

# Updates on Treating



## MDS and Secondary AML

**Mikkael A. Sekeres, MD, MS**

Chief, Division of Hematology

Sylvester Comprehensive Cancer Center

# Treating MDS | Agenda

---

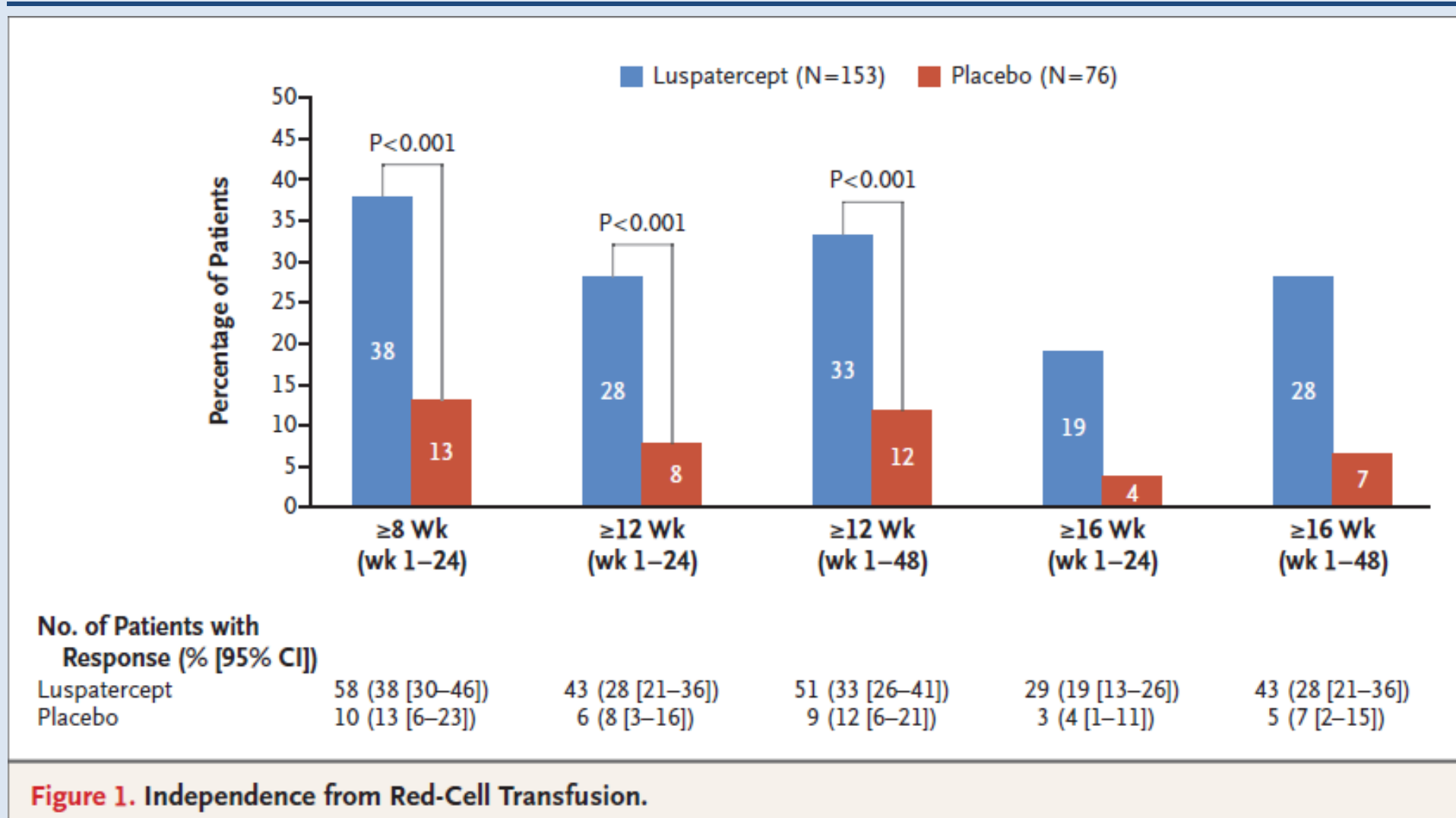
- Treating Lower-risk MDS
- Treating Higher-risk MDS
- Treating sAML

# Treating MDS | Agenda

---

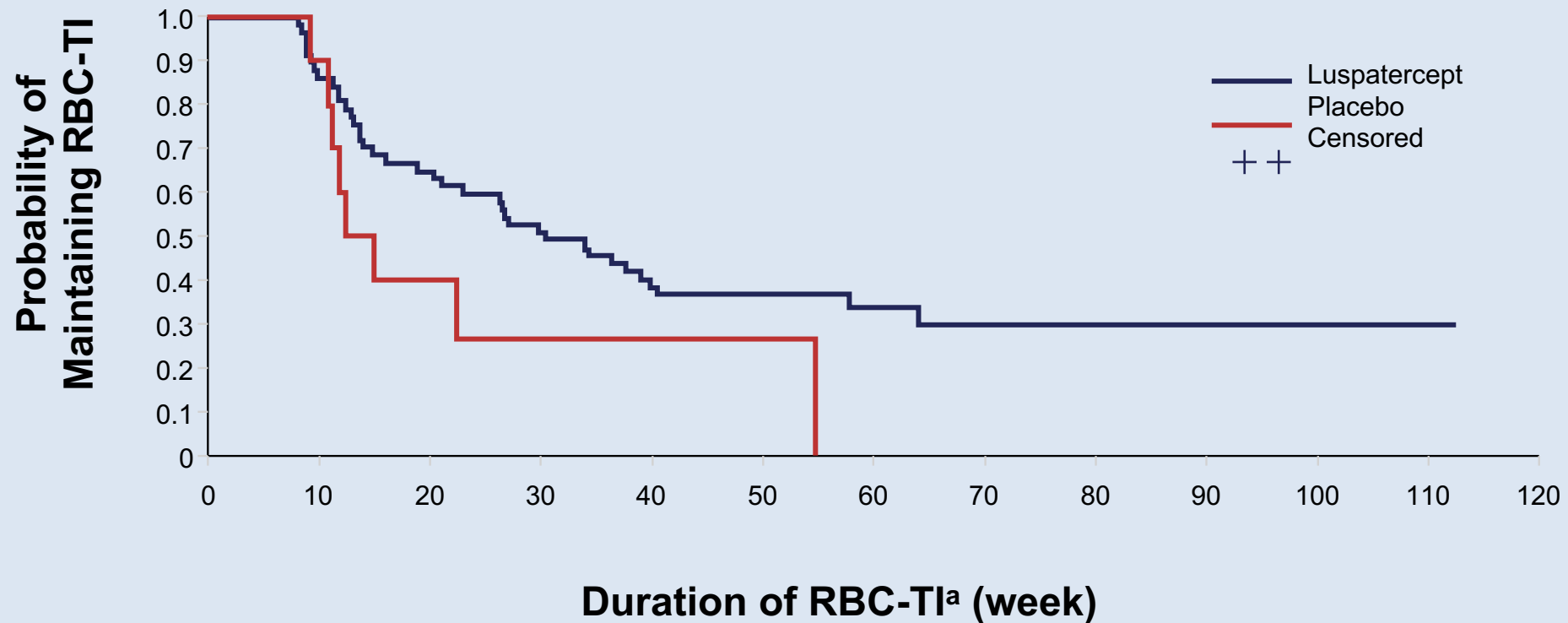
- **Treating Lower-risk MDS**
- Treating Higher-risk MDS
- Treating sAML

# MDS | Ameliorating Anemia: LUSPAT



# MDS | Ameliorating Anemia: LUSPAT

Median duration (weeks) (95% CI): 30.6 (20.6–40.6) vs 13.6 (9.1–54.9)



# MDS | Ameliorating Anemia: LUSPAT

---

## **Impact on Patient Care and Treatment Algorithm:**

- Option post-ESA
- ORR/duration similar to ESA
- Few side effects

\*\*\* But – Does TI duration/definition offset time investment for shots?  
Is it worth the \$\$? \*\*\*

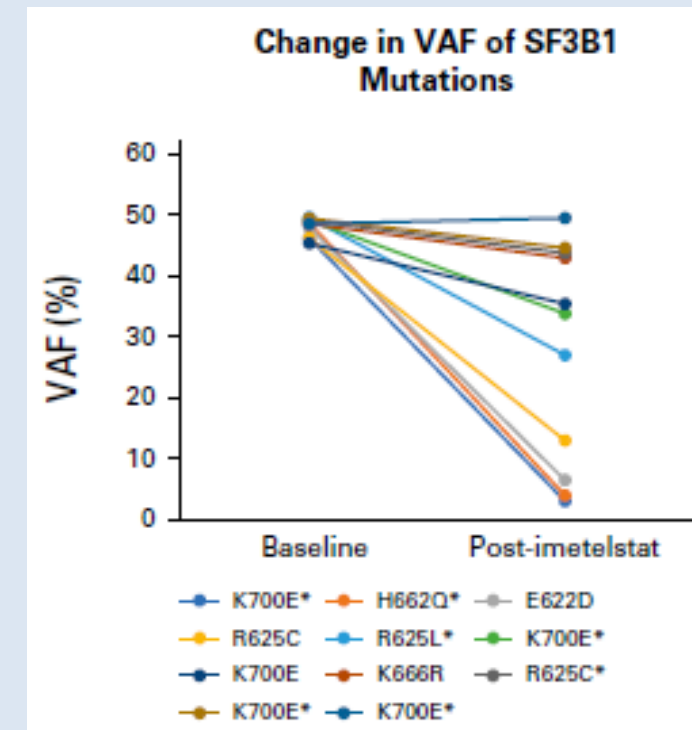
## **Implications for Future Research:**

- Compare up-front to ESA
- Is having RS such a big deal?
- Combine with ESA, LEN, HMA...

# MDS | Ameliorating Anemia: Imetel

## Imetelstat in HTB Lower-risk MDS

Parameter	Overall Population (n = 57)
8-week TI <sup>a</sup> , No. (%)	21 (37)
Median time to onset, weeks (range)	8.3 (0.1-100.6)
Median duration of TI <sup>b</sup> , weeks (range)	65 (17.0-140.9)
24-week TI <sup>a</sup> , No. (%)	13 (23)
HI-E per IWG 2006, No. (%)	37 (65)
≥ 1.5 g/dL increase in Hgb lasting ≥ 8 weeks	15 (26)
Transfusion reduction by ≥ 4 units/8 weeks	37 (65)
Response per IWG 2018, No. (%)	
Major response: 16-week TI	16 (28)
Major response: 8-week TI	21 (37)
Minor response <sup>c</sup>	28 (49)



