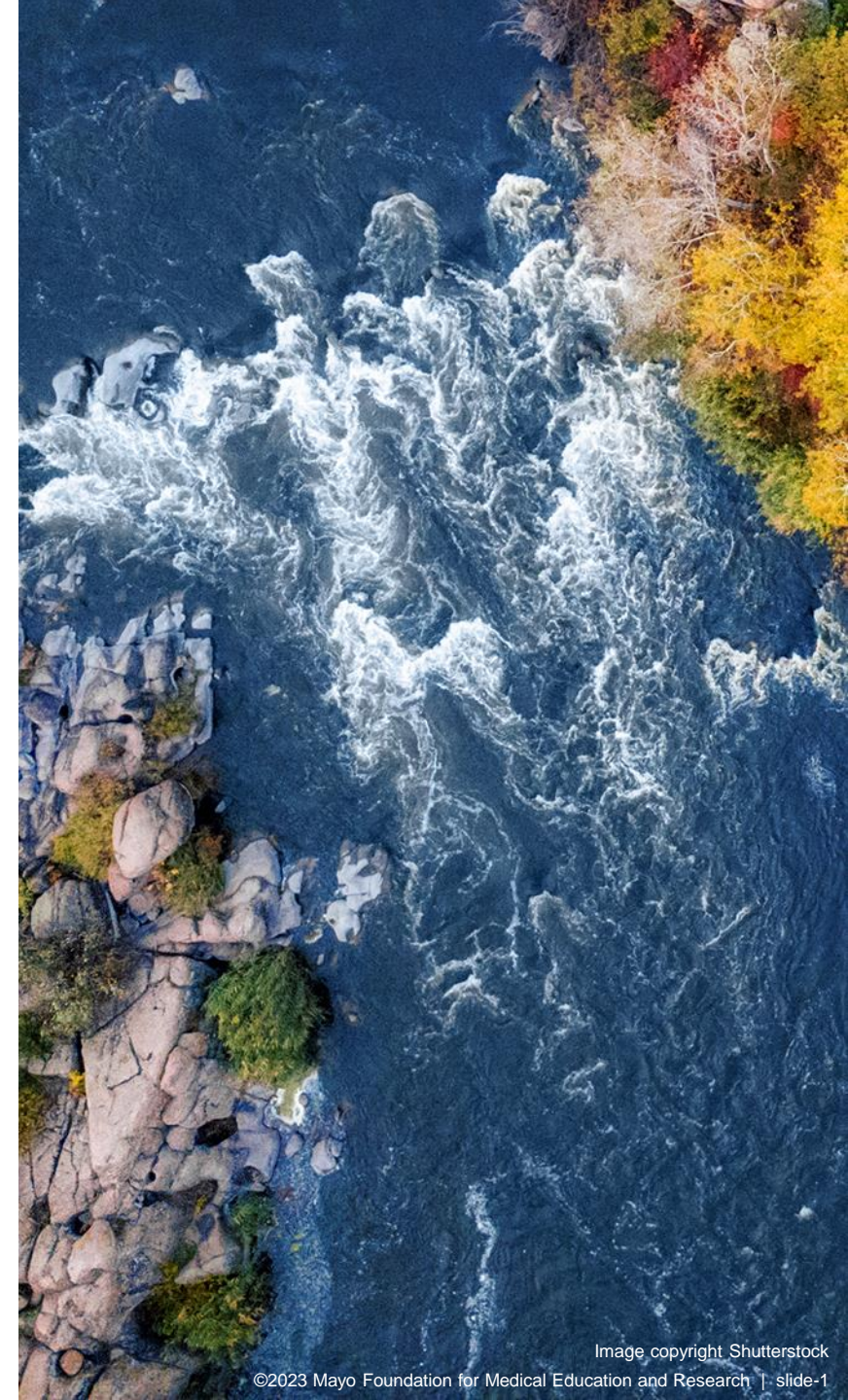


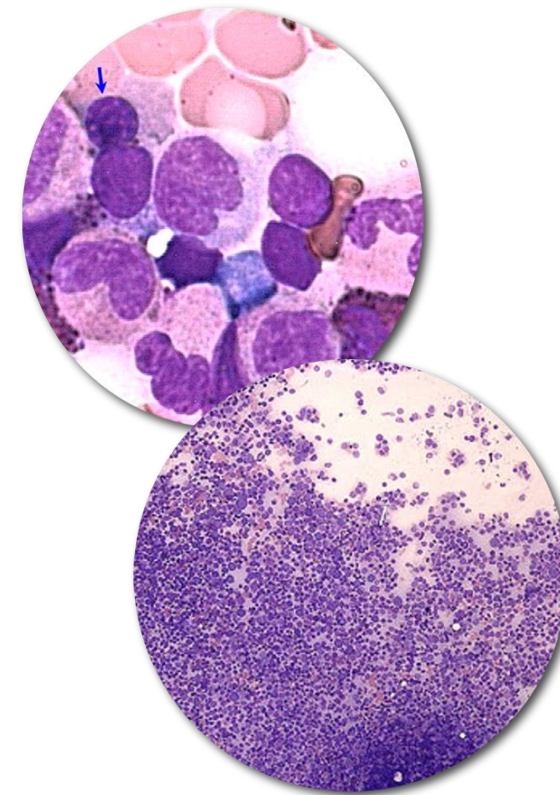
MYELODYSPLASTIC SYNDROME

- Hemant S. Murthy MD
- Division of Hematology/Oncology
- Associate Professor of Medicine
- Mayo Clinic Comprehensive Cancer Center- Florida
2023 FLASCO Fall Session
Orlando, Florida
October 21, 2023



MYELOYDYSPLASTIC SYNDROMES

- Heterogenous group of clonal malignant hematopoietic disorders sharing an ineffective production of one or more myeloid cell lines (accelerated apoptosis)
- Typically discrepancy between cellular marrow and peripheral cytopenias
- 10-30,000 cases per year in US
- Median age 76; M>F
- Ineffective hematopoiesis
 - Bleeding, anemia, infections
- Risk of transformation to AML
- Variable clinical course- need for prognostication



Bennett J, et al. The myelodysplastic syndromes. In: Abeloff MD, et al, editors. Clinical oncology. New York NY: Churchill Livingstone; 2004. pp. 2849-2881.

PATHOGENESIS OF MDS

- Arise from mutations in hematopoietic stem cells
- Genetic/ hematologic conditions
- Extrinsic factors
 - T cell dysregulation, stromal cell abnormalities, inflammasome/ inflammatory cytokines
- Familial MDS
 - Germline mutations
 - *RUNX1*, *ANKRD26*, *CEBPA*, *DDX41*, *ETV6*, *TERC*, *TERT*, *SRP72*, and *GATA2*
- Clonal hematopoiesis of indeterminate potential (CHIP)
 - mutations assoc. with MDS present, but no dysplasia, cytopenias

	Micro CHIP	CHIP	CCUS	Lower Risk MDS	Higher Risk MDS	sAML
VAF	<1%	~9%	~10-50%	~30-50%	~40-50%	~40-50%
Dysplasia	-	-	-	+	+	+
Cytopenias	-	-	+	+	+	+
BM Blast %	< 2%	< 2%	< 2%	< 2%	2-19%	20+%
Overall Risk	Background	Very Low	Low	Low	High	Very High
Treatments	N/A	Observation	Obs/BSC/GF	Obs/BSC/GF IMiD/IST	HMA/HCST	HMA/IC/HCST

PREDISPOSITIONS/ RISK FACTORS FOR MDS

- Age
 - Incidence increases with age
 - Uncommon Age <50
 - Most cases age 70's, 80's
- Sex (M>f)
- Prior chemotherapy/ radiotherapy
- Tobacco use
- Environmental
 - benzene
- Other heme disorders
 - Aplastic Anemia, PNH
- Genetic
 - Fanconi anemia
 - Shwachman-diamond syndrome
 - Diamond Blackfan anemia
 - Severe congenital neutropenia
 - Dyskeratosis congenita
- Familial MDS (rare)

MDS PRESENTATION

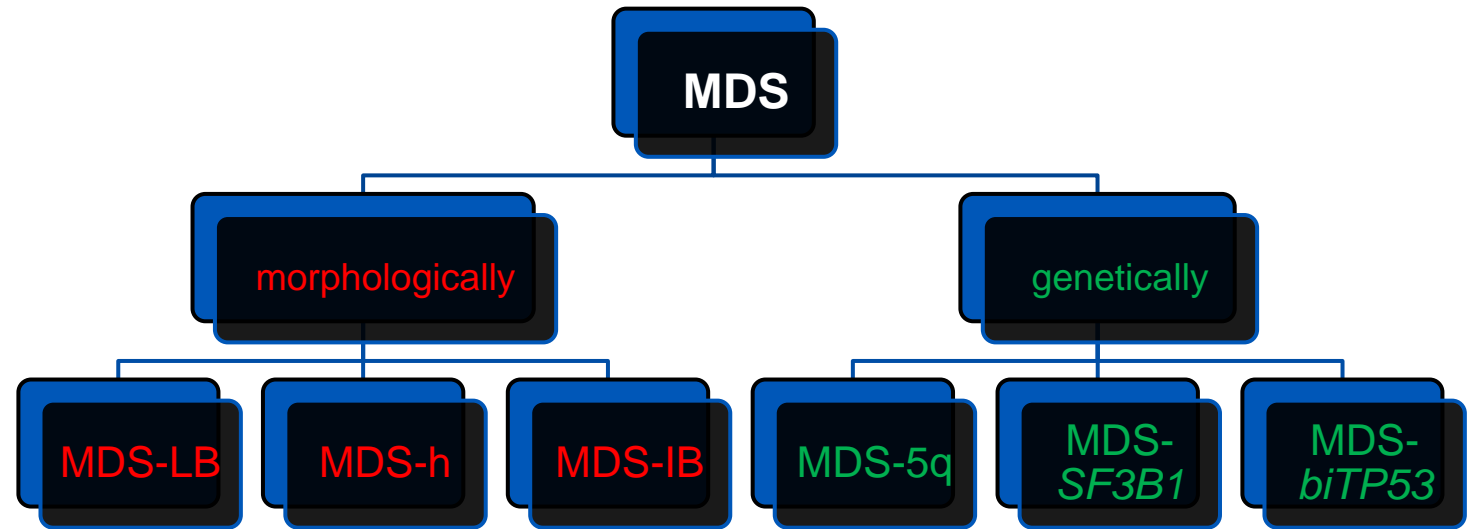
- Symptoms
 - Many present asymptomatic
 - Fatigue
 - Weakness
 - Frequent Infection
 - Bruising
- Physical signs
 - Ecchymosis
 - Pallor
- CBC
 - Anemia
 - Macrocytic (MCV >100)
 - Neutropenia
 - Pelger-huet cells
 - Thrombocytopenia

MDS WHO 2022

- Genetically defined
 - MDS-5q
 - MDS-*SF3B1m*
 - MDS-*biTP53*

- Morphologically defined

- MDS-LB low blasts
- MDS-h hypocellular
- MDS-IB increased blasts
 - IB-1: 2-4% PB blasts, 5-9% BM blasts
 - IB-2: 5-19% PB blasts, 10-19% blasts or Auer rods
 - Fibrosis: 2-19% PB blasts, 5-19% BM blasts



RISK STRATIFICATION

Factors

- CBC
- Blasts%
- Cytogenetics
- NGS
- models

Values

- Single lineage vs multiple
- $<5\%$ vs $\geq 5\%$
- Complex/-5/-7/-17
- TP53, MLL-PTD
- IPSS-molecular

R-IPSS OUTCOMES

IPSS-R Prognostic Score Values*

Prognostic Variable	0	0.5	1	1.5	2	3	4
Cytogenetics	Very Good		Good		Intermediate	Poor	Very Poor
BM Blast %	<=2		>2-<5%		5-10%	>10%	
Hemoglobin	=>10		8-<10	<8			
Platelets	=>100	50-<100	<50				
ANC	=>0.8	<0.8					

IPSS-R Cytogenetic Risk Groups*,**

Cytogenetic Prognostic Subgroups	Cytogenetic Abnormalities
Very good	-Y, del(11q)
Good	Normal, del(5q), del(12p), del(20q), double including del(5q)
Intermediate	del(7q), +8, +19, i(17q), any other single or double independent clones
Poor	-7, inv(3)t(3q)/del(3q), double including -7/del(7q), Complex: 3 abnormalities
Very Poor	Complex: >3 abnormalities

IPSS-R: Prognostic Risk Category Clinical Outcomes*

	No. Pts	Very Low	Low	Intermediate	High	Very High
Risk Score		≤1.5	>1.5-3	>3-4.5	>4.5-6	>6
Patients (%)	7012	19%	38%	20%	13%	10%
Survival***		8.8	5.3	3.0	1.6	0.8
AML/25%***, ^		NR	10.8	3.2	1.4	0.7

