

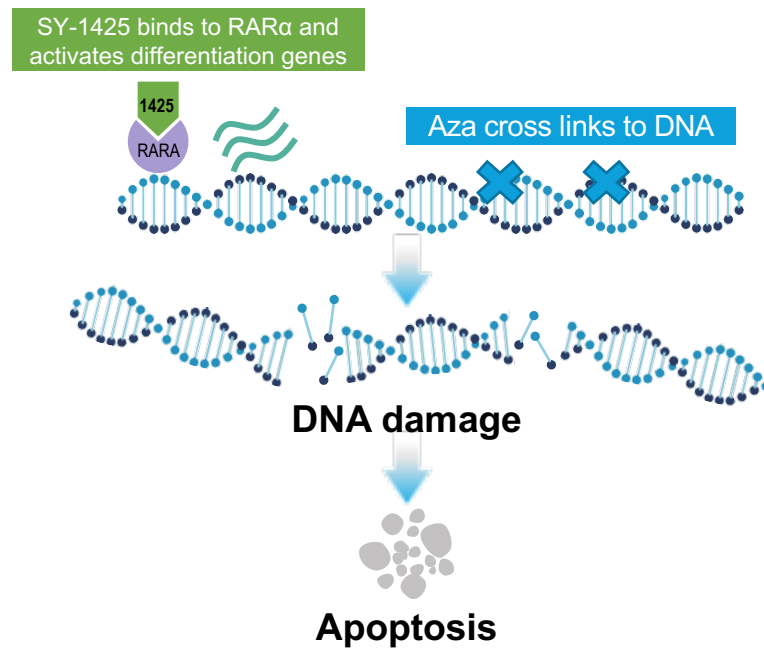
SY-1425, a Potent and Selective RAR α Agonist, in Combination with Azacitidine Demonstrates a High Complete Response Rate and a Rapid Onset of Response in RARA-positive Newly Diagnosed Unfit Acute Myeloid Leukemia

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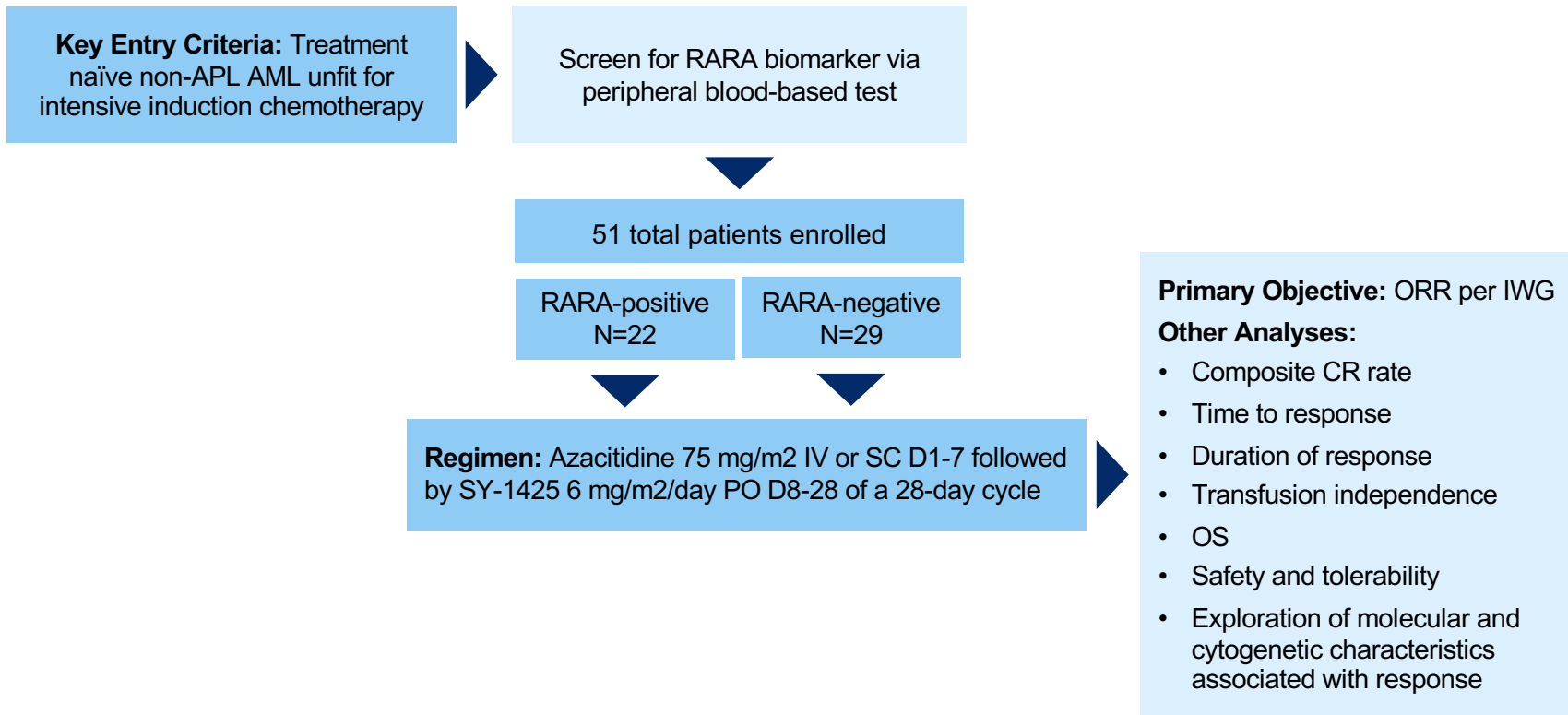
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RARA-positive AML is a Novel Patient Subset with an Actionable Target for Treatment with SY-1425, an Oral, Selective RAR α Agonist