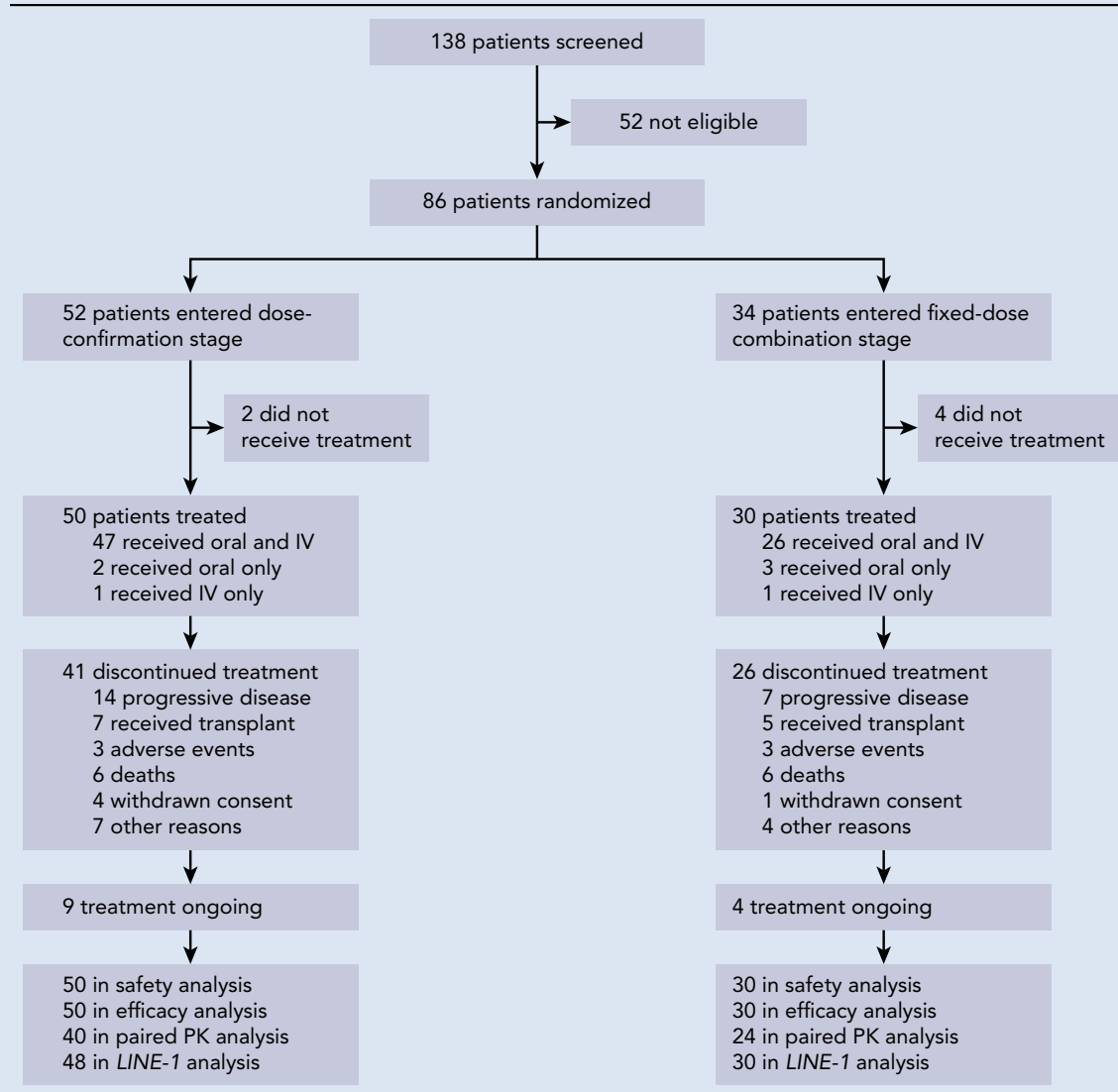
# Treating MDS | Agenda

---

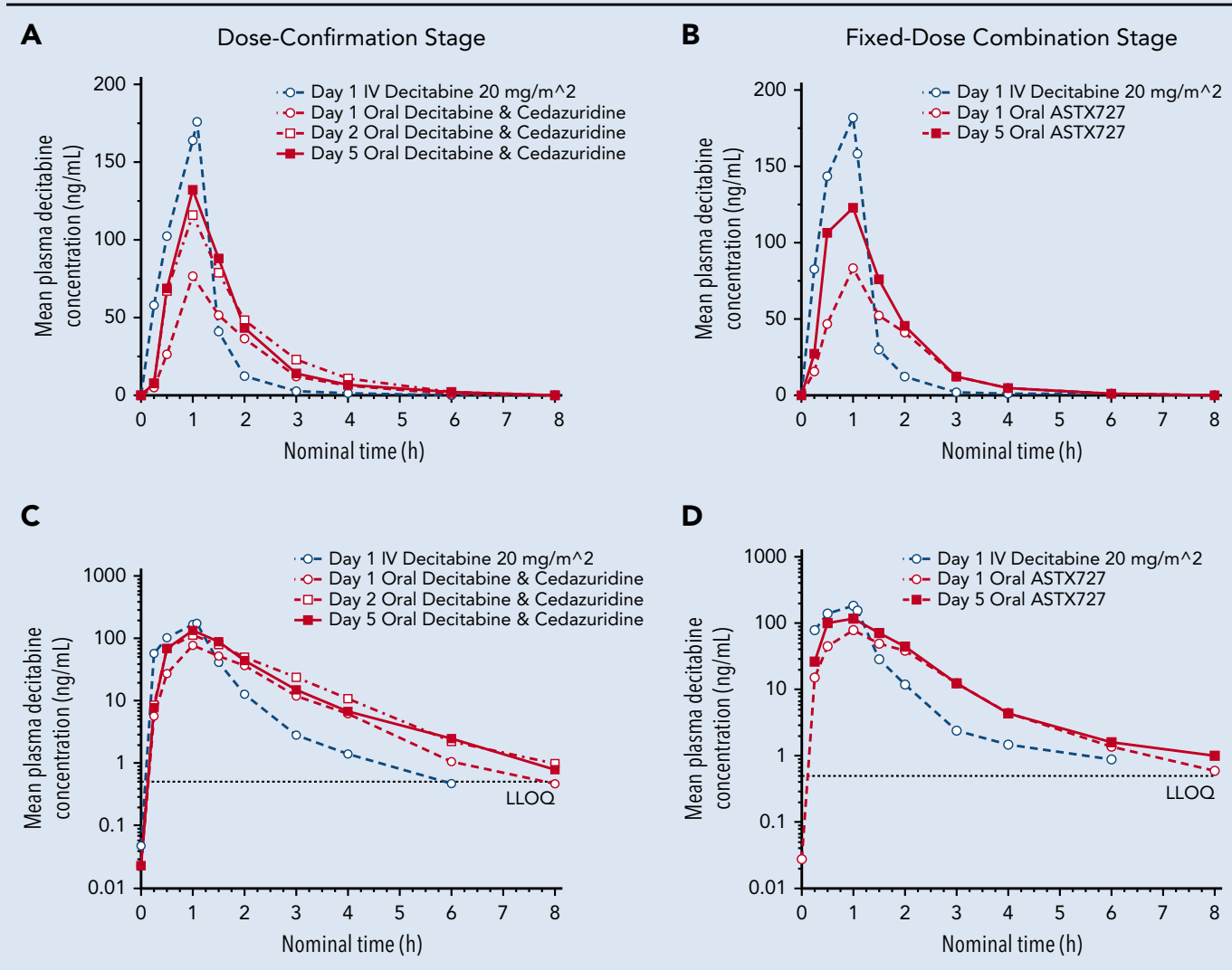
- Treating Lower-risk MDS
- **Treating Higher-risk MDS**
- Treating sAML



# Higher-risk MDS | HMAs: DAC/CED



# Higher-risk MDS | HMAs: DAC/CED



# Higher-risk MDS | HMAs: DAC/CED

## Oral Cedazuridine/Decitabine Phase 2 In Int-1, Int-2, High, CMML

Type of response	Phase 2 overall (N=80)	
	n (%)	95% CI
CR	17 (21)	13, 32
PR	0	
mCR	18 (22)	14, 33
With HI	6 (7)	3, 16
HI	13 (16)	9, 26
HI-E	8 (10)	4, 19
HI-N	2 (2)	0, 9
HI-P	11 (14)	7, 23
Overall response (CR + PR + mCR + HI)	48 (60)	48, 71
No response	32 (40)	29, 52

# Higher-risk MDS | HMAs

---

## **Impact on Patient Care and Treatment Algorithm:**

- DAC/CED has similar ORR to AZA or DAC
- Lower-risk MDS patients included in studies
- Crossover to IV occurred

\*\*\* But – Long-term follow-up pending. Impact on OS?  
Is it worth the \$\$? \*\*\*

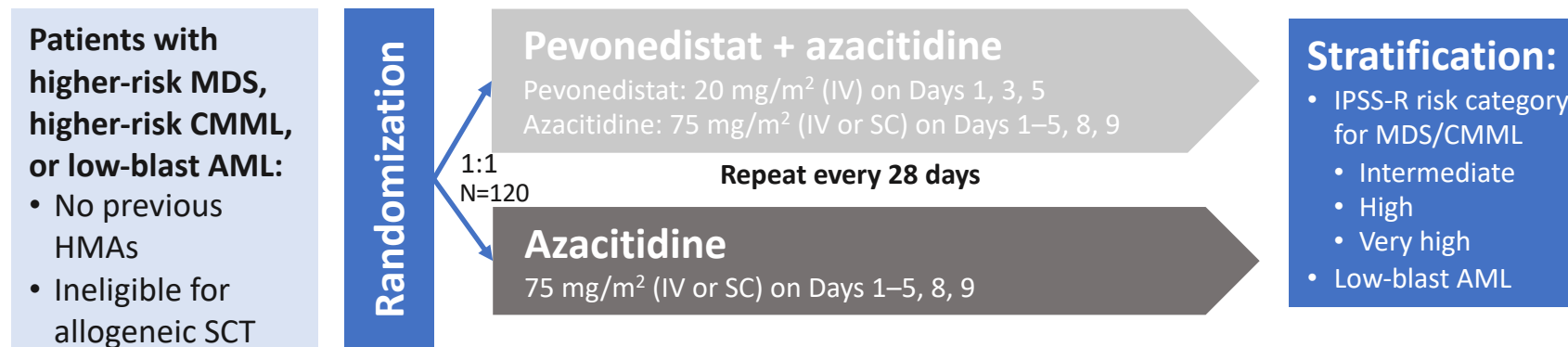
## **Implications for Future Research:**

- Identifying molecular subtypes who did particularly well
- Combine with molecularly targeted drugs
- Give as easy maintenance post-HCT, post-IC

# Higher-risk MDS | Combinations

## Study design: AZA +/- Pevonedistat

NCT02610777: Phase 2, randomized, open-label, global, multicenter study [proof of concept]



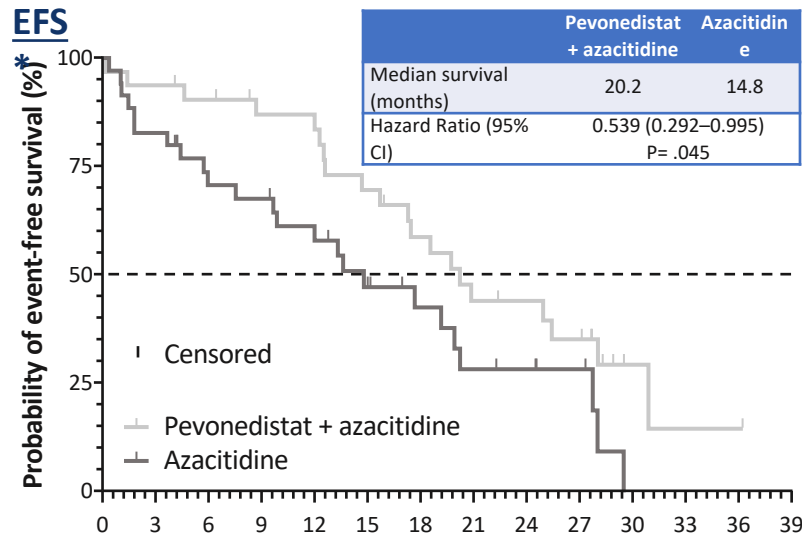
### Study endpoints

- **EFS** (defined as time to death or transformation to AML in higher-risk MDS/CMML or death in low-blast AML): Trial was powered on EFS as the original primary endpoint
- **OS**: Original secondary endpoint, changed to primary endpoint based on regulatory feedback after enrollment
- **ORR**: Secondary endpoint

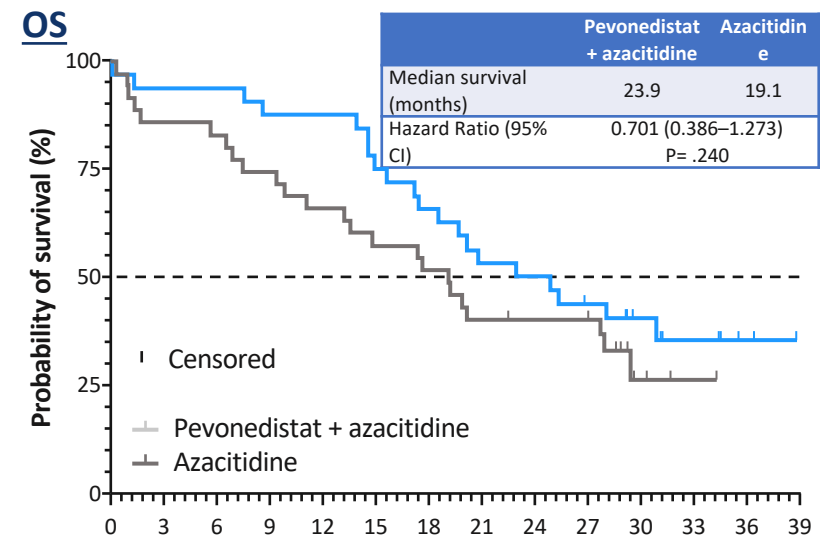
EFS, event-free survival; HMAs, hypomethylating agents; IPSS-R, Revised International Prognostic Scoring System; IV, intravenous; ORR, objective response rate; SC, subcutaneous; SCT, stem cell transplant

# Higher-risk MDS | Combinations

## EFS and OS: Higher-risk MDS



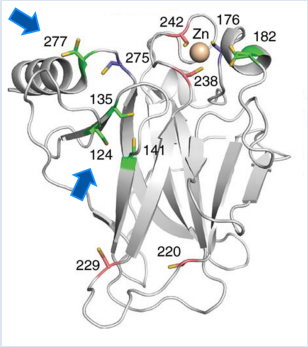
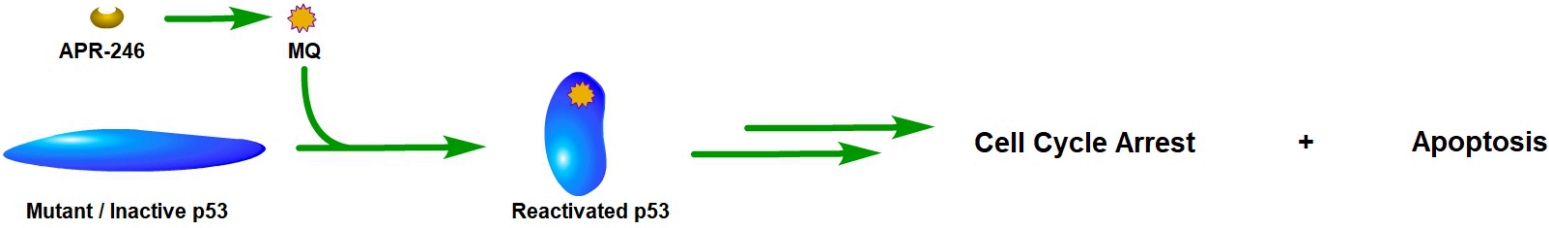
Number at risk	Months													
	0	3	6	9	12	15	18	21	24	27	30	33	36	39
Pevonedistat + azacitidine	32	30	28	25	24	20	16	11	10	8	2	1	1	0
Azacitidine	35	29	23	22	18	12	9	6	5	4	0	0	0	0



Number at risk	Months													
	0	3	6	9	12	15	18	21	24	27	30	33	36	39
Pevonedistat + azacitidine	32	30	30	28	28	24	21	17	16	13	8	5	2	0
Azacitidine	35	30	29	26	23	20	18	14	13	13	3	1	0	0

\*EFS defined as time to death or transformation to AML in higher-risk MDS/CMML or death in low-blast AML.

