)
Constipation	48 (18.6)	2 (0.8)
Cough	48 (18.6)	1 (0.4)
Hypokalemia	45 (17.4)	7 (2.7)
Vomiting	45 (17.4)	3 (1.2)
Arthralgia	41 (15.9)	5 (1.9)
Thrombocytopenia	41 (15.9)	35 (13.6)
Dizziness	40 (15.5)	1 (0.4)
Epistaxis	39 (15.1)	2 (0.8)

AEs of Interest: Primary R/R AML Set (n=125)

■ Leukocytosis

- Grade ≥ 3 leukocytosis reported in 10/125 patients (8%)
- Managed with hydroxyurea
- None were fatal

■ ECG QT prolongation

- Grade 3 QT prolongation reported in 10/125 patients (8%)
- Study drug was reduced in 1 patient and held in 5 patients (all grades)
- None were Grade 4 or fatal

- These events were managed using standard of care treatments and ivosidenib dose modifications as required

■ IDH-differentiation syndrome (IDH-DS)

- All grade reported in 12/125 patients (9.6%)
- 4/12 IDH-DS patients had co-occurring leukocytosis
- Managed with corticosteroids and diuretics, and hydroxyurea if accompanied by leukocytosis
- None were Grade 4 or fatal
- Best response for the 12 patients with IDH-DS:

Best Response	CR	CRh	CRi/CRp	MLFS	SD
n=12	2	0	3	1	6

Grade 3 = WBC > 100,000/mm³; Grade 4 = clinical manifestations of leukostasis, urgent intervention indicated

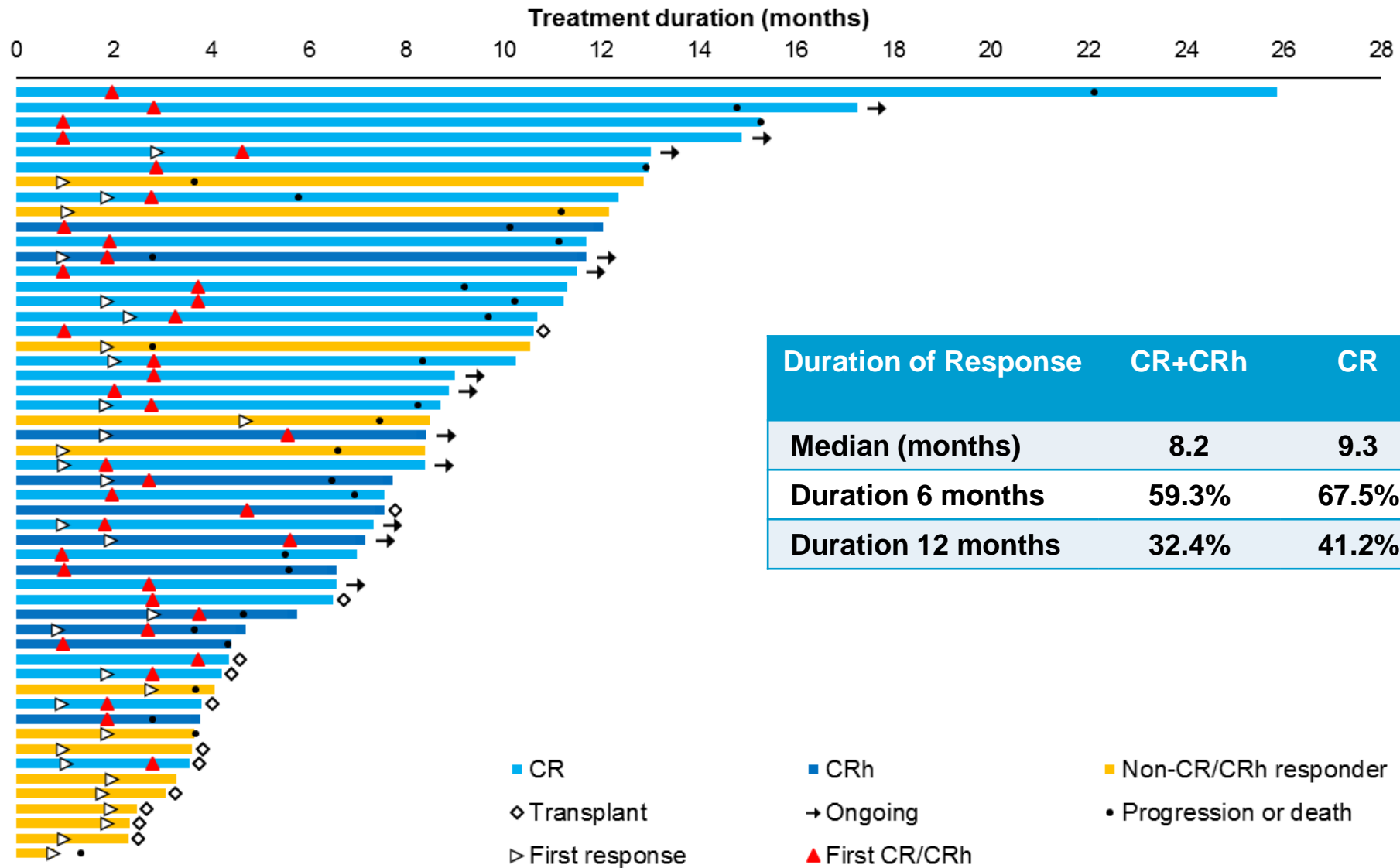
Response in R/R AML (n=125)

