

Beyond Hypomethylating Agents:



Combination Therapies in MDS

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Beyond HMAs | Agenda

- Predicting Responders to HMAs
- Combinations – Lower-risk
- Combinations – Higher-risk
- Conclusions



Beyond HMAs | Agenda

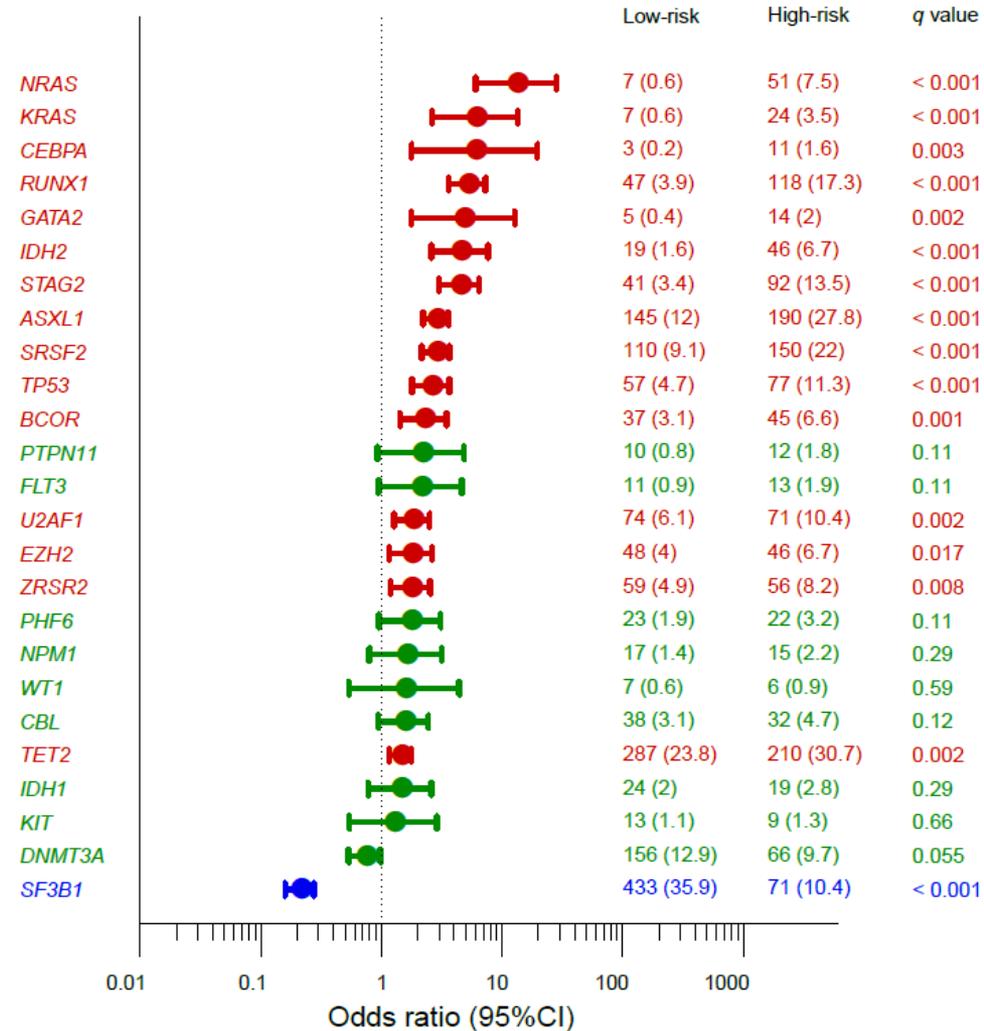
- **Predicting Responders to HMAs**
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Beyond HMAs | Mutation Risk

Driver genes can be classified into molecular subtypes differentially associated with disease severity

Low-risk MDS vs. High-risk MDS (univariate)



Makishima et al. Nat Genetics 2017; 49:204.

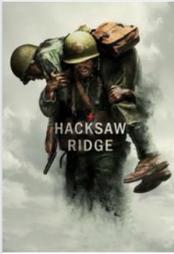
Beyond HMAs | Mutations/Response

- Somatic mutations may predict response or resistance to HMA:
 - **TET2** mutations may predict response
 - **ASXL1** may predict resistance
 - **TP53** mutations may predict response
- Challenges with these data:
 - ✓ Response might be higher but the mutation is not a biomarker
 - ✓ Genomic data are complex

Beyond HMAs | Mutations/Response



The Green Mile (1999)

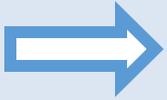


Hacksaw Ridge (2016)



Django Unchained (2012)

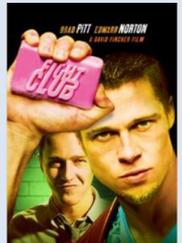
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The Lord of the Rings: The Return of the King (2003)



The Shawshank Redemption (1994)



Fight Club (1999)

?



Beyond HMAs | Mutations/Response



The Green Mile (1999)



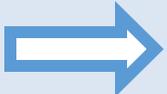
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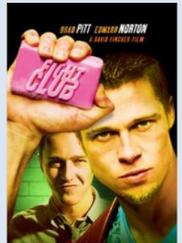
The Godfather



The Lord of the Rings: The Return of the King (2003)



The Shawshank Redemption (1994)

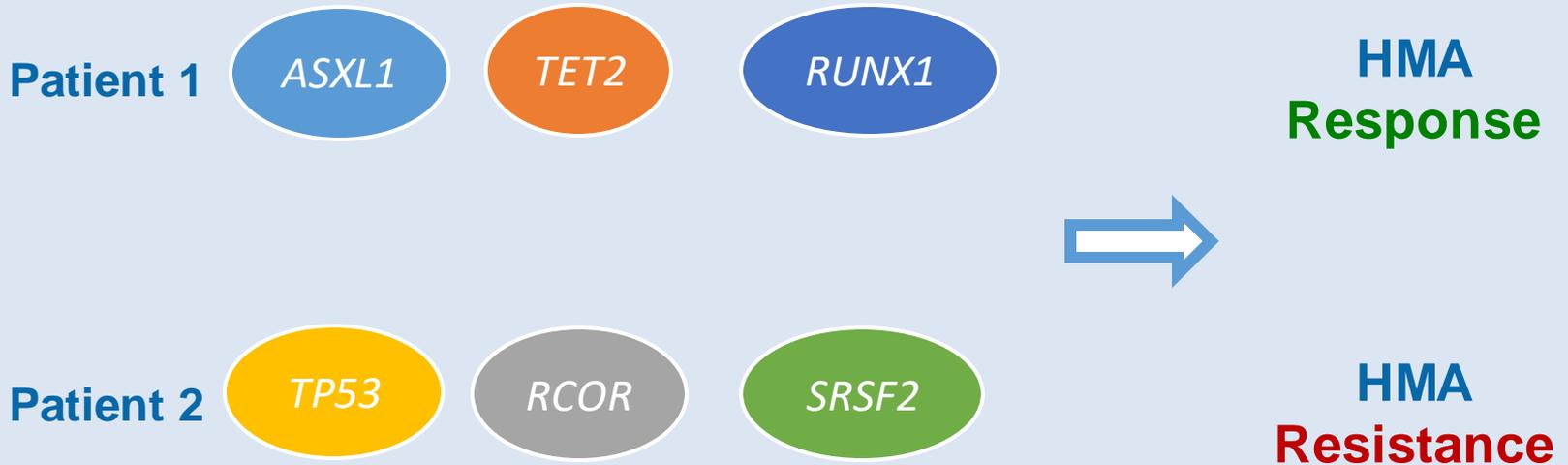


Fight Club (1999)