# Higher-risk MDS | Targeting *TP53*

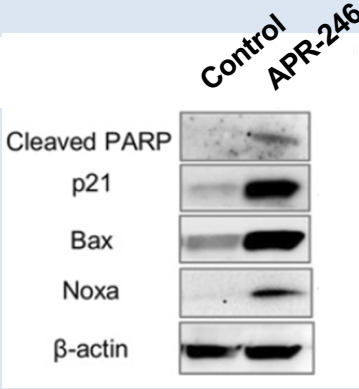


APR-246 binds covalently to p53...

p53 R175H

p53 R175H + APR-246

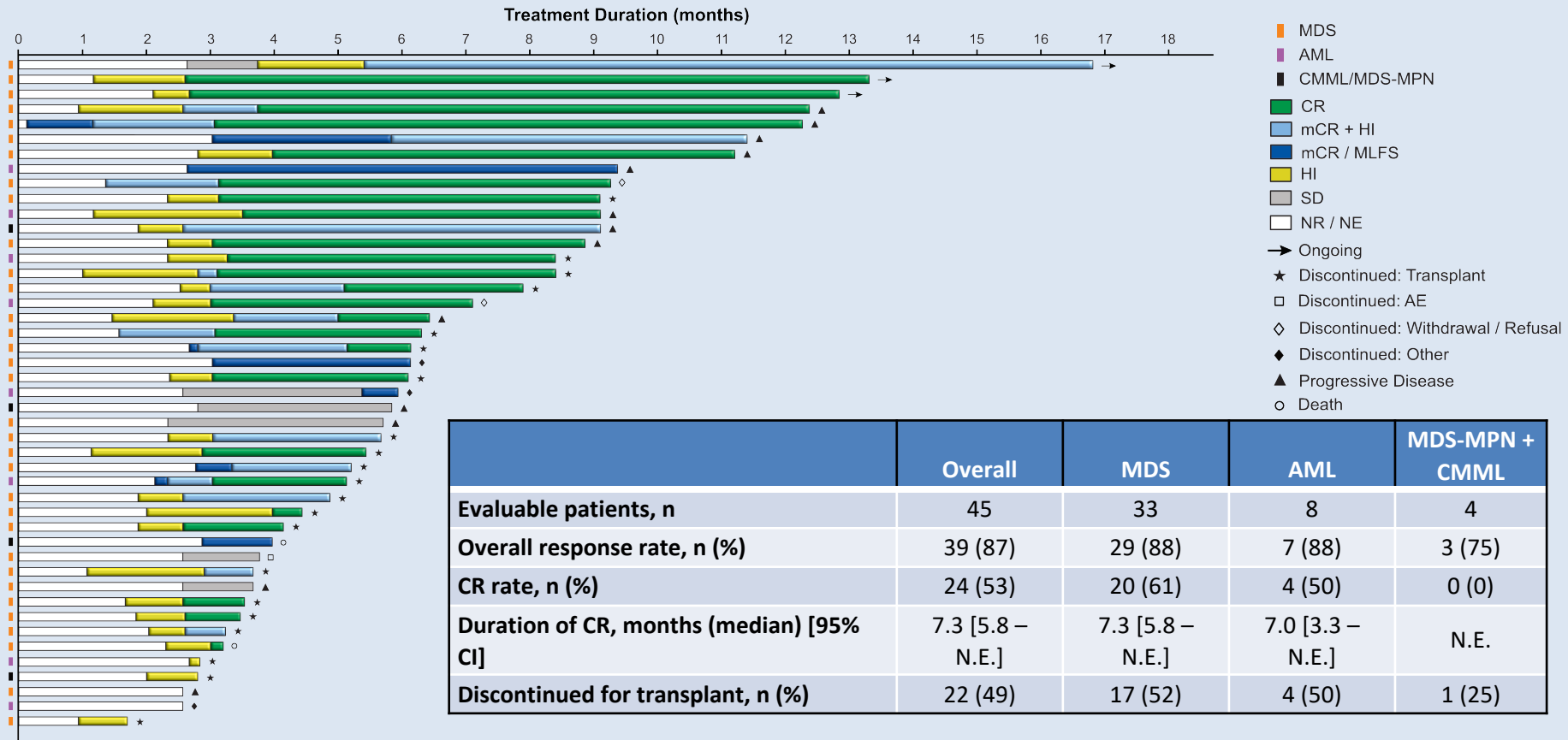
...restores wt p53 conformation & activity...



...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.

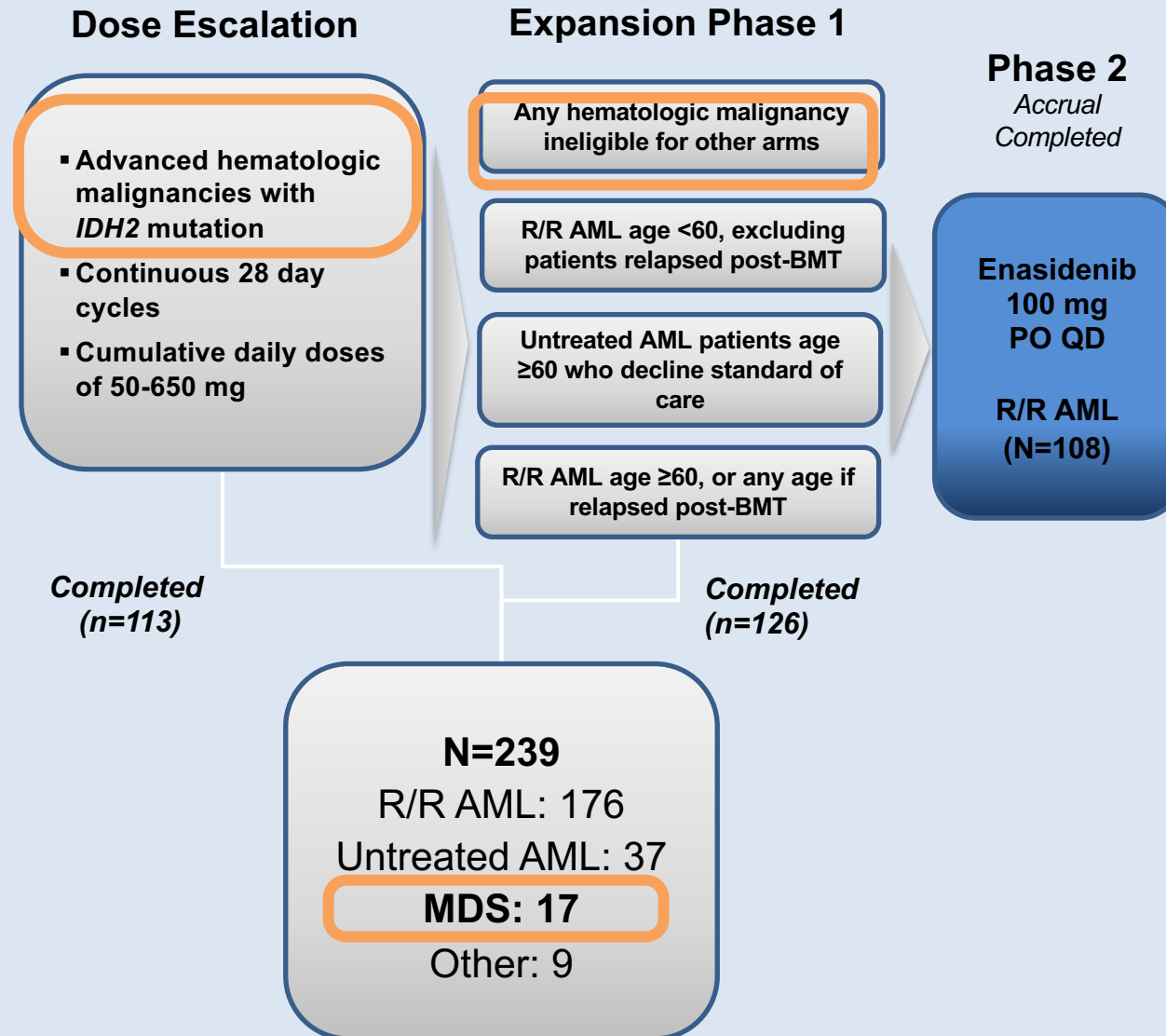
# Higher-risk MDS | Targeting *TP53*



Median duration of follow-up = 10.8 months



# Higher-risk MDS | Targeting *IDH2*



# Higher-risk MDS | Targeting *IDH2*

	MDS Patients (N=17) n (%)
<b>Overall response rate (CR + PR + mCR + HI)</b>	<b>10/17 (59)</b>
<b>Best Response</b>	
Complete remission	1/11 (9)
Partial remission	1/11 (9)
Marrow CR	3/11 (27)
<b>Any hematologic improvement (HI)<sup>†</sup></b>	<b>5/17 (29)</b>
HI-E	3/15 (20)
HI-P	4/12 (33)
HI-N	4/10 (40)

# Higher-risk MDS | Combinations

---

## Phase Ib of AZA + VEN in R/R higher-risk MDS

### Patients:

N=38

Median age = 74 years

Median 8 cycles prior HMA

Median follow-up = 6.8 months

### Responses:

CR N=3

HI N=9

TI N=13 for median of 4.1 months

Median PFS = 9.1 months

# Higher-risk MDS | Combinations

---

## Phase Ib of AZA + VEN in treatment-naïve higher-risk MDS

Ven given in escalating dose (100, 200, and 400 mg) for 14 days of a 28-day cycle

### Patients:

N=57

Median age = 71 years

Median follow-up = 13 months

### Responses:

CR 42%

Marrow CR with HI 14%

Median response duration 14.8 months

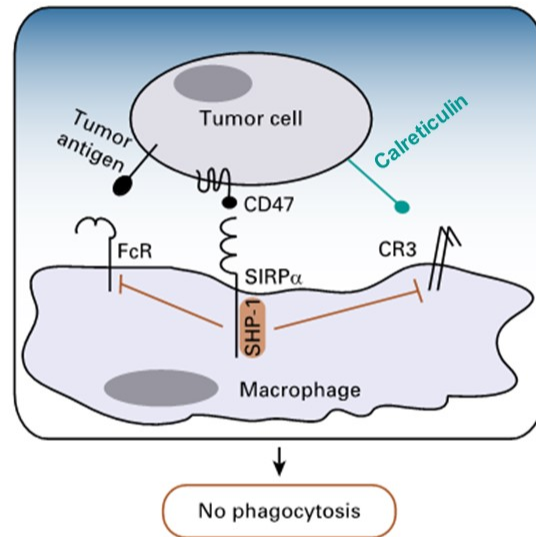
TI N=13 for median of 4.1 months

Median PFS = 17.5 months

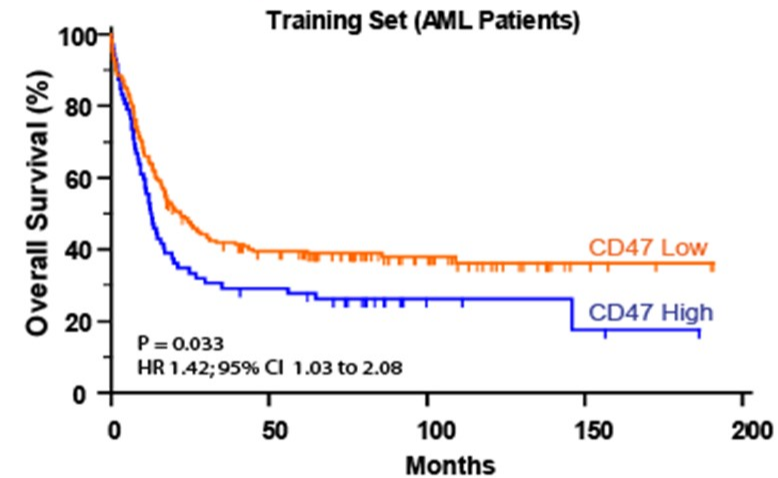
EORTC QLQ C30 improvement in fatigue and dyspnea

# Higher-risk MDS | Combinations

## CD47 Is a Major Macrophage Immune Checkpoint and ‘Do Not Eat Me’ Signal in Myeloid Malignancies Including MDS and AML



### CD47 Expression in AML Patients

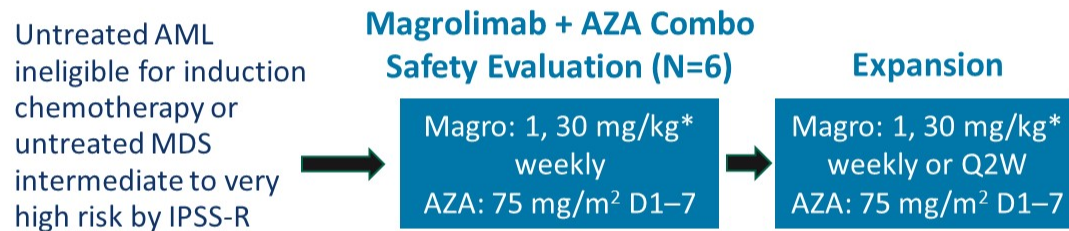


- CD47 is a “do not eat me” signal on cancers that enables macrophage immune evasion
- Increased CD47 expression predicts worse prognosis in AML patients

Figure at left adapted from Veillette A, Tang Z. *J Clin Oncol*. 2019;37(12):1012-1014, and Chao MP, et al. *Current Opin Immunol*. 2012; 24(2):225-232.  
Figure at right adapted from Majeti R, et al. *Cell*. 2009;138(2):286-299.

