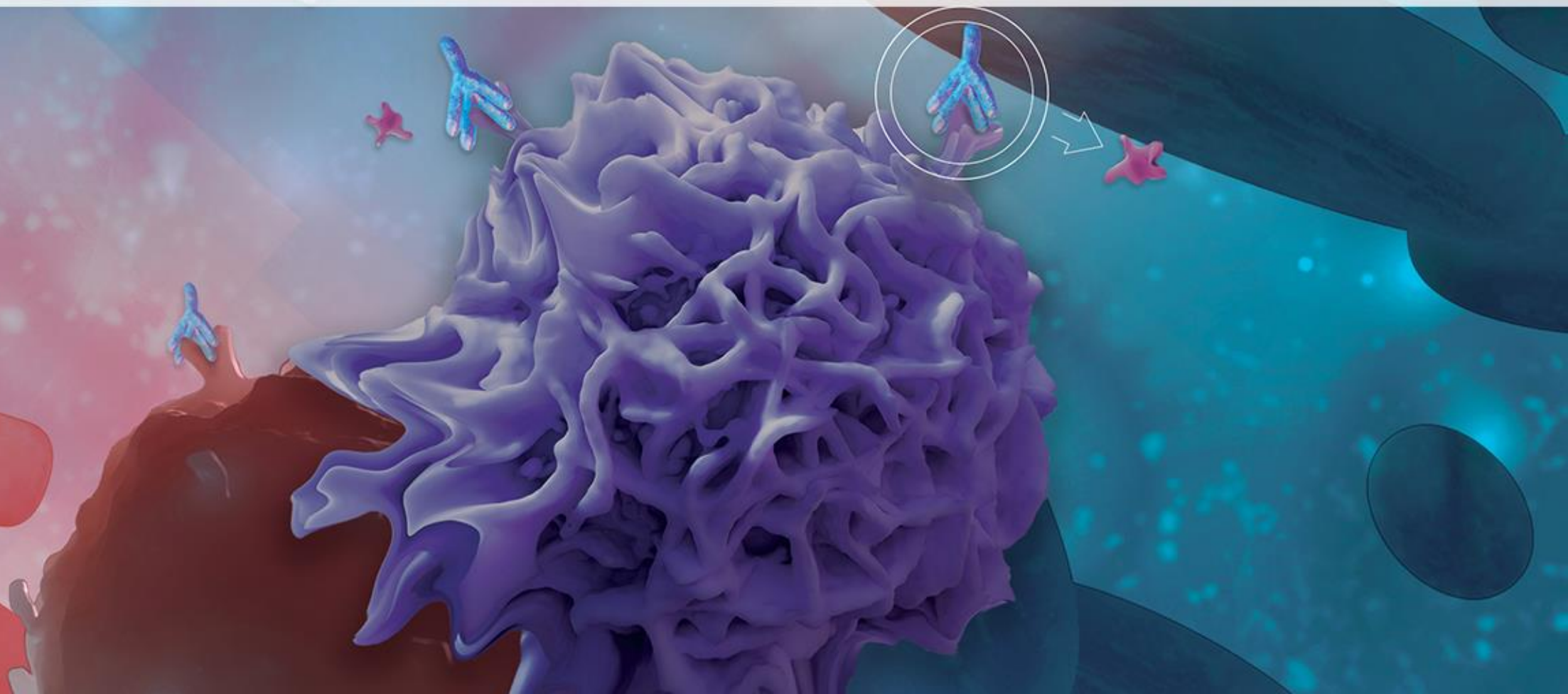


Harnessing the Power of the Immune System to Manage Higher-Risk MDS



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Disclosure of Conflicts of Interest

Consultant: Acceleron Pharma; Agios Pharmaceuticals, Inc; Bristol-Myers Squibb Co; Celgene Corp; Gilead; Keros Therapeutics; Novartis Pharmaceuticals Corp; Taiho Pharmaceutical Co, Ltd; and Takeda Oncology.

Learning Objectives

Upon completion of this activity, participants should be better able to:

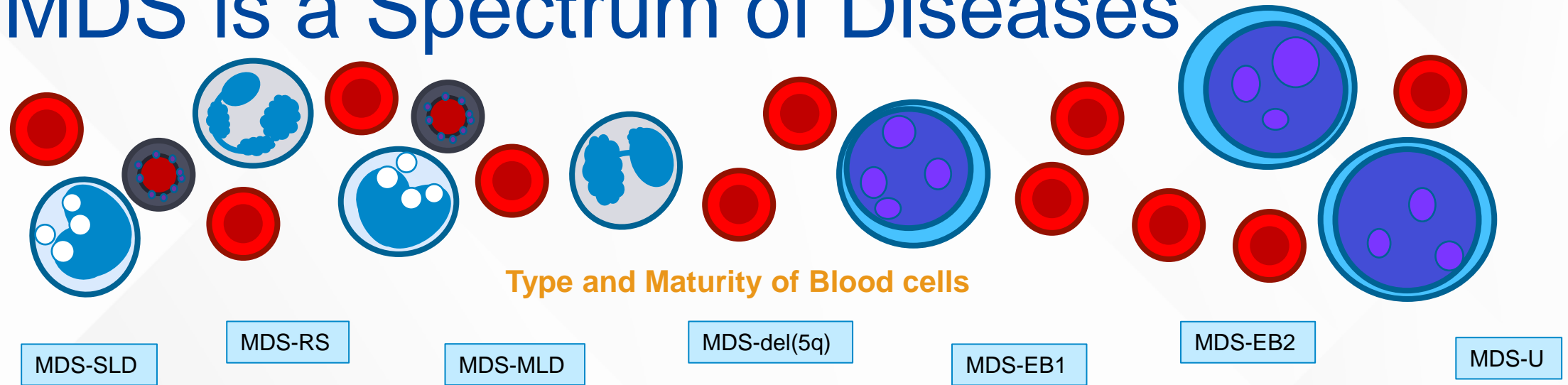
- Utilize biomarker testing and prognostic scoring systems to define higher-risk myelodysplastic syndrome (MDS) and to guide treatment
- Discuss the evolving role of the immune system in MDS, including the various pathways involved in dysregulation such as the TIM-3 pathway
- Review efficacy results of immuno-myeloid therapy targeting TIM-3 in combination with HMAs as treatment for higher-risk MDS
- Develop management plans to address adverse events related to novel and emerging therapies for MDS

What Are 'Higher-Risk' Myelodysplastic Syndromes?

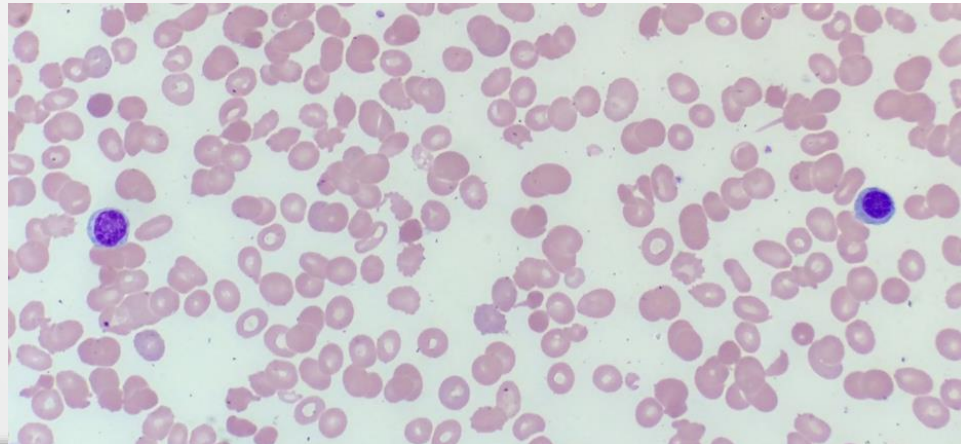
Patient Presentation

- The clinical features that can be used to identify and characterize high-risk MDS subtypes
- Risk stratification systems based on the modern MDS prognostic models, including IPSS and IPSS-R, along with IPSS-M
- Identifying higher-risk MDS subtypes based on blood counts, percentage of blast cells, cytogenetics, subclonal heterogeneity, hypermethylation of tumor suppressor genes, and unfavorable genetic mutations
- Burden of disease, diagnosis, and biomarker testing in higher risk MDS

