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A horizontal band featuring a microscopic image of blood cells, likely a peripheral smear, showing various cell types and their morphology in shades of pink and purple.

Revumenib Monotherapy in Patients with Relapsed/Refractory *KMT2Ar* Acute Leukemia: Topline Efficacy and Safety Results from the Pivotal AUGMENT-101 Phase 2 Study

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KMT2Ar Acute Leukemia

- Many patients relapse after chemotherapy and/or HSCT¹
- In adults, remission rates after relapse (CR, 5%) and median OS (2.4 months) after ≥ 2 salvage therapies remain low¹
- Outcomes in infants/children after relapse remain poor

No approved targeted therapies for *KMT2Ar* disease

OS in Adult Patients With R/R *KMT2Ar* AML After ≥ 3 rd-Line Therapy

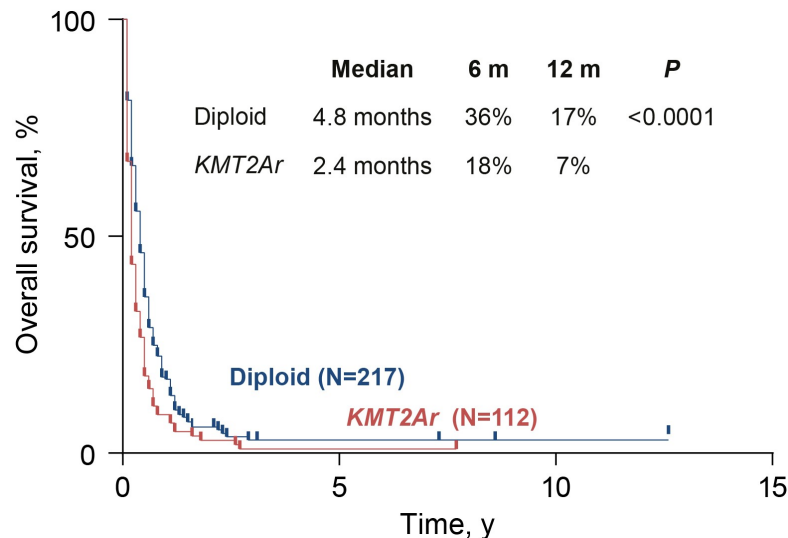
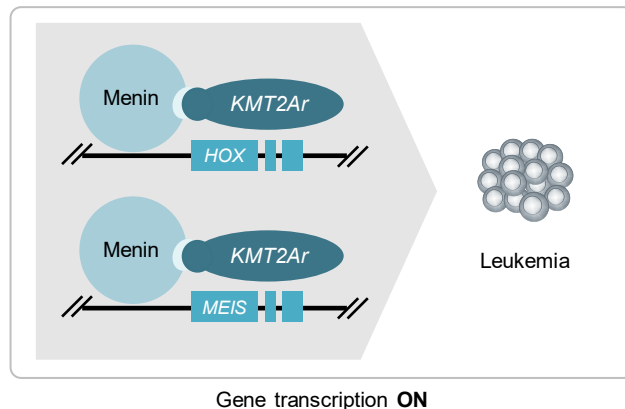


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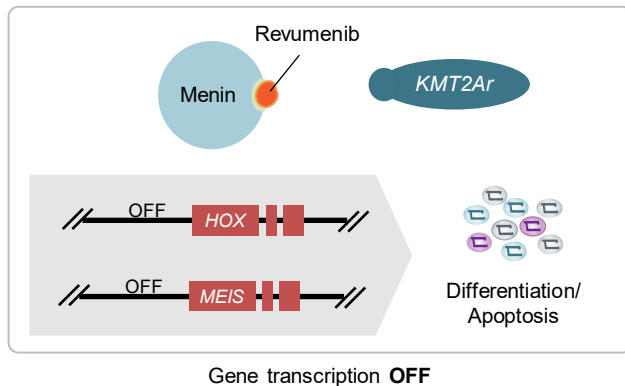
Revumenib

- The menin-KMT2A interaction is a key driver of leukemogenesis¹
- In a phase 1 study of R/R *KMT2Ar* and *NPM1m* acute leukemias, revumenib demonstrated
 - Clinically meaningful responses that were consistent across subgroups²
 - High percentage (67%) of responders proceeding to transplant²
 - Manageable safety profile²