Primary R/R AML Set (n=125)	
CR+CRh rate, n (%) [95% CI]	38 (30.4%) [22.5, 39.3]
Time to CR/CRh, median (range) months	2.7 (0.9, 5.6)
Duration of CR/CRh, median [95% CI] months	8.2 [5.5, 12.0]
CR rate, n (%) [95% CI]	27 (21.6%) [14.7, 29.8]
Time to CR, median (range) months	2.8 (0.9, 8.3)
Duration of CR, median [95% CI] months	9.3 [5.6, 18.3]
CRh rate, n (%)	11 (8.8%)
Overall Response Rate, n (%) [95% CI]	
52 (41.6%) [32.9, 50.8]	
Time to first response, median (range) months	1.9 (0.8, 4.7)
Duration of response, median [95% CI] months	6.5 [4.6, 9.3]
Best response, n (%)	
CR	27 (21.6)
CRi or CRp	16 (12.8)
MLFS	9 (7.2)
SD	44 (35.2)
PD	13 (10.4)
NA	16 (12.8)

- CRh = 6 patients with investigator assessed responses of CRi/CRp and 5 with MLFS

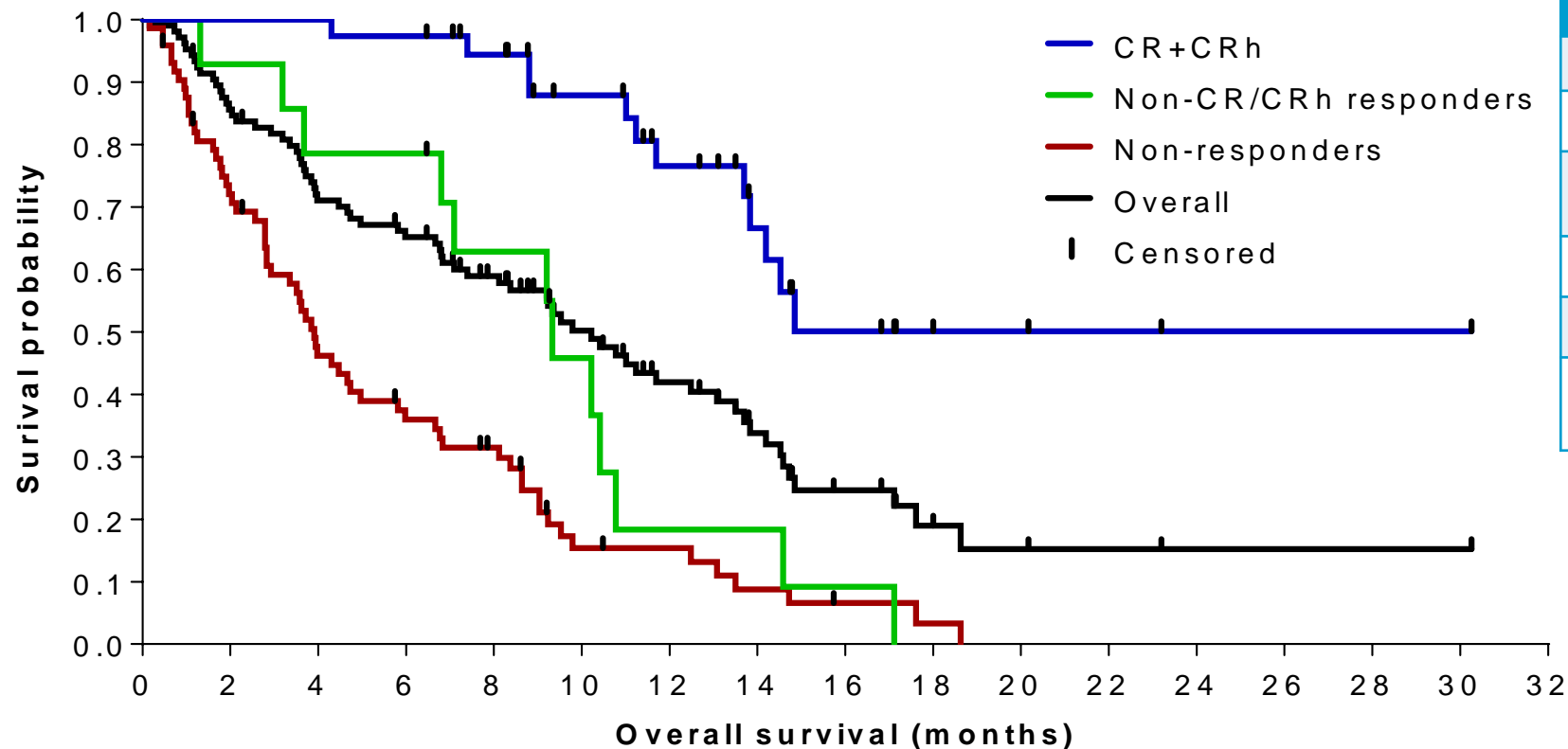
Duration of Treatment and Best Overall Response in Responders

Primary R/R AML Set (n=52)



Duration of Response	CR+CRh	CR	Overall Response
Median (months)	8.2	9.3	6.5
Duration 6 months	59.3%	67.5%	55.0%
Duration 12 months	32.4%	41.2%	24.6%

Overall Survival by Best Response in R/R AML (n=125)