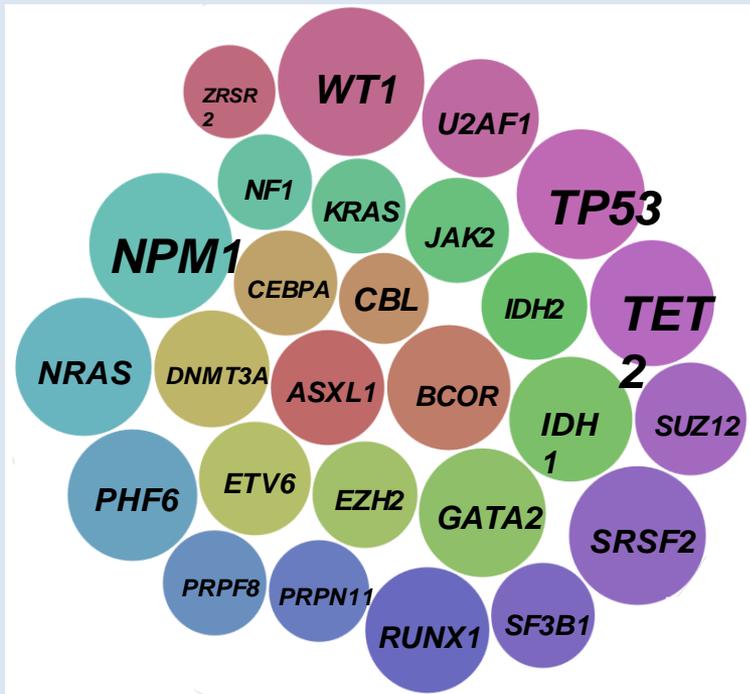
NETFLIX

Beyond HMAs | Mutations/Response

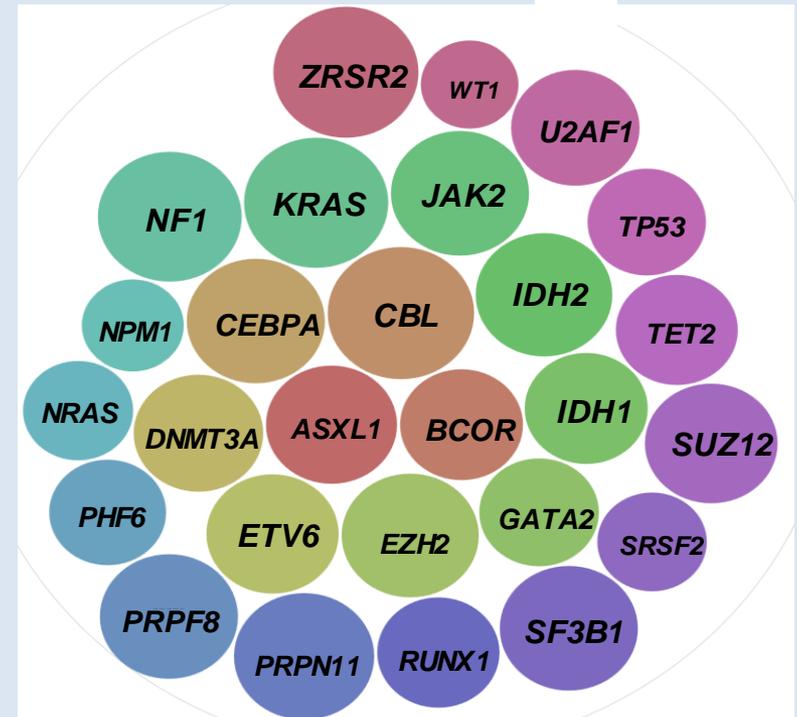


Beyond HMAs | Mutations/Response

Responders



Non-Responders



N = 433 Patients treated with HMAs → Validated in 113 Patients enrolled in S1117

Beyond HMAs | Mutations/Response

Results: Association Rules

Training

Association Rules (Resistance)
ASXL1, NF1
ASXL1, EZH2, TET2
ASXL1, EZH2, RUNX1
EZH2, SRSF2, TET2
ASXL1, EZH2, SRSF2
ASXL1, RUNX1, SRSF2
ASXL1, TET2, SRSF2
ASXL1, BCOR, RUNX1

Association Rules (Response)
TET2, RUNX1, SRSF2

31% pts
≥ 3 mutations/sample

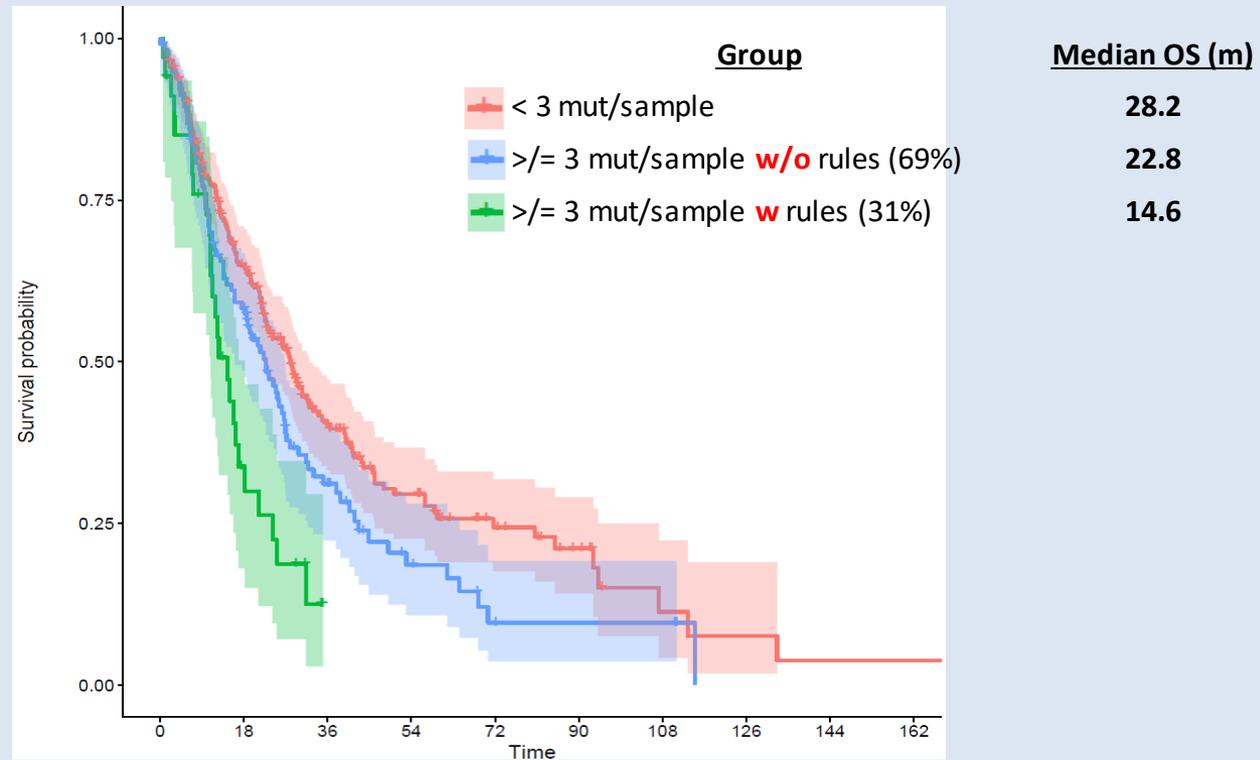
29% pts
Very Low/Low risk by IPSS-R

ORR to HMAs = **43%**

Median # mutations per
patient = **3 (range, 0-9)**

Accuracy: **87%**

Beyond HMAs | Mutations/Response



Beyond HMAs | Agenda

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Lower-risk MDS | Ameliorating Anemia: LEN