# Higher-risk MDS | Combinations

## 5F9005 Study Design: Magrolimab in Combination With AZA in MDS and AML



### Primary objectives

1. Safety of magrolimab alone or with AZA
2. Efficacy of magrolimab + AZA in untreated AML/MDS

### Secondary objectives

1. Pharmacokinetics, pharmacodynamics, and immunogenicity of 5F9
2. Additional measures of efficacy (DOR, PFS, OS)

### Exploratory objective

To assess CD47 receptor occupancy, markers of immune cell activity, and molecular profiling in AML/MDS

- A magrolimab priming dose (1 mg/kg) and dose ramp-up was utilized to mitigate on-target anemia
- Data from the expansion cohort is presented

\*Dose ramp-up from 1 mg/kg to 30 mg/kg by week 2, then 30 mg/kg maintenance dosing or 30 mg/kg Q2W starting Cycle 3+.  
IPSS-R: Revised International Prognostic Scoring System.

# Higher-risk MDS | Combinations

## Patient Characteristics (N=68): Magrolimab + AZA in Untreated (1L) MDS and AML

Characteristic	1L MDS 5F9+AZA (N=39)	1L AML 5F9+AZA (N=29)
Median age (range)	70 (47–80)	74 (60–89)
ECOG Performance Status: 0	11 (28%)	7 (24%)
1	26 (67%)	20 (69%)
2	2 (5%)	2 (7%)
Cytogenetic Risk: Favorable	0	0
Intermediate	11 (28%)	2 (7%)
Poor	25 (64%)	21 (72%)
Unknown/missing	3 (8%)	6 (21%)
WHO AML classification: MRC		19 (66%)
Recurrent genetic abnormalities		2 (7%)
Therapy related	NA	3 (10%)
Not otherwise specified		5 (17%)
WHO MDS classification:		
RS and single/multilineage dysplasia	1 (3%)	
Multilineage dysplasia	7 (18%)	NA
RS with multilineage dysplasia	3 (8%)	
Excess blasts	22 (56%)	
Unclassifiable/unknown/missing	6 (15%)	
IPSS-R (MDS): Intermediate	13 (33%)	
High	19 (49%)	NA
Very High	6 (15%)	
Unknown/missing	1 (3%)	
Therapy related MDS	12 (31%)	
Unknown/missing	1 (3%)	
Harboring a TP53 mutation	5 (13%)	13 (45%)

MRC, myelodysplasia-related changes; NA, not applicable; all patients enrolled on study are shown; WHO, World Health Organization.

- 64%–72% of MDS and AML patients were poor cytogenetic risk
- 66% of AML patients had underlying myelodysplasia (MRC)
- 31% of MDS patients were therapy related
- 45% of AML patients were TP53 mutant

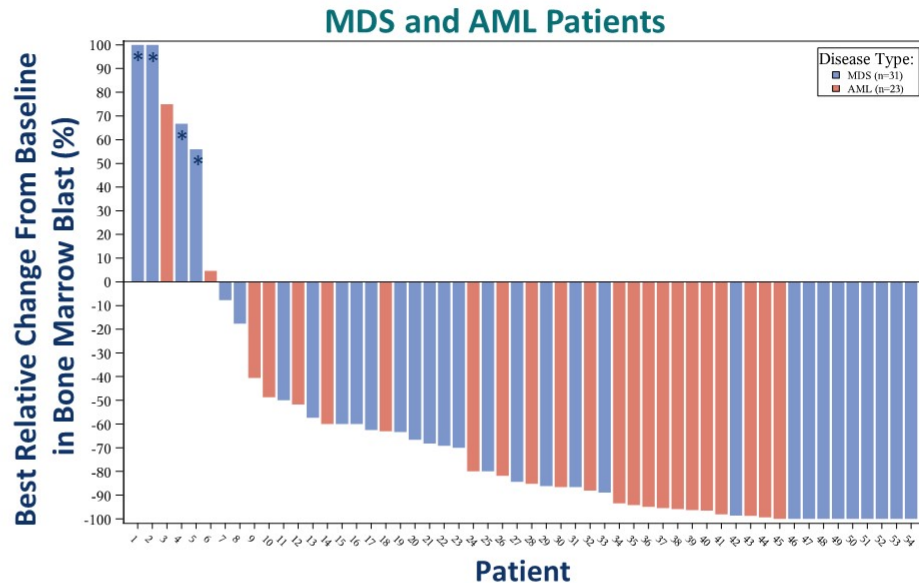


# Higher-risk MDS | Combinations

## Magrolimab + AZA Induces High Response Rates in MDS and AML

Best Overall Response	1L MDS N=33	1L AML N=25
ORR	30 (91%)	16 (64%)
CR	14 (42%)	10 (40%)
CRi	NA	4 (16%)
PR	1 (3%)	1 (4%)
MLFS/marrow CR	8 (24%) 4 with marrow CR + HI	1 (4%)
Hematologic improvement (HI)	7 (21%)	NA
SD	3 (9%)	8 (32%)
PD	0	1 (4%)

Response assessments per 2006 IWG MDS criteria and 2017 AML ELN criteria. Patients with at least 1 post-treatment response assessment are shown; all other patients are on therapy and are too early for first response assessment, except for 2 MDS patients not evaluable (withdrawal of consent) and 3 AML patients (1 AE, 2 early withdrawal).



Four patients not shown due to missing values; <5% blasts imputed as 2.5%. \*Baseline bone marrow blasts ≤5%.

- Magrolimab + AZA induces a 91% ORR (42% CR) in MDS and 64% ORR (56% CR/CRi) in AML
- Responses deepened over time with a 56% 6-month CR rate in MDS patients (assessed in all patients 6 months after initial treatment)
- Median time to response is 1.9 months, more rapid than AZA alone
- Magrolimab + AZA efficacy compares favorably to AZA monotherapy (CR rate 6-17%<sup>1,2</sup>)

1. Azacitidine USPI. 2. Fenaux P, et al. *Lancet Oncol.* 2009 ;10(3):223-232.



# Treating MDS | Combinations

---

## **Impact on Patient Care and Treatment Algorithm:**

- Modest ORR for AZA + Ven in previously treated
- Good ORR for AZA + Ven in untreated higher-risk MDS
- Good ORR for AZA + Mag in untreated Intermediate/Higher-risk MDS

\*\*\* But – Is ORR really better than AZA monotherapy in selected patients?  
Is durability of response any better?  
Is it worth the \$\$? \*\*\*

## **Implications for Future Research:**

- Identifying molecular subtypes who did particularly well
- Optimizing Ven dose and schedule
- Combine with molecularly targeted drugs (Triplet therapy)

Zeiden et al. EHA 2020;S118.

Garcia et al. ASH 2020;656a.

Sallman et al. ASCO 2020;7507a.

Courtesy of Mikkael A Sekeres, MD, MS

# Treating MDS | Agenda

---

- Treating Lower-risk MDS
- Treating Higher-risk MDS
- **Treating sAML**

# Treating sAML | CPX-351

- In total, 309 patients were randomized to receive CPX-351 (n = 153) or 7+3 (n = 156) and were included in the intent-to-treat population for efficacy analyses; the safety population included 153 and 151 patients, respectively<sup>6</sup>

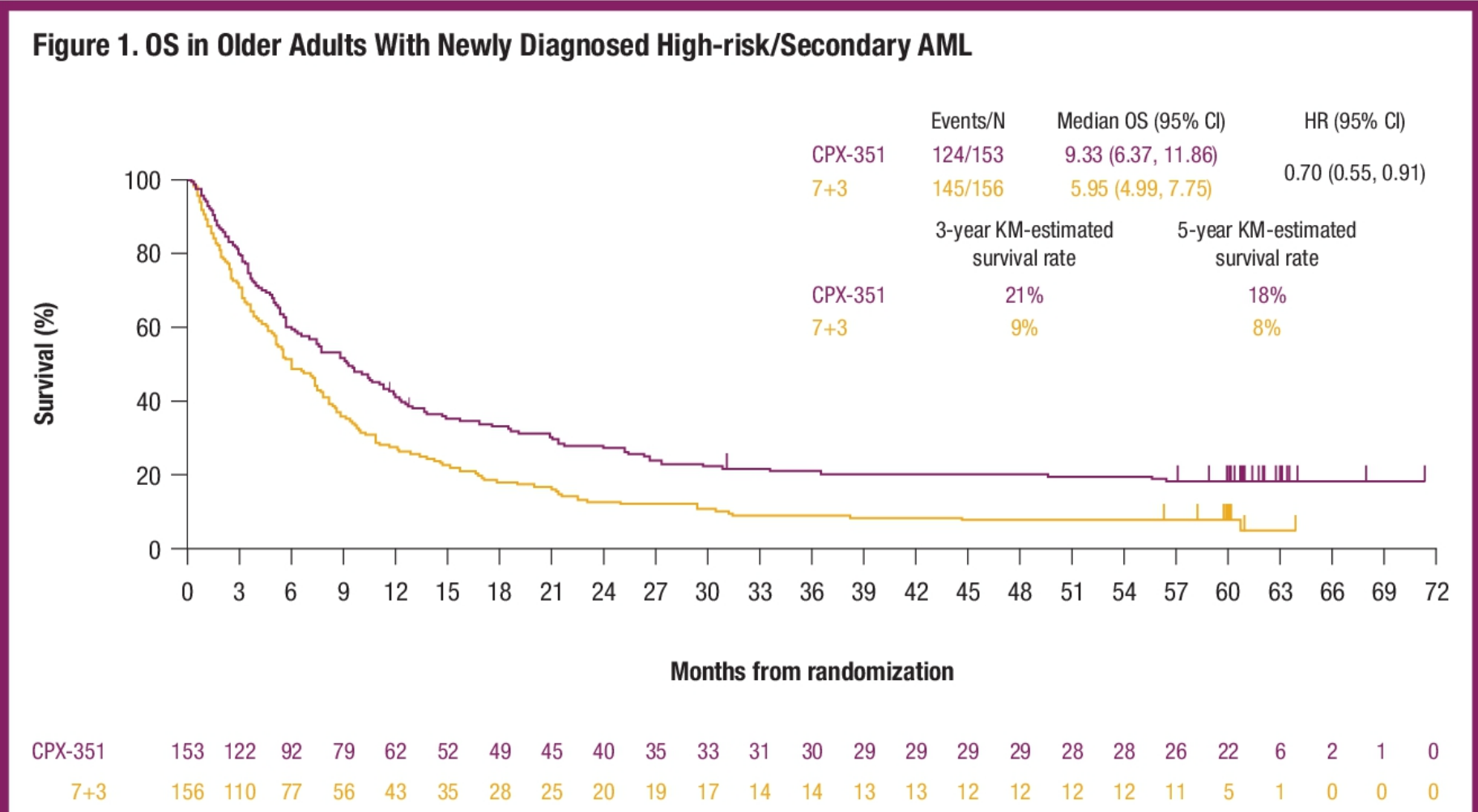
**Table 1. Baseline Characteristics in Older Adults With Newly Diagnosed High-risk/Secondary AML<sup>6</sup>**

Characteristic, n (%)	CPX-351 (n = 153)	7+3 (n = 156)
<b>Demographic characteristics</b>		
<b>Age</b>		
Mean (SD), years	67.8 (4.2)	67.7 (4.1)
60 to 69 years, n (%)	96 (63)	102 (65)
70 to 75 years, n (%)	57 (37)	54 (35)
<b>Male, n (%)</b>	94 (61)	96 (62)
<b>ECOG performance status, n (%)</b>		
0	37 (24)	45 (29)
1	101 (66)	89 (57)
2	15 (10)	22 (14)
<b>Clinical characteristics</b>		
<b>AML subtype, n (%)</b>		
t-AML	30 (20)	33 (21)
AML with antecedent MDS		
With prior HMAs	50 (33)	55 (35)
Without prior HMAs	21 (14)	19 (12)
AML with antecedent CMML	11 (7)	12 (8)
<i>de novo</i> AML with MDS karyotype	41 (27)	37 (24)
<b>Prior HMA therapy, n (%)<sup>a</sup></b>	62 (41)	71 (46)
<b>Cytogenetic risk by NCCN, n (%)</b>		
Favorable	143	146
Intermediate	7 (5)	5 (3)
Unfavorable	64 (45)	58 (40)
	72 (50)	83 (57)
<b>Median bone marrow blasts (range), %</b>	35 (5, 93)	35 (3, 97)
<b>WBC count &lt;20,000/<math>\mu</math>L, n (%)</b>	131/153 (86)	131/155 (85)

AML, acute myeloid leukemia; SD, standard deviation; ECOG, Eastern Cooperative Oncology Group; t-AML, therapy-related AML; MDS, myelodysplastic syndrome; HMA, hypomethylating agent; CMML, chronic myelomonocytic leukemia; NCCN, National Comprehensive Cancer Network; WBC, white blood cell.

<sup>a</sup>Includes patients in the prespecified randomization strata of antecedent MDS with prior HMA exposure as well as patients in other strata (eg, t-AML, antecedent CMML) who had previously received HMAs.