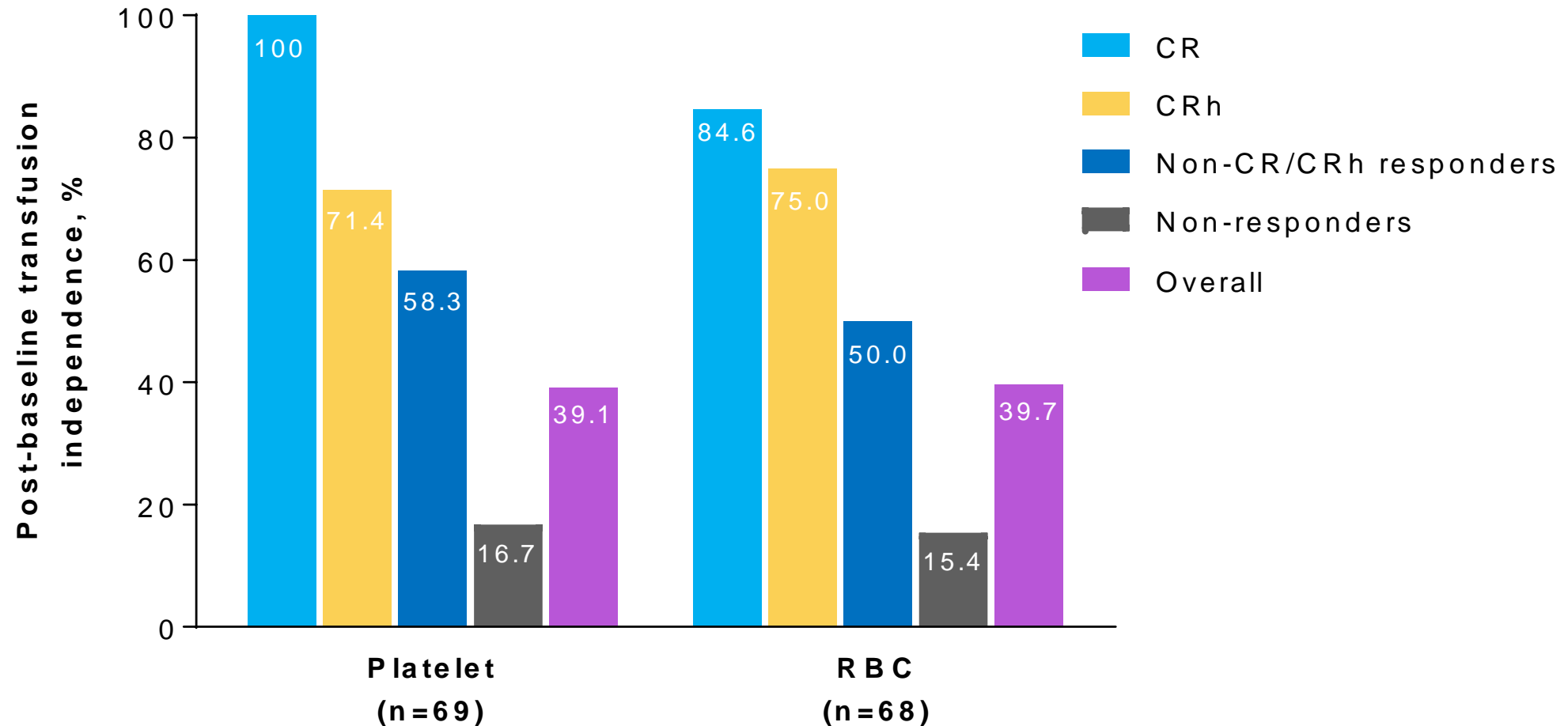
Months	
Overall Survival, median [95% CI]	
CR+CRh	NE [13.8, NE]
Non-CR/CRh responders	9.3 [3.7, 10.8]
Non-responders	3.9 [2.8, 5.8]
All	8.8 [6.7, 10.2]
Overall follow-up, median (range)	14.8 (0.2–30.3)

Number of patients at risk:

38	38	38	37	32	25	19	13	8	5	4	3	1	1	1	1	CR+CRh
14	13	11	11	8	5	2	2	1	0	Non-CR/CRh responders						
73	51	32	24	19	8	7	4	2	1	0	Non-responders					

Non-responders = all others including those with best responses of SD, PD or not evaluable

Transfusion Independence was Observed Across all Response Categories in Primary R/R AML Set Patients Who Were Dependent at Baseline



Post-baseline transfusion independence defined as no transfusion for at least one 56-day period.