MUTATIONS TIPS

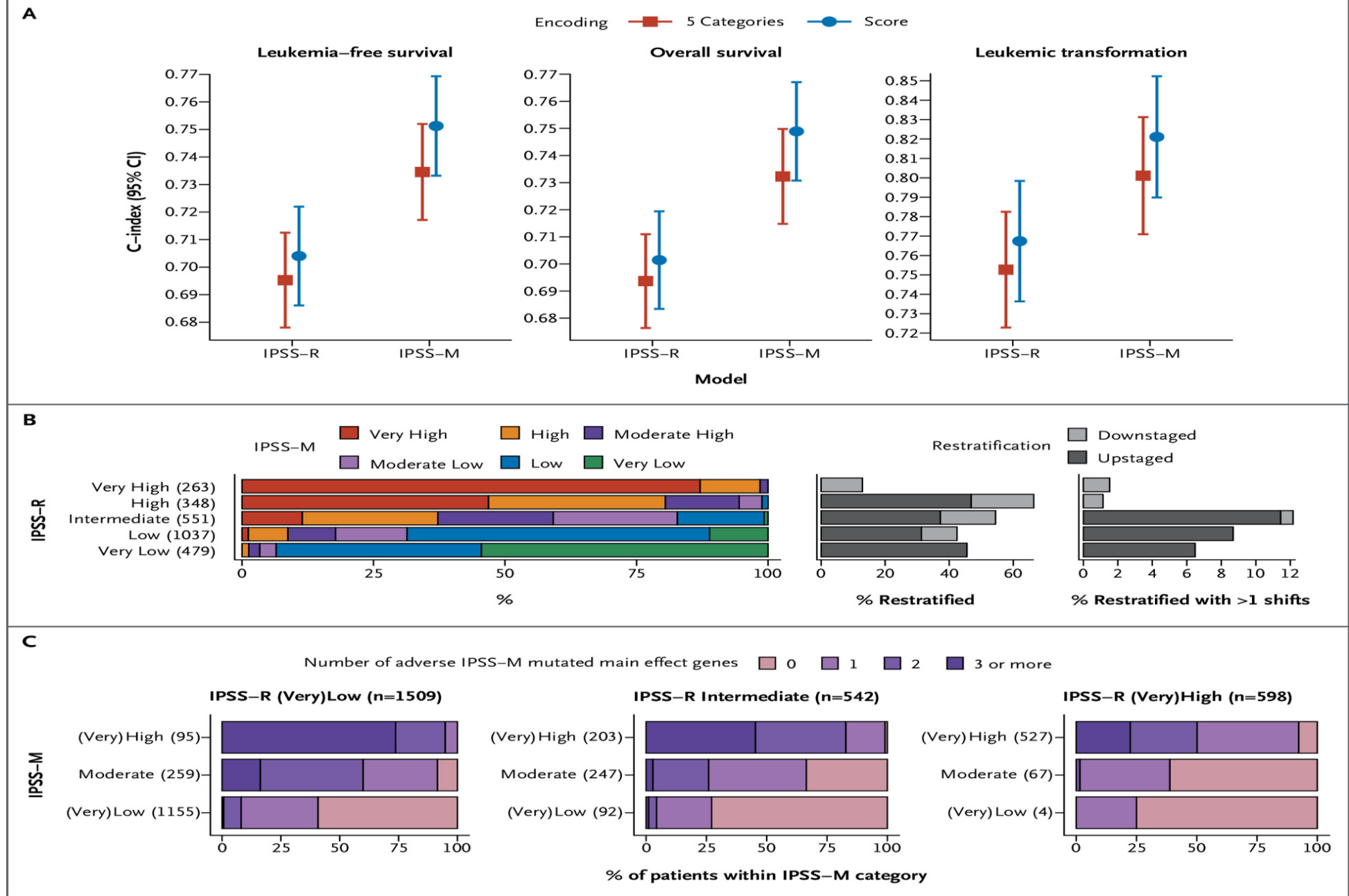
Gene	Correlation	Rx
<i>SF3B1</i>	Ring sideroblast	luspatercept
<i>IDH1</i>	Cbc ~	Ivosidenib, HMA+VEN
<i>IDH2</i>	Cbc ~	Enasidenib, HMA+VEN
<i>FLT3</i>	AML transformation	Gilteritinib
<i>NPM1</i>	AML-defining	CTX vs HMA+VEN
<i>RUNX1</i>	AML transformation	HMA+VEN
<i>DDX41</i>	Germline ?, cbc ~	HMA+VEN, LEN
<i>STAT3</i>	LGL	ISA
<i>PIGA1</i>	PNH	Complement inhibitor
<i>UBA1</i>	VEXAS	HMA, JAKi
<i>TP53</i>	T-MN	? PO DAC

@AlkaliDr



MOL-IPSS

- Clinical-molecular prognostic model (IPSS-Molecular [IPSS-M]), pretreatment diagnostic or peridiagnostic samples from 2957 patients with MDS were profiled for mutations in 152 genes
- *TP53*^{multihit}, *FLT3* mutations, and *MLL*^{PTD} top genetic predictors of adverse outcomes.
- *SF3B1* mutations were associated with favorable outcome
- Compared with the IPSS-R, the IPSS-M improved prognostic discrimination across all clinical end points and restratified 46% of patients



Low-grade MDS

LUSPATERCEPT

SF3B1m,
MDS-RS,
EPO >500

ESA

EPO <500

TPO mimetic

Low PLT

LEN

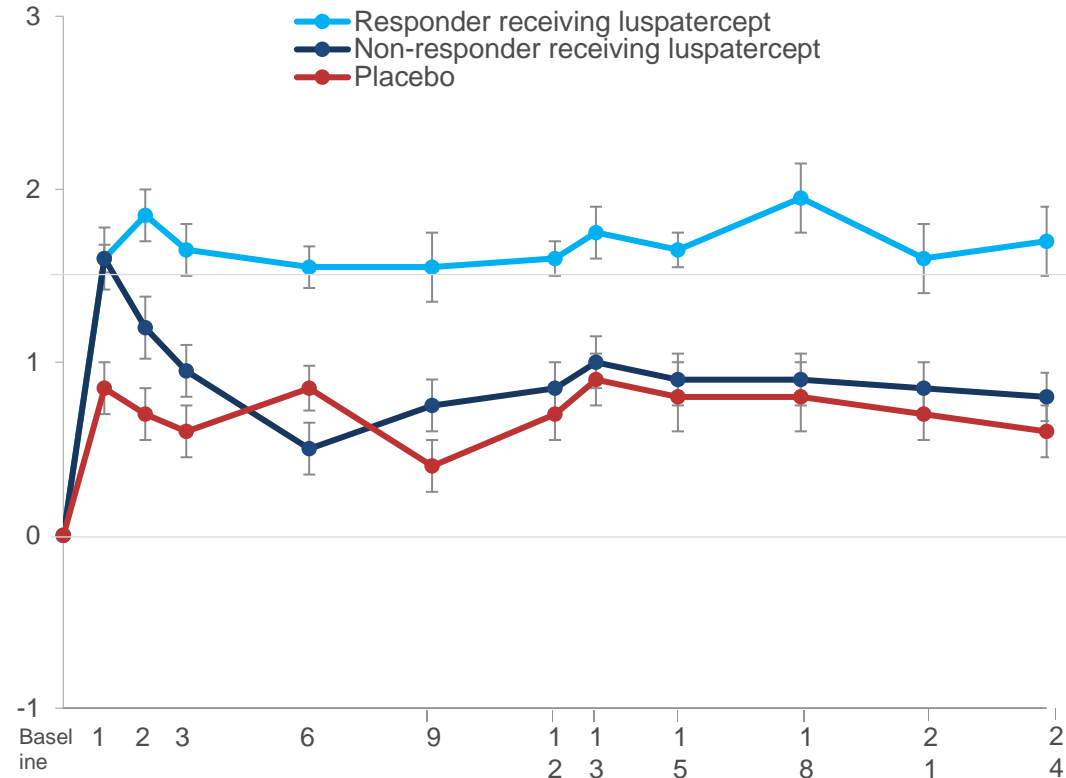
Del 5q

HMA
3-7 days

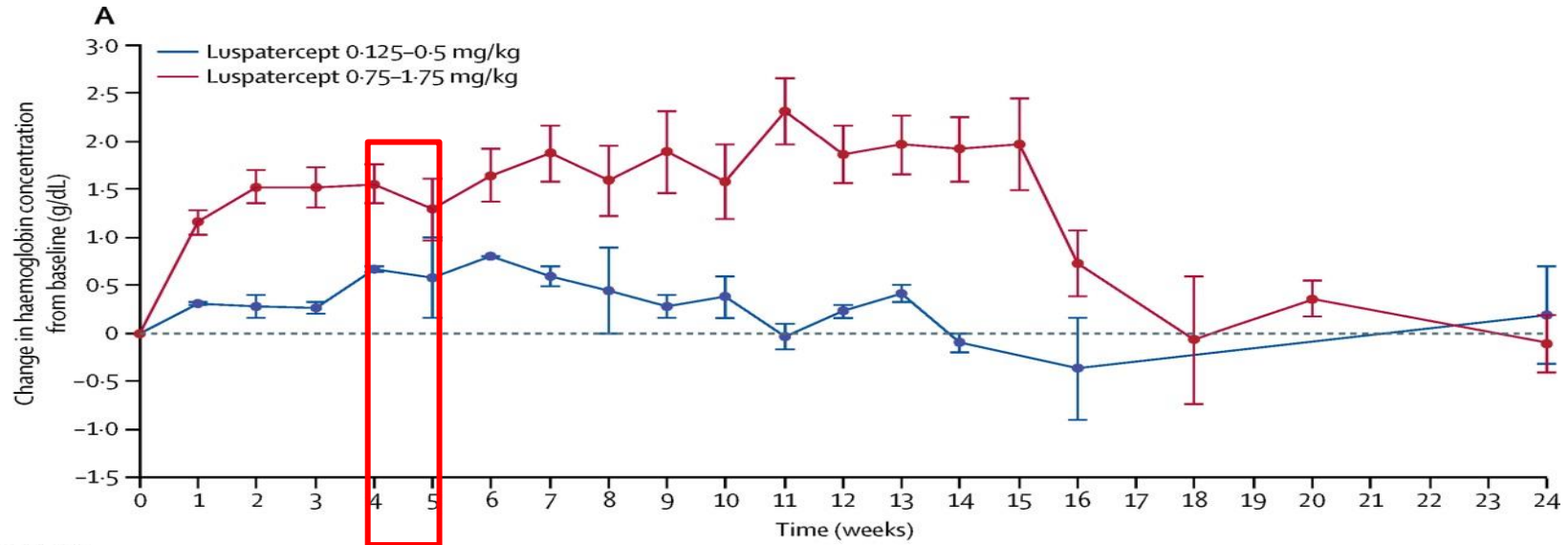
Failure to
prior lines, >1
cytopenia

LUSPATERCEPT

- TGFb ligand and Smad Fusion protein inhibitor
- In lower-risk, Ring sideroblast -positive MDS:
 - Luspatercept better in reducing rbc transfusion
 - More transfusion independence
 - Better Hb increase than placebo
 - Erythroid responses were durable, with approximately 40% of patients achieving RBC-TI sustained at 12 months of treatment
 - Well tolerated in this patient population
- FDA approved for thalassemia, MDS

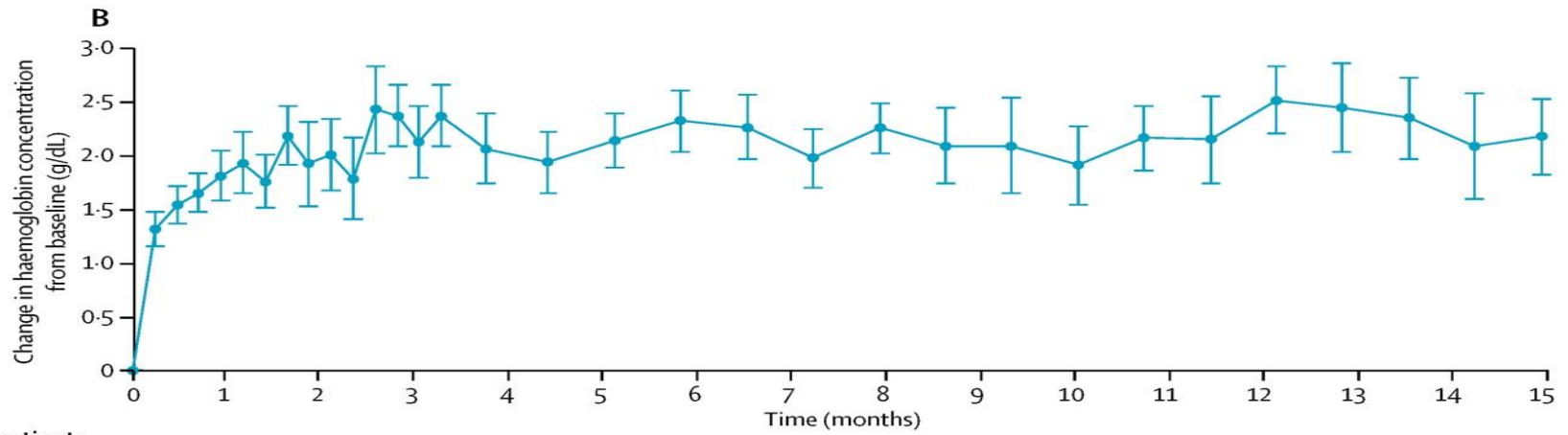


Courtesy Alan List



Number of patients

Luspatercept 0.125-0.5 mg/kg	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2
Luspatercept 0.75-1.75 mg/kg	17	16	16	16	16	17	14	17	16	13	16	13	16	13	12	8	4	4	4	3	3	3	3	3	3	3	3	3	3	3	3



Number of patients

Luspatercept 1.0-1.75 mg/kg	13	13	10	13	10	10	13	13	10	11	9	13	13	13	12	12	11	11	10	11	11	10	9	9	9	10	8
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MDS MEDALIST

- MDS-RS*
- >18 years
- IPSS-R:
 - VL, L, Int-1
- **TD** (2/8wks)
- ESA
 - Failure* or
 - not likely to respond