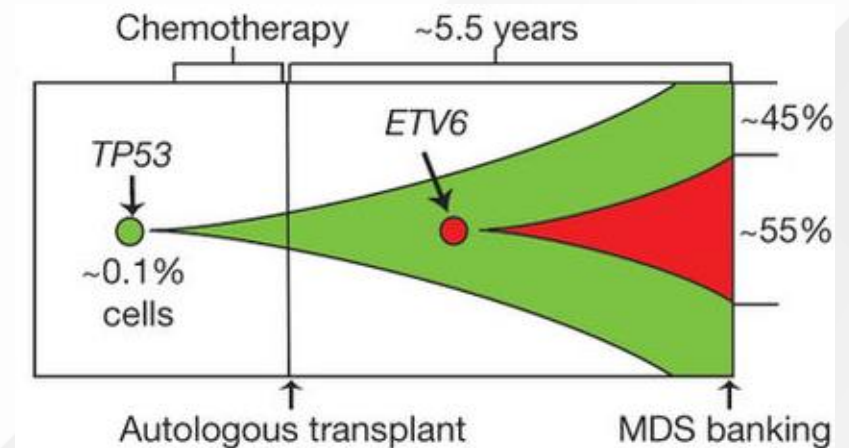
MDS is a Spectrum of Diseases



Variation in Blood Counts



Variation in Driver Mutations



Characterizing MDS

Newly Diagnosed MDS
Evaluate Type and Depth of Cytopenias

CBC count and
differential

Full cytogenetic
analysis

Bone marrow
core/aspirate

Molecular
diagnostics

Risk Stratification:

IPSS-R

Consideration of LR-PSS, WPSS, IPSS-M, Molecular Diagnostics

IPSS-R Calculation

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	-	-	-	-	-

Cytogenetics

Very good: -Y, del(11q)

Good: normal, del(5q), del(12p), del(20q)
double clone w/ del(5q)

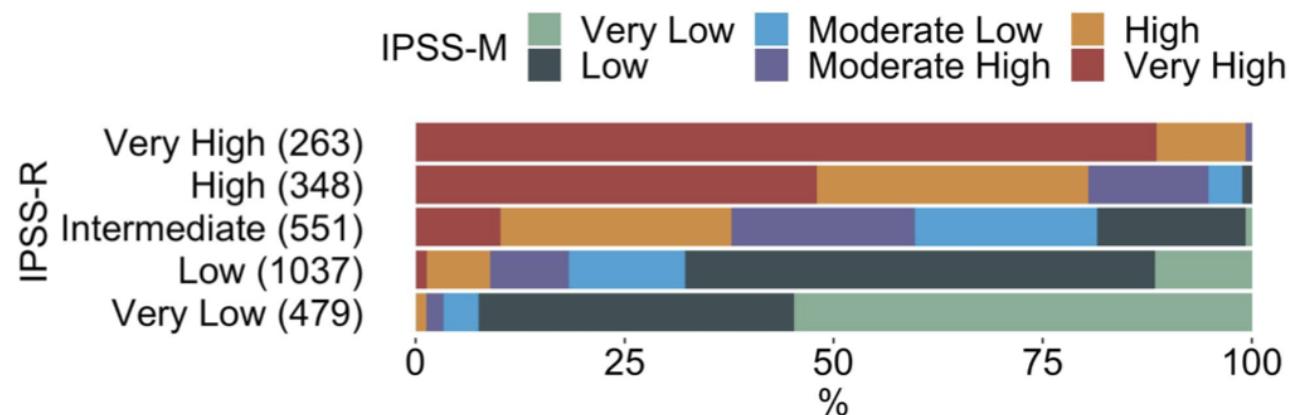
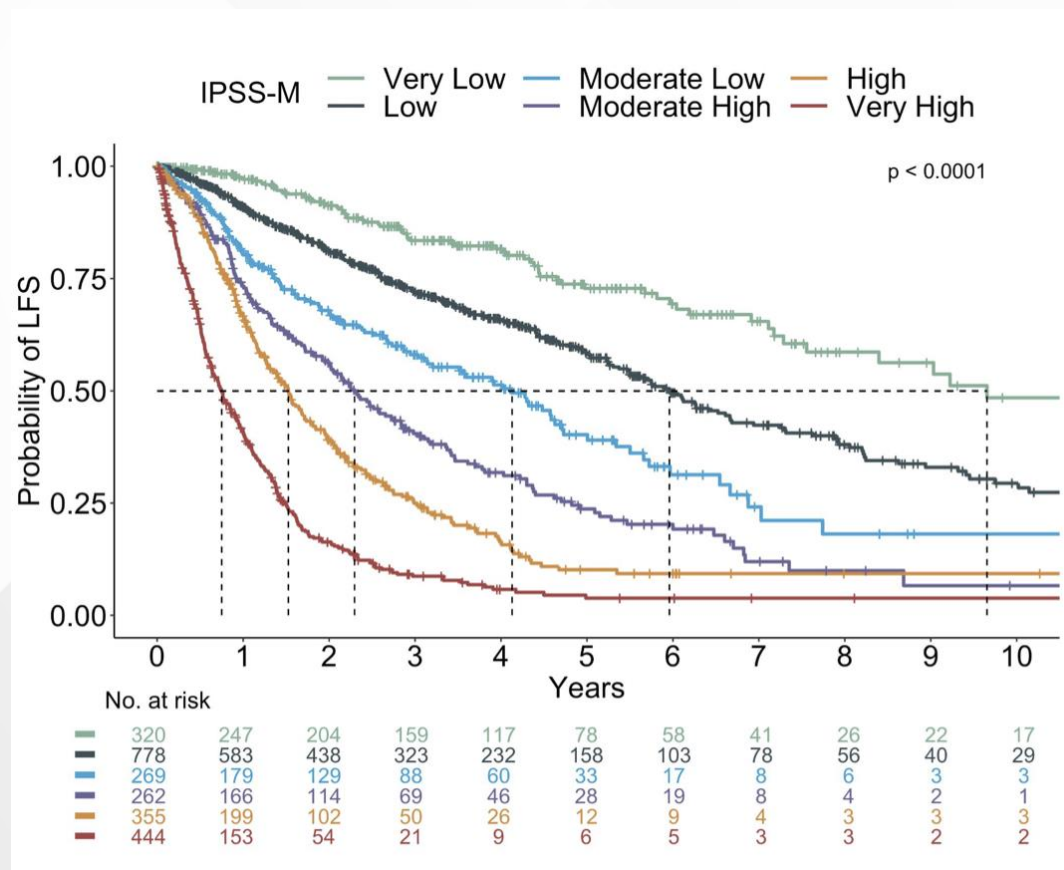
Intermediate: del(7q), +8, +19, i(17q), other
single/double clone

Poor: -7, inv(3)/t(3q)/del(3q), double clone
including -7/del(7q), complex w/ 3 abnl

Very poor: Complex (>3 abnormalities)

	Very Low	Low	Intermediate	High	Very High
SCORE	≤1.5	>1.5-3	>3-4.5	>4.5-6	>6
OS (years)	8.8	5.3	3.0	1.6	0.8
25% AML (years)	NR	10.8	3.2	1.4	0.73

IPSS-M



- Poor LFS, OS, and AML transformation:
 - TP53 multi-hit mutations (7%)
 - MLL PTD (2.5%)
 - FLT3 mutations (1%)
- More favorable clinical course:
 - SF3B1, depending on commutations

MDS Treatment Is Based on Disease Risk

Risk Stratification by IPSS or IPSS-R Blood Counts, Blasts, and Karyotype

Risk for Serious or Life-threatening
Complication related to MDS:

Infection
Bleeding

Risk for Progression to Acute Myeloid
Leukemia

IPSS and IPSS-R Risk do not always match the risk of the WHO disease subtype

Defining Higher-Risk MDS

Goal: Identify patients whose disease, left untreated, is at high risk of:

- Death (most often from infection/bleeding/cardiac disease)
- or
- Leukemic progression within months (generally <18 months)