Exposure-adjusted Incidence of Febrile Neutropenia and Grade \geq 3 Infections Primary R/R AML Set (n=125)

	Best Response				Overall (n=125)
	CR (n=27)	CRh (n=11)	Non-CR/CRh responders (n=14)	Non- responders (n=73)	
All Grade Febrile Neutropenia^a	2.6 [1.2, 5.4]	3.8 [1.2, 11.8]	3.9 [1.3, 12.1]	14.2 [10.0, 20.0]	6.9 [5.1, 9.2]
Grade \geq 3 Infections^b	2.6 [1.2, 5.4]	6.4 [2.6, 15.3]	14.4 [8.0, 25.9]	23.0 [17.6, 30.2]	11.5 [9.2, 14.4]

Incidence rate reported as 100 patients / month [95% CI]^c

^aPreferred term, including febrile bone marrow aplasia preferred term

^bBased on MedDRA V20.0 System Organ Class of infection and infestations

^cCalculated as total number of specific AEs / total person exposure time in months x 100 for all patients with the same best overall response

Response in Untreated AML and MDS

Characteristic	Untreated AML Arm 2 ^a (n=34)	MDS Arm 3 ^b (n=12)
Women / men, n	15 / 19	3 / 9
Age in years, median (range)	76.5 (64–87)	72.5 (52–78)
ECOG PS at screening, n (%)		
0	8 (23.5)	4 (33.3)
1	20 (58.8)	6 (50.0)
Prior MDS, n (%)	18 (52.9)	NA
Response		
Overall Response Rate, n (%) [95% CI]	19 (55.9) [37.9, 72.8]	11 (91.7) [61.5, 99.8]
Duration of response, median [95% CI] months	9.2 [1.9, NE]	NE [2.3, NE]
Duration of CR, median [95% CI] months	NE [5.6, NE]	NE [2.8, NE]
Best response, n (%)		
CR	7 (20.6)	5 (41.7)
CRi/CRp	7 (20.6)	n/a
PR	1 (2.9)	n/a
MLFS/mCR	4 (11.8)	6 (50.0)
SD	10 (29.4)	0
PD	3 (8.8)	1 (8.3)
NA	2 (5.9)	0

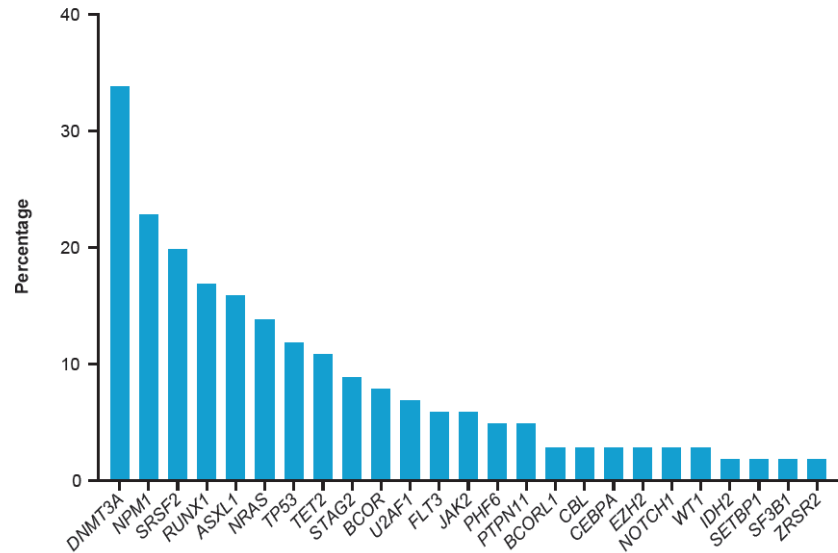
^aUntreated AML patients not eligible for standard of care therapies in expansion Arm 2 and from dose escalation whose starting dose was 500 mg QD

^bMDS patients in expansion Arm 3 and from dose escalation whose starting dose was 500 mg QD

Data cutoff: 12 May 2017. CR, complete remission; CRi, CR with incomplete hematologic recovery; CRp, CR with incomplete platelet recovery; mCR, marrow CR; MLFS, morphologic leukemia-free state; NA, not assessed; NE, not estimable; PD, progressive disease; PR, partial response; SD, stable disease

Baseline Mutation Analysis: Poster 2684, Stone RM et al.