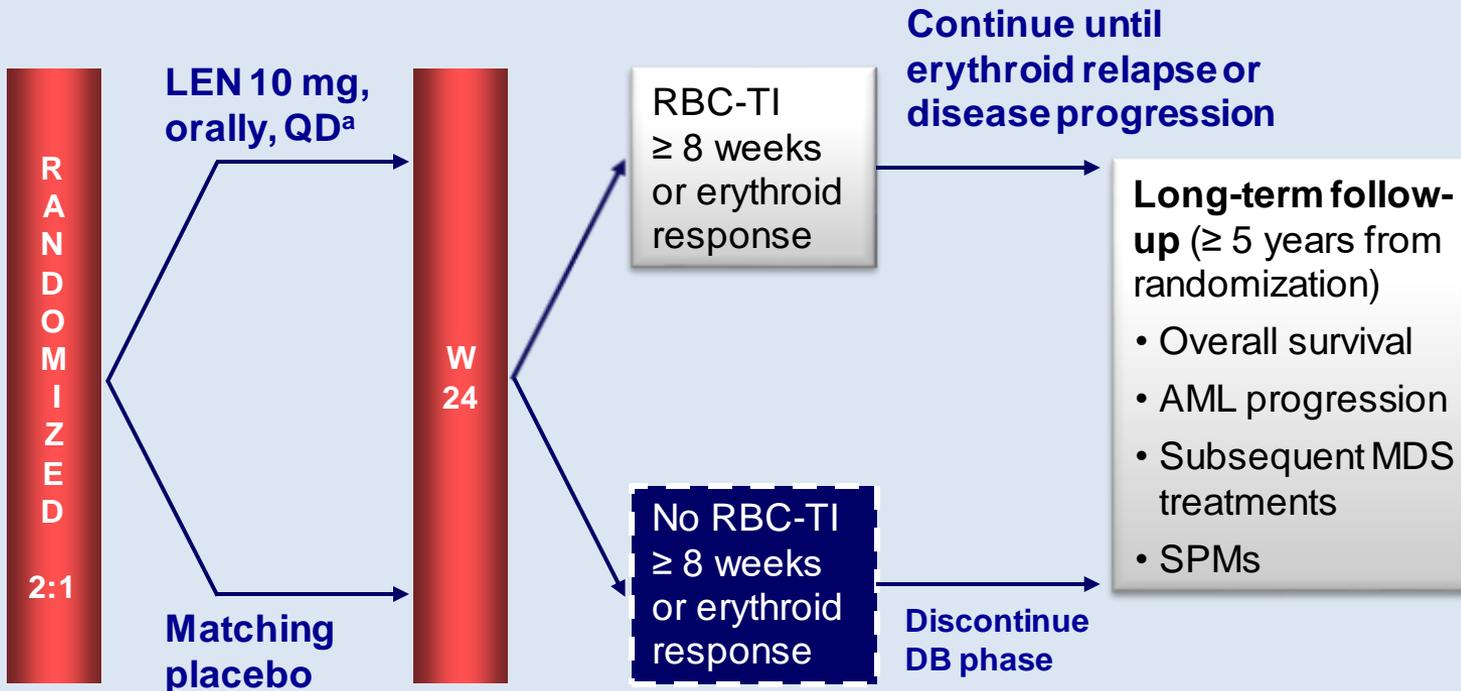
Pretreatment

Double-blind (DB) treatment

Off-treatment

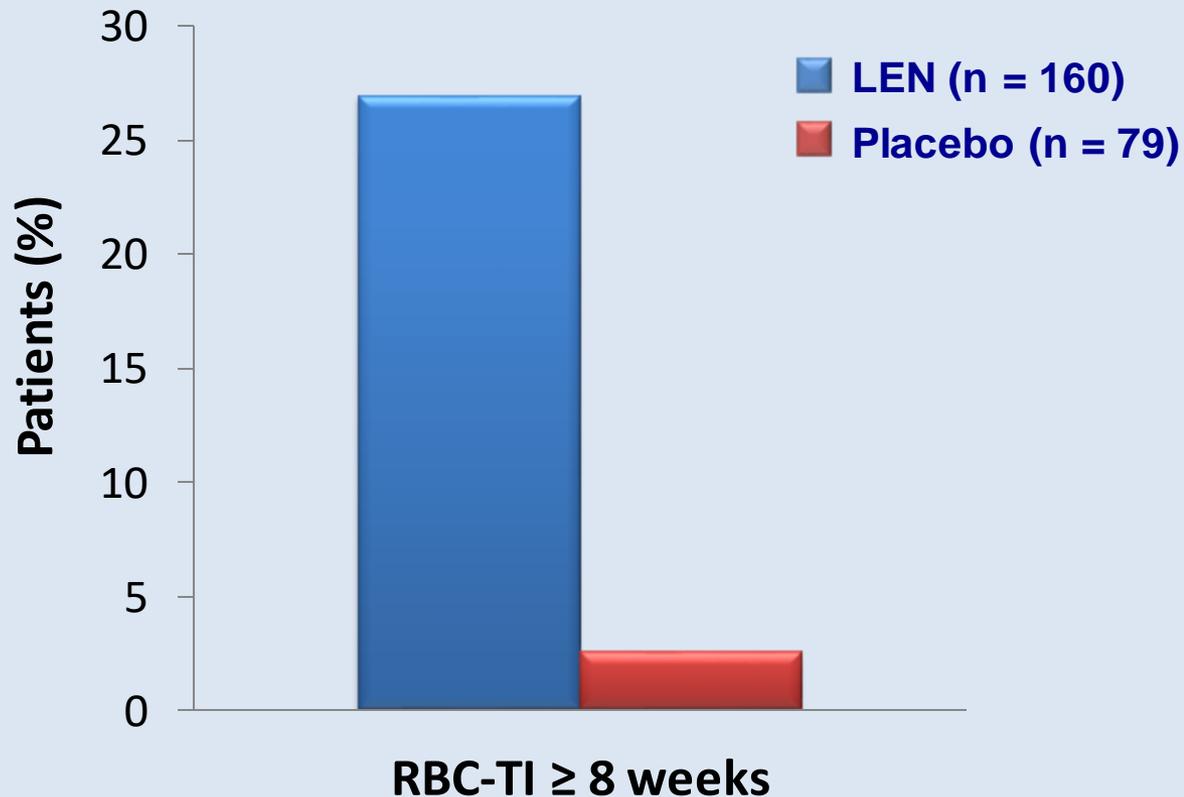
Key inclusion criteria

- Centrally reviewed IPSS Low or Int-1-risk MDS with karyotypes other than del(5q)
- RBC-TD
- Unresponsive or refractory to ESAs



Lower-risk MDS | Ameliorating Anemia: **LEN**

Significantly more LEN patients achieved RBC-TI ≥ 8 weeks versus placebo ($P < 0.001$)

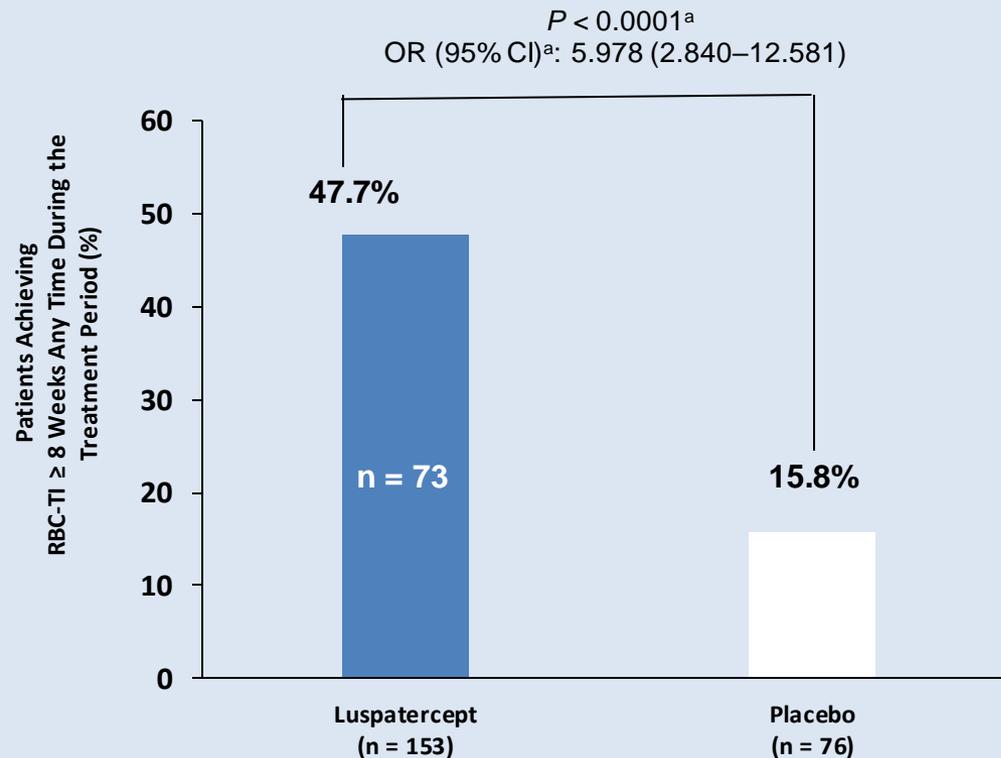


- Randomized Phase III low risk non-del(5q) ESA resistant or high endogenous EPO level (>500).
- Len 10 mg/day * 21 versus Len plus EPO 60K/week
- 205 patients randomized- 14 not treated due to EPO shortage
- Median transfusion burden 4 U/ 8 weeks
- 93% prior ESA; 18% prior DNMTi

LEN +/- EPO

	Lenalidomide	Lenalidomide plus EPO
n	96	99
Major Erythroid Response	11 (11.5%)	28 (28.3%)
After 16 weeks of therapy		
n	64	73
Major Erythroid Response	10 (16%)	28 (39%)
CROSSOVER To COMBO		
Major Erythroid Response		11/44
Duration of MER		
Median (months)	13	24