- Subset of non-APL AML patients characterized by overexpression of the *RARA* gene
 - Novel blood-based biomarker test identifies patients for treatment with SY-1425, with typical 2 to 3-day turnaround time^{1,2}
 - Approximately 30% of AML patients are RARA-positive
- Preclinical synergy of SY-1425 with azacitidine (Aza) supported development of the combination in RARA-positive myeloid malignancies³
- Early data of SY-1425/Aza demonstrated high CR rate and rapid onset of responses in RARA-positive newly diagnosed (ND) unfit AML^{4,5}
- Unmet need for new well-tolerated therapies remains, for example, one-third of ND unfit AML patients do not respond to upfront treatment with venetoclax/Aza, and a majority of responders eventually relapse⁶



Study SY-1425-201: A Phase 2, Multi-center, Open-label Trial



Baseline Demographics and Patient Characteristics

Characteristic	RARA-positive (N=22)	RARA-negative (N=29)
Median age, years (range)	77 (60-91)	76 (64-86)
Male, n (%)	13 (59)	19 (66)
Diagnosis, n (%)		
De novo AML	16 (73)	13 (45)
Secondary AML	6 (27)	16 (55)
Evolved from antecedent hematologic malignancy	6 (27)	13 (45)
Associated with treatment from prior malignancy	0 (0)	3 (10)
AML cytogenetic risk, n (%)		
Intermediate	16 (73)	18 (62)
Poor	6 (27)	10 (34)
Not done	0 (0)	1 (3)
Baseline bone marrow blasts, n (%)		
≤ 30%	7 (32)	11 (38)
>30%	15 (68)	18 (62)

- Elderly patient population
- Large proportion with high blast counts and poor risk features