IPSS

INT-2 Risk
High Risk

IPSS-R

Score >3.5
Intermediate
High
Very High

Disease
History

Progression after
prior therapies; eg,
formerly low risk

*

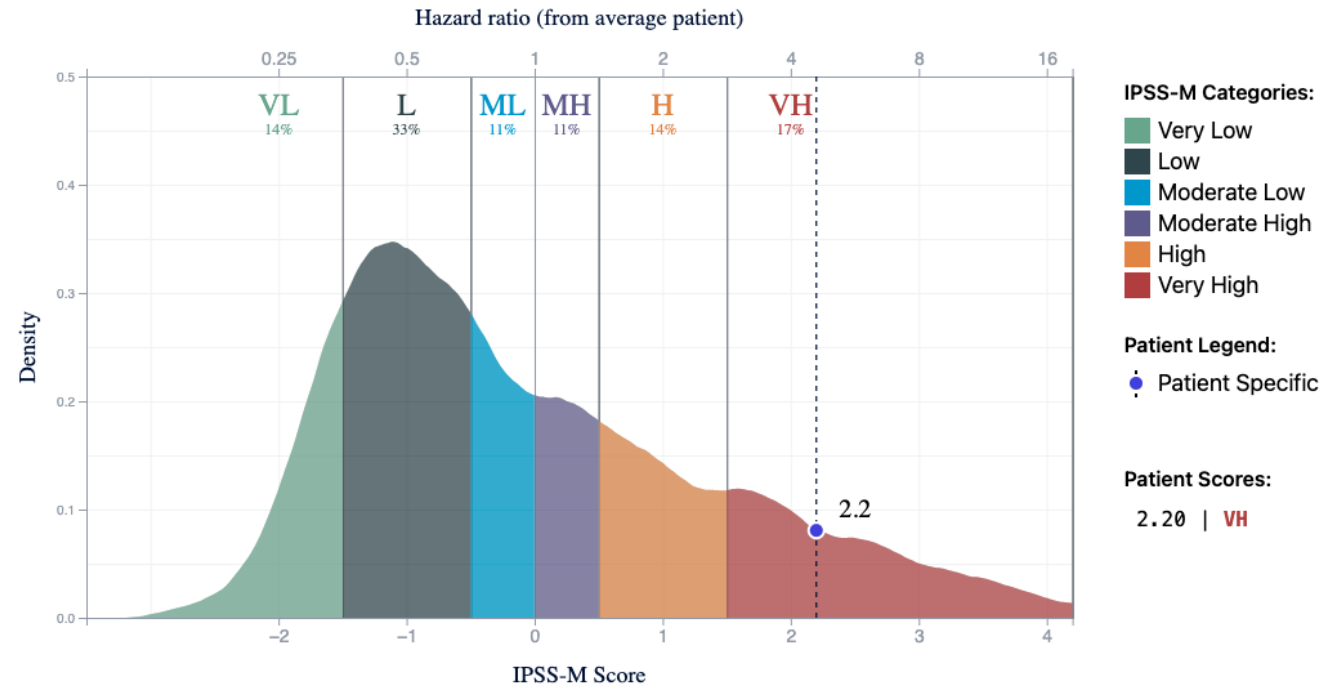
Molecular

TP53
EZH2, RUNX1, ASXL1
“AML-like” mutations

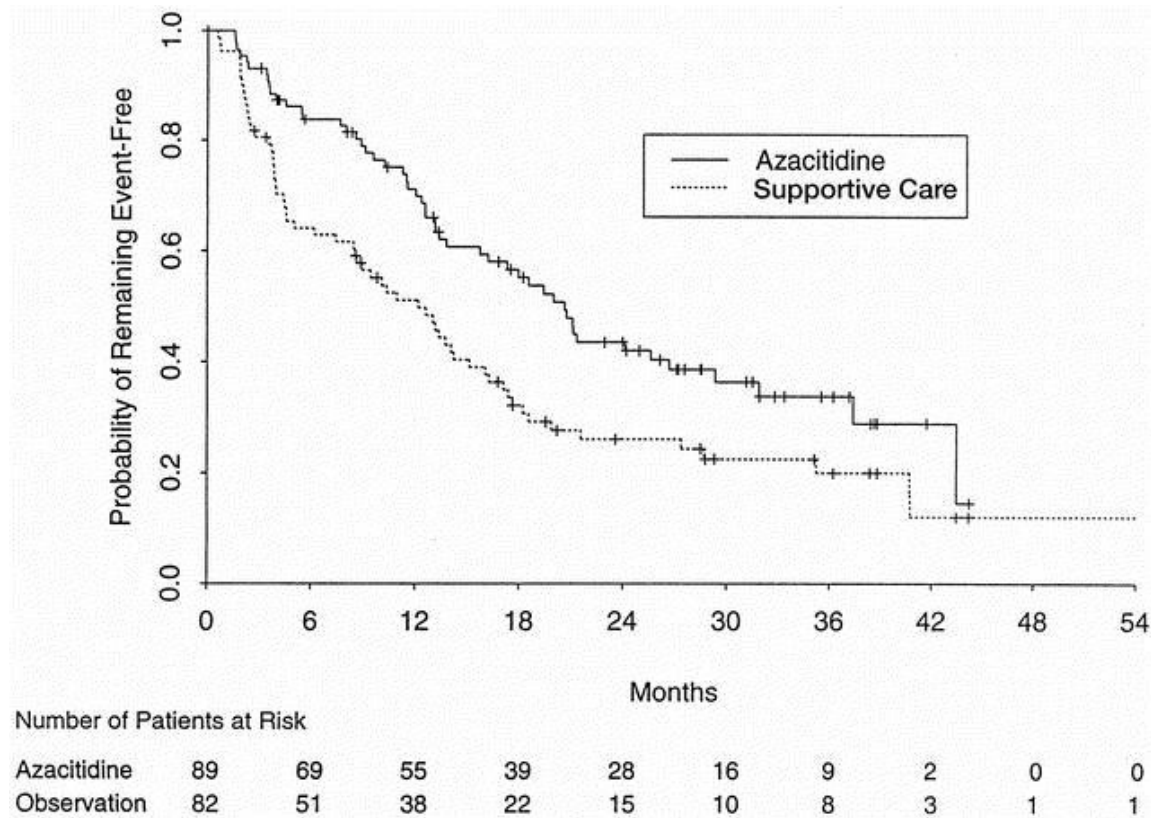
Higher-Risk MDS Case

Patient JB:

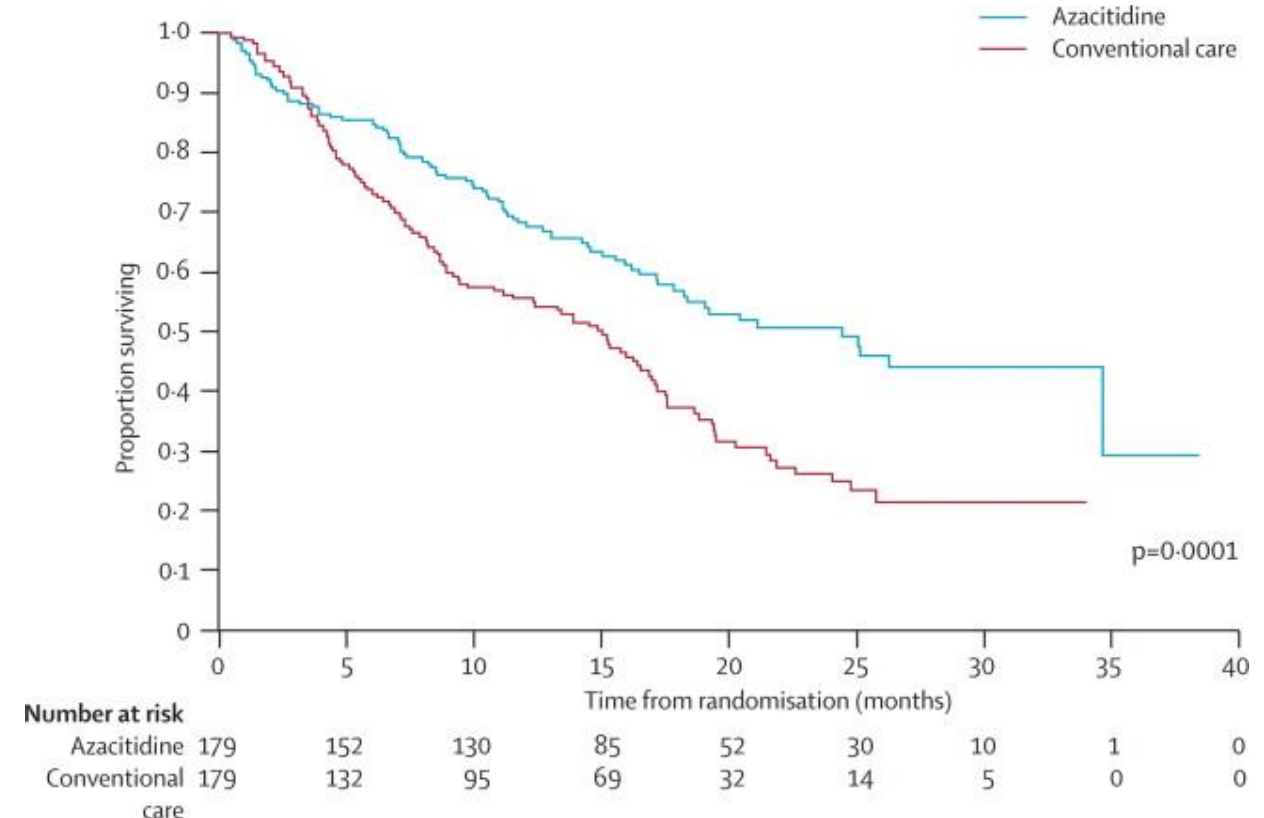
- 80-year-old woman with progressive anemia
- CBC and differential with WBC 2, ANC 0.6, Hgb 8 g/dL, platelets 45
- Bone marrow biopsy: hypercellular, 12% CD34+ blasts, no ring sideroblasts
- Cytogenetics: 46,XX,del7q
- Molecular studies: mutations in *BCOR*, *CBL*, *U2AF1*