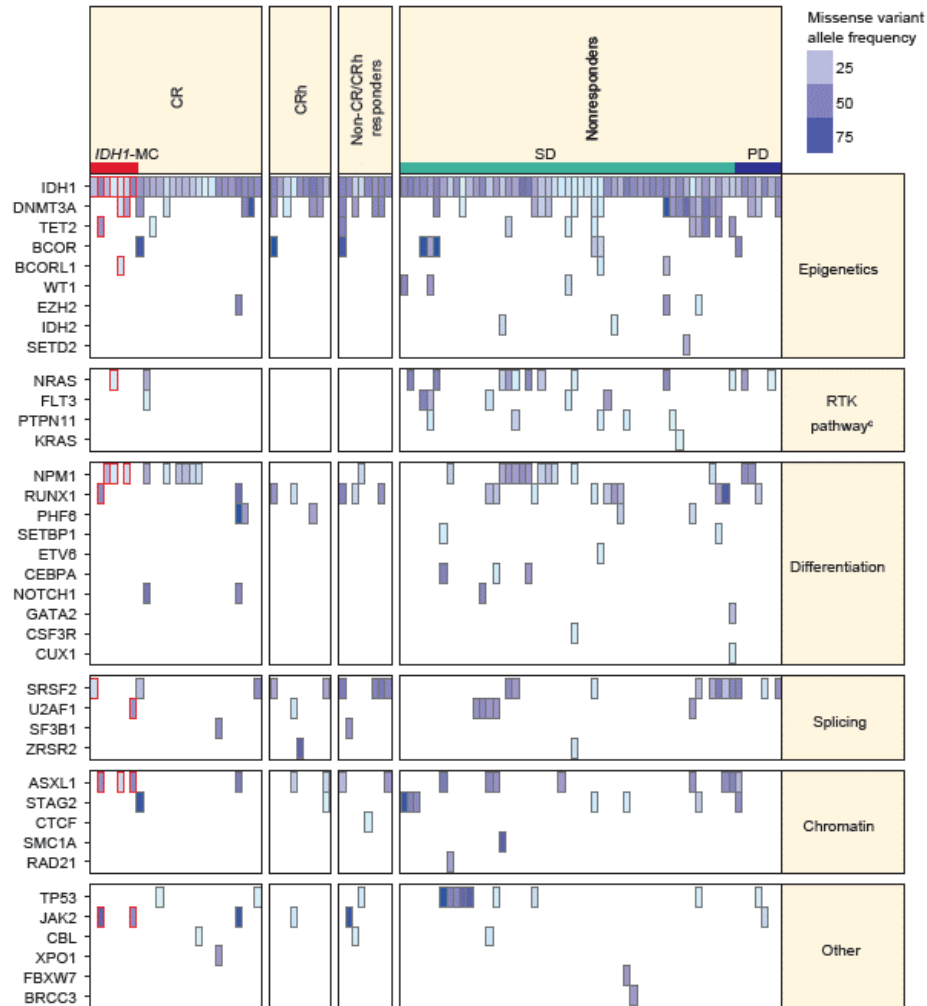
Figure 5. Most frequent (n≥2) co-occurring mutations at baseline in mLDH1 R/R AML patients (bone marrow)



Top five most frequently co-occurring mutations (*DNMT3A*, *NPM1*, *SRSF2*, *ASXL1*, *RUNX1*) in AG120-C-001 are consistent with published AML literature^{5,11,12}

- No specific single gene mutation was significantly predictive of clinical response or resistance to treatment with ivosidenib in the R/R AML patients presented. However, RTK pathway mutations were associated with a lack of response