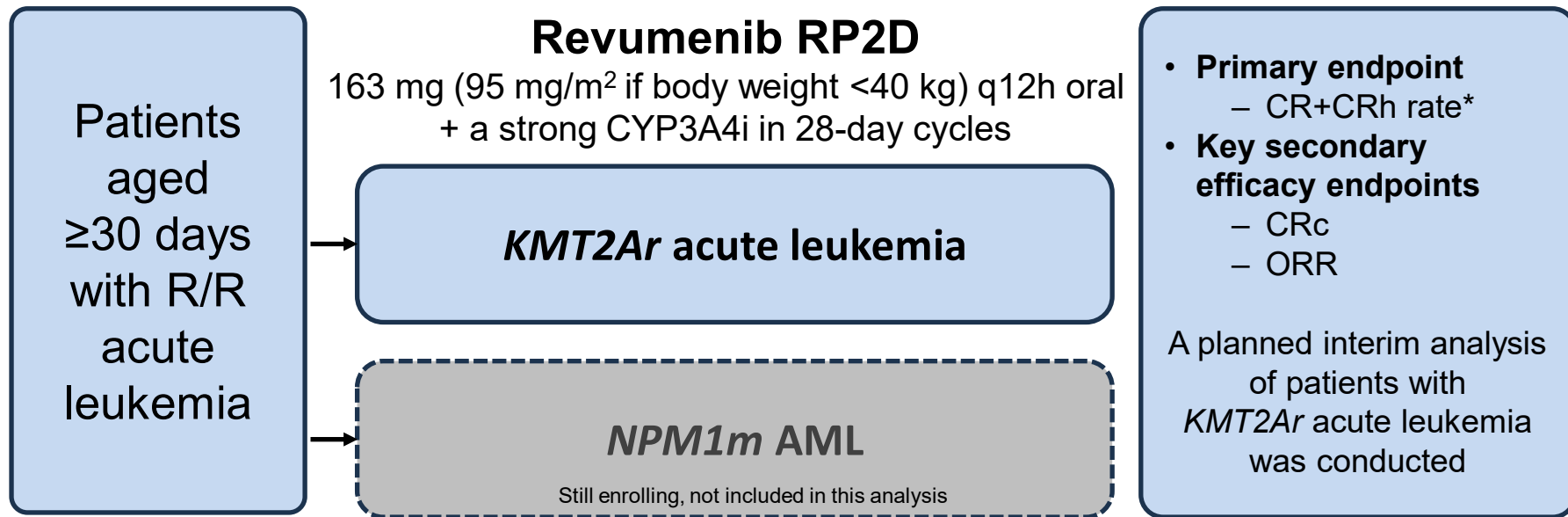
KMT2Ar acute leukemia



Menin inhibition with revumenib



AUGMENT-101 Phase 2 Study Design



*CR+CRh rate >10% in adult evaluable population considered lower efficacy bound

Patient Demographics

Parameter	Efficacy population (n=57)	Safety population (n=94) ^a
Median age, y (range)	34.0 (1.3–75)	37.0 (1.3–75)
Age <18 y, n (%)	13 (23)	23 (25)
Age ≥18 y, n (%)	44 (77)	71 (76)
Sex, n (%)		
Female	33 (58)	56 (60)
Race, n (%)		
White	43 (75)	68 (72)
Non-White	10 (18)	14 (15)
Unknown	4 (7)	12 (13)

Data cutoff: July 24, 2023. ^aDefined as patients with *KMT2Ar* acute leukemia having received at least 1 dose of revumenib.

Baseline Characteristics

Parameter	Efficacy population (n=57)	Safety population (n=94) ^a
Leukemia type, n (%)		
AML	49 (86)	78 (83)
ALL	7 (12)	14 (15)
MPAL/Other	1 (2)	2 (2)
Co-mutations ^b , n (%)		
<i>FLT3</i>	5 (9)	7 (7)
<i>RAS</i>	9 (16)	12 (13)
<i>p53</i>	4 (7)	5 (5)
Primary refractory, n (%)	14 (25)	18 (19)
Number of prior lines of therapy, median (range)	2 (1–11)	2 (1–11)
1, n (%)	17 (30)	25 (27)
2, n (%)	14 (25)	28 (30)
≥3, n (%)	26 (46)	41 (44)
Prior venetoclax, n (%)	41 (72)	61 (65)
Prior HSCT, n (%)	26 (46)	47 (50)

Data cutoff: July 24, 2023. ^aDefined as patients with *KMT2Ar* acute leukemia having received at least 1 dose of revumenib. ^bIn patients that had co-mutation status reported.