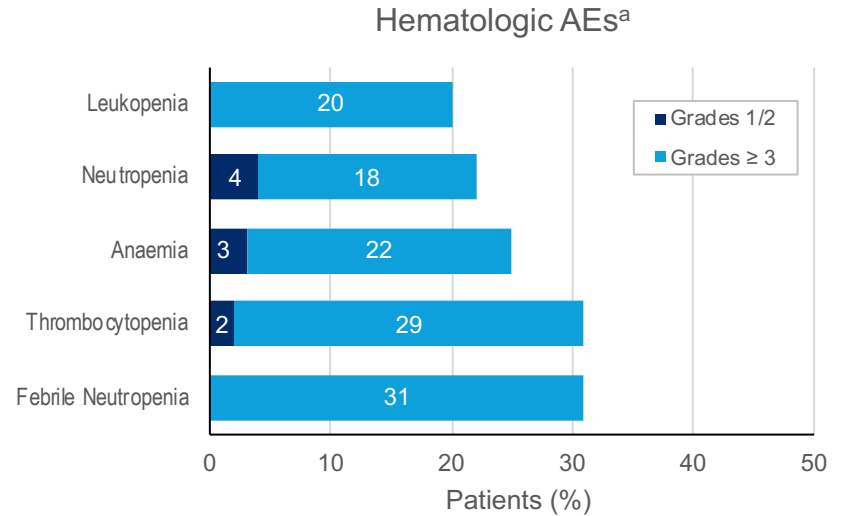
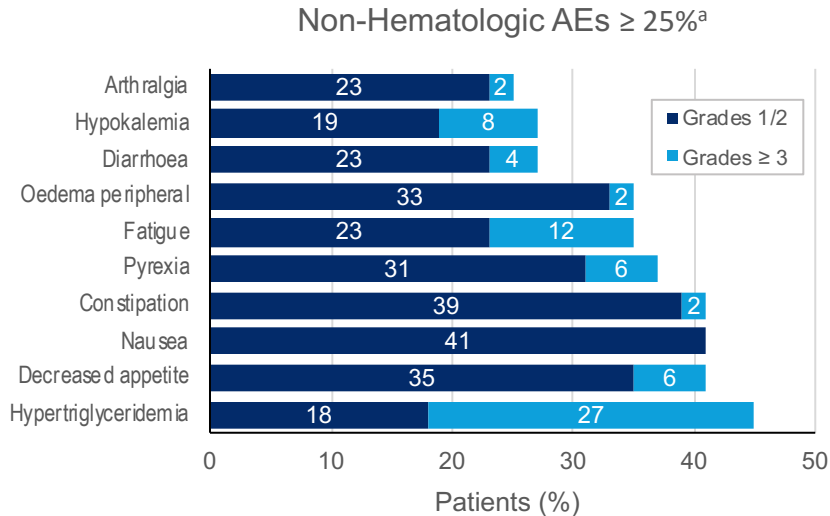
Patient Disposition

Characteristic	Enrolled Population N=51
Discontinued treatment, n (%)	46 (90)
AE ^a	16 (31)
PD	14 (27)
Treatment failure	3 (6)
Withdrawal of consent	3 (6)
Lack of clinical benefit	1 (2)
Death	1 (2)
Other	8 (16)

^aIncludes 2 patients who discontinued treatment prior to first dose of SY-1425. Of the 16 patients who discontinued due to AE, 2 were due to AEs assessed as related to study drug.

Safety Summary

- Combination generally well tolerated with no increased toxicity relative to either single agent SY-1425 or Aza in AML
- Myelosuppression comparable to reports of single agent Aza in this population
- Majority of non-hematologic AEs are low grade and reversible
- SAEs were reported for 42 patients; the most frequent (occurring in ≥ 5 pts) included febrile neutropenia (14 pts), pyrexia (6 pts), pneumonia (6 pts) and sepsis (5 pts)



^aIncludes all enrolled patients, N=51.

RARA-positive Patients Have a High Complete Remission Rate with a Rapid Time to Response

Best IWG Response ¹	RARA-positive n (%)	RARA-negative n (%)
Response Evaluable, N ^a	18	28
ORR	12 (67)	12 (43)
CR/CRi	11 (61)	9 (32)
CR	9 (50)	7 (25)
CRm	4 (22)	3 (11)
CRc	4 (22)	3 (11)
CRi	2 (11)	2 (7)
MLFS	1 (6)	1 (4)
PR	0 (0)	2 (7)

^a Only response evaluable patients are included, defined as all patients who completed one cycle of treatment with at least one post-baseline response evaluation or discontinued earlier due to disease progression, and who have not had any major protocol violations. There were 4 non-response evaluable RARA-positive patients (2 discontinued prior to first dose of SY-1425 and 2 discontinued prior to completion of cycle 1 due to AE not related to study drug) and 1 non-response evaluable RARA-negative patient (discontinued due to clinical progression without post-baseline response evaluation).