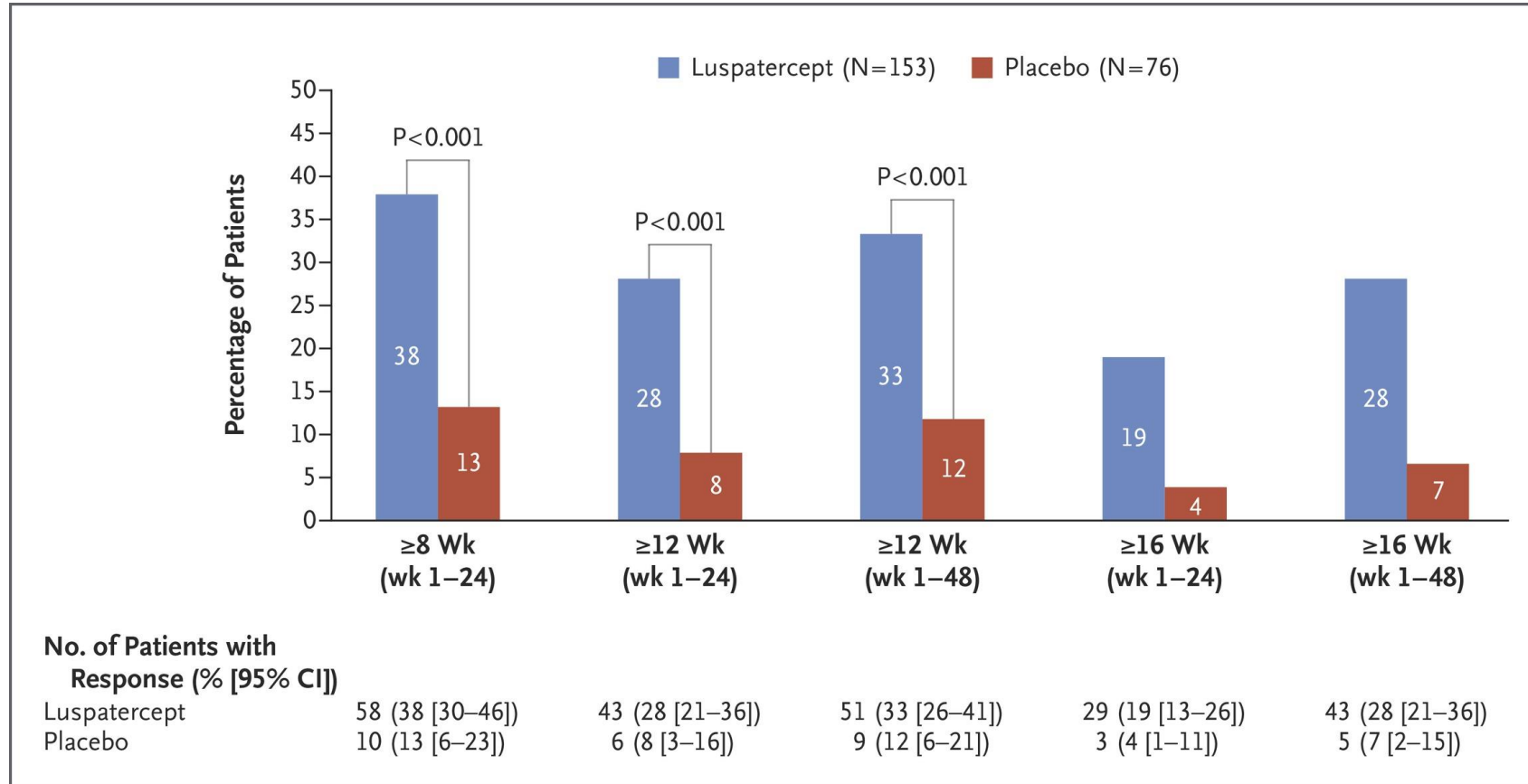
R



Luspatercept SC 1mg/Kg q 21 days
N=153

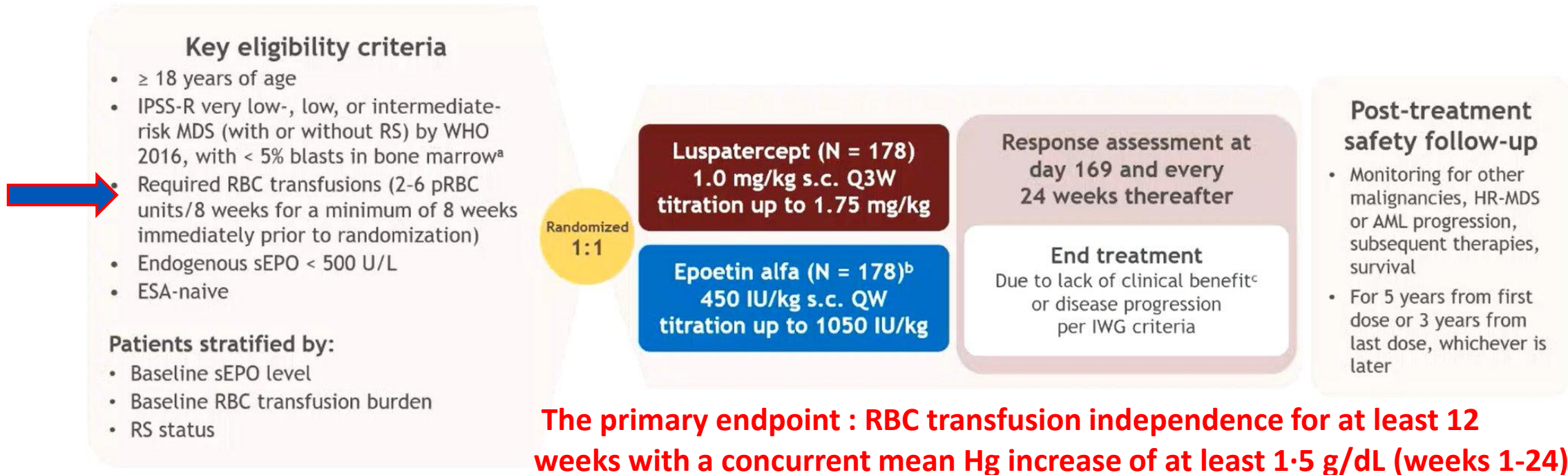
Placebo SC q 21 days
N= 76

MDS MEDALIST



The COMMANDS study

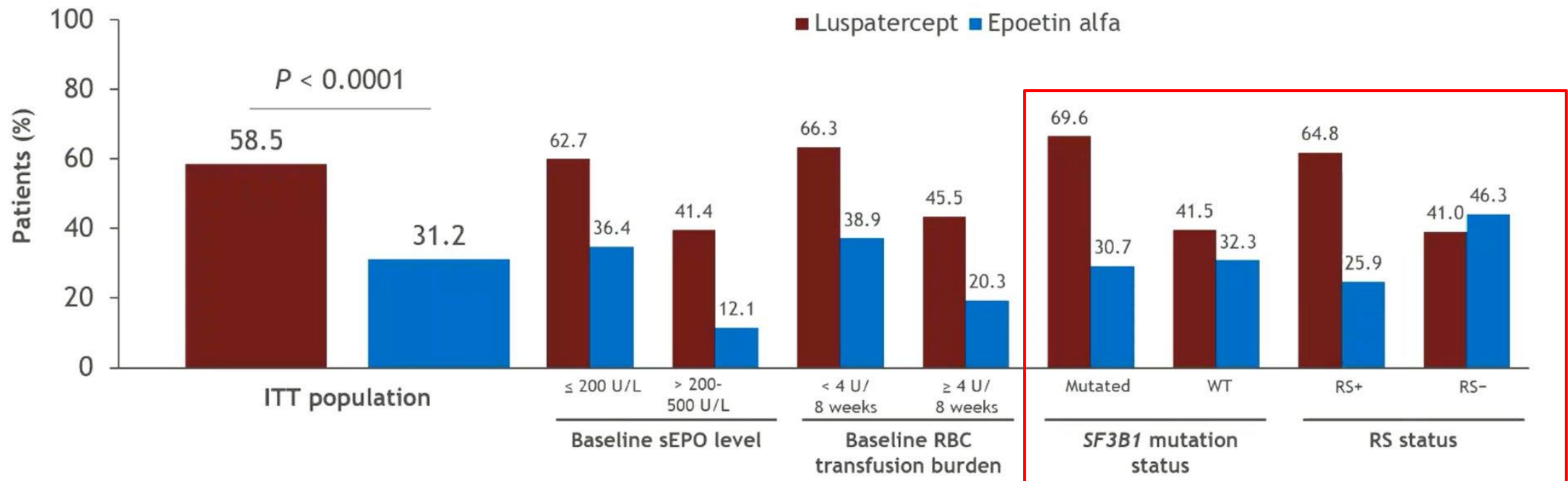
The COMMANDS study (NCT03682536) is a phase 3, global, open-label, randomized trial comparing the efficacy and safety of luspatercept versus epoetin alfa for the treatment of anemia due to IPSS-R LR-MDS in ESA-naive patients who require RBC transfusions



^aMDS with del(5q) were excluded. ^b2 patients randomized to the epoetin alfa arm withdrew consent prior to receiving their first dose; ^cClinical benefit defined as transfusion reduction of ≥ 2 pRBC units/8 weeks versus baseline; AML, acute myeloid leukemia; ESA, erythropoiesis-stimulating agent; IPSS-R, Revised International Prognostic Scoring System; IWG, International Working Group; LR-MDS, lower-risk MDS; MDS, myelodysplastic syndromes; pRBC, packed RBC; QW, once weekly; Q3W, every 3 weeks; RBC, red blood cell; RS, ring sideroblasts; s.c., subcutaneously; sEPO, serum erythropoietin; WHO, World Health Organization.

Primary endpoint: luspatercept superior to epoetin alfa

- Of 301 pts included in the efficacy analysis, 86 (58.5%) patients receiving luspatercept and 48 (31.2%) epoetin alfa achieved the primary endpoint
 - Achievement of the primary endpoint favored luspatercept or was similar to epoetin alfa for all subgroups analyzed



This prespecified interim analysis included 301 patients who had either completed 24 weeks of treatment or discontinued prior to completing 24 weeks of treatment.

Della Porta MG, et al. EHA 2023 [Abstract #S102]

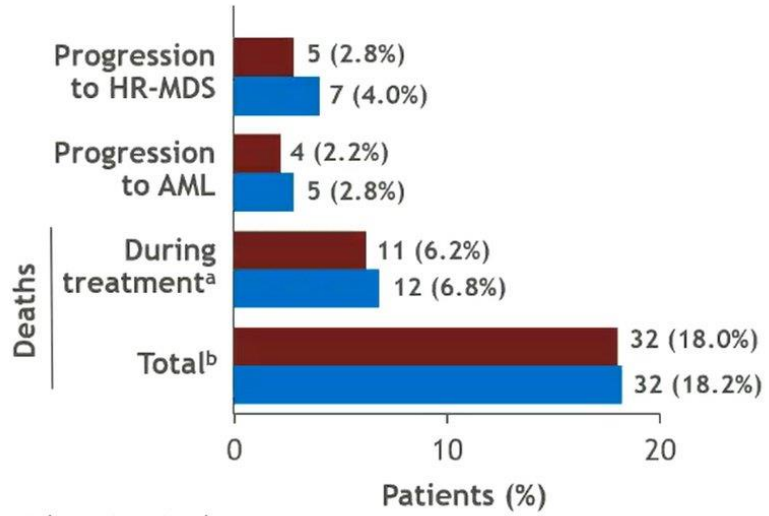
Safety profile of luspatercept manageable and comparable to previous studies

- Exposure to luspatercept was ~2 times longer compared with epoetin alfa, providing a longer reporting period for AEs

Patients, n (%)	Luspatercept (N = 178)		Epoetin alfa (N = 176)	
	Any grade	Grade 3/4	Any grade	Grade 3/4
Heme-related TEAEs				
Anemia	17 (9.6)	13 (7.3)	17 (9.7)	12 (6.8)
Thrombocytopenia	11 (6.2)	7 (3.9)	3 (1.7)	1 (0.6)
Neutropenia	9 (5.1)	7 (3.9)	13 (7.4)	10 (5.7)
Leukocytopenia	2 (1.1)	0	3 (1.7)	0
TEAEs of interest				
Fatigue	26 (14.6)	1 (0.6)	12 (6.8)	1 (0.6)
Diarrhea	26 (14.6)	2 (1.1)	20 (11.4)	1 (0.6)
Peripheral edema	23 (12.9)	0	12 (6.8)	0
Asthenia	22 (12.4)	0	25 (14.2)	1 (0.6)
Nausea	21 (11.8)	0	13 (7.4)	0
Dyspnea	21 (11.8)	7 (3.9)	13 (7.4)	2 (1.1)
TEE	8 (4.5)	5 (2.8)	5 (2.8)	1 (0.6)

TEAEs of any grade
 164 (92.1%) luspatercept
 150 (85.2%) epoetin alfa

Treatment duration, median (range), weeks
 41.6 (0-165) luspatercept
 27.0 (0-171) epoetin alfa



Safety data are not exposure-adjusted.
^a11 deaths in each arm led to treatment discontinuation. One additional death occurred in the epoetin alfa arm after treatment discontinuation due to an AE; the death occurred during the 42-day safety follow up, which was considered a death during treatment but not counted as a death leading to treatment discontinuation. ^bDeaths during treatment period and post-treatment period. TEE, thromboembolic event.
 Della Porta MG, et al. EHA 2023 [Abstract #S102]

LUSPATERCEPT

- SC Q21 days
- Adjust dose based on response
- Hold if Hg >11.5
- FDA approved
- Works more if
 - Frontline
 - RS+
 - *SF3B1*+
 - Thrombocytosis+
 - LT

IMELSTAT

- First-in-class telomerase inhibitor
 - Targets cells with high telomerase activity and human telomerase reverse transcriptase expression, both of which have been reported in MDS
- IMERGE- global phase 2 clinical trial on LR-MDS resistant to ESA
 - Treatment with imetelstat achieved >1 year transfusion independence in 29% of RBC TD, ESA-R/R LR-MDS patients who were non-del(5q) and lenalidomide/HMA-naïve.
 - 24-week TI predicted a likelihood to achieve TI >1 year
-