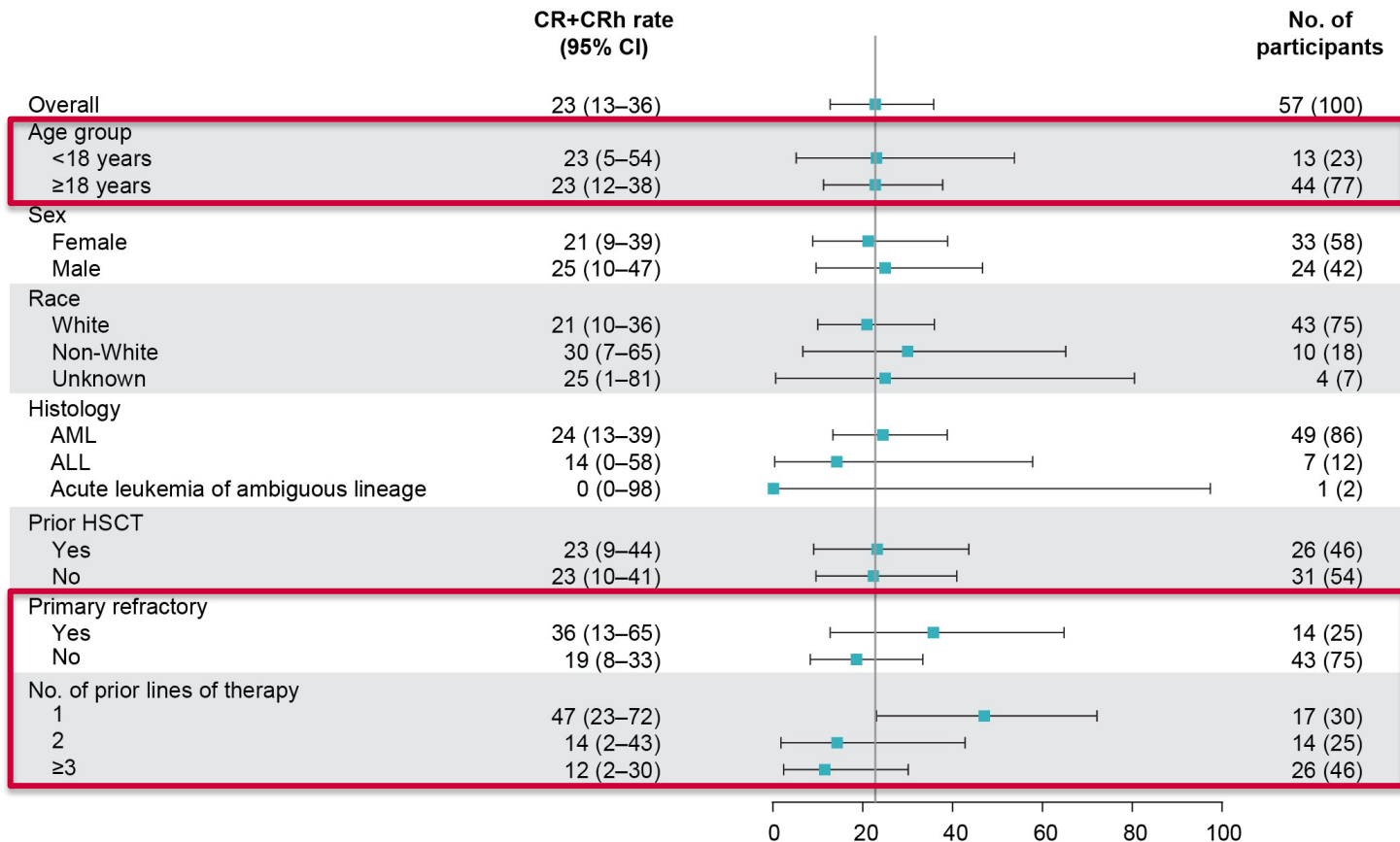
Response

Parameter	Efficacy population (n=57)
ORR, n (%)	36 (63)
CR+CRh rate, n (%)	13 (23)
95% CI	12.7–35.8
P value, 1-sided	0.0036
CRc	25 (44)
95% CI	30.7–57.6
Negative MRD status ^a	
CR+CRh	7/10 (70)
CRc	15/22 (68)