- RARA-positive patients:
 - High CR/CRi response rate
 - Deep CR with 8/9 (89%) CRm or CRc
 - Rapid time of onset of initial complete response^b with median 1.2 months
 - Median duration of complete response^b 10.8 months (95% CI: 2.9, 15.2)
- RARA-negative patients:
 - Response rates comparable to historical response rates for single agent Aza²⁻⁴
 - Median time to initial complete response^b delayed relative to RARA-positive patients at 3.0 months
 - Median duration of complete response^b 10.3 months (95% CI: 3.1, NE)

^bComplete response includes CR, CRi, CRh

Responses Observed in RARA-positive Patients Irrespective of Mutation or Cytogenetic Risk

	Patients with IWG Response												Patients without IWG Response					
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18
IWG response	Achieved IWG response																	
TP53				Presence of the indicated molecular mutation														
ASXL1					Presence of the indicated molecular mutation		Presence of the indicated molecular mutation											
RUNX1															Presence of the indicated molecular mutation			
NPM1	Presence of the indicated molecular mutation		Presence of the indicated molecular mutation		Presence of the indicated molecular mutation				Presence of the indicated molecular mutation	Presence of the indicated molecular mutation								Presence of the indicated molecular mutation
FLT3															Presence of the indicated molecular mutation			Presence of the indicated molecular mutation
CEBPA																		Presence of the indicated molecular mutation
IDH1																		Presence of the indicated molecular mutation
IDH2																		Presence of the indicated molecular mutation
DNMT3A															Presence of the indicated molecular mutation			
TET2				Presence of the indicated molecular mutation	Presence of the indicated molecular mutation													
BCORL1	Presence of the indicated molecular mutation																	
BCOR					Presence of the indicated molecular mutation													
EZH2					Presence of the indicated molecular mutation										Presence of the indicated molecular mutation			
KRAS															Presence of the indicated molecular mutation			
CBL				Presence of the indicated molecular mutation	Presence of the indicated molecular mutation													
PHF6					Presence of the indicated molecular mutation													
Cytogenetic Risk	Intermediate	Intermediate	Intermediate	Intermediate	Intermediate	Intermediate	Intermediate	Poor	Intermediate	Poor	Poor	Intermediate	Intermediate	Intermediate	Intermediate	Intermediate	Intermediate	Intermediate

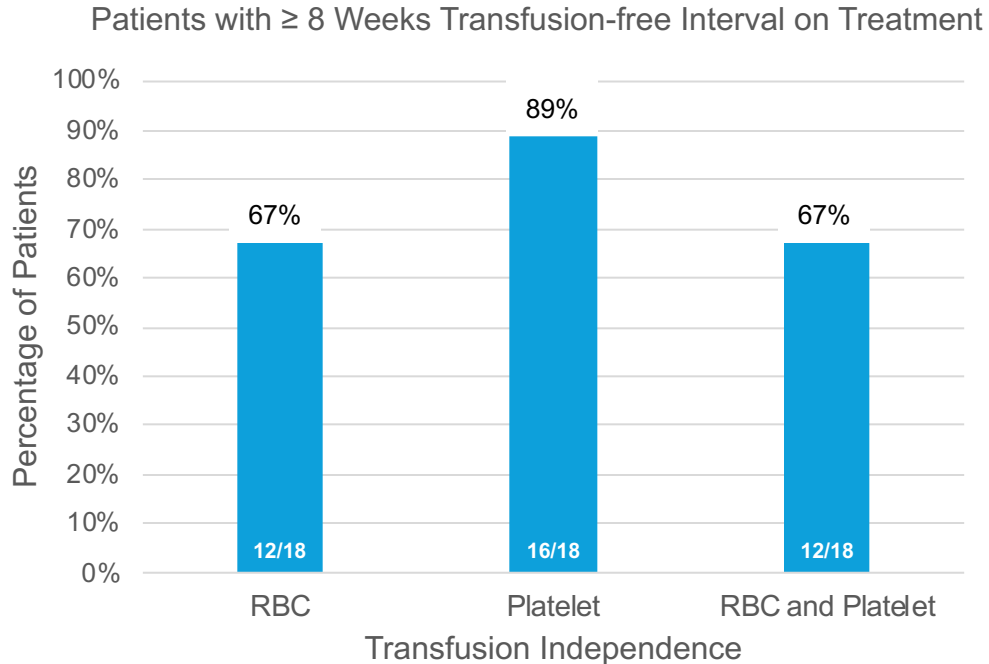
- Achieved IWG response
- Presence of the indicated molecular mutation