IMERGE PHASE 3: STUDY DESIGN

- International, double-blind, randomized phase III trial

Patients with low-risk or intermediate 1-risk MDS
(IPSS-R); R/R to ESA or EPO >500 mU/mL
(ESA ineligible); RBC transfusion dependent
(≥4 U/8 wk over 16 wk prestudy); non-del(5q);
no prior lenalidomide or HMAs
(N = 178)

2:1

Imetelstat

7.5 mg/kg IV Q4W
(n = 118)

Placebo

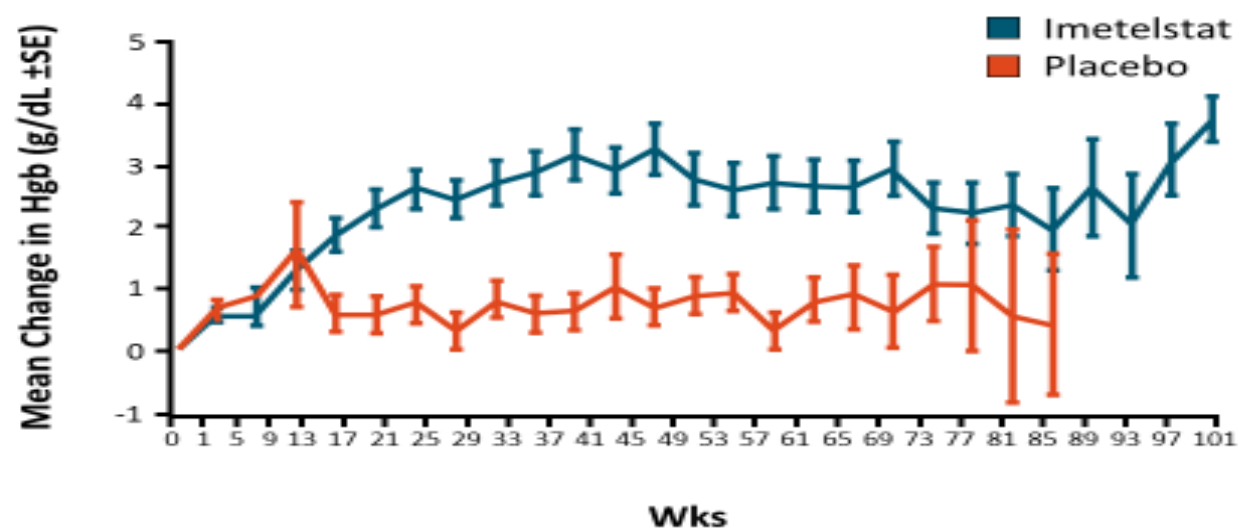
(n = 60)

- **Primary endpoint:** 8-wk RBC-TI
- **Key secondary endpoints:** 24-wk RBC-TI, TI duration, HI-E, safety
- **Key exploratory endpoints:** changes in VAF, PRO (FACIT-Fatigue)

IMERGE3

Measure, n (%)	Imetelstat N=118	Placebo N=60	<i>p</i> ^a
8-wk TI	47 (39.8)	9 (15.0)	<0.001
TI duration, median wks (95% CI) ^b	51.6 (26.9– 83.9)	13.3 (8.0– 24.9)	<0.001 ^d
24-wk TI	33 (28.0)	2 (3.3)	<0.001
HI-E ^c	50 (42.4)	8 (13.3)	<0.001

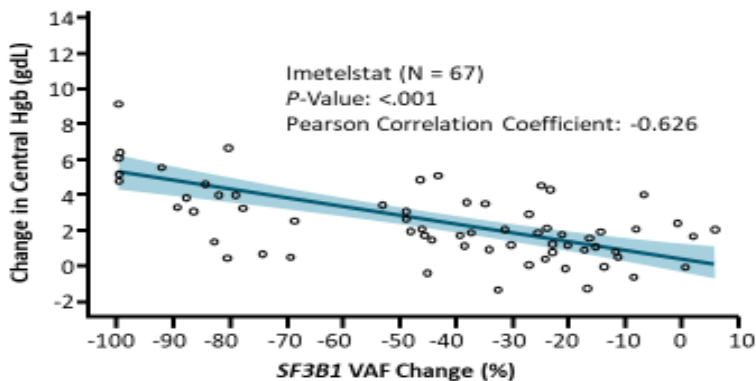
Entire Cohort: Mean Change in Hb



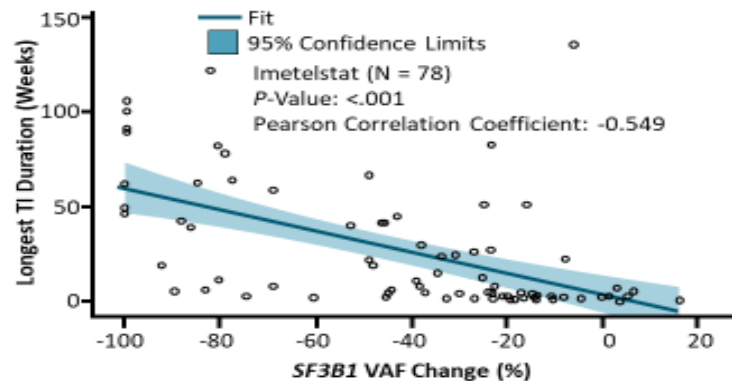
- No new safety signals were identified.
- The most common Grade 3/4 AEs were thrombocytopenia and neutropenia, with similar rates of Grade ≥ 3 bleeding and infections on imetelstat and placebo.
- In pts treated with imetelstat, cytopenias were manageable, of short duration, and >80% were reversible to Grade ≤ 2 within 4 wks.

IMPACT OF TREATMENT ON REDUCTION OF VAF

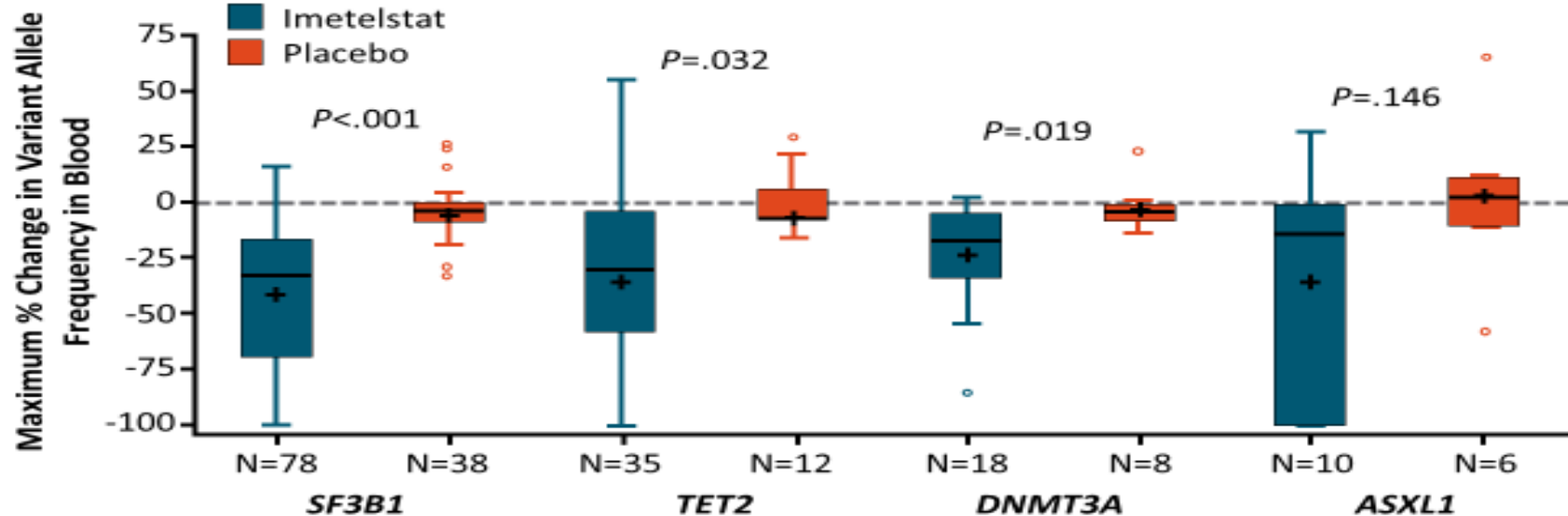
Maximum Increase in Hb vs Maximum Reduction in SF3B1 VAF



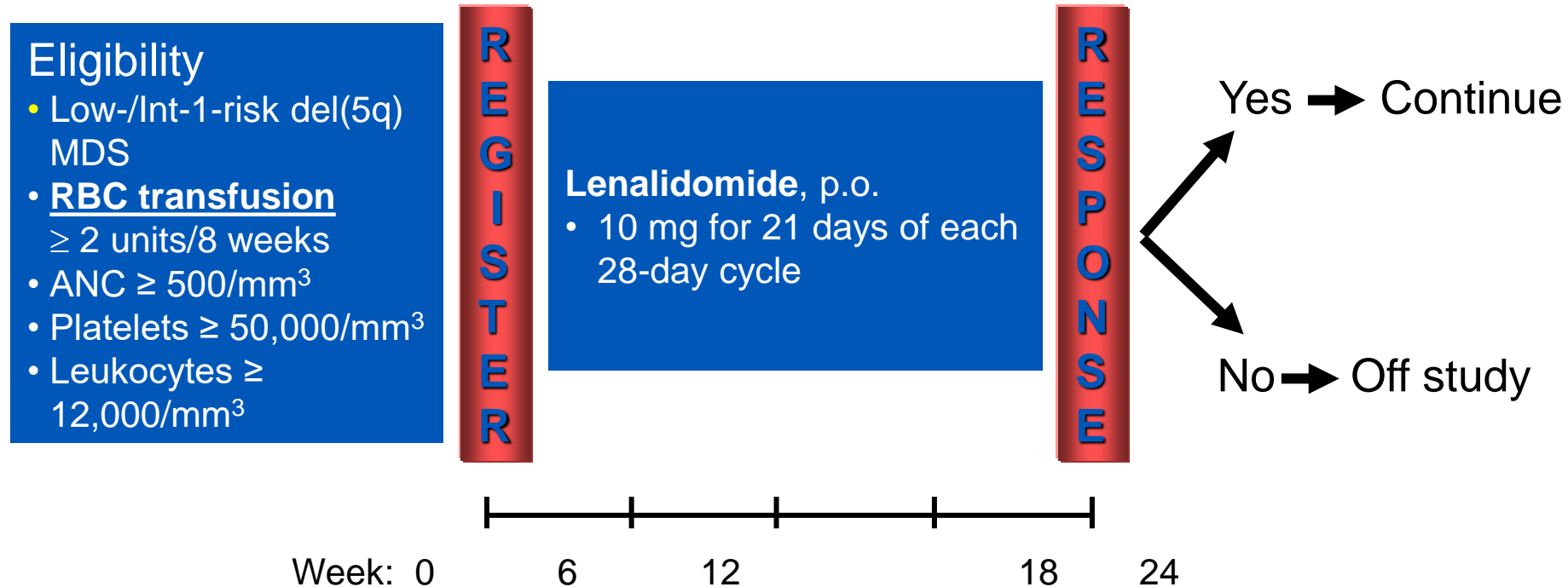
Longest RBC-TI Duration vs Maximum Reduction in SF3B1 VAF



TET2, DNMT3A, or ASXL21 VAF reductions correlated with longer RBC-TI duration



MDS-003: STUDY DESIGN



Primary objective: erythroid response* (RBC-TI)

Secondary objectives: RBC-TI duration, cytogenetic response, tolerability

*As defined by the IWG 2000 criteria: Cheson BD, et al. Blood 2000;96:3671–4. ANC, absolute neutrophil count; p.o., orally; RBC, red blood cell; TI, transfusion independence.