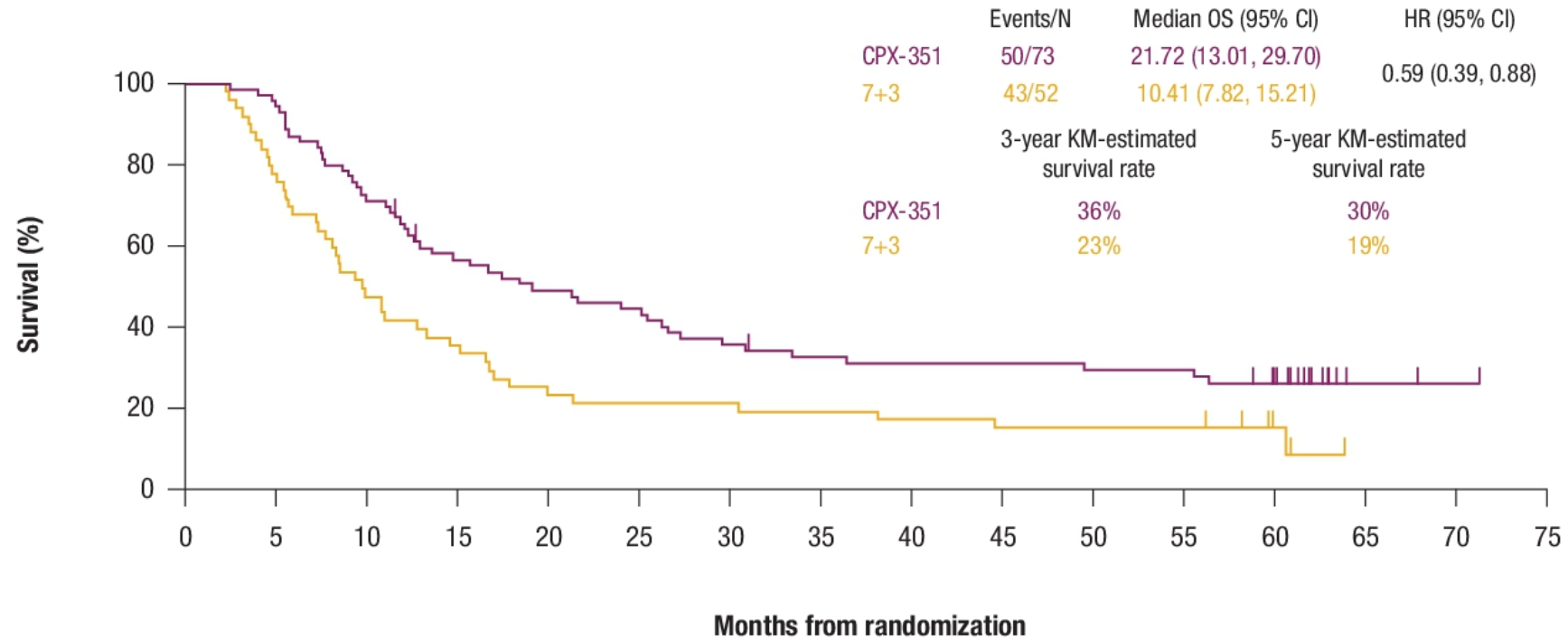
# Treating sAML | CPX-351



# Treating sAML | CPX-351

**Figure 3. OS in Older Adults With Newly Diagnosed High-risk/Secondary AML Who Achieved CR or CRi**

**A. All Patients Who Achieved CR or CRi**



CPX-351	73	69	54	41	36	33	27	24	23	23	22	22	18	2	1	0
7+3	52	41	26	20	15	13	13	12	11	10	10	10	3	0	0	0

# Treating sAML | CPX-351

---

	<i>N</i> = 52
Best response	
CR	
<i>n</i> (%)	15 (29)
95% CI	17.1–43.1
CRi	
<i>n</i> (%)	8 (15)
95% CI	6.9–28.1
CR + CRi	
<i>n</i> (%)	23 (44)
95% CI	30.5–58.7
Time to CR or CRi	
<i>n</i>	23
Median (range), d	37.0 (15–72)
Mean (SD), d	41.5 (15.24)

---

# Treating sAML | CPX-351

---

## **Impact on Patient Care and Treatment Algorithm:**

- Continued improved OS among patients with sAML
- Particularly improved OS for those achieving CR/CRi and in those undergoing HCT.
- Suggests deeper responses.

\*\*\* But – did comparison arm perform as well as historically?  
Was drug intensity similar?  
Is it worth the \$\$? \*\*\*

## **Implications for Future Research:**

- Identifying molecular subtypes who did particularly well
- Outpatient therapies
- Combine with molecularly targeted drugs

# Treating sAML | Magrolimab

---

## Phase Ib of AZA + Mag in treatment-naïve AML

### Patients:

N=52

Median age = 73 years

TP53 + in 65%

Median follow-up = ???

34 evaluable for response

### Responses:

CR 44%, 48% among TP53+

CRi 12%, 19% among TP53+

Median response duration 9.9 months

Median OS for TP53+ = 12.9 months, for non-TP53 18.9 months



# Treating sAML | Magrolimab

---

## **Impact on Patient Care and Treatment Algorithm:**

- Good ORR for AZA + Mag in AML
- Good responses in TP53+
- Response duration good, c/w other single arm combo studies

\*\*\* But – Is ORR really better than AZA monotherapy in selected patients?  
Is durability of response any better?  
Is it worth the \$\$? \*\*\*

## **Implications for Future Research:**

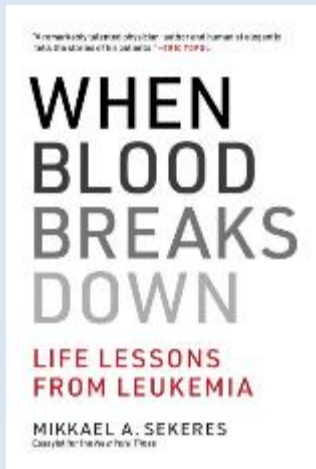
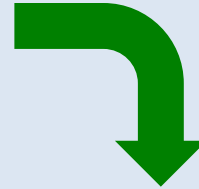
- Identifying molecular subtypes who did particularly well
- Combine with molecularly targeted drugs (Triplet therapy)
- Is it better than AZA + Ven???

# Thanks!!!



**MDS** | The National  
Myelodysplastic  
Syndromes  
Natural History  
Study

Sponsored by the National Heart, Lung, and Blood Institute  
in collaboration with the National Cancer Institute



**UNIVERSITY  
OF MIAMI**

# Appendix 1

---

- Additional Recent Data Set Provided by Dr Sekeres

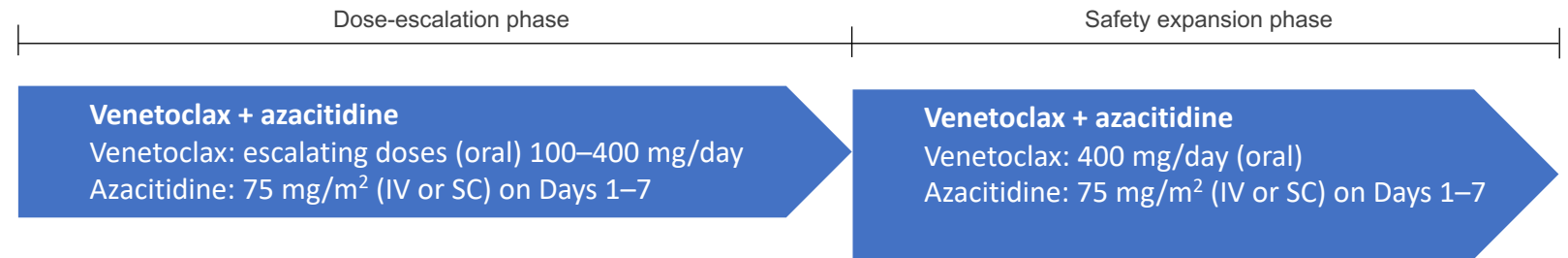
# Venetoclax/azacitidine has been evaluated in a Phase I study

## Ongoing Phase 1b, open-label, dose-escalation,\* multicenter study<sup>1,2</sup>



**Higher-risk MDS**  
No prior therapy, not eligible for intensive chemotherapy or HSCT

N=82



### Primary endpoints:

- Safety
- RP2D



### Secondary endpoints include:

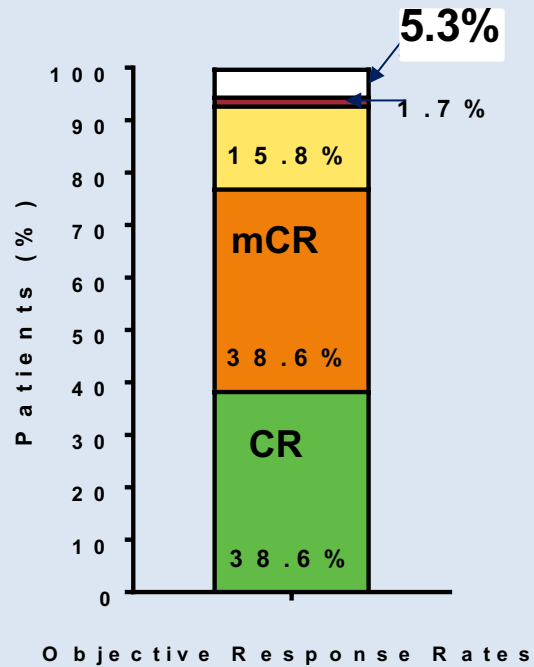
- ORR
- PFS
- TTR
- DoR
- OS

1. Wei AH, *et al. Blood* 2019;134 (Suppl. 1):Abstract #568; 2. ClinicalTrials.gov, NCT02942290

\*Originally a 3-arm, randomized study; amended to dose-escalation safety study after two deaths

TTR, time to response

# 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

# Appendix 2

---

- AML Overview slides
- Prognostic/Predictive Tools slides
- Additional Historic Data Sets Provided

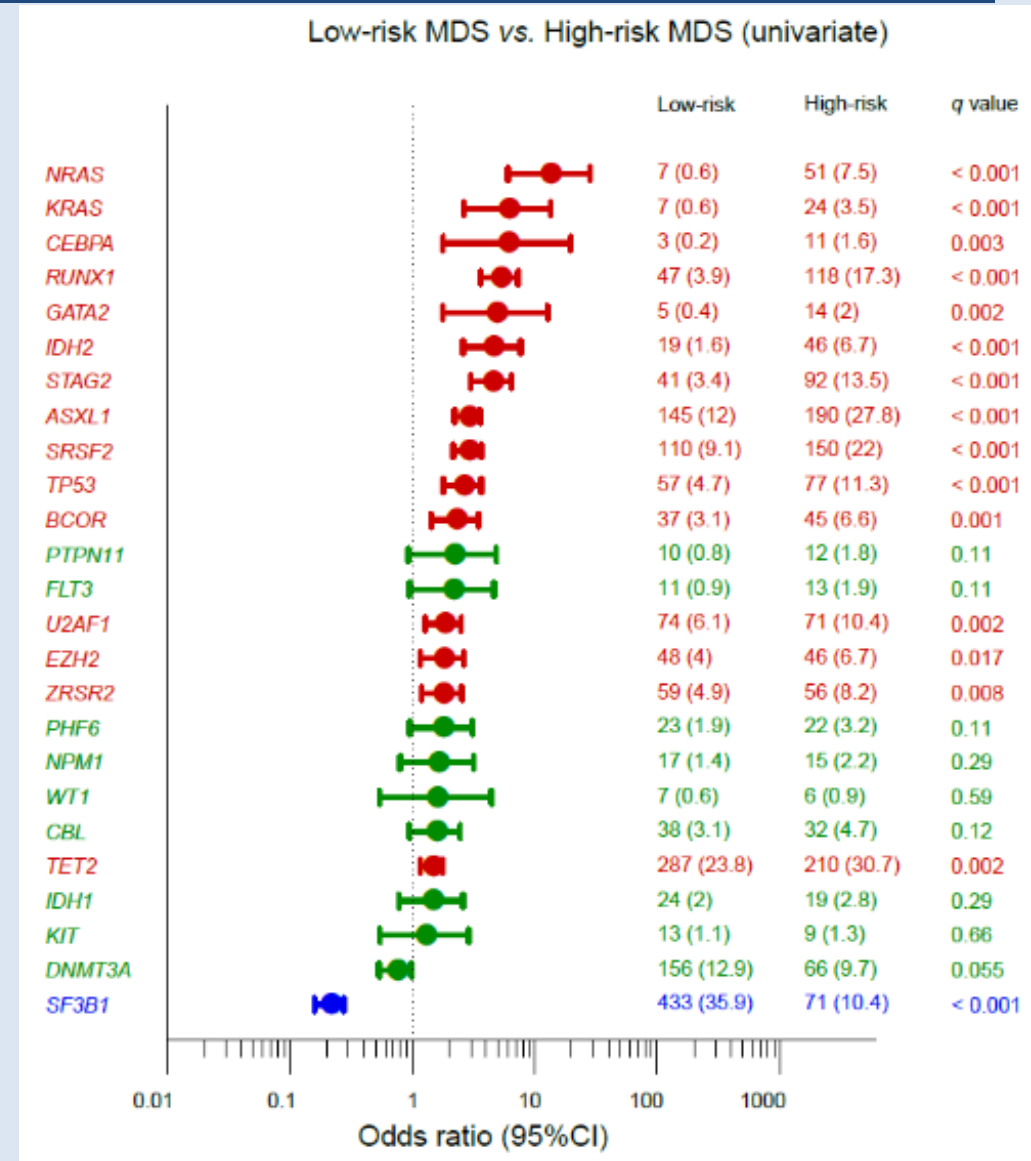
# Treating MDS | Agenda

---

- **Prognostic/Predictive tools**
- Treating Lower-risk MDS
- Treating Higher-risk MDS
- Treating sAML