Cytogenetic Risk^a

- Intermediate
- Poor

*Data shown for the 18 response evaluable patients
^aCytogenetic risk per NCCN AML guidelines 2018

Transfusion Independence

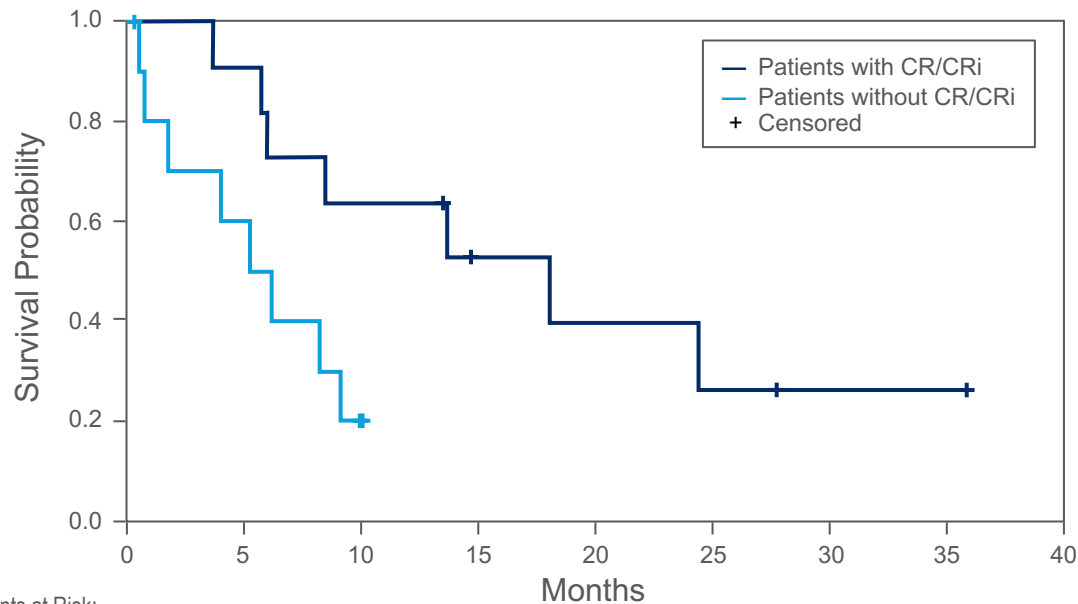


- High proportion of RARA-positive patients achieved or maintained transfusion independence:
 - 67% (12/18) of patients were free of both RBC and platelet transfusions for a ≥ 8 -week interval on treatment
 - 86% (6/7) of patients dependent on transfusions at baseline converted to transfusion independence during treatment

Patients on treatment ≥ 56 days evaluable for transfusion independence.

Transfusion independence defined as not requiring RBC or platelet transfusions during any 56-day post baseline period.

Overall Survival in RARA-positive Patients Stratified by Response Status



- RARA-positive patients with CR/CRi (N=11):
 - Median OS 18.0 months (95% CI: 5.7, NE)
- RARA-positive patients without CR/CRi (N=11)^a
 - Median OS 5.6 months (95% CI: 0.4, 9.0)
- Total enrolled RARA-positive patients (N=22):
 - Median OS 8.4 months (95% CI: 5.2, 18.0)

^a RARA-positive patients without CR/CRi included 4 non-response evaluable patients (2 discontinued prior to first dose of SY-1425 and 2 discontinued prior to completion of cycle 1 due to AE not related to study drug).