Fenaux et al, Luspatercept: RBC-TI \geq 8 weeks Achieved any time during treatment period



Beyond HMAs | **LEN + LUS**

The L2 Regimen

Dose Level	Lenalidomide Schedule	Luspatercept Schedule
1	5 mg PO days 1-21	1.0 mg/kg day 1
2	5 mg PO days 1-21	1.33 mg/kg SC day 1
3	5 mg PO days 1-21	1.75 mg/kg SC day 1

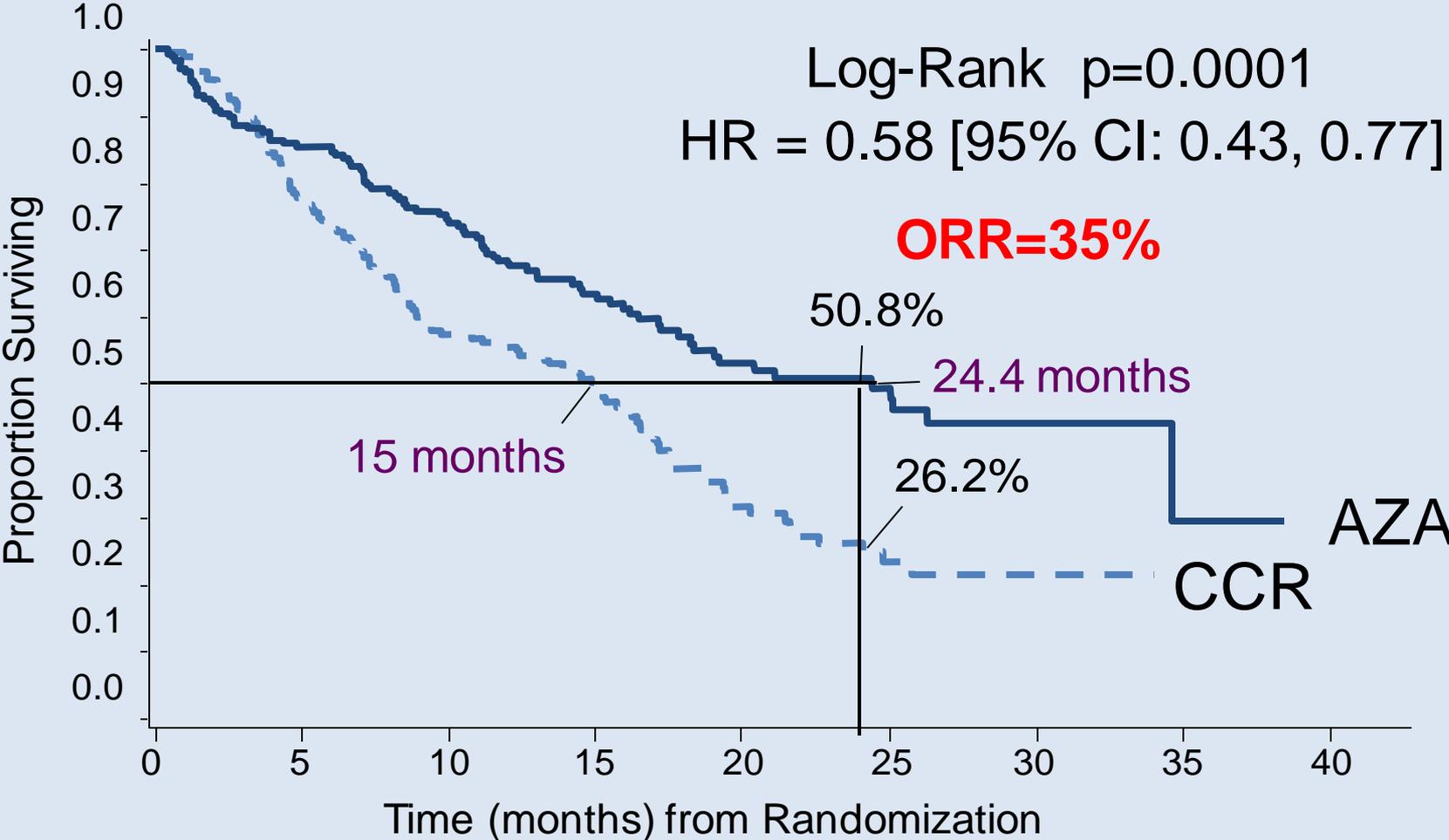
Phase IB/II
Lower-risk, Non-Del(5q)
N = 40

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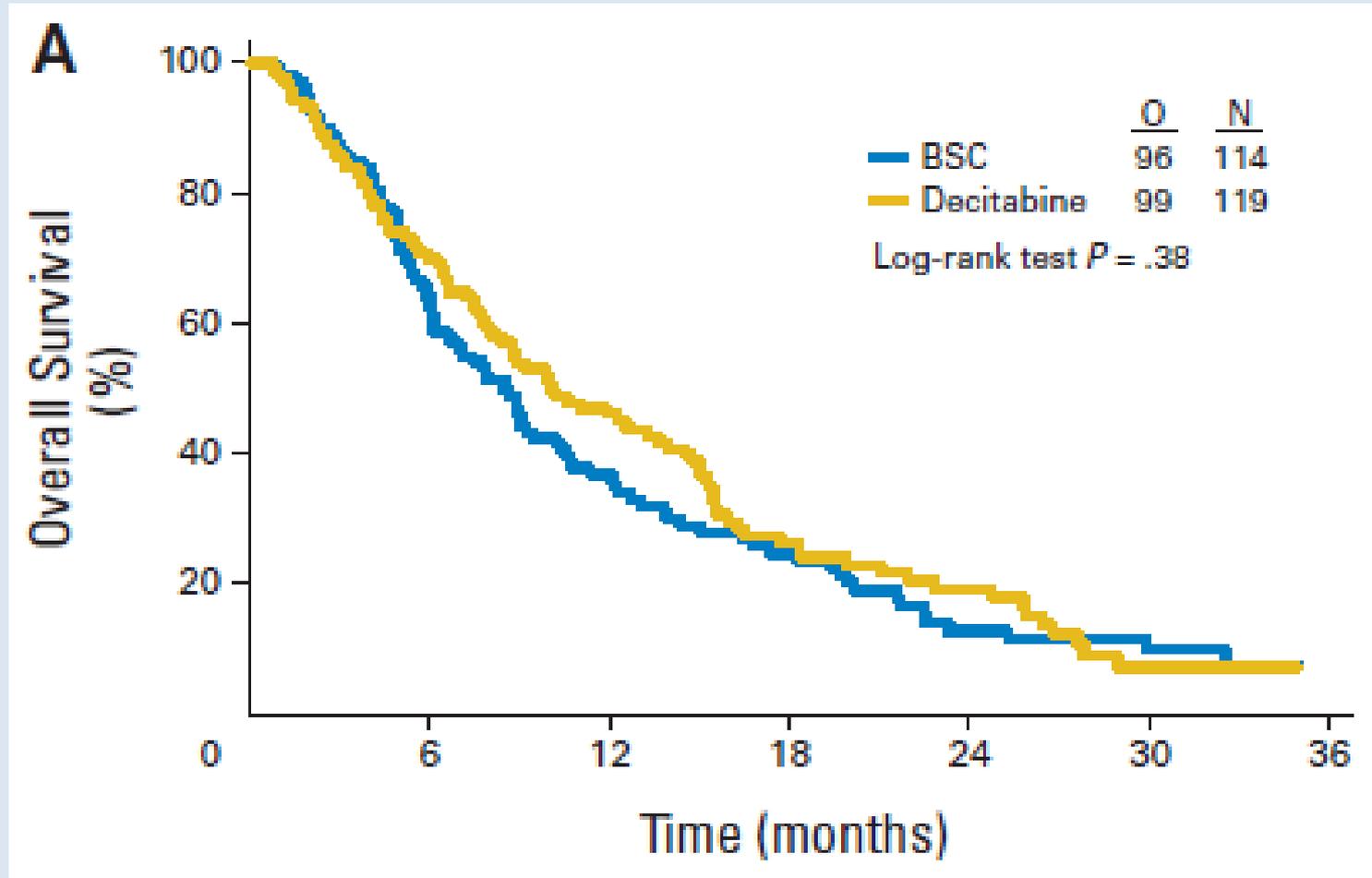


Beyond HMAs | AZA



Fenaux P, et al. Lancet Oncology 2009;10:223-232.