~~MDS-003: ERYTHROID RESPONSE AT 24 WEEKS~~

Variable

N = 148 (ITT)

Erythroid response, n (%)

**RBC-TI (≥ 56 consecutive days
without transfusion and Hb levels
rose by 1 g/dL)**

99 (67)

$\geq 50\%$ decrease in transfusions

13 (9)

**Total (RBC-TI + $\geq 50\%$ decrease in
transfusions)**

112 (76)

Median Hb increase, g/dL (range)

5.4 (1.1–11.4)

**Median time to response, weeks
(range)**

4.6 (1–49)

- RBC-TI achieved by 67% of lenalidomide-treated del(5q) patients**

LEN IN MDS 5Q TP53+

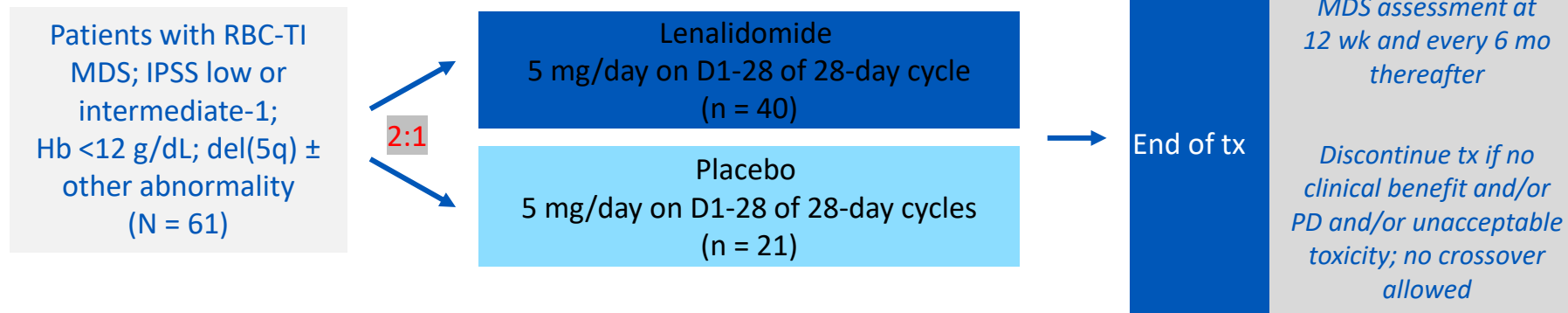
Table 1. Patient characteristics of all patients analyzed for TP53 mutations

	All	TP53 ^{WT}	TP53 ^{Mut}	
Number of patients	67	59	8	
Gender, n (%)				
Male	13 (18)	11 (19)	2 (25)	NS
Female	54 (82)	48 (81)	6 (75)	
Median age (range)	70 (40–87)	70 (40–84)	74 (55–87)	NS
WHO subtype, n (%)				
MDS-U	2 (3)	2 (3)	0	NS
RCUD (RA, RT)	13 (19)	12 (20)	1	
RCMD	43 (64)	37 (63)	6 (75)	
RCMD-RS	9 (13)	8 (14)	1	
FAB subtypes, n (%)				
RA	55 (77)	47 (80)	7 (88)	NS
RARS	9 (13)	8 (13)	1 (12)	
Missing	4 (6)	4 (7)	0	
IPSS risk groups, n (%)				
Low	42 (63)	35 (59)	7 (87)	NS
Intermediate-1	21 (31)	20 (34)	1 (13)	
Missing	4 (6)	4 (7)	0	
IPSS-R risk groups, n (%)				
Very low	8 (10)	6 (10)	2 (25)	NS
Low	49 (73)	44 (75)	5 (62)	
Intermediate	6 (9)	5 (8)	1 (13)	
Missing	4 (6)	4 (7)	0	
Mean blood counts (s.d.)				
Hb (g/dl)	8.9 (1.3)	8.9 (1.3)	9.2 (1.5)	NS
ANC (1 × 10 ⁹ /l)	3.12 (3.1)	3.11 (3.1)	3.65 (2.9)	NS
Platelets (1 × 10 ⁹ /l)	333 (265)	311 (192)	501 (575)	NS
Ferritin (µg/l)	1381 (1306)	1405 (1394)	1227 (683)	NS
Mean medullary blast count, % (Std)	2.0 (1.4)	2.3 (1.3)	1.6 (1.6)	NS
Median number of PRBC transfusions 8 weeks before enrollment, n (range)	2 (0–10)	2 (0–8)	5 (0–10)	NS
Median time from first diagnosis to treatment, months (range)	30 (1–181)	22 (1–181)	14 (2–39)	NS
Median number of Len cycles, n (range)	15 (1–49)	16 (1–49)	6 (1–28)	NS

	TP53 wt	TP53 mu	P value
N	59	8	
Median cycles	16	6	0.38
TI	75%	50%	0.2
TTR	4 months	4 months	
event	25%	63%	0.045
progression	15%	25%	0.6
Median time to 25% AML	NR	18 m ←	
OS, months	NR	43.2	0.002

SINTRA-REV: STUDY DESIGN

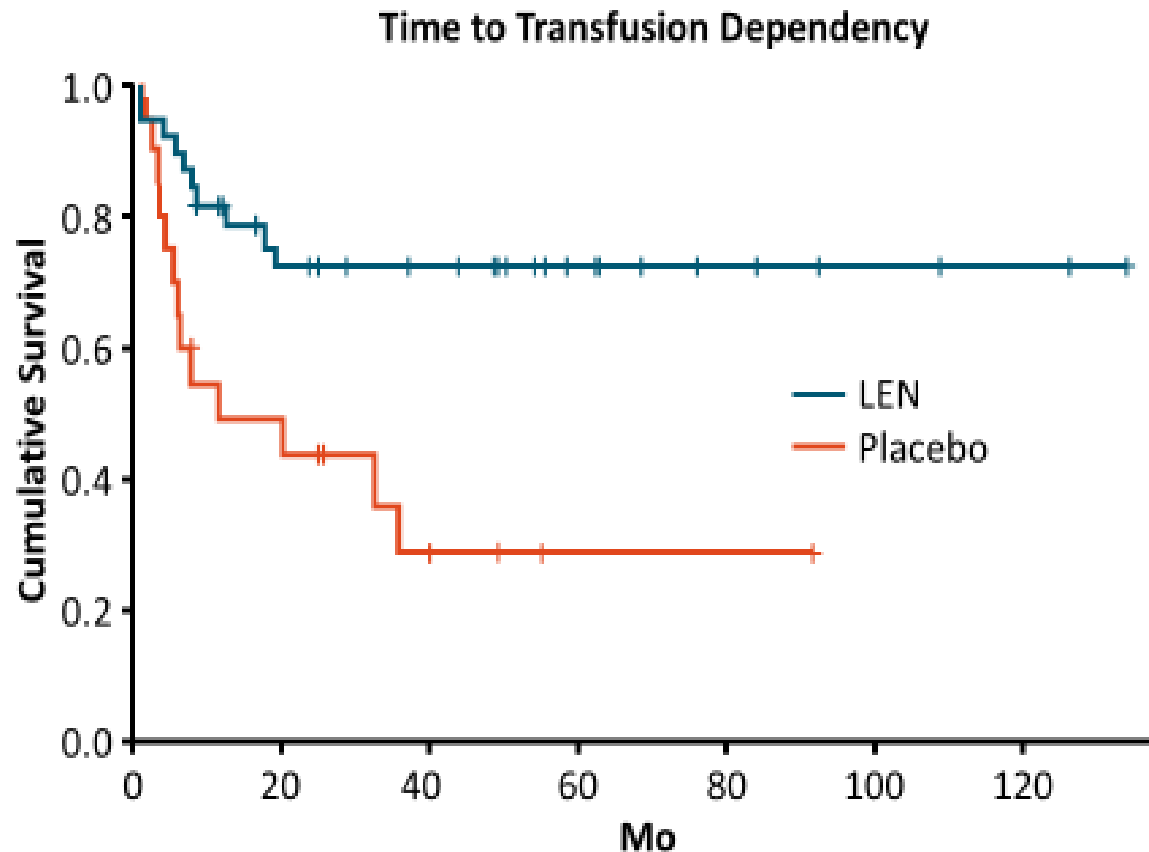
- Multicenter, randomized, double-blind, placebo-controlled phase III trial



Primary endpoint: time to transfusion dependence

Secondary endpoints: erythroid response, cytogenetic response, duration of RBC-TI, change in Hb, bone marrow response, neutrophil/platelet changes, safety, OS, EFS, AML transformation, clonal evolution

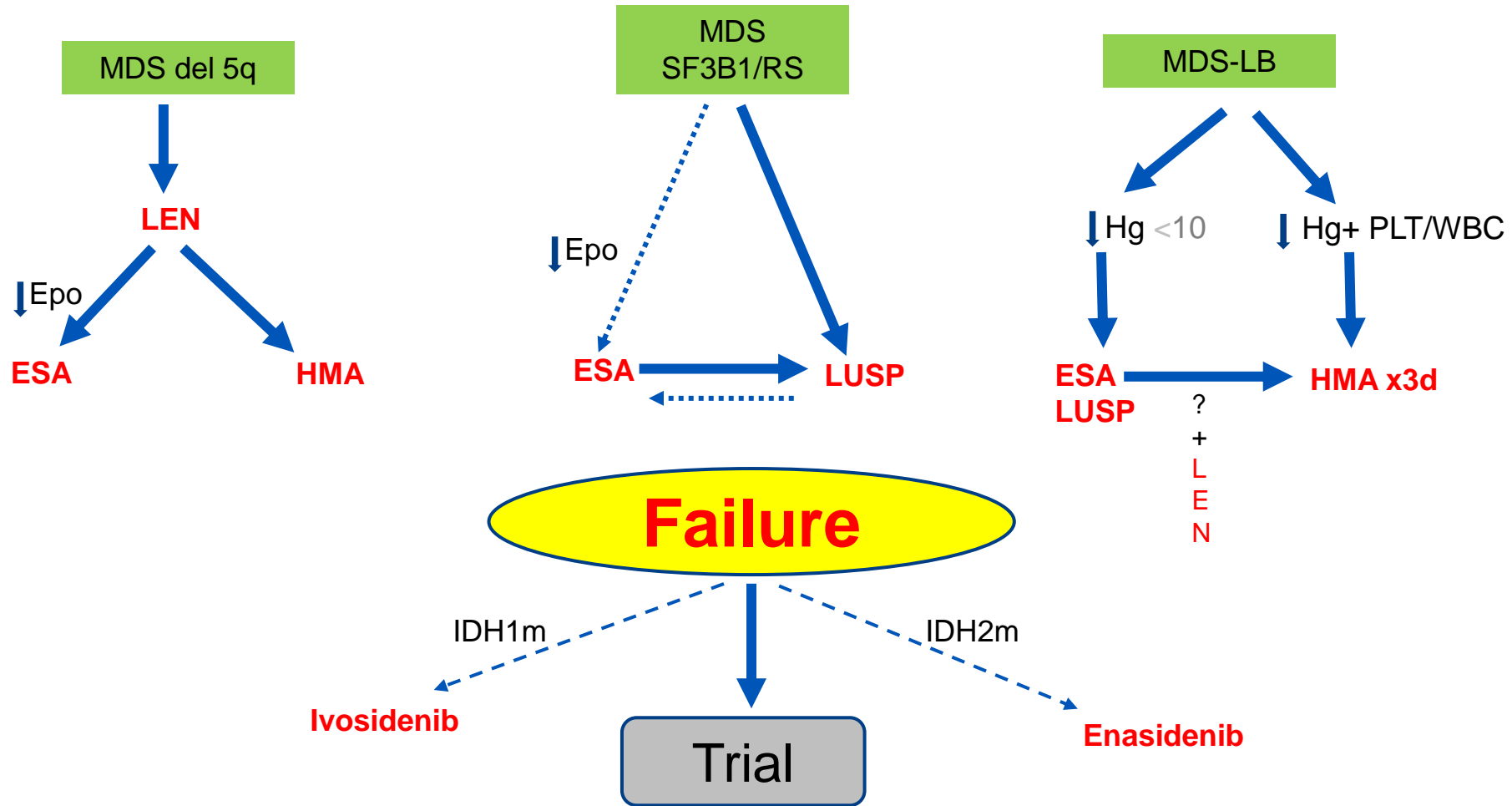
SINTRA-REV: TIME TO TRANSFUSION DEPENDENCY IN ITT POPULATION (PRIMARY ENDPOINT)



	Len (n = 40)	Placebo (n = 21)	P
Events, n (%)	10 (25)	13 (65)	.005
HR (95% CI)	0.302 (0.132-0.692)		
Median time to TD, mo	NR	11.6	.003

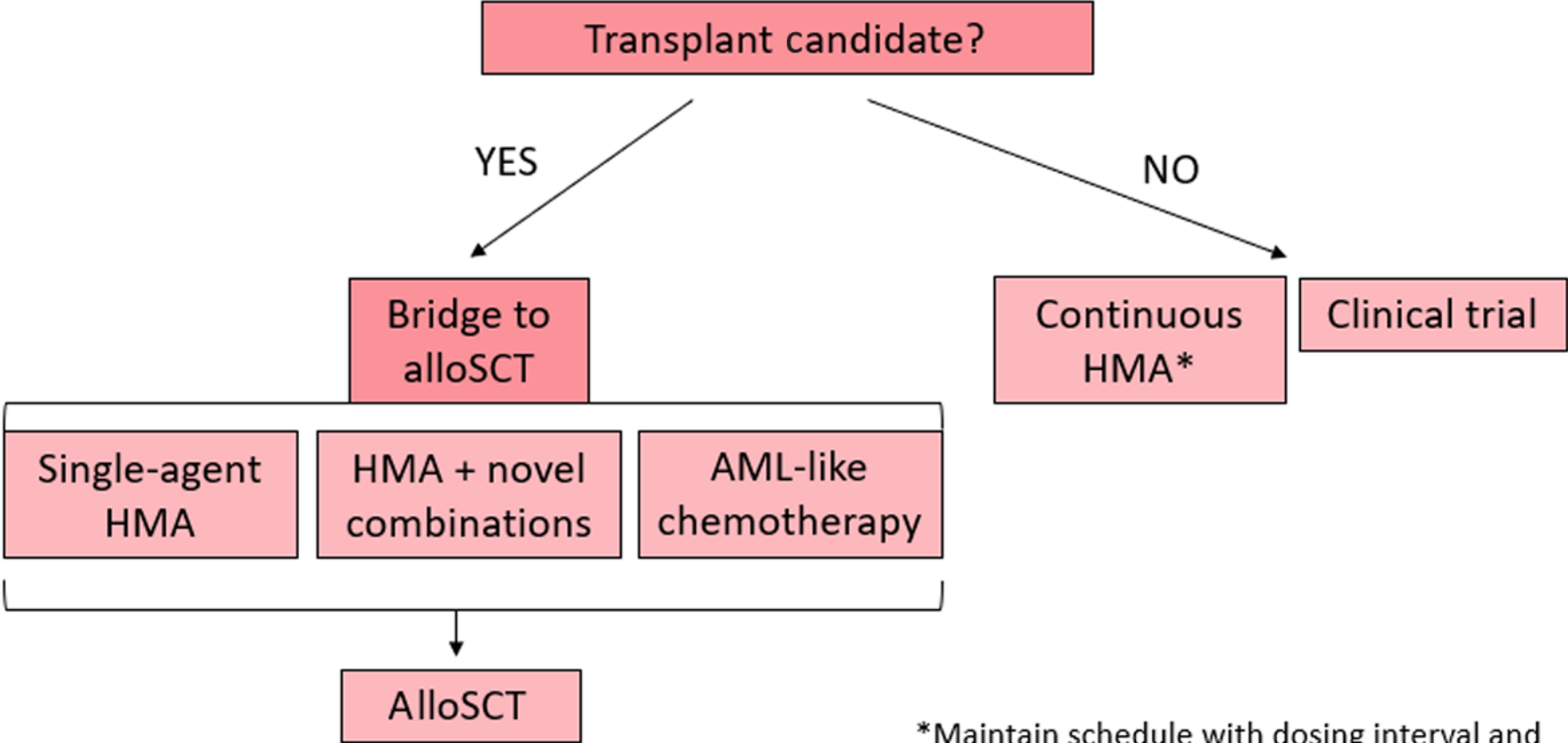
- Median follow-up: 5.05 yr (range: 0.3-11)