Patients at Risk:

Patients with CR/CRi 11

10

7

4

3

2

1

0

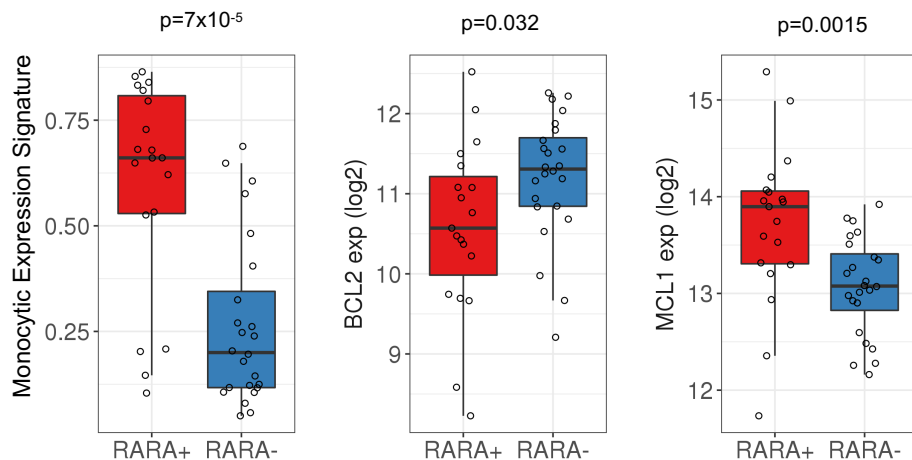
Patients without CR/CRi 11

6

0

RARA-positive ND Unfit AML Patients Including Those with Response to SY-1425 Plus Aza are Enriched for Features Associated with Venetoclax Resistance

Analyses of Patient Samples from Clinical Trial



- Multiple recent studies report venetoclax resistance is associated with a monocytic phenotype¹⁻³
- A monocytic expression signature was developed using 9 well-established monocytic and primitive gene expression markers⁴
- ~80% of RARA-positive ND unfit AML trial patients have monocytic phenotype associated with venetoclax resistance, which includes lower BCL2 and higher MCL1 expression⁴
- Majority of RARA-positive ND unfit AML patients who achieved CR/CRi with SY-1425/Aza have this monocytic phenotype⁴

Selection of RARA-positive Newly Diagnosed Unfit AML Patients with Elevated *RARA* Gene Expression Enriches for Features Associated with Primary Resistance to Venetoclax and Clinical Response to SY-1425, a Potent and Selective RAR α Agonist, plus Azacitidine (abstract # 137323) to be presented in Session 616 AML: Novel Therapy, excluding Transplantation: Poster III on Mon, Dec 7

Conclusions

- SY-1425/Aza demonstrates high CR rates including the majority with molecular and cytogenetic CRs in RARA-positive AML, a novel subset of AML characterized by *RARA* overexpression
 - Rapid onset of response
 - Responses observed across cytogenetic risk groups and mutations
 - Majority achieved or maintained transfusion independence
 - Median OS for responders was 18.0 months, suggesting clinically meaningful benefit
- SY-1425/Aza was generally well-tolerated with no evidence of increased toxicity relative to either as a single agent
 - Rates of myelosuppression were comparable to single-agent Aza
- ~80% of RARA-positive ND unfit AML trial patients have monocytic phenotype associated with venetoclax resistance, which includes lower BCL2 and higher MCL1 expression¹
 - Majority of RARA-positive ND unfit AML patients who achieved CR/CRi with SY-1425/Aza have this monocytic phenotype, suggesting the potential for combination treatment with SY-1425 to address significant unmet need in ND unfit AML, including in those who may be resistant to venetoclax¹
- Further development is warranted in RARA-positive AML and other myeloid malignancies