Beyond HMAs | DAC



Median OS 10.1 vs. 8.5 months

ECOG E1905

Eligible patients (no prior azacitidine):

- **MDS** (*higher risk; if IPSS low/INT-1 risk, then platelets $<50 \times 10^9/L$ or ANC <500)*
- **CMML** with WBC $<12 \times 10^9/L$
- **AML** with multilineage dysplasia and WBC $\leq 30 \times 10^9/L$ for ≥ 4 weeks

Azacitidine SC 50 mg/m² x 10 days every 28 days

Primary Endpoint:

IWG 2000 responses with hematological normalization
(CR+PR+trilineage HI)

**Azacitidine SC 50 mg/m² x 10 days every 28 days, plus
Entinostat (MS-275) 4 mg/m² PO days 3 and 10 each cycle**

Randomize 1:1

E1905 study results

Response / AE	Arm A (n=68) (azacitidine monotherapy)	Arm B (n=68) (combination therapy)
Complete response (CR)	12%	7%
Partial response (PR)	9%	7%
Trilineage hematological improvement (tHI)	10%	10%
Qualifying Response (CR+PR+tHI)	31%	24% (p=NS)
<i>Other hematological improvement</i>	12%	19%
<i>Any response</i>	43%	44%
Grade IV thrombocytopenia	44%	63% (p=0.07)
Grade III/IV fatigue	13%	23% (p=0.13)

Aza + Pracinostat in MDS: Study Design



- **102 evaluable patients: one-to-one randomization**
- **Azacitidine: 75 mg/m² 7 days I.V./sq every 28 days**
- **Pracinostat or placebo P.O. 60 mg 3 days/week for 3 weeks**
- **Cycles repeated every 28 days until disease progression, lack of benefit, or intolerance**

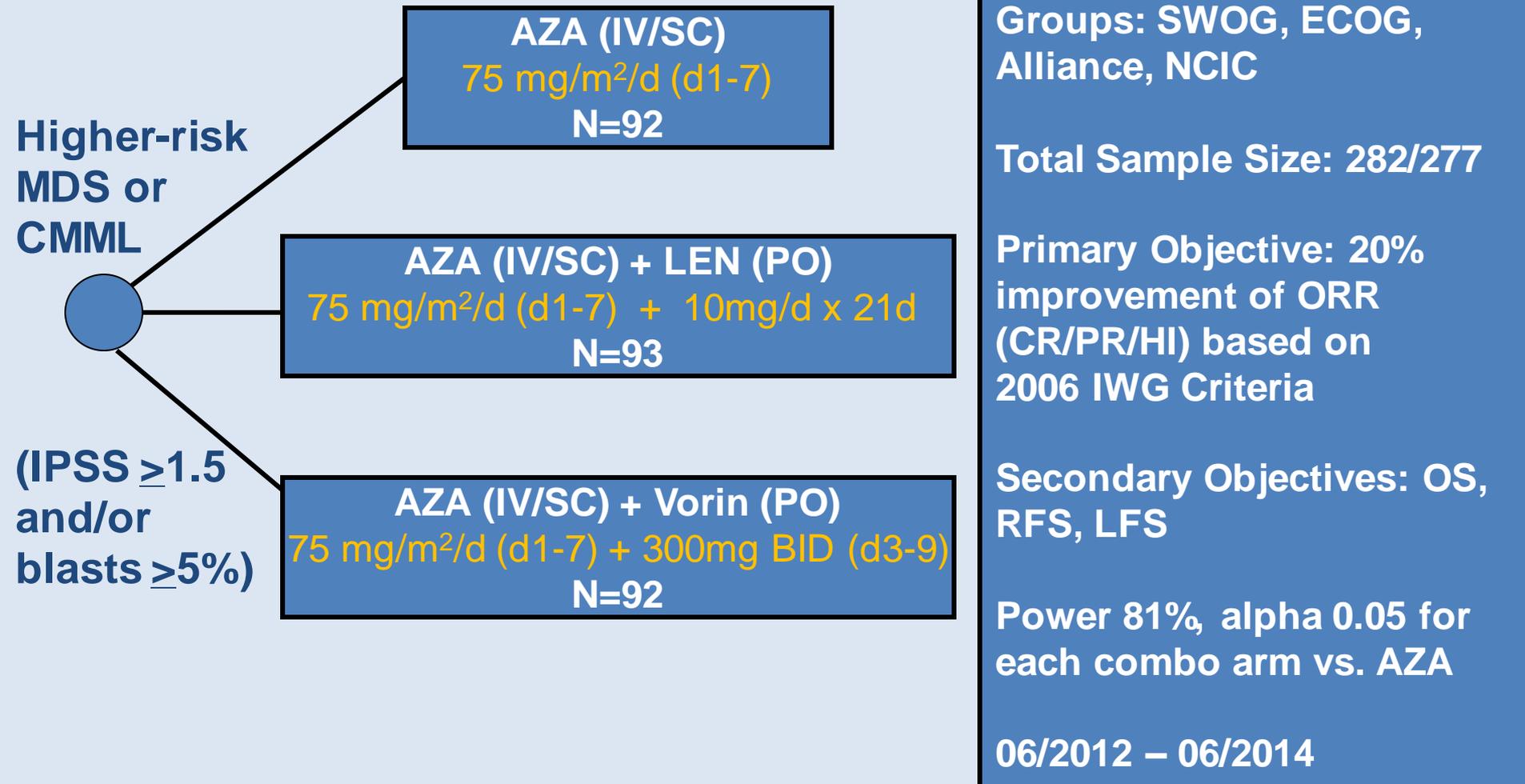
Aza + Pracinostat in MDS: Summary of Response

	Pracinostat	Placebo
CR, within 180 days	18%	33%
Best Response		
Complete Remission	20%	33%
Partial Remission	0%	0%
Marrow CR	28%	22%
Stable Disease	26%	29%
Progressive Disease	6%	6%
Not evaluable	22%	10%

Aza + Pracinostat in MDS: Summary of Response

RESPONSE	Pracinostat	Placebo
Hematological improvement	35%	55%
Erythroid response (HI – E)	28%	45%
Platelet response (HI – P)	31%	53%
Neutrophil response (HI – N)	26%	39%
Clinical benefit rate (CR + PR + HI + mCR)	53%	63%
Cytogenetic response	42%	55%
Cytogenetic CR	24%	29%
Cytogenetic PR	18%	26%

Higher-risk MDS | Combinations



Higher-risk MDS | Combinations

Variable Median and N (%)	AZA n=92 (33%)	AZA+LEN n=93 (34%)	AZA+VOR n=92 (33%)	Total n=277 (100%)
Age (yrs, range)	69 (42, 88)	70 (51, 86)	70 (28, 93)	70 (28, 93)
Female	31 (34)	32 (34)	22 (24)	81 (31)
CMML	18 (20)	19 (20)	16 (18)	53 (19)
tMDS	7 (8)	6 (6)	5 (5)	19 (7)
Baseline ANC (x10³)	2 (0, 110)	1 (0, 336)	2 (0, 36)	2 (0, 336)
Baseline Platelet count (x10³)	70 (8, 4000)	75 (3, 452)	62 (3, 1462)	68 (3, 4000)
Baseline Median Blast %	8 (0, 22)	10 (0, 20)	10 (1, 18)	9 (0, 22)