# Predictive Tools | Mutation Risk

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



Courtesy of Mikkael A Sekeres, MD, MS

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



# MDS | Machine Learning

Data

Random Survival Forest

Important Variables

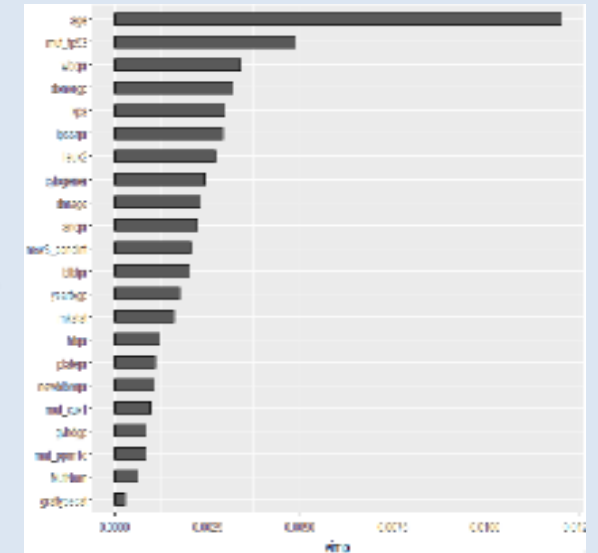
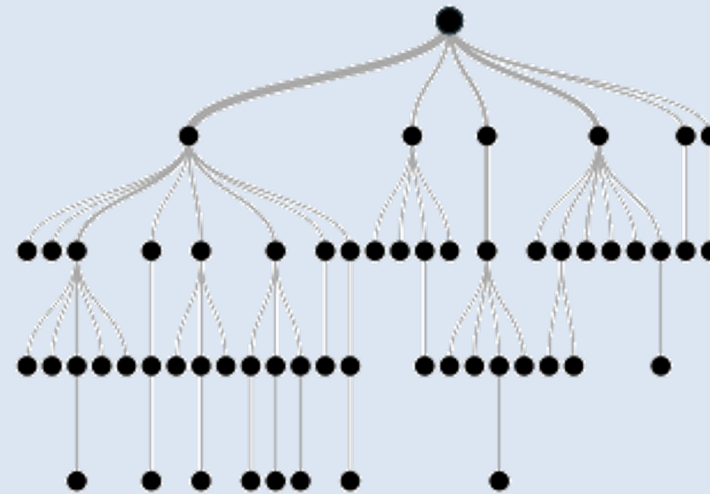
Demographic

Clinical

Genomic

Training  
CC + MLL

Validation  
Moffitt

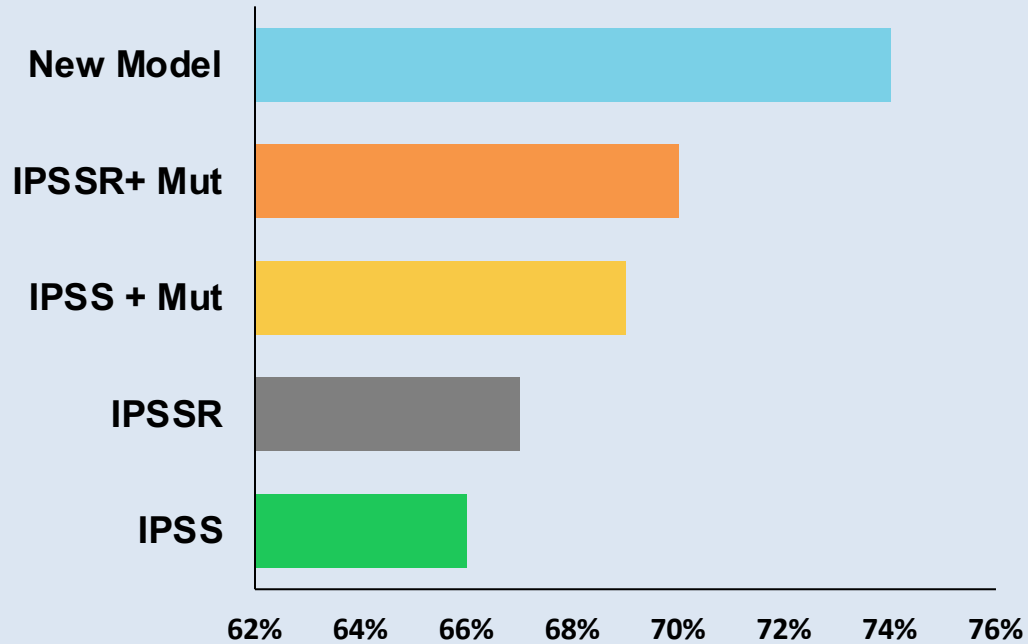


# MDS | Machine Learning

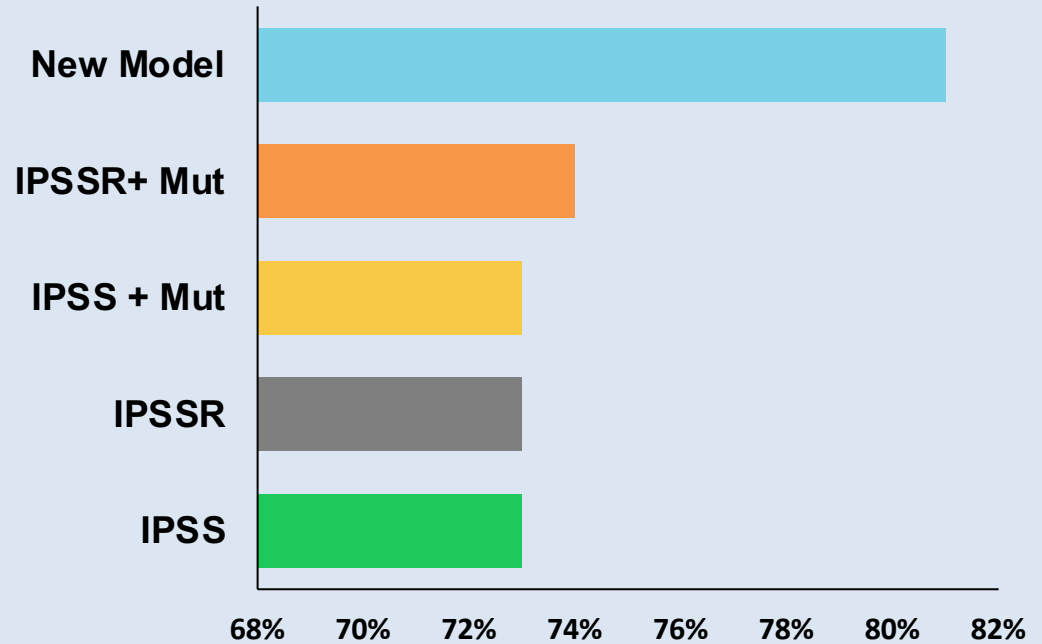
Parameter	Training No. (%) / [range]	Validation No. (%) / [range]	<i>p</i>
<b>Total</b>	1471	831	
<b>Median age, years</b>	71 [19-99]	69 [4-93]	NS
<b>Clinical Variables</b>			
Median WBC, 10 <sup>9</sup> /L	4.2 [0.6-82.6]	4 [0.1-25.6]	NS
Median ANC, 10 <sup>9</sup> /L	2.1 [0-65.1]	2 [0-8.5]	NS
Median Hb, g/dL	9.9 [3.9-15.6]	10 [3.4-17.1]	NS
Median Plts, 10 <sup>9</sup> /L	120 [4-975]	117 [7-1280]	NS
Median BM Blasts %	4 [0-19]	3 [0-19]	NS
<b>2008 WHO Category</b>			
RCMD / RCUD	578 (38)	350 (42)	NS
RARS	209 (11)	128 (15)	
RAEB-1 / RAEB-2	573 (37)	302 (36)	
MDS-U	49 (9)	18 (2)	
MDS with del (5q)	62 (5)	33 (4)	