Low grade MDS



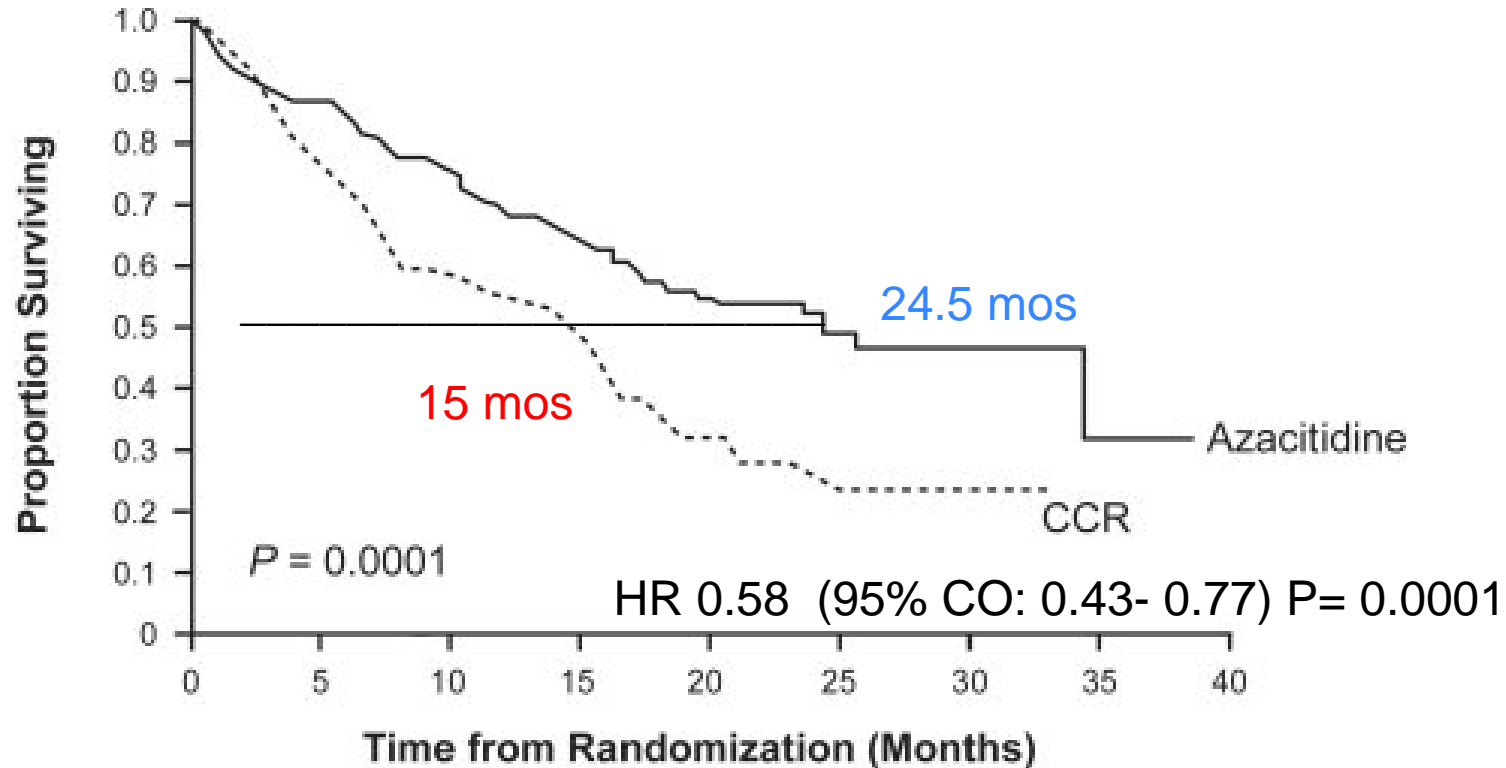
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Current Treatment Algorithm in HR-MDS



*Maintain schedule with dosing interval and intensity for first 4-6 cycles for maximum benefit

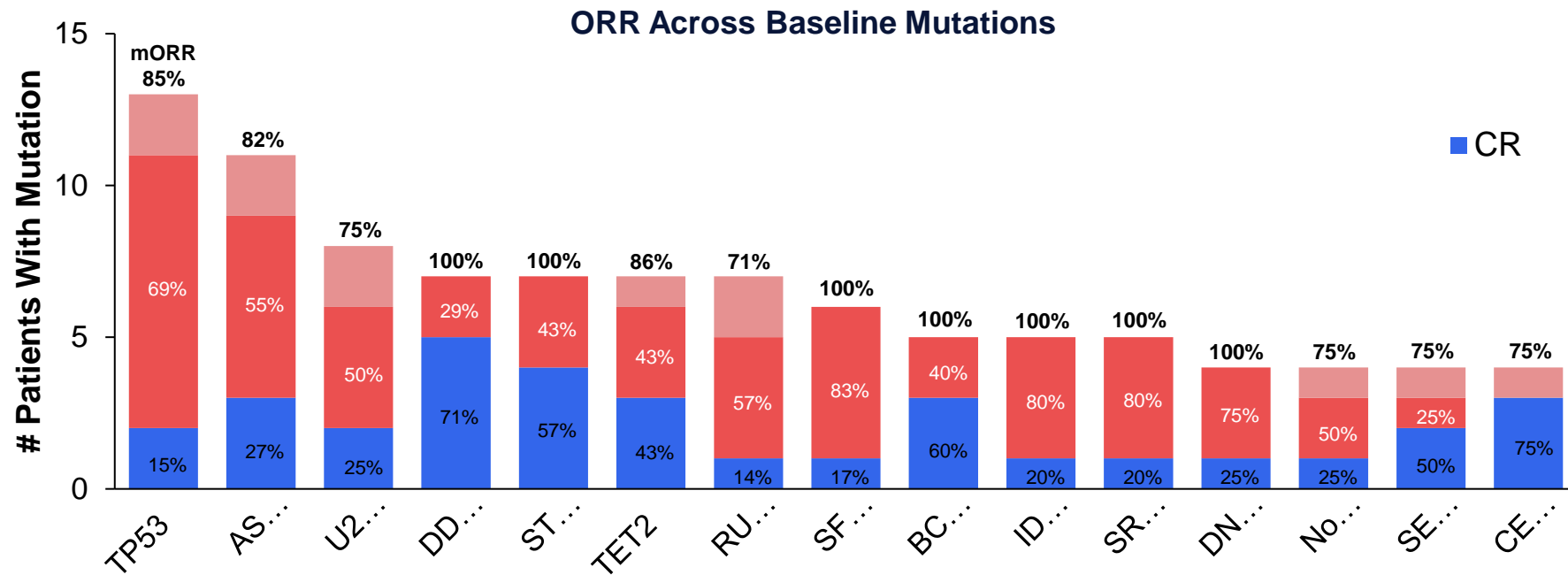
AZA vs CCR in high risk MDS



CCR: 7+3, LDAC, BSC

Fenaux et al Lancet Oncology 2009

Azacitidine and Venetoclax in HR-MDS



To transplant or not.....

Biologic Assignment Trial of Reduced-Intensity Hematopoietic Cell Transplantation Based on Donor Availability in Patients 50-75 Years of Age With Advanced Myelodysplastic Syndrome

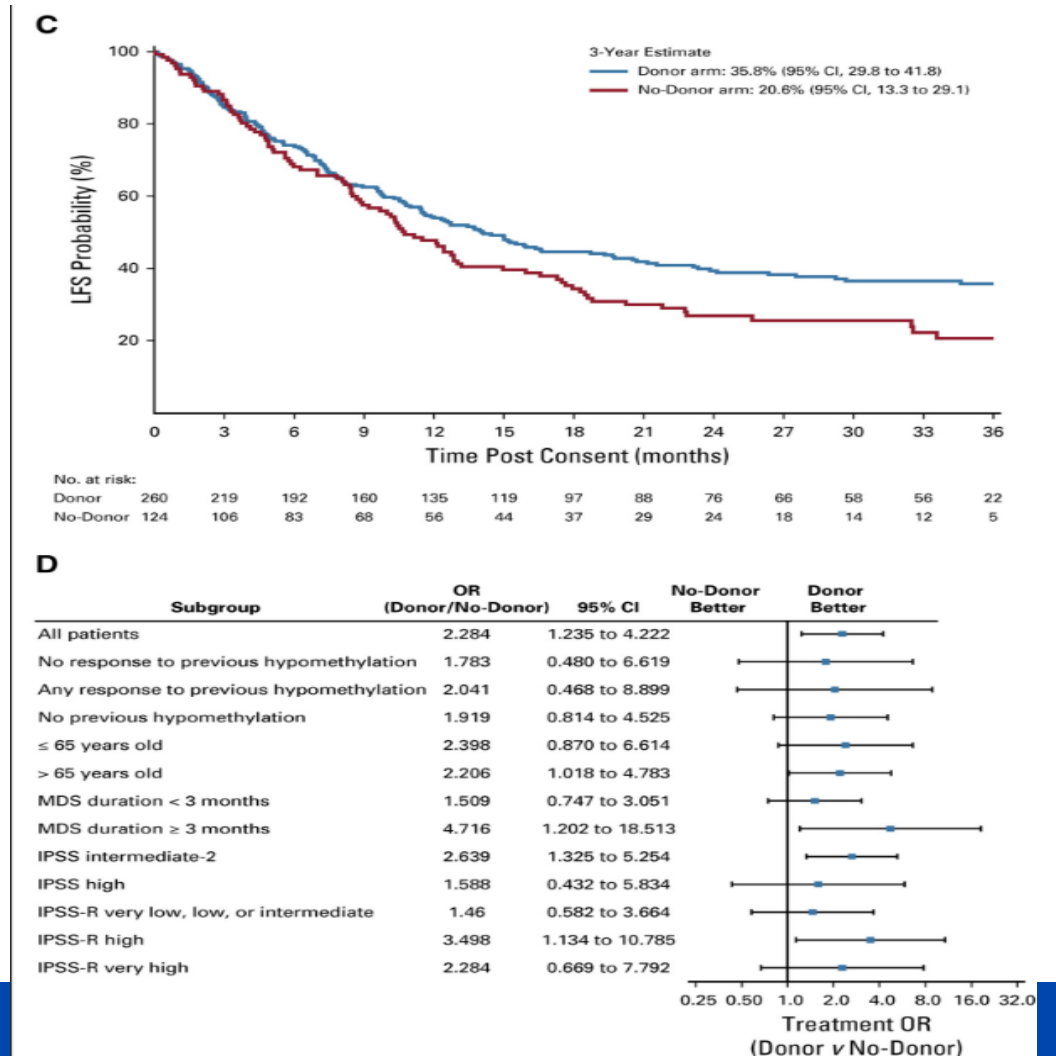
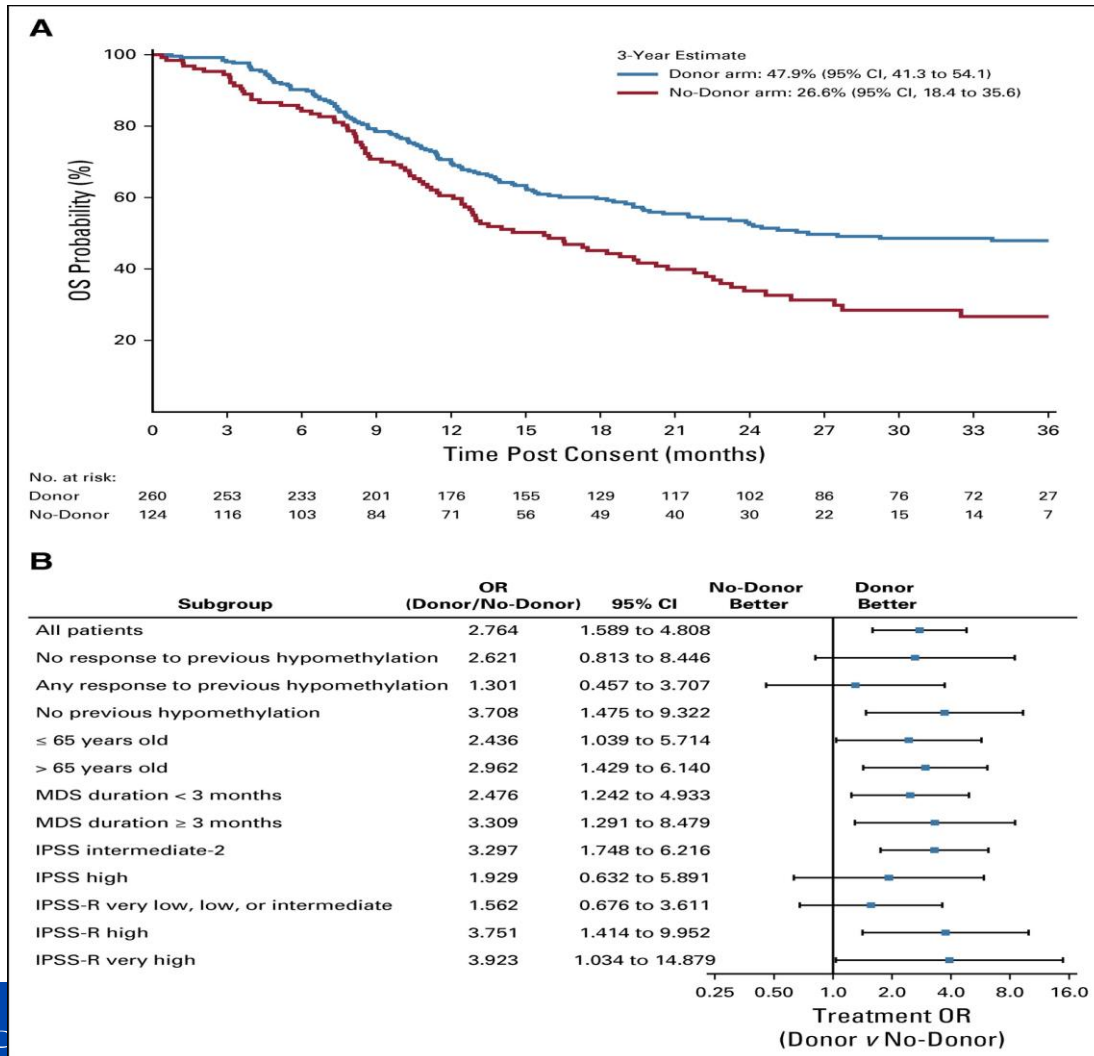
Che
up