Higher-risk MDS | Combinations

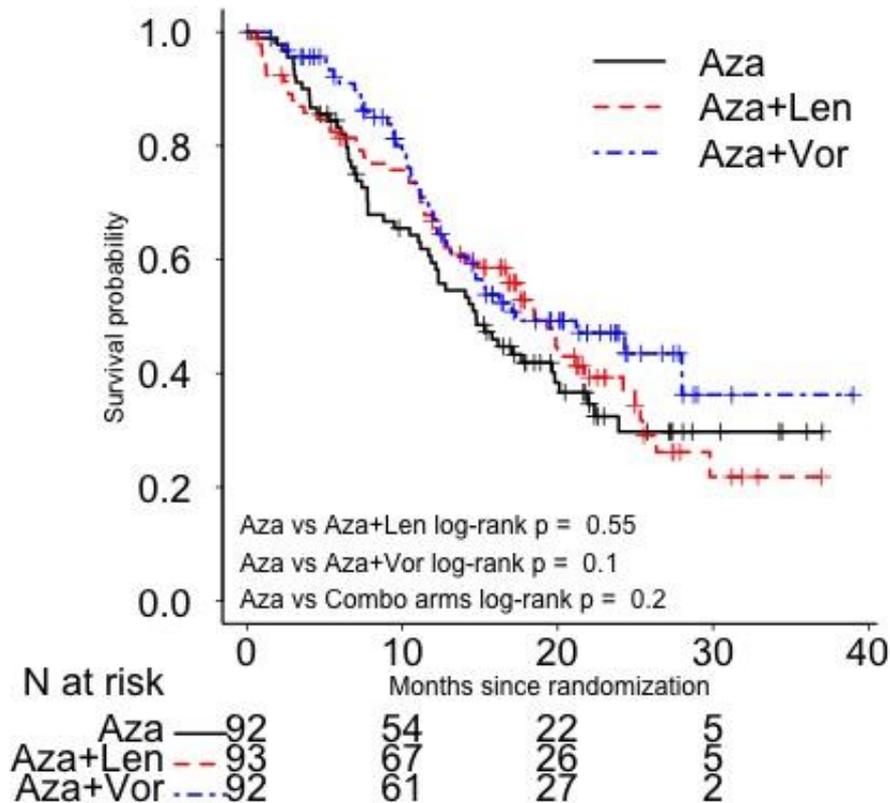
Toxicity Variable	AZA	AZA+LEN (P-value vs. AZA)	AZA+VOR (P-value vs. AZA)	Total n=271
Febrile neutropenia (n)	10	13 (.66)	12 (.51)	36
GI (n)	4	12 (.10)	14 (.02)	28
Rash (n)	3	14 (<.01)	1 (1)	17
Off Tx due to Toxicity/Side Effect/Complication	8%	20% (.05)	21% (.03)	18%
Non-protocol defined dose modifications	24%	43% (.002)	42% (.01)	33%

Higher-risk MDS | Combinations

Response Variable	AZA	AZA+LEN (P-value vs. AZA)	AZA+VOR (P-value vs. AZA)	Total n=277
Median Tx Duration (Wks)	25	24	20	22
Overall Response Rate (%)	38	49 (.16)	27 (.16)	38%
CR/PR/Hi (%)	24/0/14	24/1/ 25	17/1/9	22/1/16%
CMML ORR (%)	5 (28)	13 (68) (.02)	2 (12) (.41)	37%
ORR Duration (median)	10 months	14 months (.41)	15 months (.31)	14 months
CMML ORR Duration (median)	15 months	14 months (.87)	24 months (.69)	15 months

Higher-risk MDS | Combinations

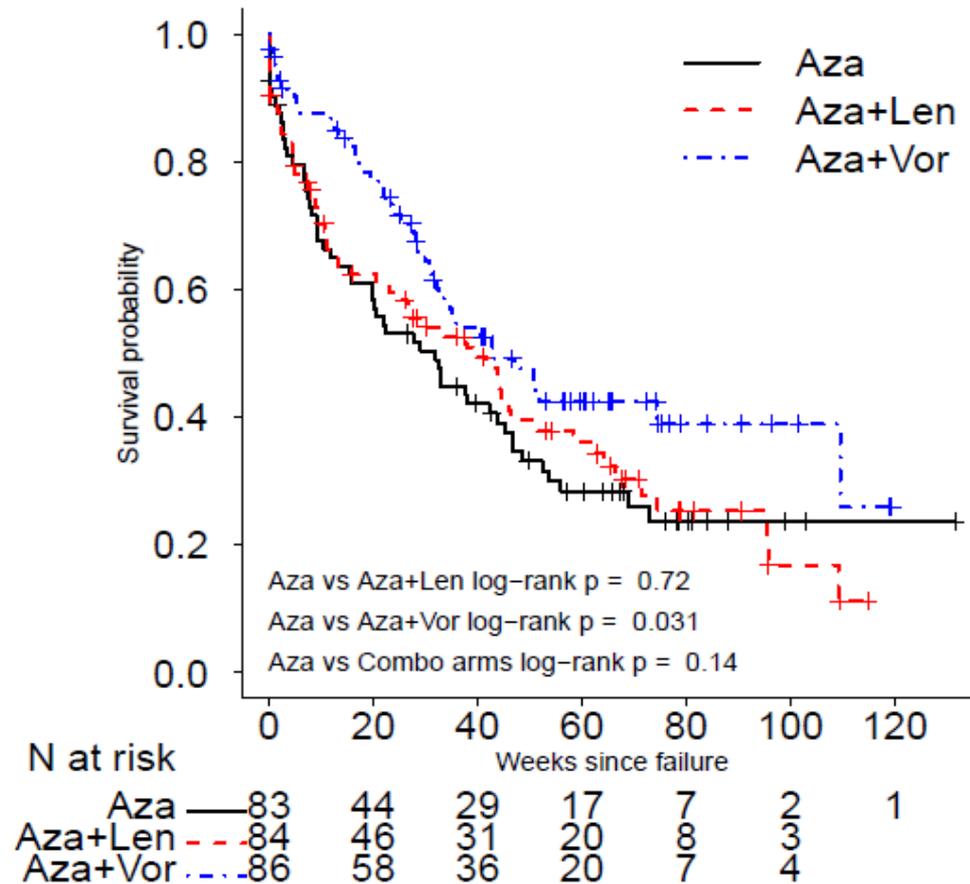
Overall Survival



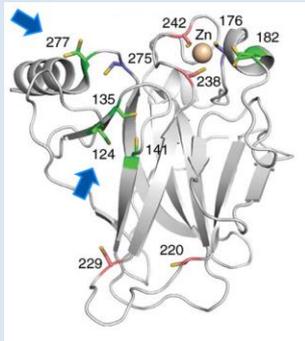
Comparisons are between combination arms and AZA monotherapy

Higher-risk MDS | Combinations

Overall Survival After Failure



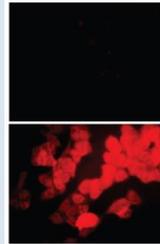
Higher-risk MDS | Targeting *TP53*



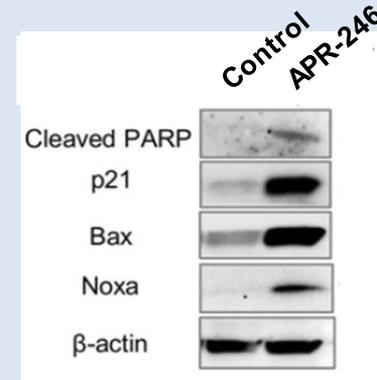
APR-246 binds covalently to p53...

p53
R175H

p53
R175H
+
APR-246



...restores wt p53 conformation & activity...