Higher-Risk MDS: Hypomethylating Agents

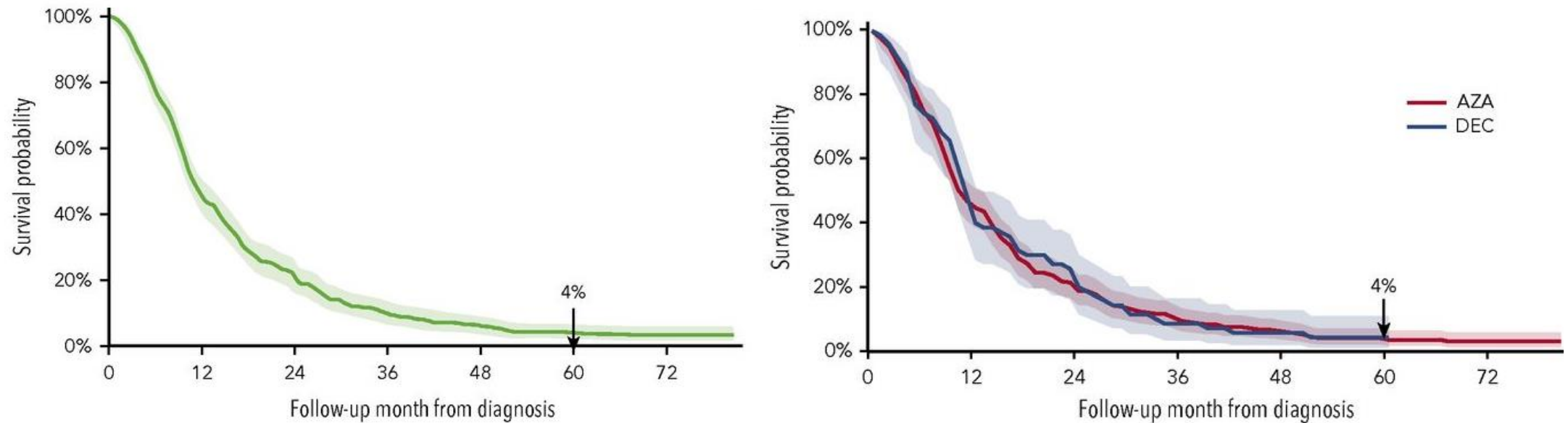


CALGB 9221: OS 20 mo (AZA) vs. 14 mo (Conventional Care)



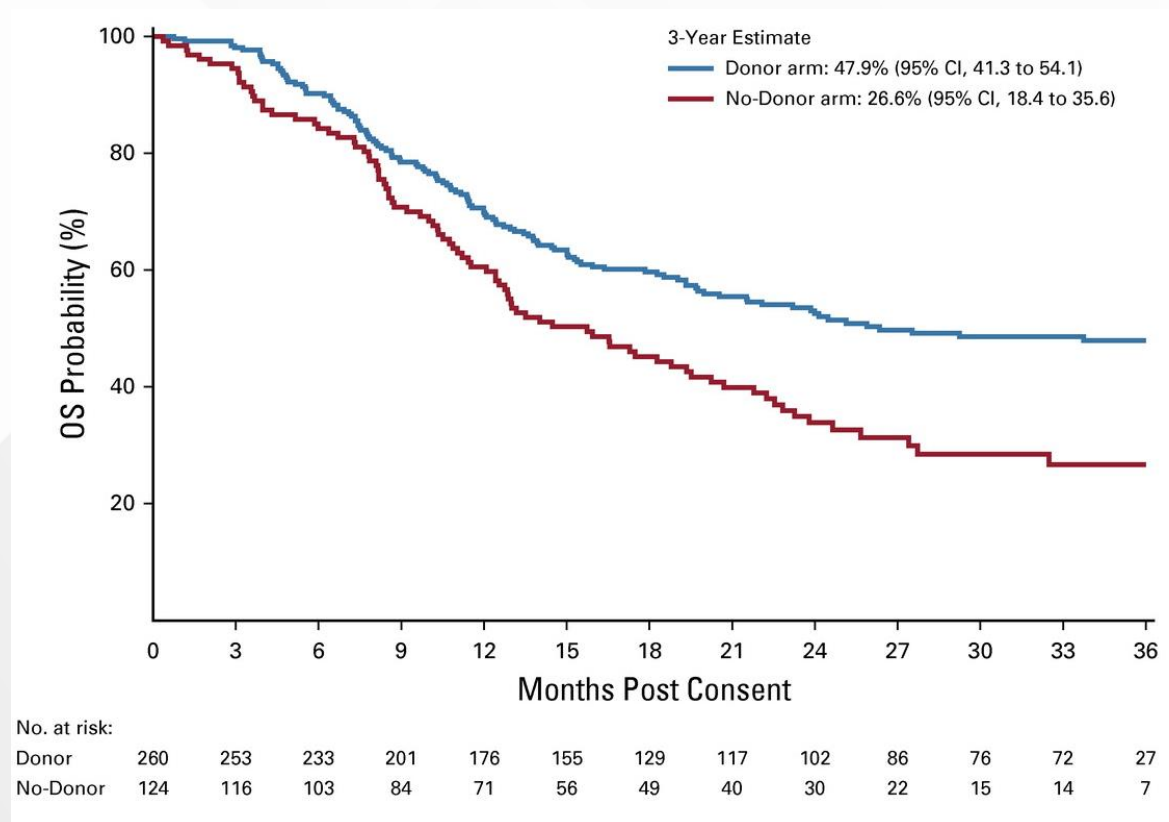
AZA001: OS 21.1 mo (AZA) vs. 15.0 mo (Conventional Care)