Ryotaro Nakamura, MD¹; Wael Saber, MD, MS²; Michael J. Martens, PhD²; Alyssa Ramirez, BS³; Bart Scott, MD⁴; Betul Oran, MD⁵; Eric Leifer, PhD⁶; Roni Tamari, MD⁷; Asmita Mishra, MD⁸; Richard T. Maziarz, MD⁹; Joseph McGuirk, DO¹⁰; Peter Westervelt, MD, PhD¹¹; Sumithira Vasu, MBBS¹²; Mrinal Patnaik, MBBS¹³; Rammurti Kamble, MD¹⁴; Stephen J. Forman, MD¹; Mikkael A. Sekeres, MD, MS¹⁵; Frederick Appelbaum, MD⁴; Adam Mendizabal, PhD³; Brent Logan, PhD²; Mary Horowitz, MD, MS²; and Corey Cutler, MD, MPH¹⁶; on behalf of the Blood and Marrow Transplant Clinical Trials Network

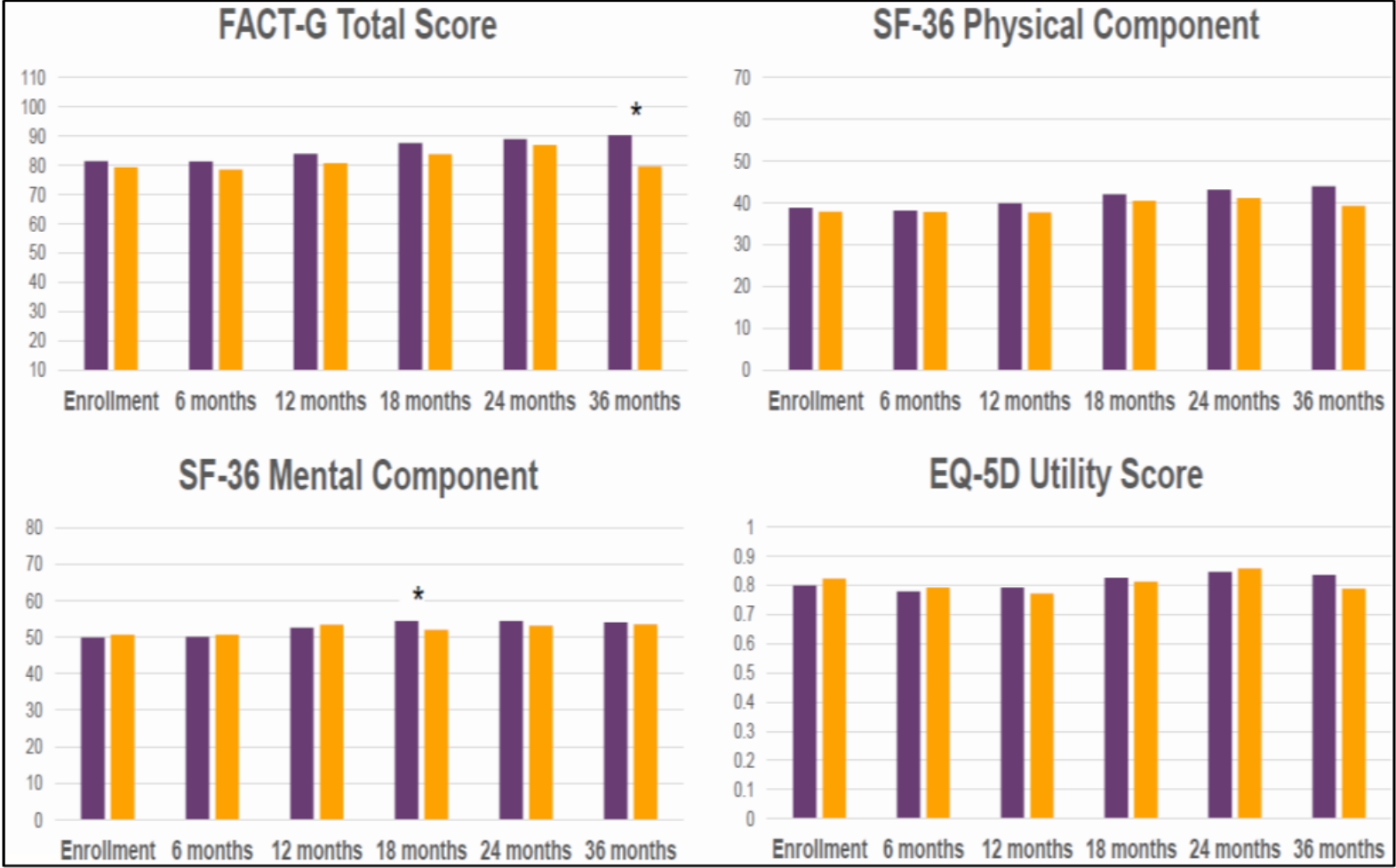
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Clinical outcomes

(Donor arm=260 and No donor arm=124 patients)



Quality of life similar in both groups

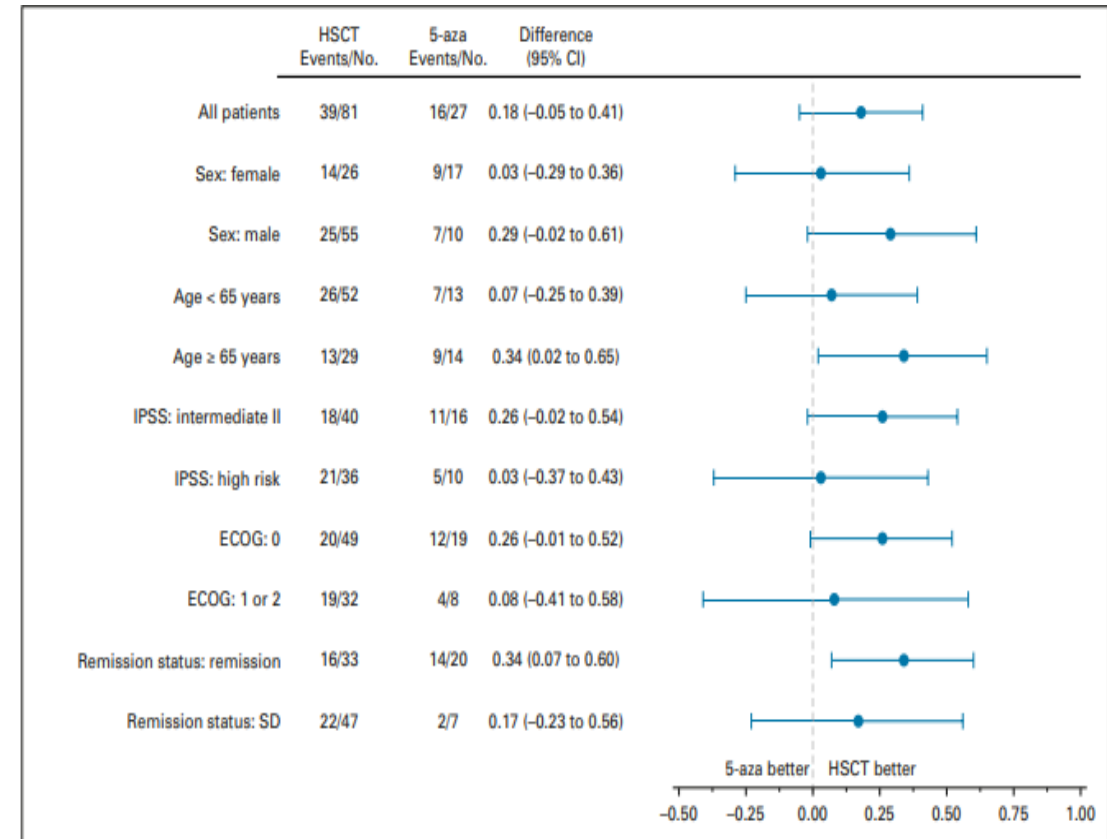
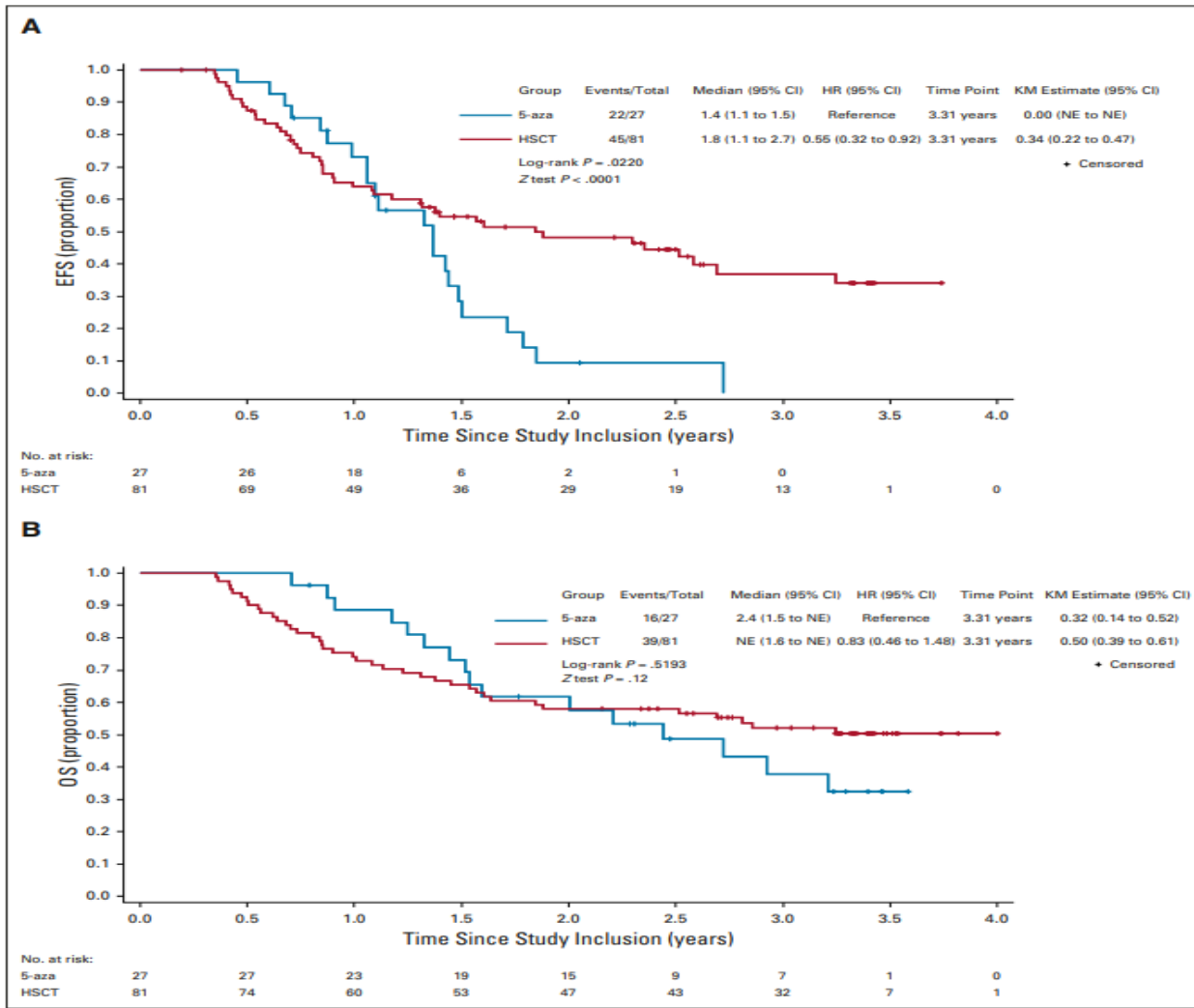


Comparison Between 5-Azacytidine Treatment and Allogeneic Stem-Cell Transplantation in Elderly Patients With Advanced MDS According to Donor Availability (VidazaAllo Study)