# MDS | Machine Learning

## Overall Survival



## Leukemia-Free Survival



# Predictive Tools | Mutations/Response

## 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

## Results: Association Rules

**31%** pts

≥ 3 mutations/sample

**29%** pts

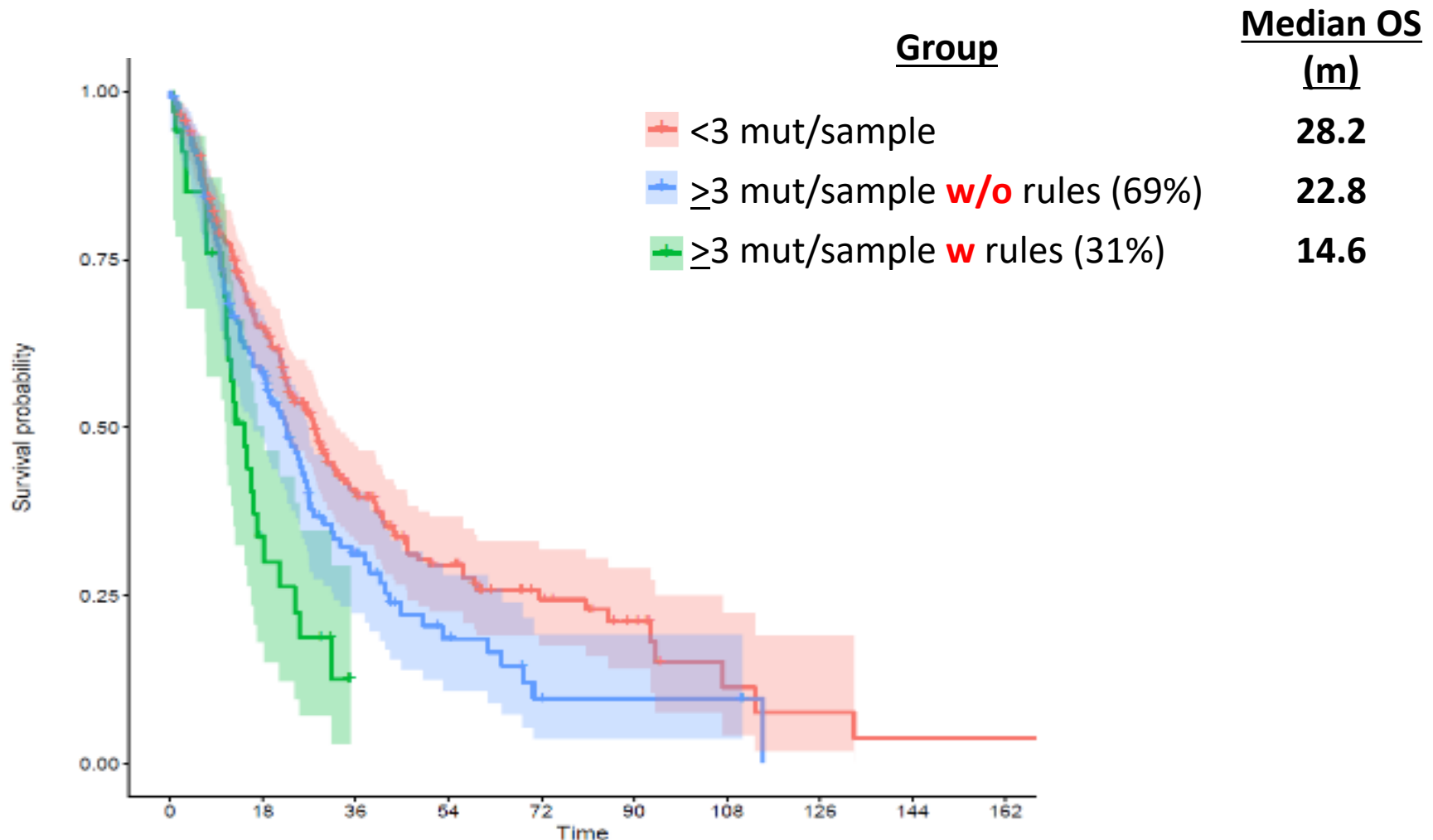
Very Low/Low risk by IPSS-R

ORR to HMAs = **43%**

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

Accuracy: **87%**

# Predictive Tools | Mutations/Response



# Treating MDS | Disease Biology

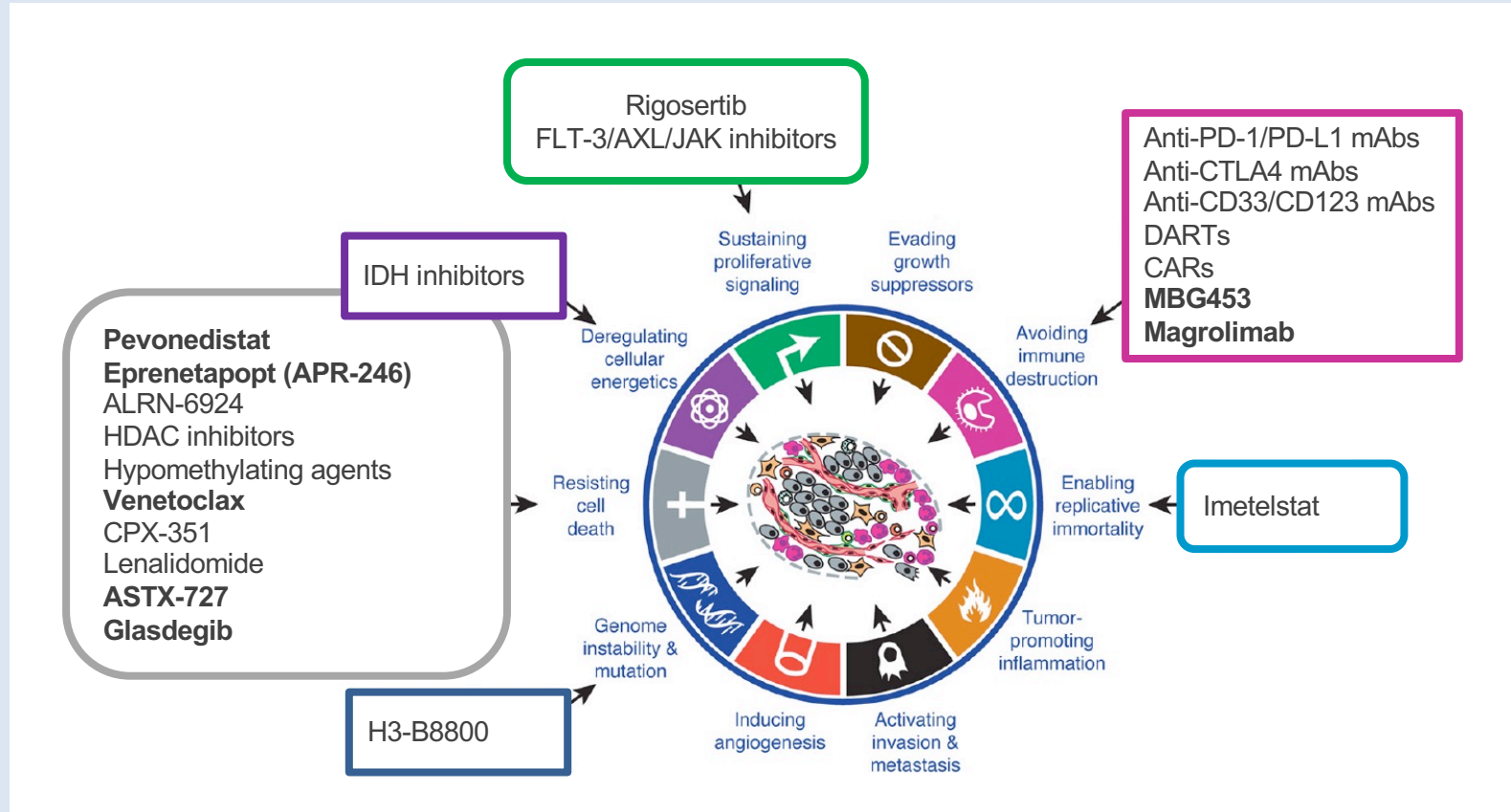
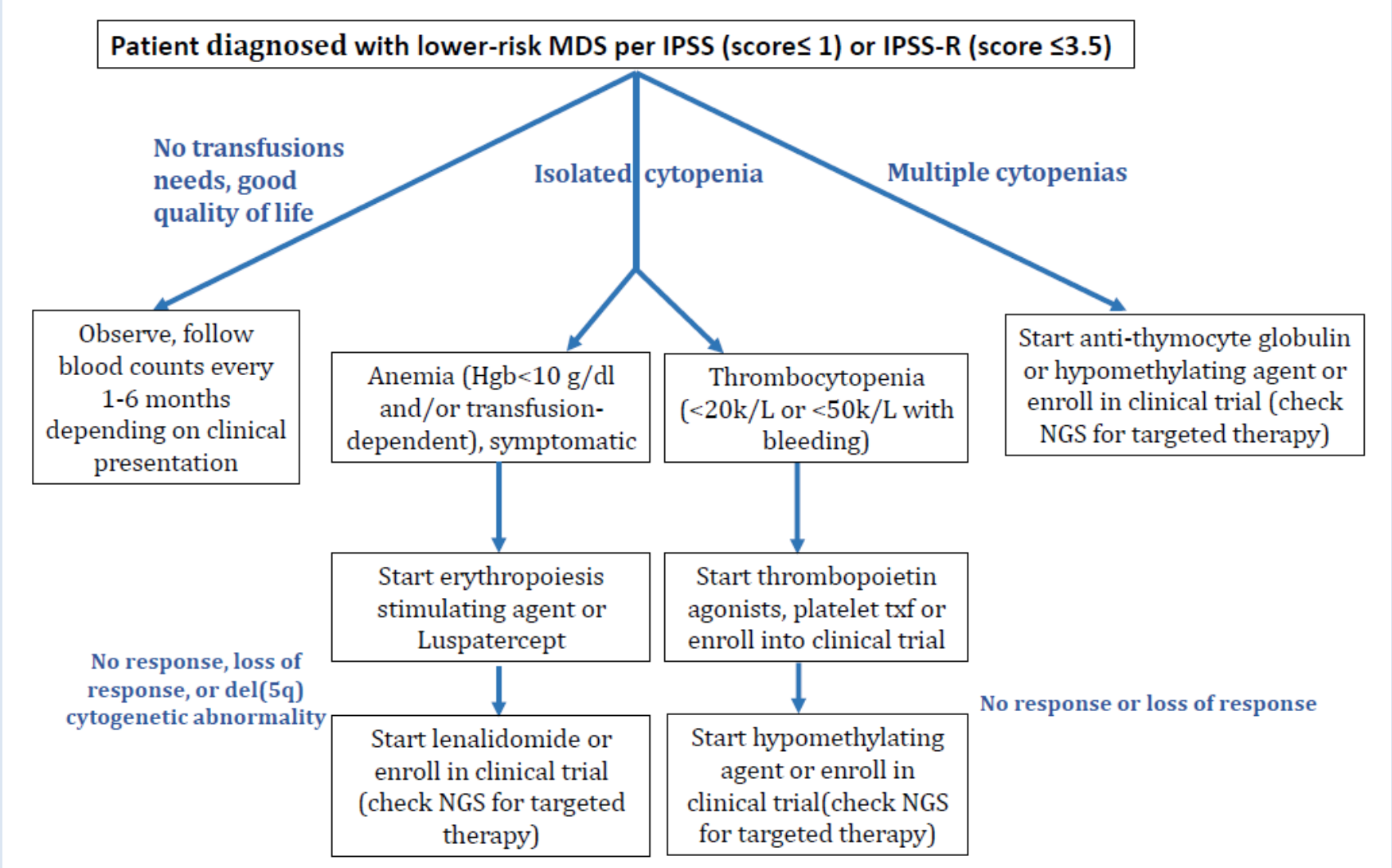


Figure adapted from Hanahan D, Weinberg RA. *Cell* 2011;144:646–74

Courtesy of Mikkael A Sekeres, MD, MS

BH3, bcl homology domain 3; CTLA4, cytotoxic T-lymphocyte-associated protein 4; DARTs, dual affinity retargeting agents; EGFR, epidermal growth factor receptor; HGF, hepatocyte growth factor; mAb, monoclonal antibody; PARP, poly adenosine diphosphate ribose polymerase; VEGF, vascular endothelial growth factor

# MDS | Treatment – Lower-risk



# Lower-risk MDS | Modifying MLD: HMA

---

- Regimens:
  - DAC 20 mg/m<sup>2</sup> IV D1-3 every 4 weeks
  - AZA 75 mg/m<sup>2</sup> IV/SC D1-3 every 4 weeks
- 113 pts with LR-MDS treated and evaluable for response
- Median duration of follow-up = 14 months (range: 2-30 months)
- Randomized follow-up study [NCT02269280](#)



# Lower-risk MDS | Modifying MLD: HMA

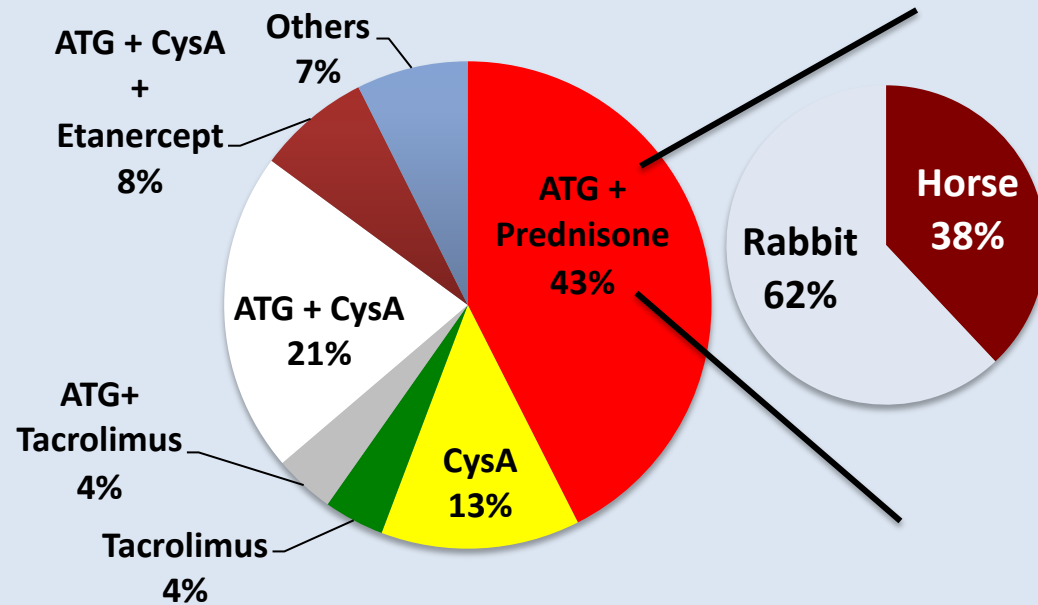
---

Response	N (%)
CR	33 (36)
mCR	8 (9)
HI	13 (14)
<b>ORR</b>	<b>54 (59)</b>
SD	31 (34)
PD	6 (7)

- Median time to best response: 2 months (range: 1-20)
- Median number of cycles received: 9 (range: 2-32)