Parameter	Efficacy population (n=57)
Best response, n (%)	
CR	10 (18)
CRh	3 (5)
CRi	1 (1.8)
CRp	11 (19)
MLFS	10 (18)
PR	1 (1.8)
PD	4 (7)
No response	14 (25)
Other ^b	3 (5)

Data cutoff: July 24, 2023. ^aMRD done locally; not all patients had MRD status reported. ^bIncludes patients without postbaseline disease assessment.

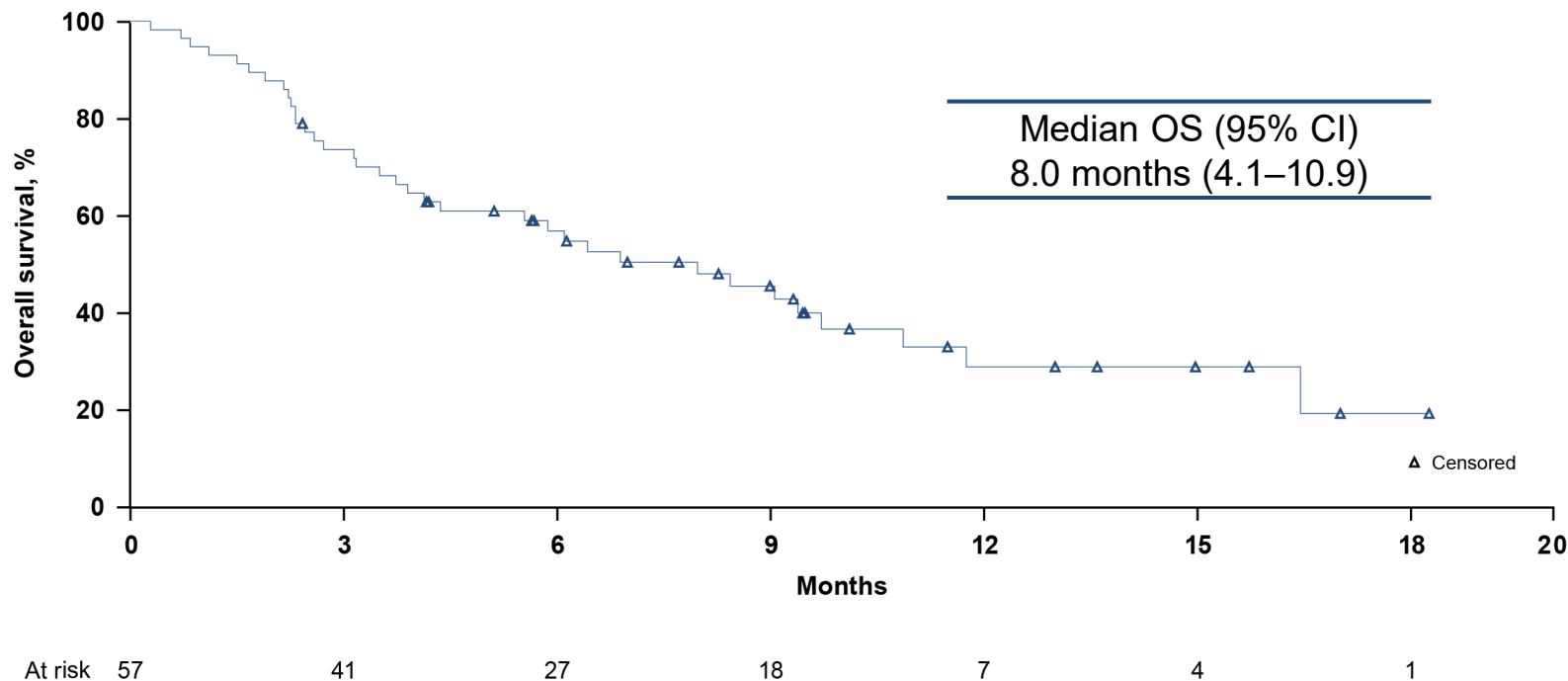
Subgroup Analyses



Responses Observed Across *KMT2A* Rearrangements

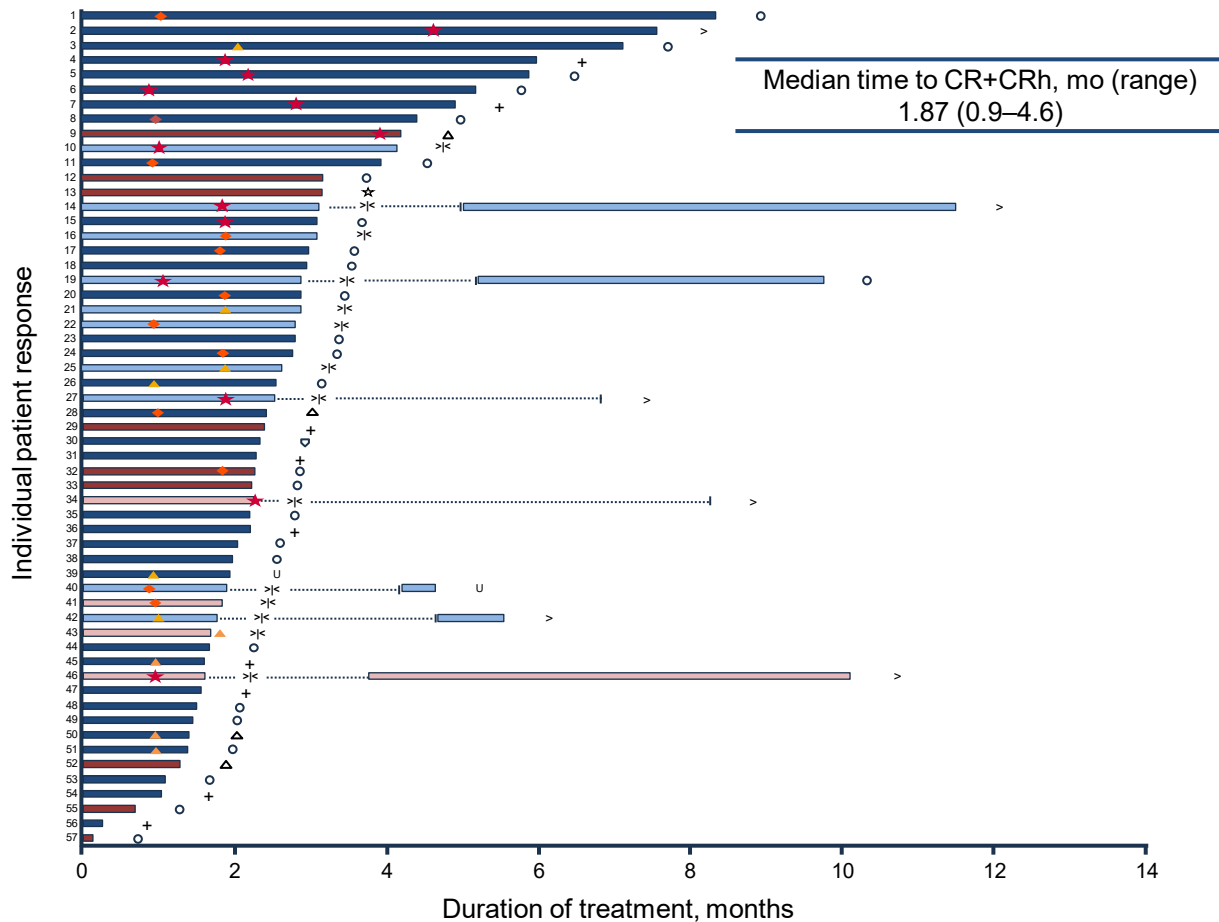
<i>KMT2A</i> rearrangement/ translocation	Summary of ORR		Summary of CR+CRh rate	
	n/N	ORR (95% CI)	n/N	CR+CRh rate (95% CI)
9;11	10/11	91 (58.7–99.8)	2/11	18 (2.3–51.8)
11;19	7/13	54 (25.1–80.8)	2/13	15 (1.9–45.4)
10;11	5/7	71 (29.0–96.3)	2/7	29 (3.7–71.0)
6;11	5/7	71 (29.0–96.3)	2/7	29 (3.7–71.0)
4;11	2/2	100 (15.8–100.0)	0/2	0 (0.0–84.2)
1;11	0/2	0 (0.0–84.2)	0/2	0 (0.0–84.2)
11;16	1/1	100	0/1	0
11;22	1/1	100	1/1	100
Unknown <i>KMT2A</i> fusion partner	5/13	39 (13.9–68.4)	4/13	31 (9.1–61.4)