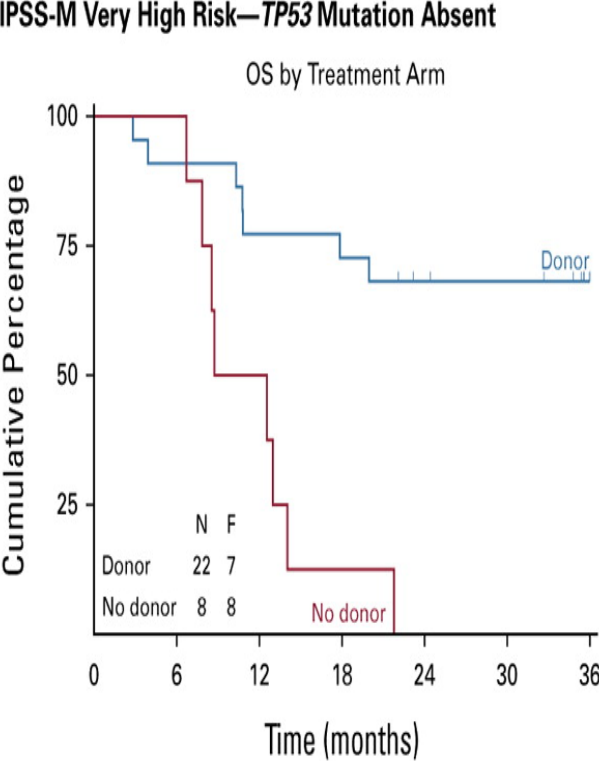
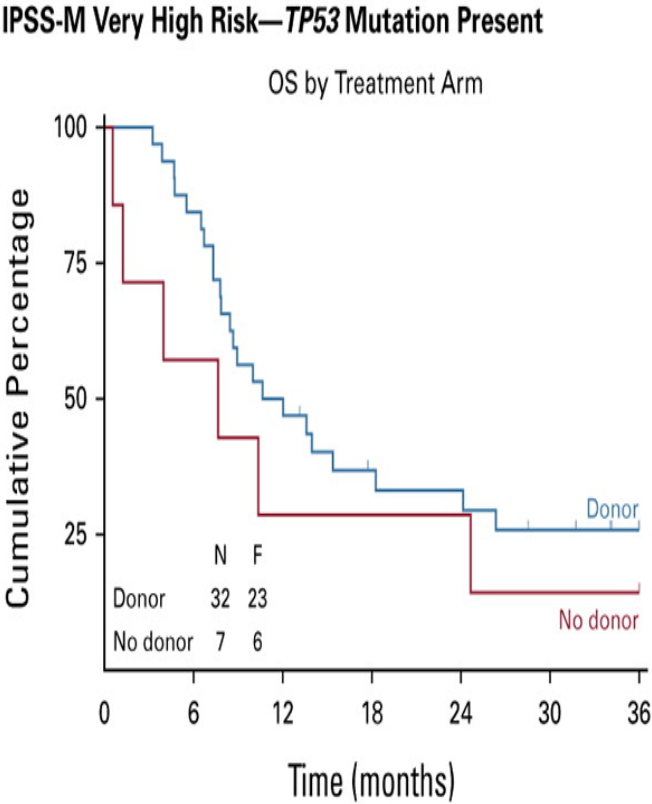
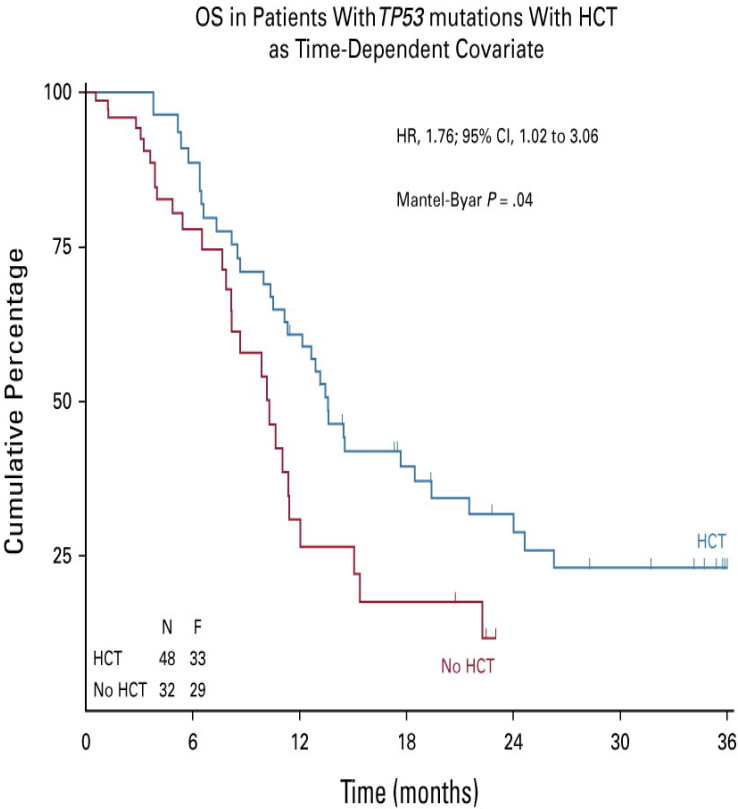
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J Clin Oncol 39:3318-3327. © 2021 by American Society of Clinical Oncology

Clinical outcomes (HCT arm=81 and continuous 5-Aza arm=27)



Does transplant work even with high risk mutations such as TP53?



No. at risk:

Eprenetapopt (APR-246) + HMA following Allo-hct

- 33 patients (14 AML, 19 MDS)
- Median age was 65 (range, 40-74) years.
- median number of eprenetapopt cycles was 7 (range, 1-12).
- With a median follow-up of 14.5 months, the
 - median RFS was 12.5 months
 - 1-year RFS probability was 59.9% (95% CI, 41 to 74).
 - median follow-up of 17.0 months,
 - median overall survival (OS) was 20.6 months (95% CI, 14.2 to not estimable)
 - 1-year OS probability was 78.8% (95% CI, 60.6 to 89.3).

Conclusions

- Many recent advances in Prognostication and Treatment of MDS
 - M-IPSS
 - CHIP, CCUS
- New indications/ therapies for LR-MDS
 - Luspatercept, Imelstat
- Allo-hct with proven survival benefit in HR-MDS in older adults
 - Effective strategy even in TP53 myeloid neoplasm

Allo-HCT for high risk MDS (BMT CTN 1102)

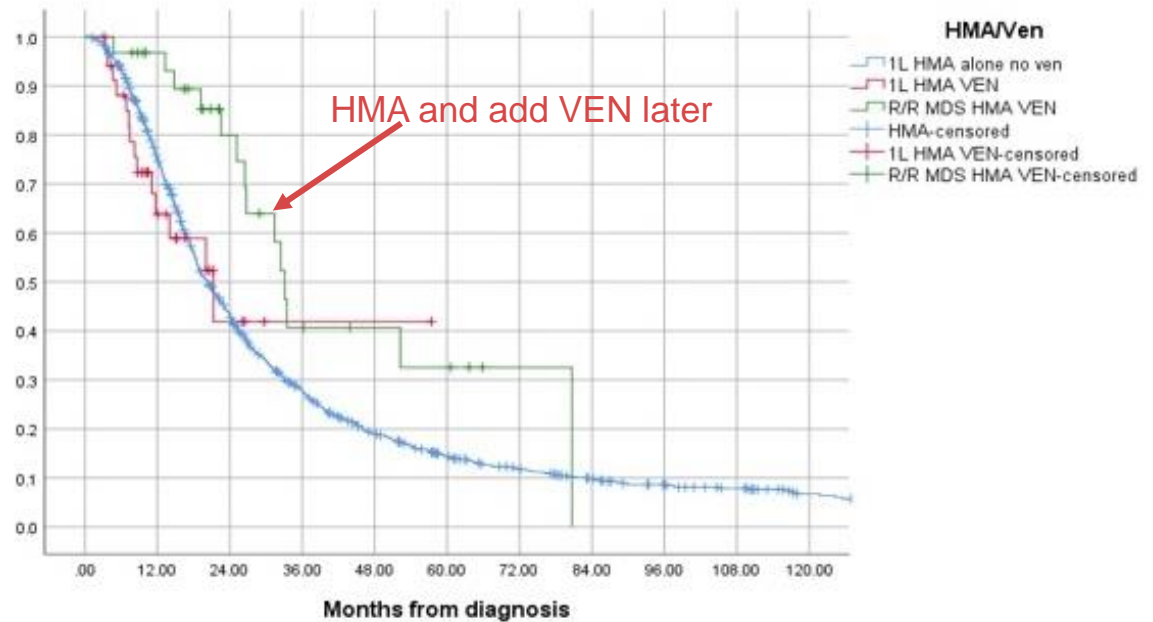
- Allo-HCT with reduced intensity conditioning (RIC) in patients 50 to 75 years old with HR-MDS comparing patients with and without a (HLA)-matched donor.
- 384 total patients
 - 67.7% (260 patients) had a matched donor and
 - 32.3% (124 patients) did not have a matched donor and instead received HMA and/or supportive care
 - HLA-matched donor experienced
 - increased OS (absolute improvement in 3-year OS was 21.3% [95% CI, 10.2-31.8; $P = .0001$])
 - improved leukemia-free survival (absolute improvement in 3-year LFS was 15.2% [95% CI, 13.3-29.1; $P = .003$])
- Interestingly, OS was significantly shorter in patients without response to HMA prior to alloSCT (HR, 1.64; $P = .0097$)

Assessing the Role of Venetoclax in Combination with Hypomethylating Agents in Higher Risk Myelodysplastic Syndromes

- Higher risk MDS at Moffitt Cancer Center (INT, H, VH)
- Compare: first line single agent HMA, first line HMA/Ven combination, HMA, but add Ven after HMA failure (R/R MDS HMA/Ven)

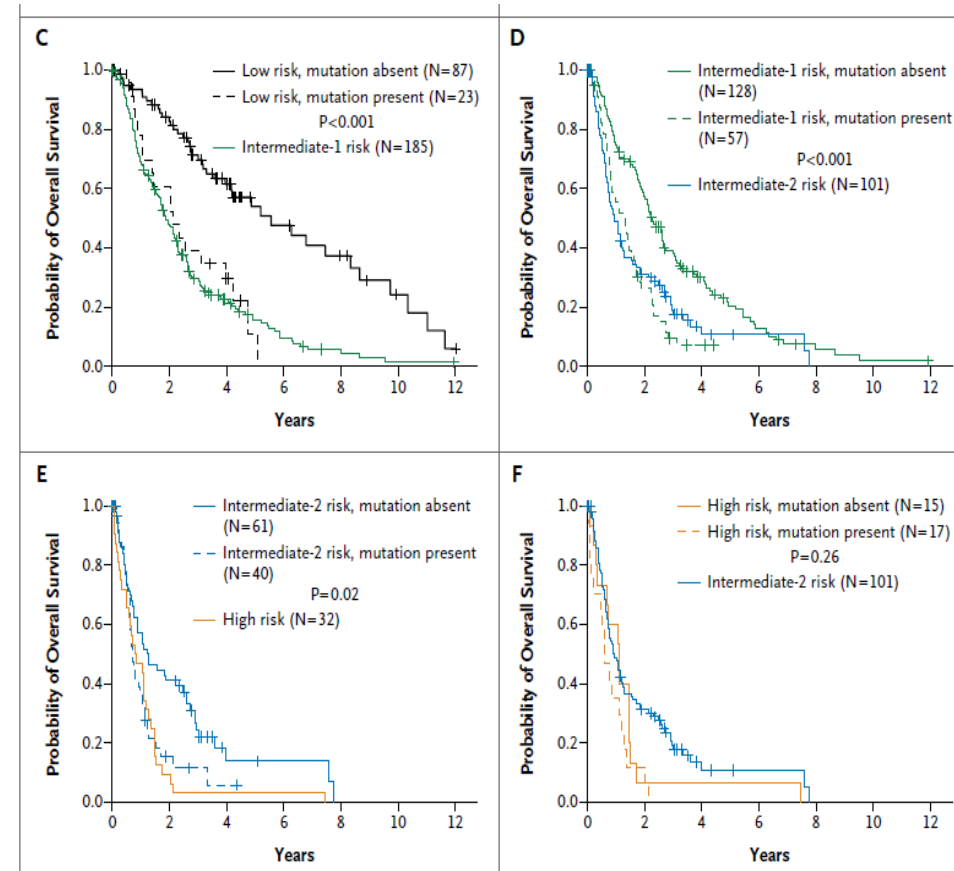
Table-1 Baseline characteristics comparing HMA alone versus HMA Ven first line therapy in higher risk MDS

		HMA 1L	HMA/Ven 1L	P value
n		1127	35	
Age	mean	68.4	67.8	.76
Gender	Male	66%	71%	.5
Race	White	90%	97%	.66
t-MDS		24%	23%	.86
WHO 2016	MDS-SLD/MLD	18%	4%	.04
	MDS-RS	6%	4%	
	MDS-EB1	33%	9%	
	MDS-EB2	39%	78%	
R-IPSS	Intermediate	31%	17%	.22
	High	31%	37%	
	Very High	38%	46%	
Myeloblasts	Mean (%)	8	13	< .005
Hgb	Mean (g/dl)	9	9	1.0
WBC	Mean	4	10.6	< .005
ANC	Mean	1.8	4.1	< .005
platelets	platelets	96	100	.8
Somatic Mutations (n= 546 sequenced)	<i>SF3B1</i>	5%	0	.3
	<i>TET-2</i>	16%	23%	.3
	<i>IDH-1</i>	3%	3%	.7
	<i>IDH-2</i>	5%	14%	.056
	<i>ASXL-1</i>	21%	46%	.002
	<i>TP53</i>	27%	34%	.6
	<i>NRAS</i>	4%	11%	.07



TP53 Mutations Predict Prognosis in MDS

- p53 is a critical tumor suppressor protein that responds to cellular stress by inducing cell cycle arrest, senescence and/or apoptosis to eliminate precancerous and tumor cells.
- Mutations in the *TP53* gene (m*TP53*) are found in 5-10% of MDS and AML patients and predict for the most inferior OS (median OS 6-12 months).
- Confers poor prognosis in allo-SCT



Thank you