# Lower-risk MDS | Modifying MLD: ATG

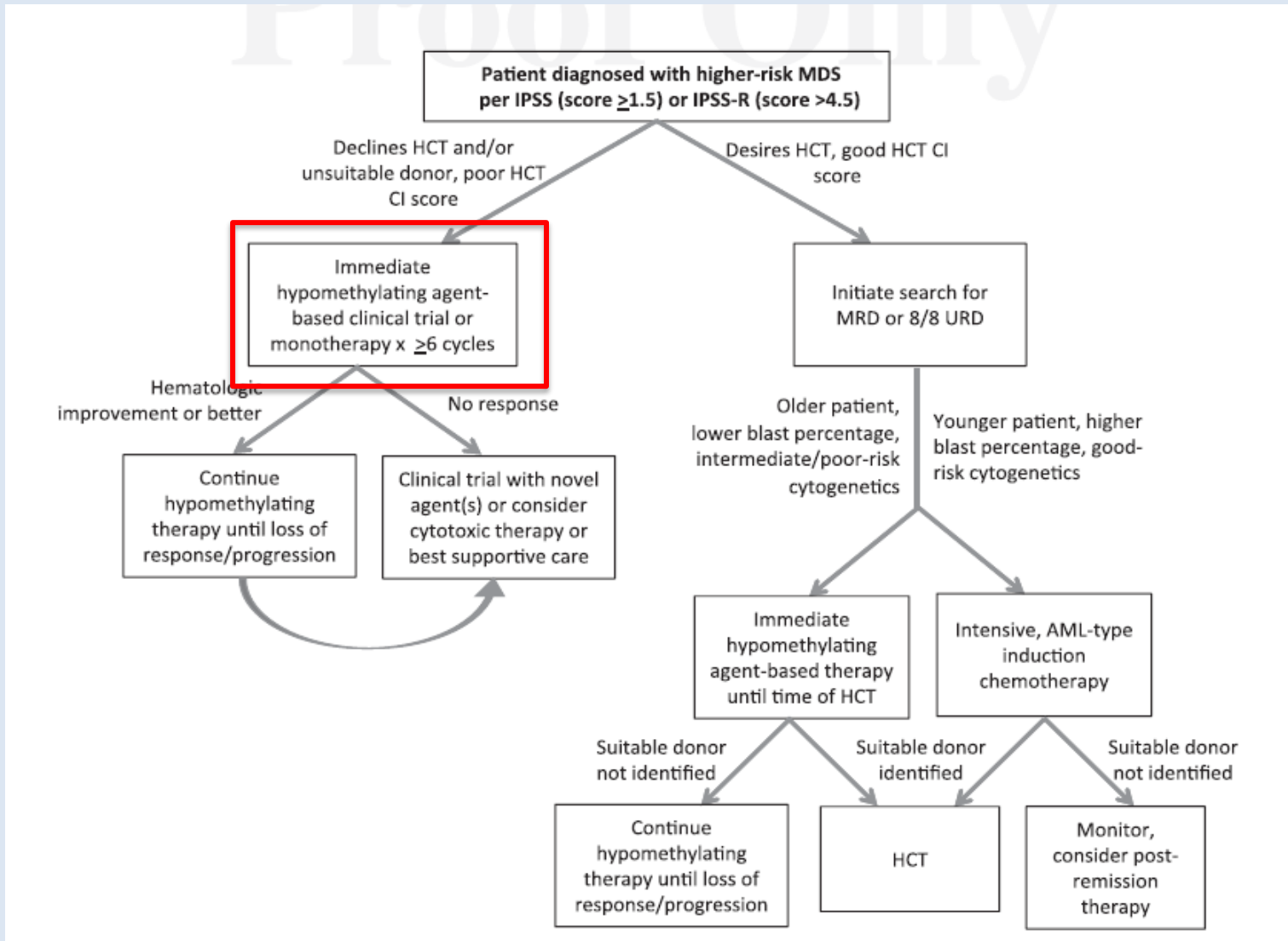
## 166 patients treated with ATG



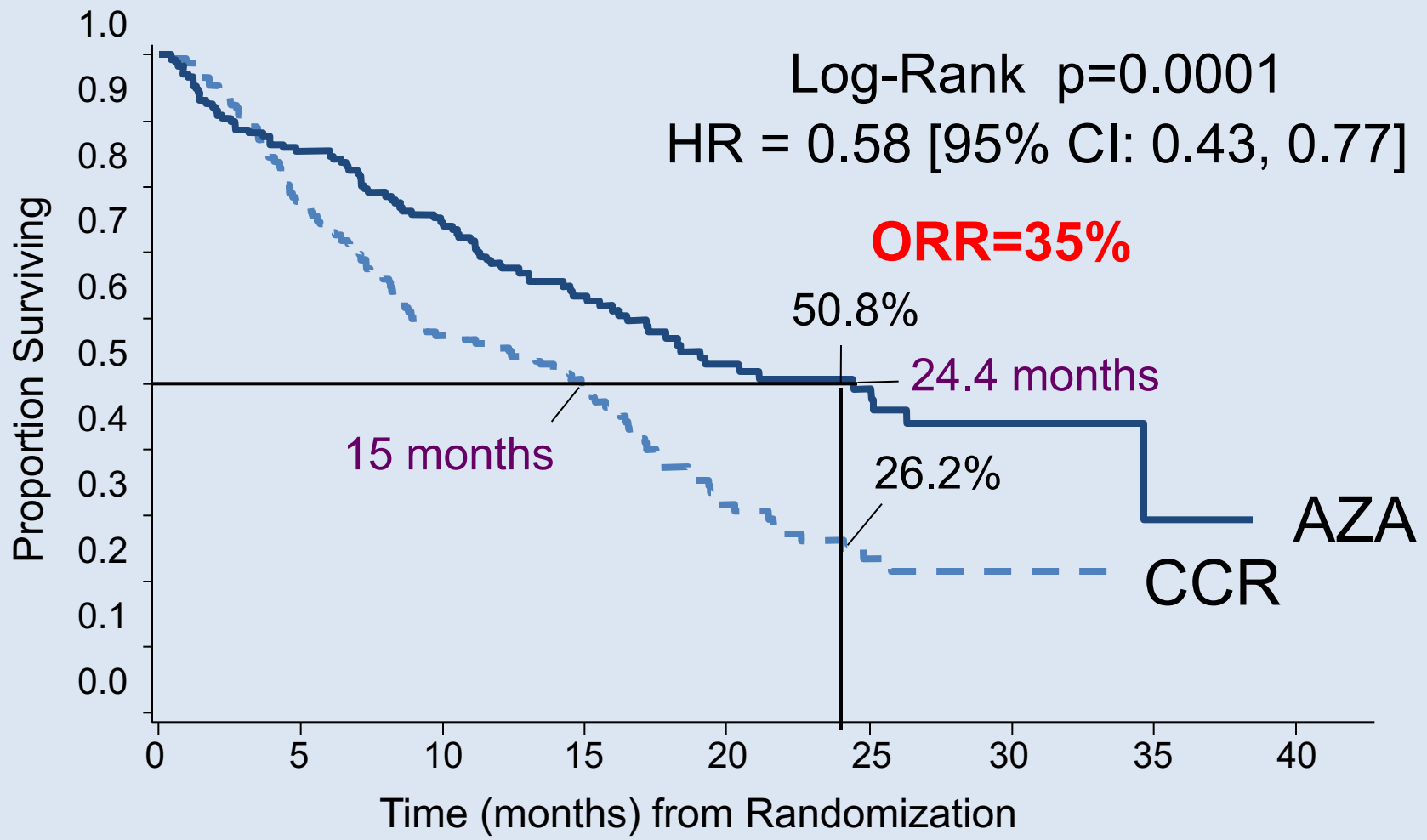
Response	%	95%CI
CR	11.2	6.5-18.4
PR	5.6	2.5-11.6
HI	32.0	24.1-41.0
SD	39.2	30.7-48.4
PD	12.0	7.1-19.3
<b>ORR</b>	<b>48.8</b>	<b>39.8-57.9</b>

## Type of IST used (N=217) and responses

# Higher-risk MDS | HMA and HCT



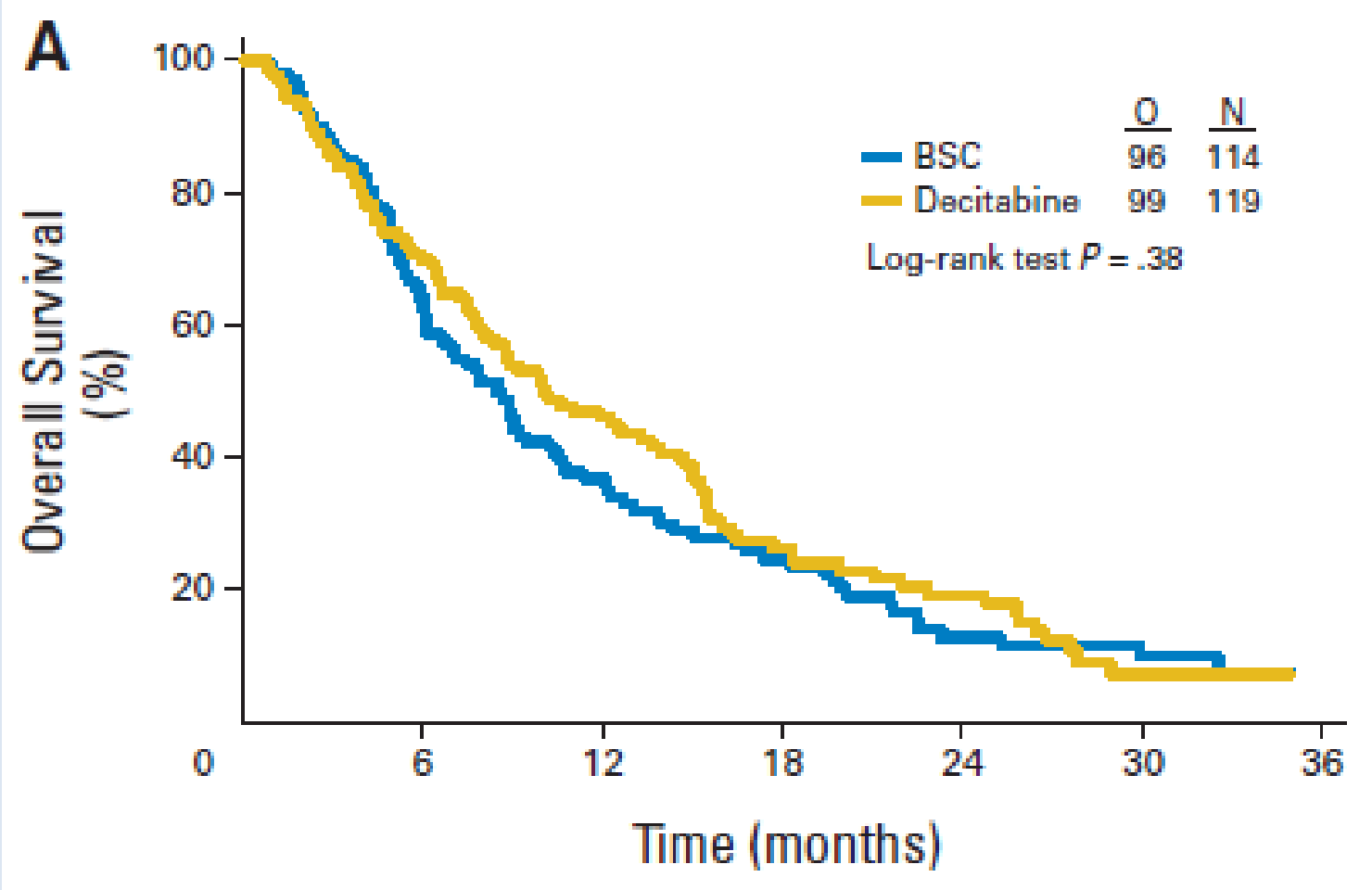
# Treating MDS | AZA



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

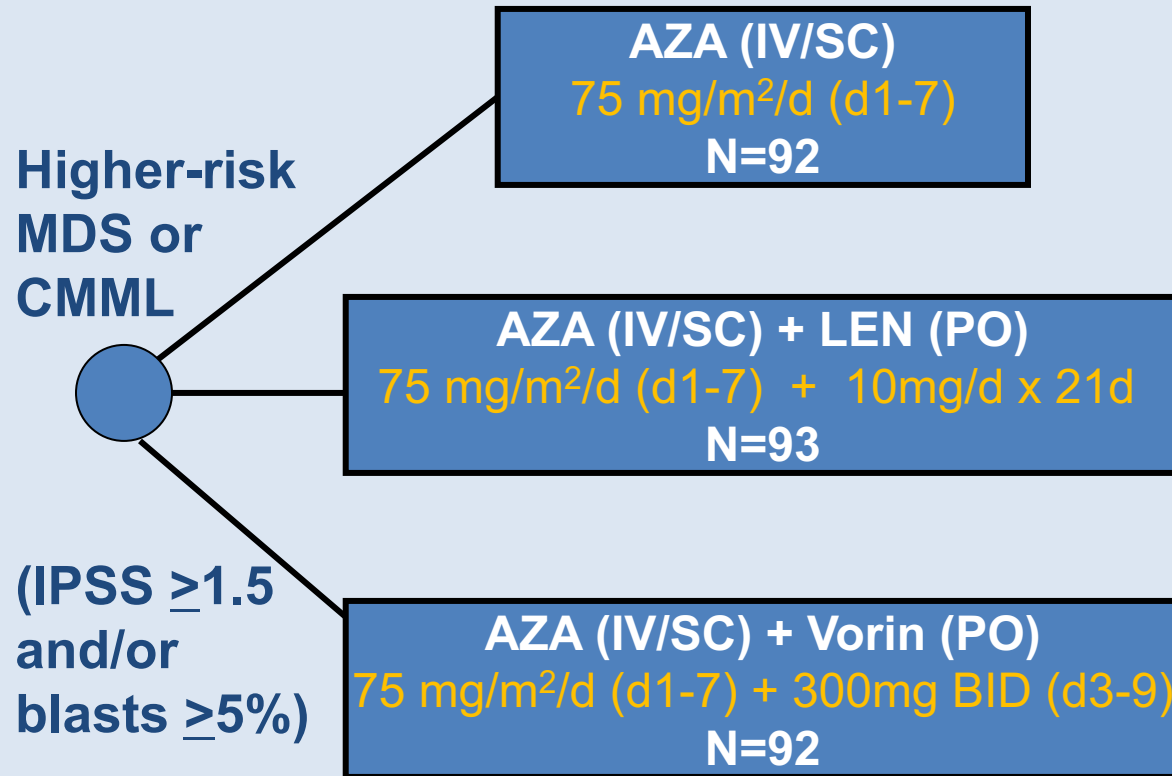
Courtesy of Mikkael A Sekeres, MD, MS

# Treating MDS | DAC



Median OS 10.1 vs. 8.5 months

# Higher-risk MDS | Combinations



Groups: SWOG, ECOG, Alliance, NCIC

Total Sample Size: 282/277

Primary Objective: 20% improvement of ORR (CR/PR/Hi) based on 2006 IWG Criteria

Secondary Objectives: OS, RFS, LFS

Power 81%, alpha 0.05 for each combo arm vs. AZA

06/2012 – 06/2014

# 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 (x10 <sup>3</sup> )	2 (0, 110)	1 (0, 336)	2 (0, 36)	2 (0, 336)
Baseline Platelet count (x10 <sup>3</sup> )	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)	<b>14 (.02)</b>	28
Rash (n)	3	<b>14 (&lt;.01)</b>	1 (1)	17
Off Tx due to Toxicity/Side Effect/Complication	8%	<b>20% (.05)</b>	<b>21% (.03)</b>	18%
Non-protocol defined dose modifications	24%	<b>43% (.002)</b>	<b>42% (.01)</b>	33%

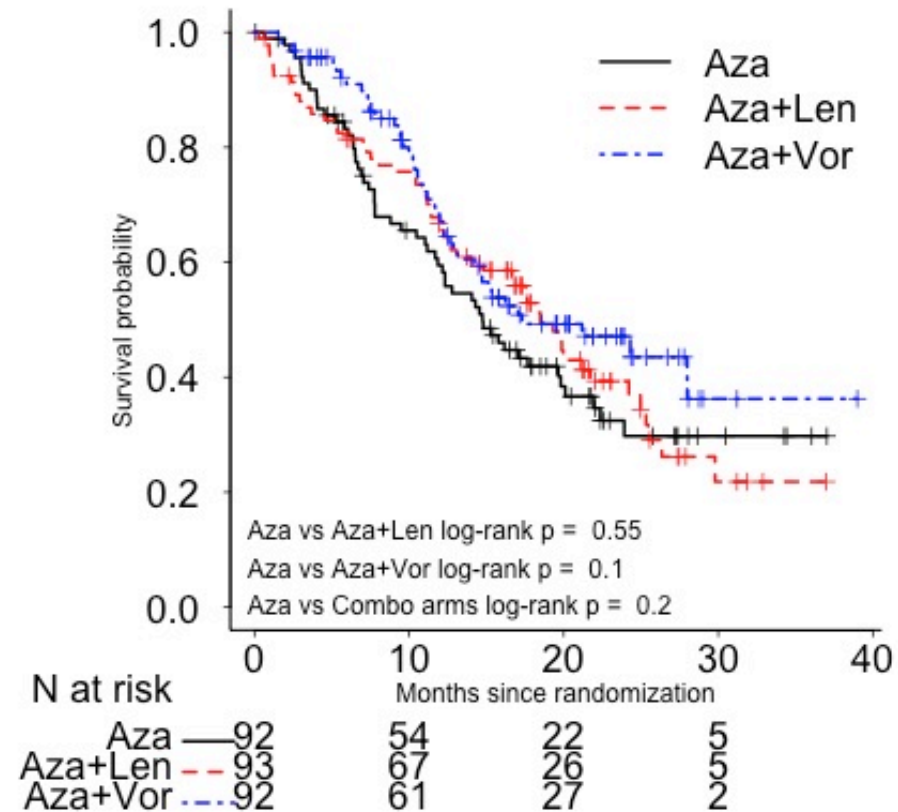


# Higher-risk MDS | Combinations

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 (%)	<b>38</b>	<b>49 (.16)</b>	<b>27 (.16)</b>	38%
CR/PR/Hi (%)	24/0/14	24/1/ <b>25</b>	17/1/9	22/1/16%
CMML ORR (%)	5 (28)	<b>13 (68) (.02)</b>	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