Overall Survival



Duration of Treatment

- Adult
- Adult, underwent HSCT
- Pediatric
- Pediatric, underwent HSCT
- ★ CR/CRh
- ◆ CRp/CRI
- ▲ MLFS
- Progressive disease
- > Ongoing at data cutoff
- + Adverse event
- △ Subject withdrew consent for treatment
- >|< HSCT
- ☆ Noncompliance
- Patient did not achieve at least a PR after 4 cycles
- U Prohibited concomitant medication



Duration of Response

Parameter	Patients achieving CR+CRh (n=13)
Median duration of CR+CRh, months (95% CI)	6.4 (3.4–NR)
Proceeded to HSCT, n (%)	14/36 (39)
Proceeded to HSCT in CR or CRh	6/14 (43)
Proceeded to HSCT in MLFS or CRp	8/14 (57)
Restarted revumenib post HSCT, n (%)	7/14 (50)*

Data cutoff: July 24, 2023

*3 additional patients remained eligible to initiate revumenib after HSCT at the time of data cutoff.

Revumenib Safety Profile

	Safety population (n=94) ^a
All terms	TEAEs
Any grade, n (%)	93 (99)
≥Grade 3, n (%)	86 (92)
Serious AE, n (%)	72 (77)
AEs leading to:	
Dose reduction	9 (10)
Discontinuation	12 (13)
Death	14 (15)

Data cutoff: July 24, 2023. ^aDefined as patients with *KMT2Ar* acute leukemia having received at least 1 dose of revumenib.

Revumenib Safety Profile (cont)

Any grade TEAEs that occurred in ≥25% patients

All terms, n (%)	Safety population (n=94) ^a
Nausea	42 (45)
Febrile neutropenia	36 (38)
Diarrhea	33 (35)
Vomiting	29 (31)
Differentiation syndrome	26 (28)
Hypokalemia	26 (28)
Epistaxis	25 (27)
QTc prolongation	24 (26)

Grade ≥3 TEAEs that occurred in ≥10% patients

All terms, n (%)	Safety population (n=94) ^a
Febrile neutropenia	35 (37)
Decreased neutrophil count	15 (16)
Decreased white blood cell count	15 (16)
Decreased platelet count	14 (15)
Anemia	17 (18)
Differentiation syndrome	15 (16)
QTc prolongation	13 (14)
Sepsis	11 (12)
Hypokalemia	10 (11)

Data cutoff: July 24, 2023. ^aDefined as patients with *KMT2Ar* acute leukemia having received at least 1 dose of revumenib.

No patients discontinued due to differentiation syndrome, QTc prolongation, or cytopenias

Conclusions

- Revumenib is effective and safe in pediatric and adult patients with R/R *KMT2Ar* acute leukemia
- Durable MRD-negative remissions were observed in responders
- High rates of transplants among responders
- Discontinuations and dose reductions due to adverse events were low
- Study was stopped early after meeting the primary efficacy endpoint at the predefined interim analysis
 - A New Drug Application for *KMT2Ar* leukemia has been initiated under the FDA Real-Time Oncology Review program based on these data

The independent NPM1m cohort continues to enroll at all sites



Acknowledgements

- All study patients, their families, and caregivers for participating in this study
- Study teams at the individual sites
- Syndax Pharmaceuticals, Inc., for funding the study