HMA in Myelodysplastic Syndromes

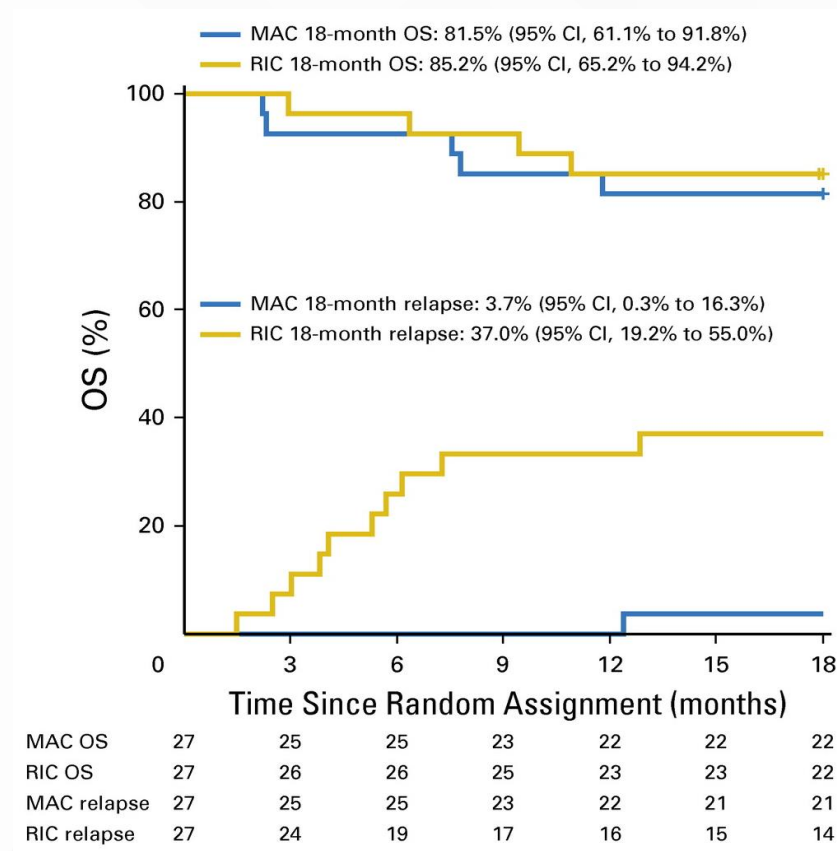


SEER-Medicare, all patients, irrespective of MDS disease risk

Role of Transplant

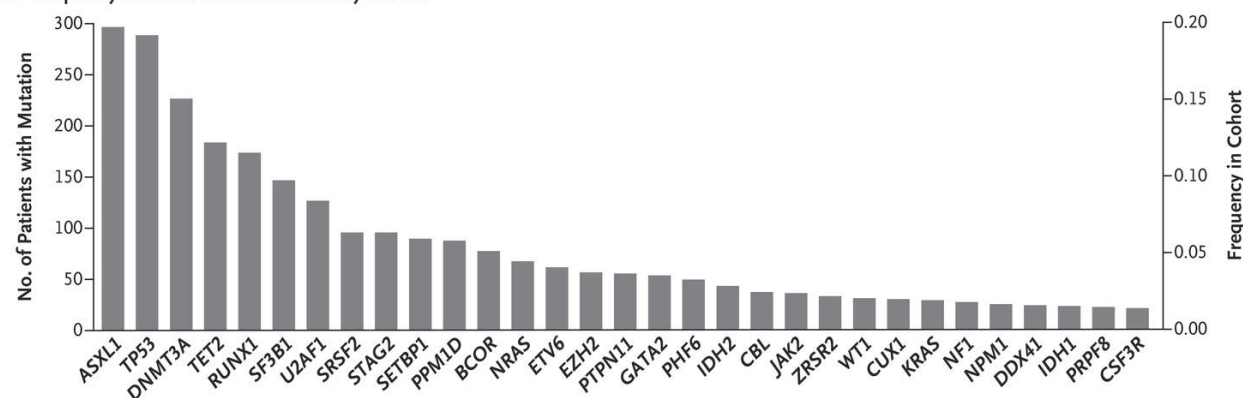


MAC vs RIC in MDS

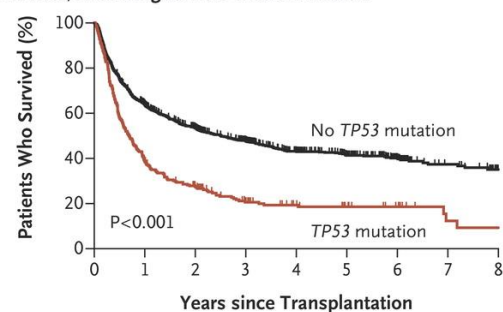


Mutations and Transplant

A Frequency of Driver Mutations in Study Cohort



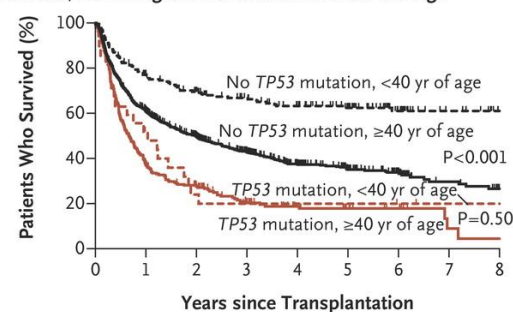
B Overall Survival, According to *TP53* Mutation Status



No. at Risk

No <i>TP53</i> mutation	1224	757	529	370	261	183	109	53	32
<i>TP53</i> mutation	289	109	66	39	26	20	14	6	5

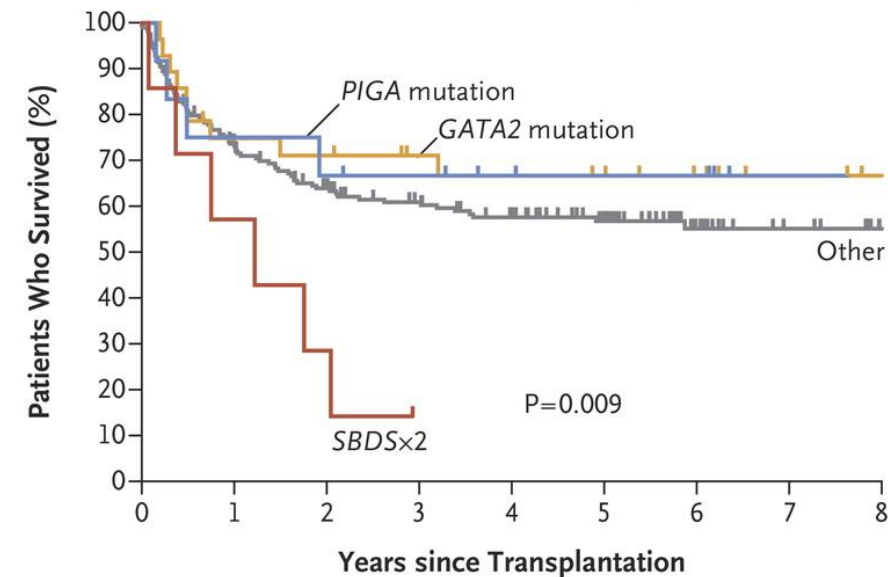
C Overall Survival, According to *TP53* Mutation Status and Age



No. at Risk

No <i>TP53</i> mutation									
<40 yr of age	214	159	133	115	100	78	42	23	13
≥40 yr of age	1010	598	396	255	161	105	67	30	19
<i>TP53</i> mutation									
<40 yr of age	27	14	7	5	5	5	4	4	3
≥40 yr of age	262	95	59	34	21	15	10	3	2

B Overall Survival among Patients <40 Yr of Age, According to Mutation Status

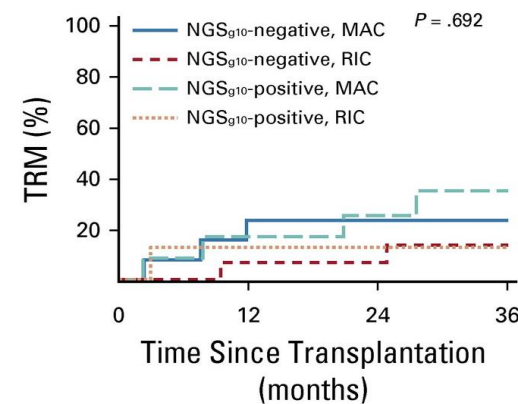
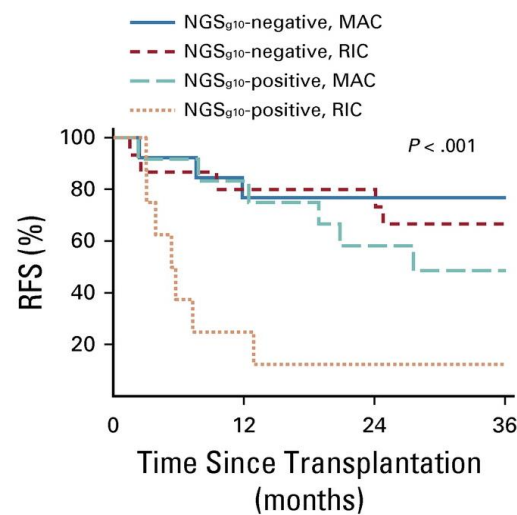
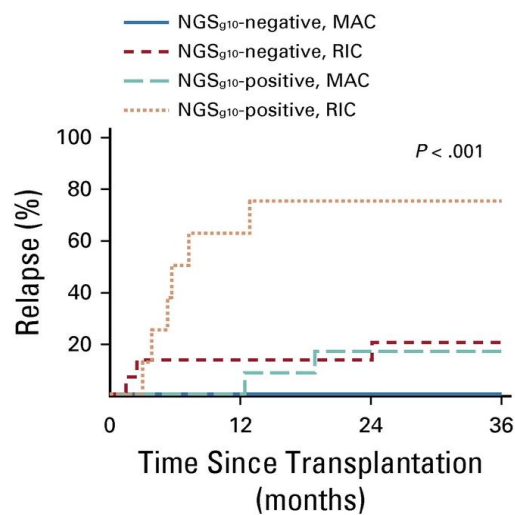