Figure 6. Co-occurring mutations at baseline (NGS, bone marrow, n=101)^{a,b}



^aIn this heatmap, each column corresponds to a single R/R AML patient, arranged by best overall response to ivosidenib. Detected known or likely oncogenic mutations are denoted by boxes and shaded by VAF

^bNo specific single gene mutation was significantly predictive of clinical response or resistance to treatment with ivosidenib in the R/R AML patients presented

^cRTK pathway genes assayed in this 95-gene NGS Rapid Heme Panel include: *FLT3* (TKD and ITD), *KRAS*, *NRAS*, *BRAF*, *KIT*, *MAP2K1*, *PTPN11*, and *RET*. In this dataset, RTK pathway mutations were detected in *NRAS*, *FLT3* (TKD only), *PTPN11*, and *KRAS*. Mutations in the RTK pathway occurred less frequently in patients who achieved CR or CRh as a best response relative to those who did not achieve CR or CRh (p=0.003 by Fisher's exact test)

Longitudinal Mutant IDH1 Analysis: Poster 2684, Stone RM et al.

Figure 3. Ivosidenib treatment reduced *mIDH1* VAF in BMMCs and neutrophils from patients with best overall response of CR or CRh (R/R AML)

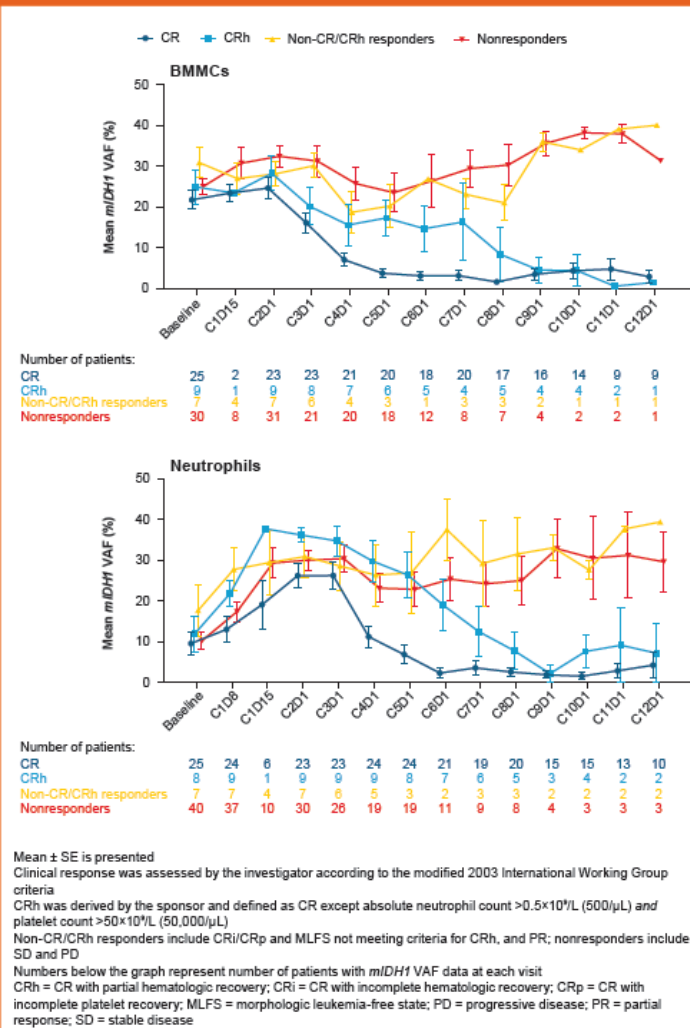
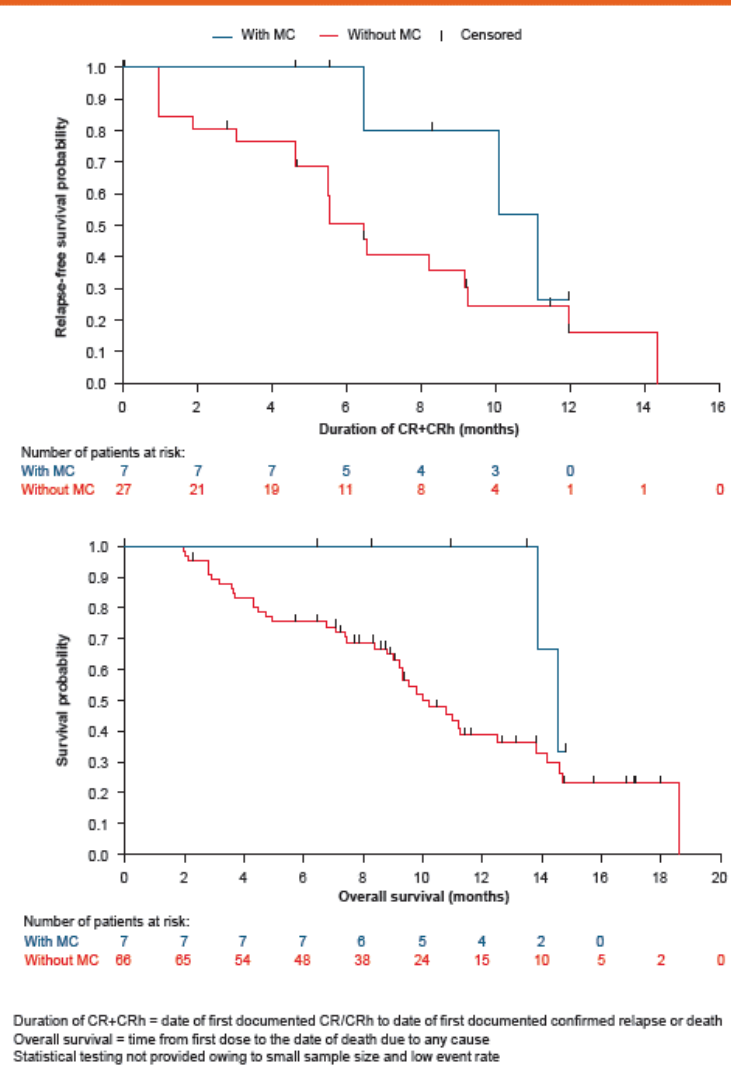