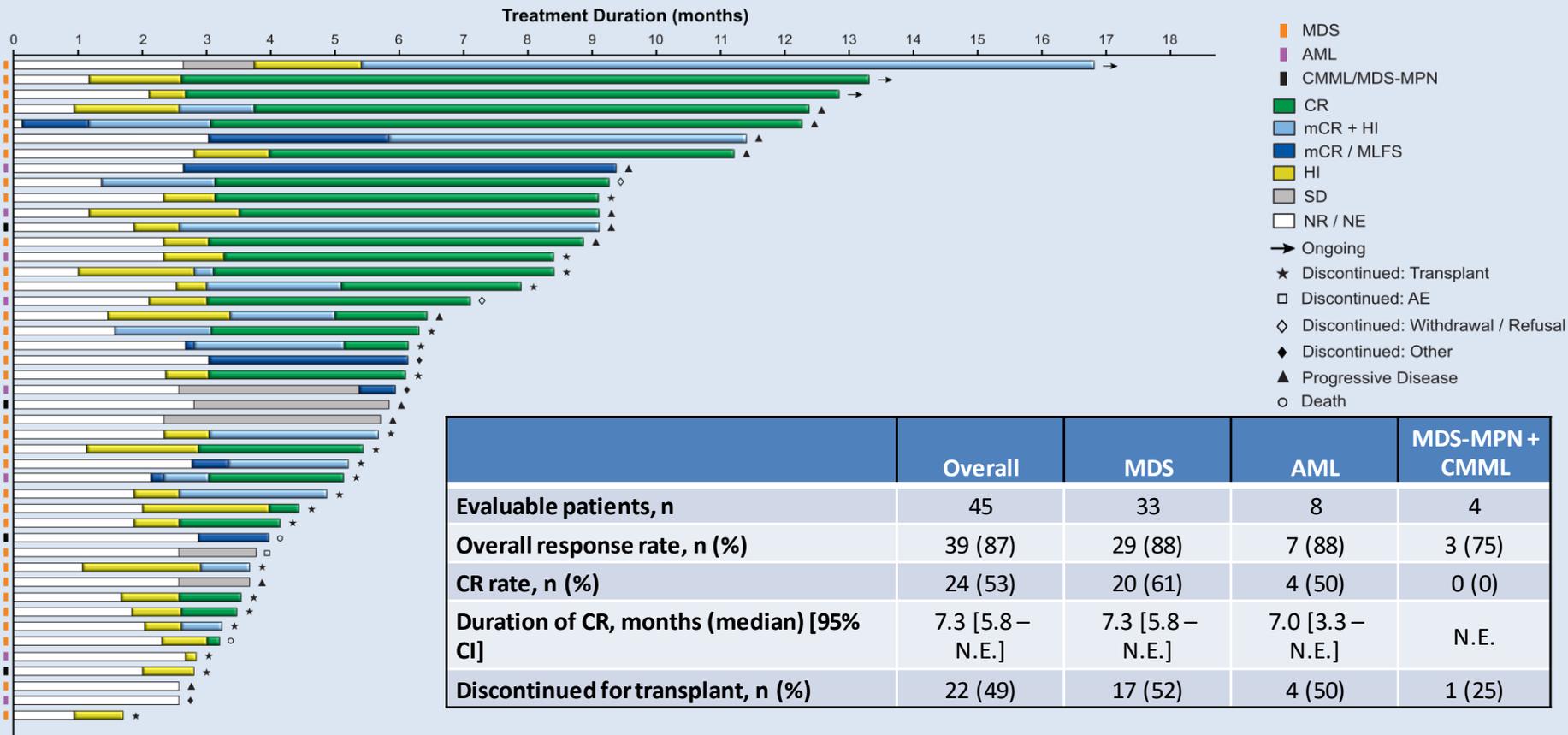


...and triggers cell cycle arrest and apoptosis

A. Fersht et al. (2010) Prot. Sci; Q. Zhang et al, (2018) Cell Death Disease; H. Furukawa et al, (2018) Cancer Sci.

Sallman et al, Cluzeau et al. ASH 2019, Abstract 676-7.

Higher-risk MDS | Targeting *TP53*



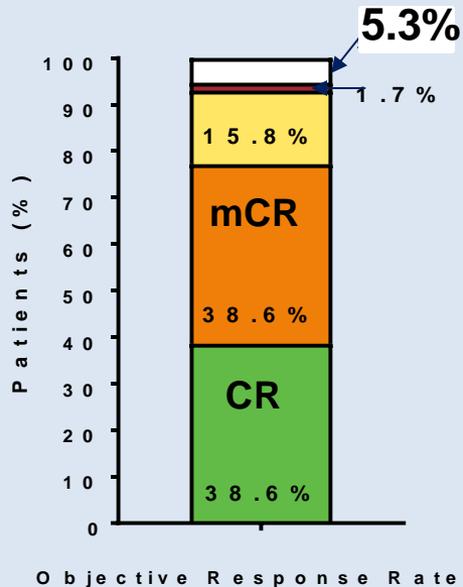
	Overall	MDS	AML	MDS-MPN + CMML
Evaluable patients, n	45	33	8	4
Overall response rate, n (%)	39 (87)	29 (88)	7 (88)	3 (75)
CR rate, n (%)	24 (53)	20 (61)	4 (50)	0 (0)
Duration of CR, months (median) [95% CI]	7.3 [5.8 – N.E.]	7.3 [5.8 – N.E.]	7.0 [3.3 – N.E.]	N.E.
Discontinued for transplant, n (%)	22 (49)	17 (52)	4 (50)	1 (25)

Median duration of follow-up = 10.8 months

Wei et al, Abstract 568 – AZA plus Venetoclax for HR-MDS

- Phase 1b study
- Untreated *de novo* MDS, IPSS Int-2 or high risk, not planning intensive chemo or transplant
- Ven days 1-14 (400 mg/day, no ramp up)
 - Prophylactic antimicrobials required
- 57 patients
 - Med Age 71 (26-85);
 - IPSS-R very high risk: 60%

Wei et al, Abstract 568 – AZA plus Venetoclax for HR-MDS: Response Rates



Median time to CR, months (range)	2.2 (1.2-11.1)
12-mo estimate of DoR after CR, % (95% CI)	83.3 (2.3, 97.5)
mCR with HI (HI-E, HI-P or HI-N), n/N (%)	10/22 (45.5)

■ Complete Remission
 ■ Marrow Complete Remission
 ■ Stable Disease
■ Progressive Disease
 Non-Evaluable

Excludes patients of arm C (Aza only); Objective response rate (ORR) includes [complete remission (CR) + marrow complete remission (mCR) + partial remission (PR)]; # of patients with PR=0; per IWG (Cheson et al., *Blood*2006;108:419-425)
 DoR: Duration of response; HI: hematological improvement; HI-E: hematologic improvement in erythroids; HI-N: hematologic improvement in neutrophils; HI-P: hematologic improvement in platelet count; n: patients with favorable outcomes; N: patients eligible for evaluating outcomes

Abstract 569 – AZA plus Magrolimab for HR-MDS

- CD47 is a macrophage immune checkpoint and “Do Not Eat Me” signal in MDS
- Magrolimab targets CD47 and synergizes with AZA in preclinical models
- Phase 1b study
- *Untreated* MDS, IPSS-R intermediate or higher risk disease
 - Magro dose ramp up to 30mg/kg weekly in C1 to mitigate on-target anemia
 - AZA given at standard 75mg/m² D1-7 doses
- 35 patients
 - Med Age 70 (47-80);
 - IPSS-R high or very high risk disease: 65%
- Safety profile consistent with AZA monotherapy; on-target anemia seen but mitigated with ramp up (median Hgb drop 0.4 g/dL with first dose)

AZA plus Magrolimab for HR-MDS – Preliminary Efficacy

Best Overall Response	1L MDS N=24
CR	12 (50%)
CRi	-
PR	0
MLFS/ marrow CR	8 (33%) 4 with marrow CR + HI
Hematologic improvement (HI)	2 (8%)
SD	2 (8%)
PD	0

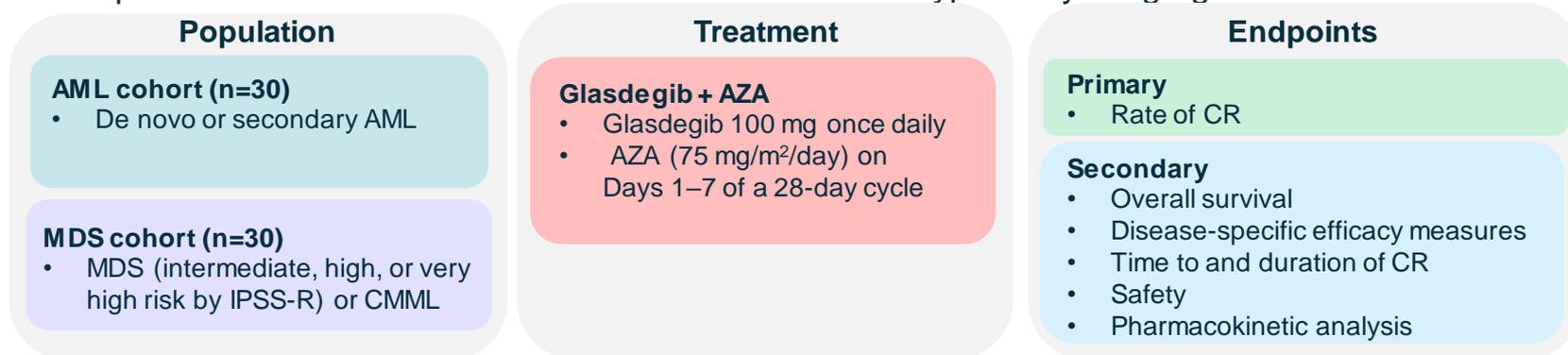
Parameter	1L MDS N=24
RBC transfusion independence ¹	4/9 (44%)
Complete cytogenetic response in responders ²	5/19 (26%)
MRD negativity in responders	5/22 (23%)
Median duration of response (months)	Not reached (0.03+ – 9.76+)
Median follow-up [range] (months)	6.4 [2.0 – 14.4]

Med OS not reached
5 patients received allogeneic HSCT



BRIGHT MDS & AML 1012 Study Design

- BRIGHT MDS & AML 1012 (NCT02367456) is an ongoing open-label, multicenter, phase 1b trial
- Key eligibility criteria:
 - Patients were aged ≥ 18 years
 - Newly diagnosed AML, higher-risk MDS, and CMML
 - Clinical indication for treatment with AZA for AML or MDS
 - No prior treatment with a Smoothened inhibitor and/or a hypomethylating agent