No. at Risk

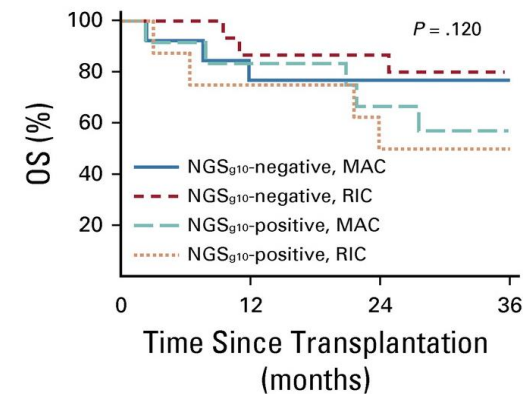
<i>PIGA</i> mutation	12	10	9	8	6	5	5	2	1
<i>GATA2</i> mutation	28	21	20	17	15	13	10	6	5
Other	198	140	111	97	85	66	32	19	11
<i>SBDS</i> ×2	7	5	3	1	0	0	0	0	0

Does pre-transplant therapy/response impact post-transplant outcomes?

RIC vs. MAC Transplant



No. at risk				
NGS _{g10} -negative	MAC	13	10	10
	RIC	15	12	10
NGS _{g10} -positive	MAC	12	10	7
	RIC	8	2	1



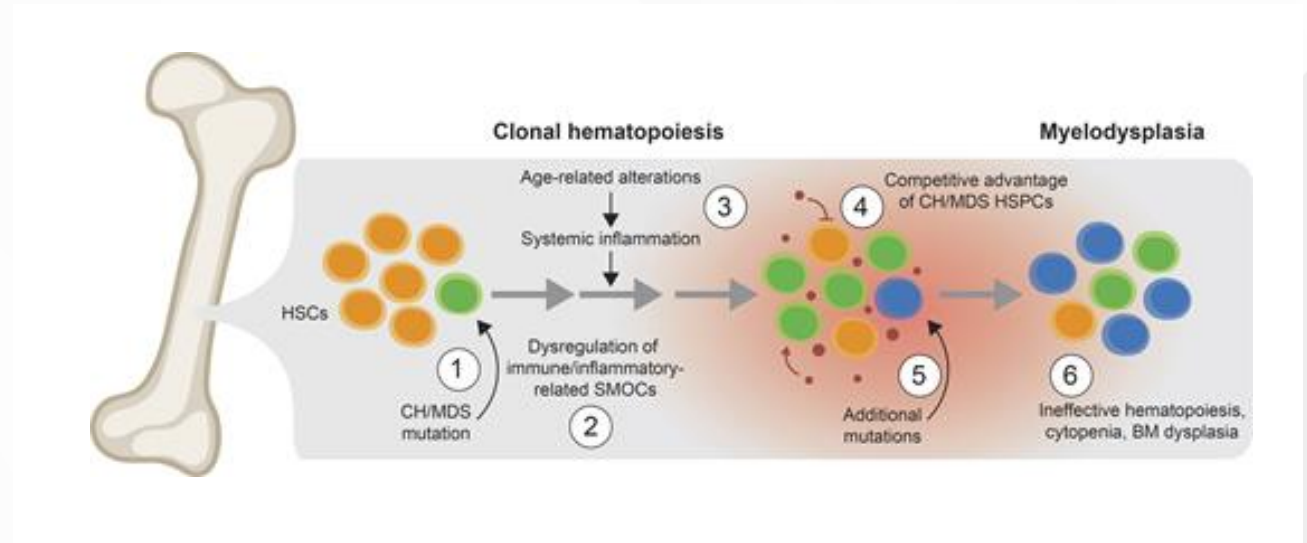
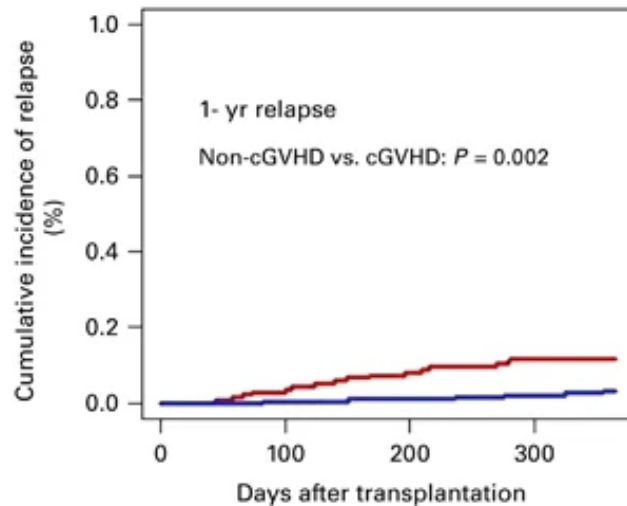
No. at risk				
NGS _{g10} -negative	MAC	13	10	10
	RIC	15	13	12
NGS _{g10} -positive	MAC	12	10	8
	RIC	8	6	4

Innovative Therapeutics for High-Risk MDS

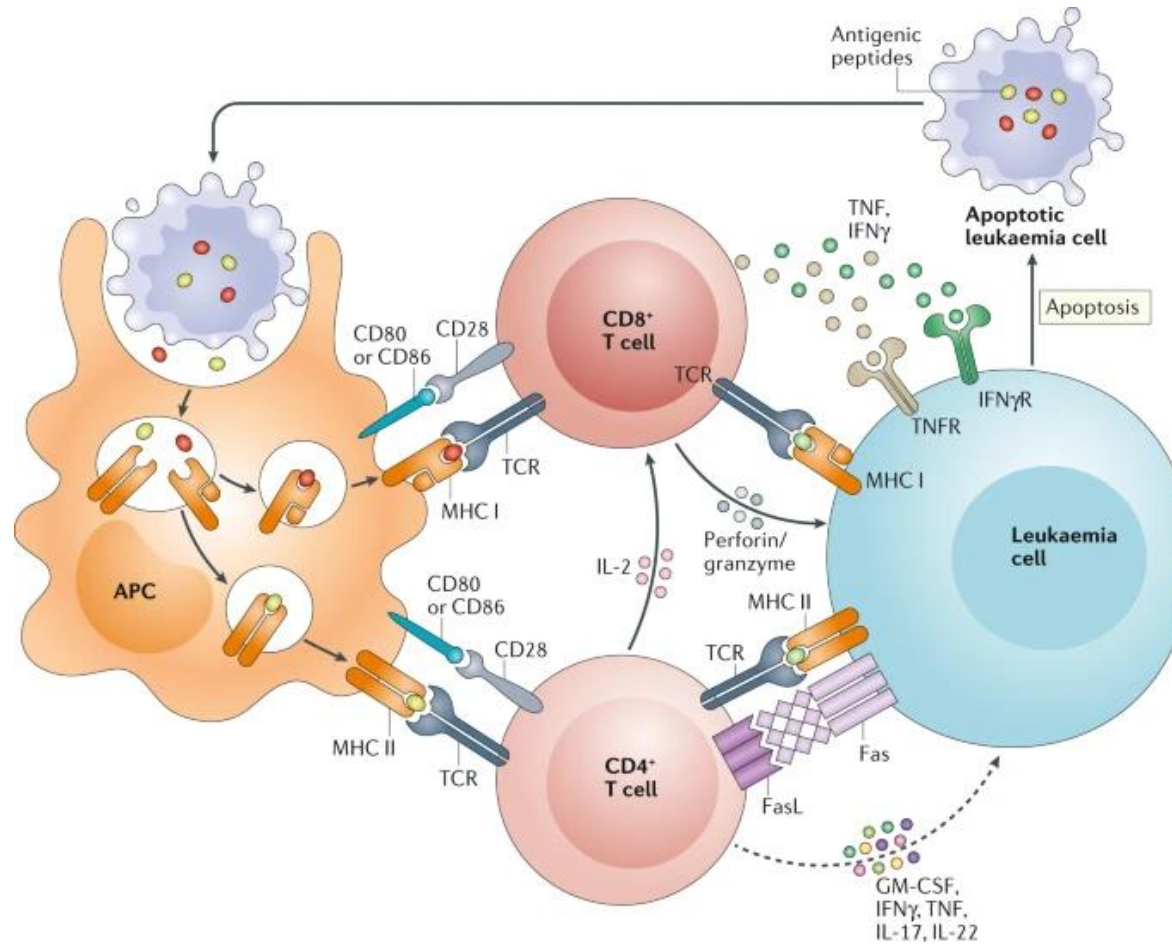
Rationale for Use and Integration Into Treatment Plans