Figure 4. Patients with *IDH1*-MC had improved duration of CR+CRh and overall survival (R/R AML, BMMCs)



- Ivosidenib reduced *mIDH1* allele burden in both BMMCs and neutrophils in R/R AML patients in the expansion phase who achieved CR or CRh
- MRD-negative CR was observed in 7 of 25 (28%) R/R AML patients who achieved CR
 - Patients with MRD-negative CR had improved duration of CR compared to patients with CR with persistent MRD in this limited dataset
 - Patients with MRD-negative CR had improved overall survival compared to all other R/R AML patients with persistent MRD

Conclusions

- Ivosidenib was well tolerated; most AEs were grade 1–2 in severity
- In patients with R/R AML, most of whom had received multiple prior AML treatments, ivosidenib induced durable responses, with CR+CRh and ORR rates of 30.4% and 41.6% respectively, and corresponding durations of 8.2 and 6.5 months, with additional benefits:
 - Transfusion independence across best response categories
 - Decreased frequency of febrile neutropenia and infections in responders
- Durable responses were observed in patients with untreated AML, with CR and ORR rates of 20.6% and 55.9% and corresponding durations of NE and 9.2 months
- Ongoing AML studies:
 - Phase 1 of ivosidenib or enasidenib with AZA: presented this morning
 - AGILE: Global Phase 3, ivosidenib+AZA vs placebo+AZA in 1st-line AML
 - Phase 1 of ivosidenib or enasidenib in combination with standard AML induction and consolidation therapy, to be presented next

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