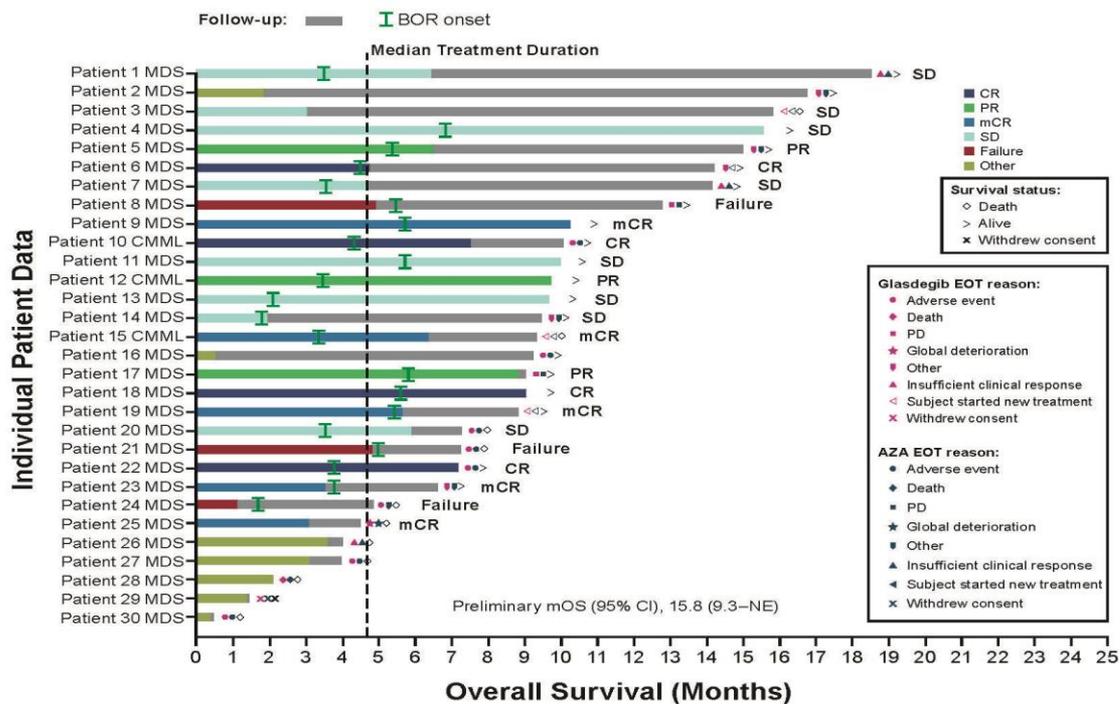


Study start date: April 28, 2015. Data cut-off: September 11, 2019.

AML=acute myeloid leukemia; AZA=azacitidine; CMML=chronic myelomonocytic leukemia; CR=complete remission; IPSS-R=Revised International Prognostic Scoring System; MDS=myelodysplastic syndrome



MDS Cohort: Overall Survival With Best Overall Response



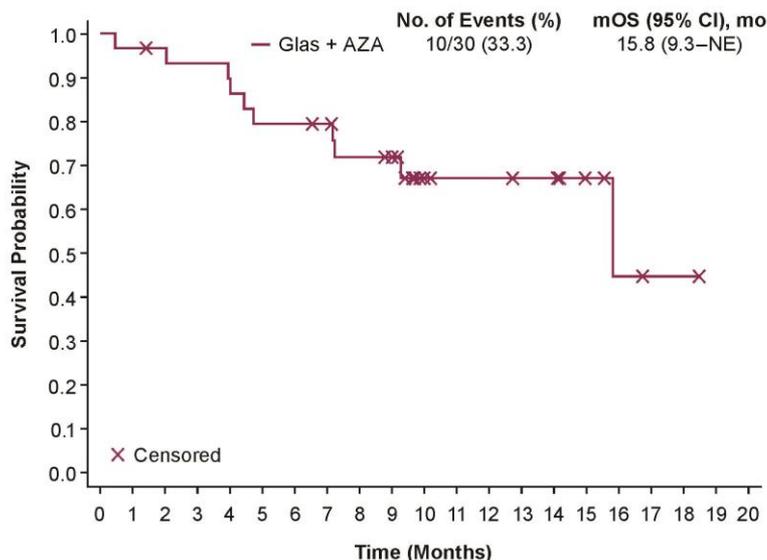
Response	n (%)
CR	4 (13.3)
PR	3 (10.0)
HI without CR or PR	3 (10.0)
CR + PR + HI without CR or PR	10 (33.3)
mCR	5 (16.7)
SD	8 (26.7)

AZA=azacitidine; BOR=best overall response; CMML=chronic myelomonocytic leukemia; CR=complete remission; EOT=end of treatment; HI=hematologic improvement; mCR=marrow complete remission; MDS=myelodysplastic syndrome; PD=progressive disease; PR=partial remission; SD=stable disease

MDS Cohort: Preliminary Overall Survival

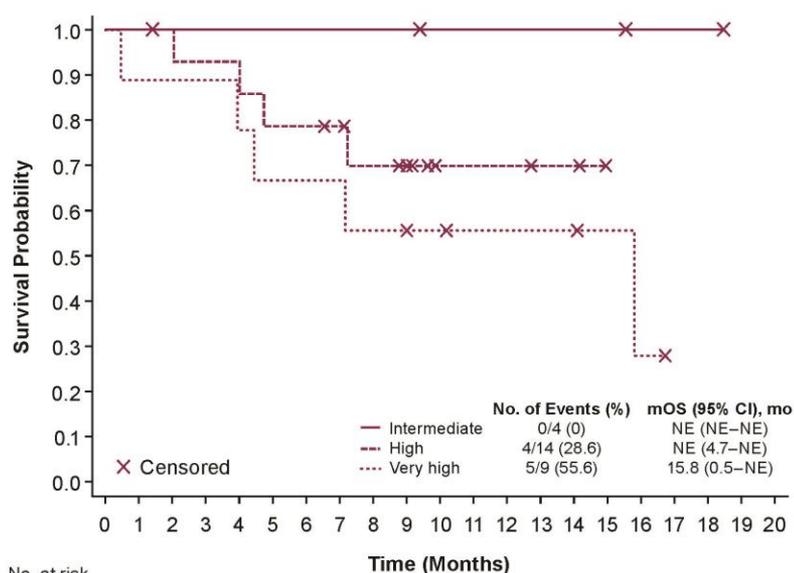


All Patients



No. at risk 30 29 28 27 26 23 23 22 19 18 9 8 8 7 7 4 2 1 1 0

IPSS-R Genetic Risk Category*



No. at risk
Intermediate 4 4 3 3 3 3 3 3 3 2 2 2 2 2 2 1 1 1 0
High 14 14 14 13 13 11 11 10 8 7 3 3 3 2 2 0
Very high 9 8 8 8 7 6 6 6 5 5 4 3 3 3 3 2 1 0

*Risk category based on 27 MDS patients alone.

AZA=azacitidine; CI=confidence interval; Glas=glasdegib; IPSS-R=Revised International Prognostic Scoring System; MDS=myelodysplastic syndrome; mo, months; mOS=median overall survival; NE=not evaluable

Beyond HMAs | Other Combos

AZA + Checkpoint inhibitors

AZA + Other HDAC-I

AZA + Pevonedistat

AZA + IDHi (or IDHi alone)

AZA + mAb (or mAb alone)

AZA + Rigosertib

(or DAC + any of the above)



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- Predicting Responders to HMAs
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- Combinations – Higher-risk
- **Conclusions**



Beyond HMAs | Conclusions

- Predicting response to HMAs coming of age
- Combos in lower-risk MDS focused on anemia
- Combos in higher-risk take different mechanism of action approaches or on genetics

Our drugs fail our patients! Our patients don't fail our drugs.



Thanks!!!



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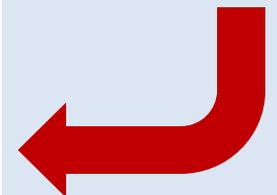


WHEN
BLOOD
BREAKS
DOWN

LIFE LESSONS
FROM LEUKEMIA

MIKKAEL A. SEKERES
Essayist for the New York Times

April
2020!



And Our Patients!!!