Altered Immunity in MDS

- How the altered immune system may play a therapeutic role in MDS
- How an altered immune system may impact MDS pathogenesis



Allogeneic Transplant and MDS/AML

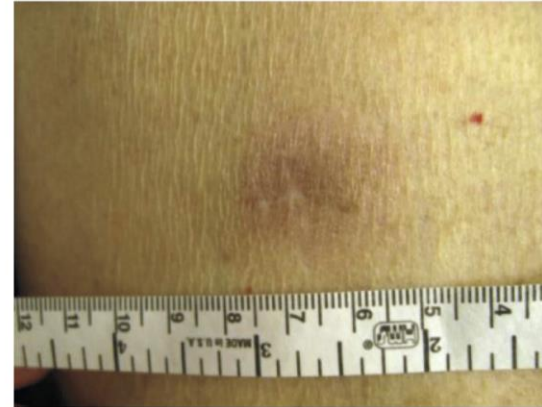


Immune Checkpoint Inhibition in MDS/AML

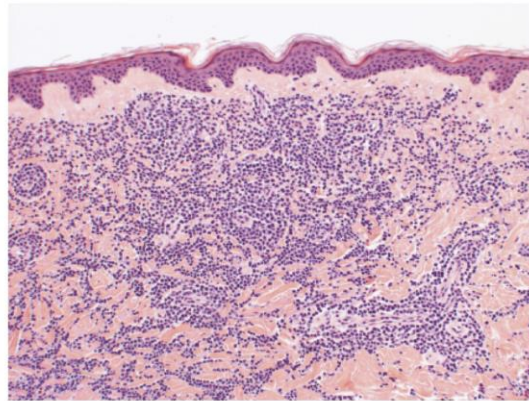
A Before Treatment



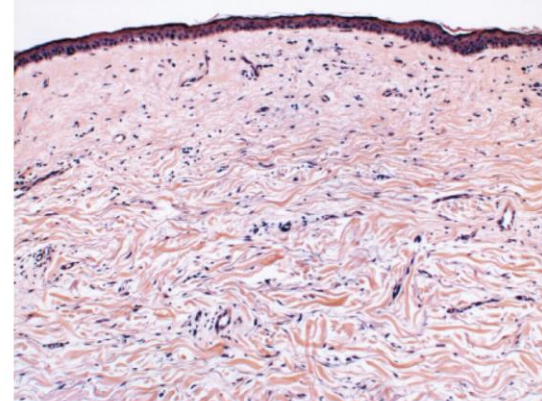
B After Treatment



C Before Treatment



D After Treatment



Challenges with Canonical ICI in MDS

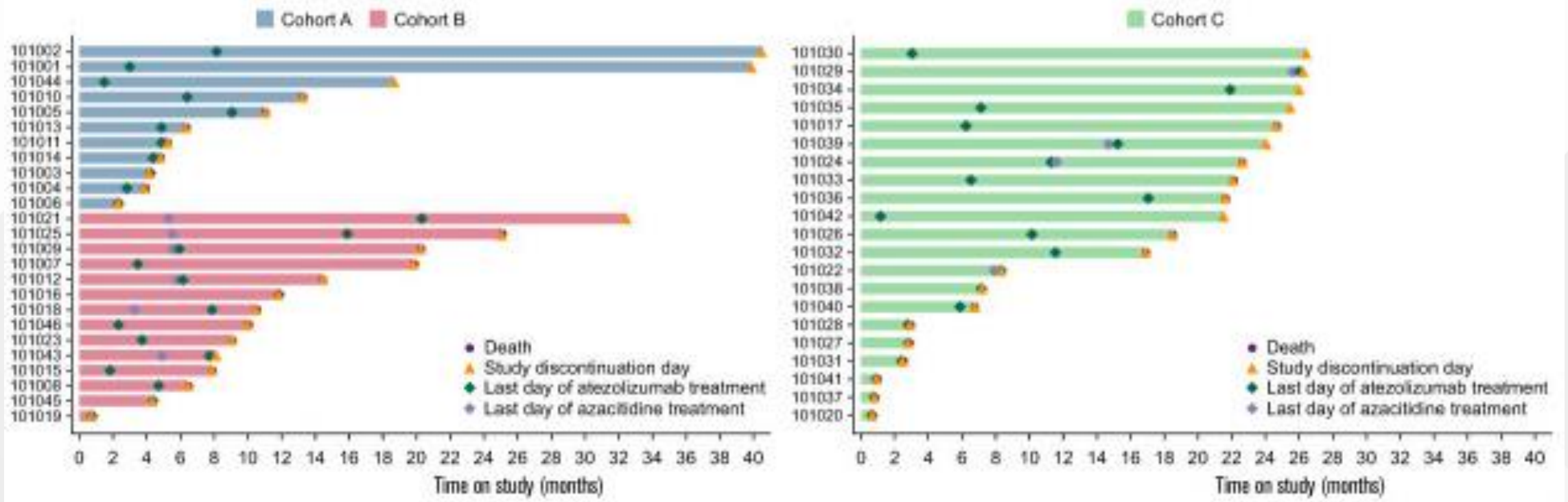
Response	Arm A (azacitidine + durvalumab) (N = 42)		Arm B (azacitidine) (N = 42)		<i>P</i>
	No. (%)	95% CI	No. (%)	95% CI	
ORR (CR + PR + mCR + HI)	26 (61.9)	47.22-76.59	20 (47.6)	32.51-62.72	.1838
CR	3 (7.1)	0.00-14.93	4 (9.5)	0.65-18.40	
mCR	15 (35.7)	21.22-50.21	8 (19.0)	7.17-30.92	
PR	0		0		
HI only	8 (19.0)	7.17-30.92	8 (19.0)	7.17-30.92	
SD	6 (14.3)		3 (7.1)		

Median OS (11.6 months vs 16.7 months; *P* = .74)

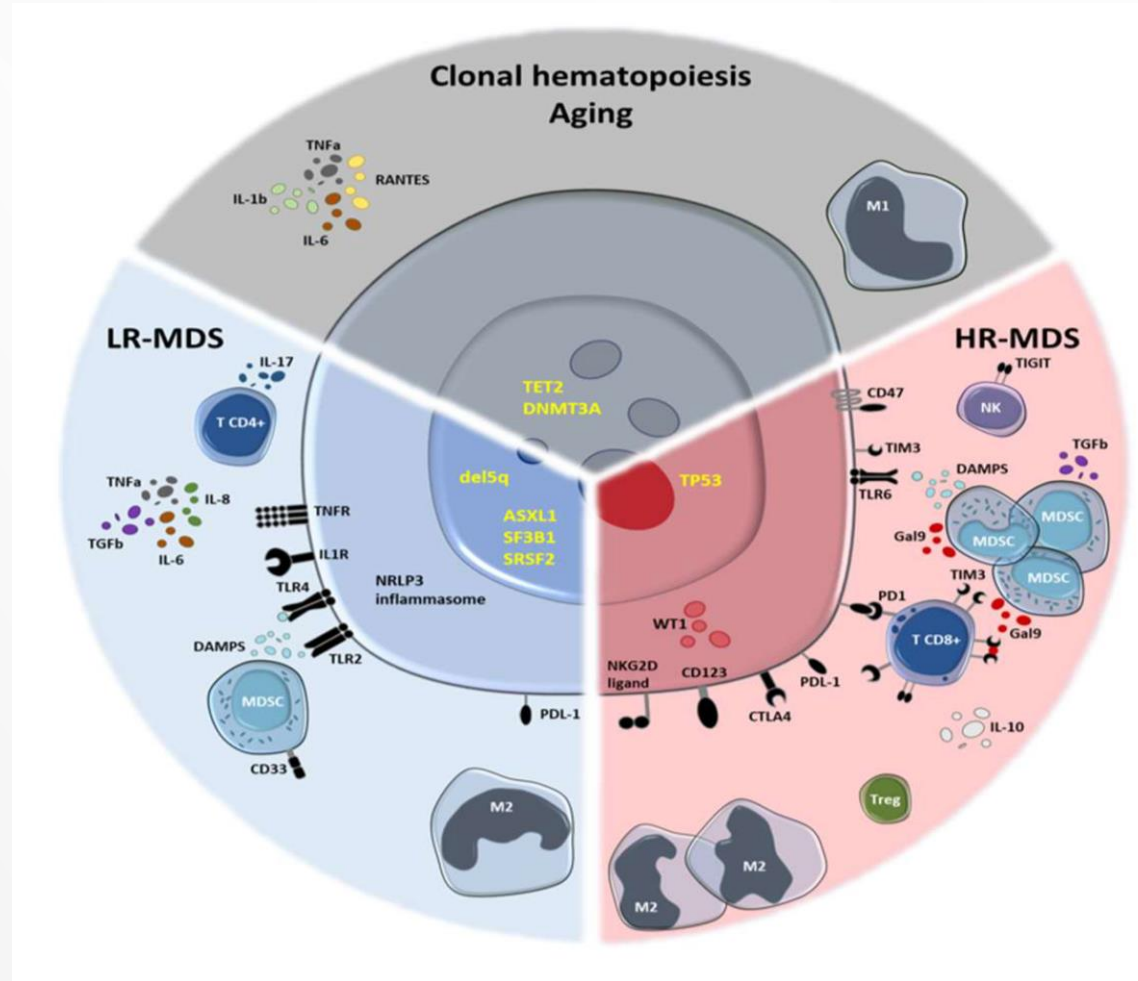
Median PFS (8.7 months vs 8.6 months; *P* = .93)

Challenges with Canonical ICIs in MDS

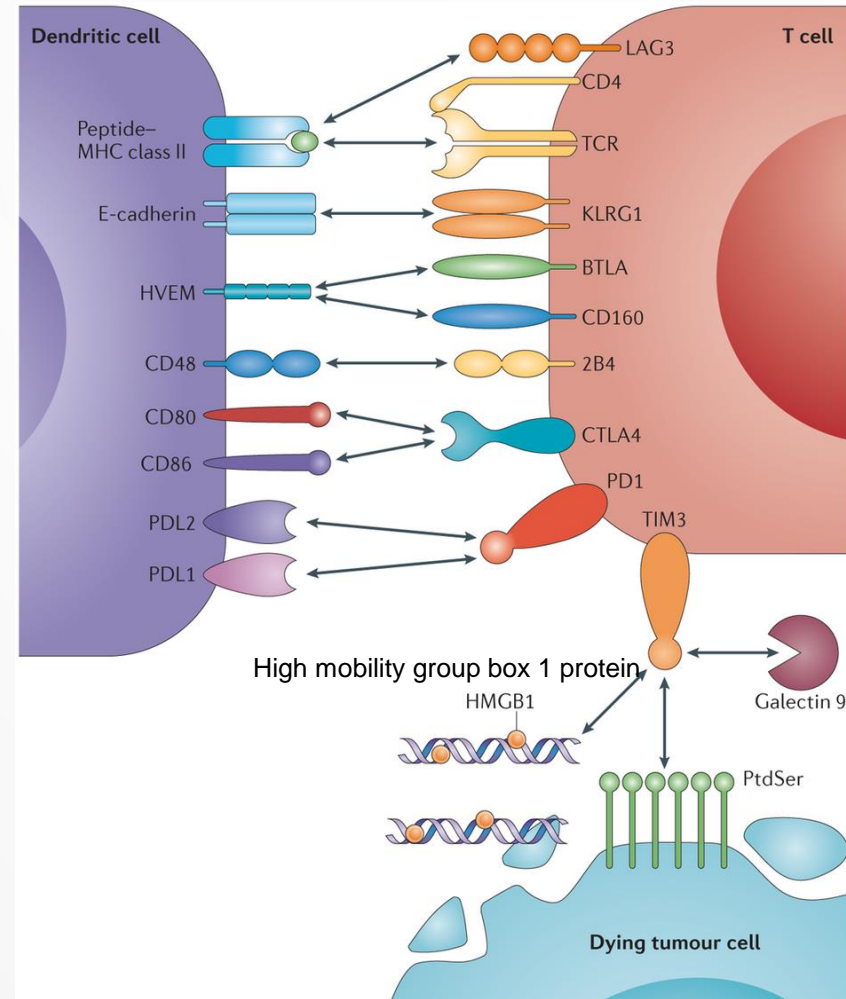
- The combination of atezolizumab plus azacitidine in HMA-naïve patients was associated with high early mortality rates, which led to early study termination
- Atezolizumab alone or in combination with azacitidine had limited clinical activity in patients with MDS previously exposed to HMAs, although this was without excessive or unexpected toxicity



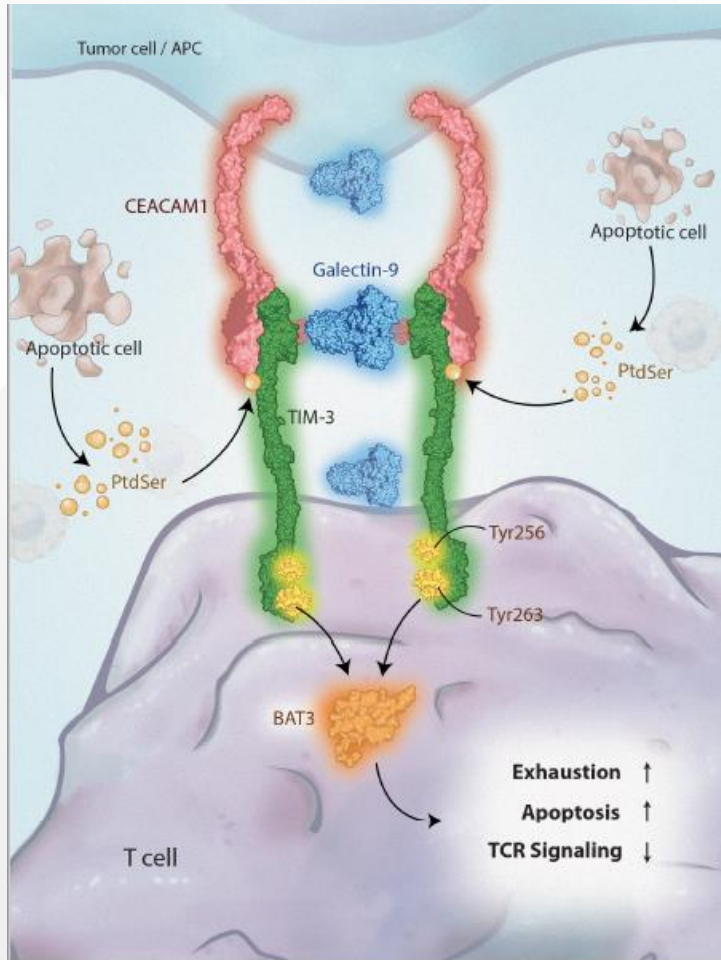
Immune Key Hubs Involved in Early Stages, Low-Risk MDS, and High-Risk MDS



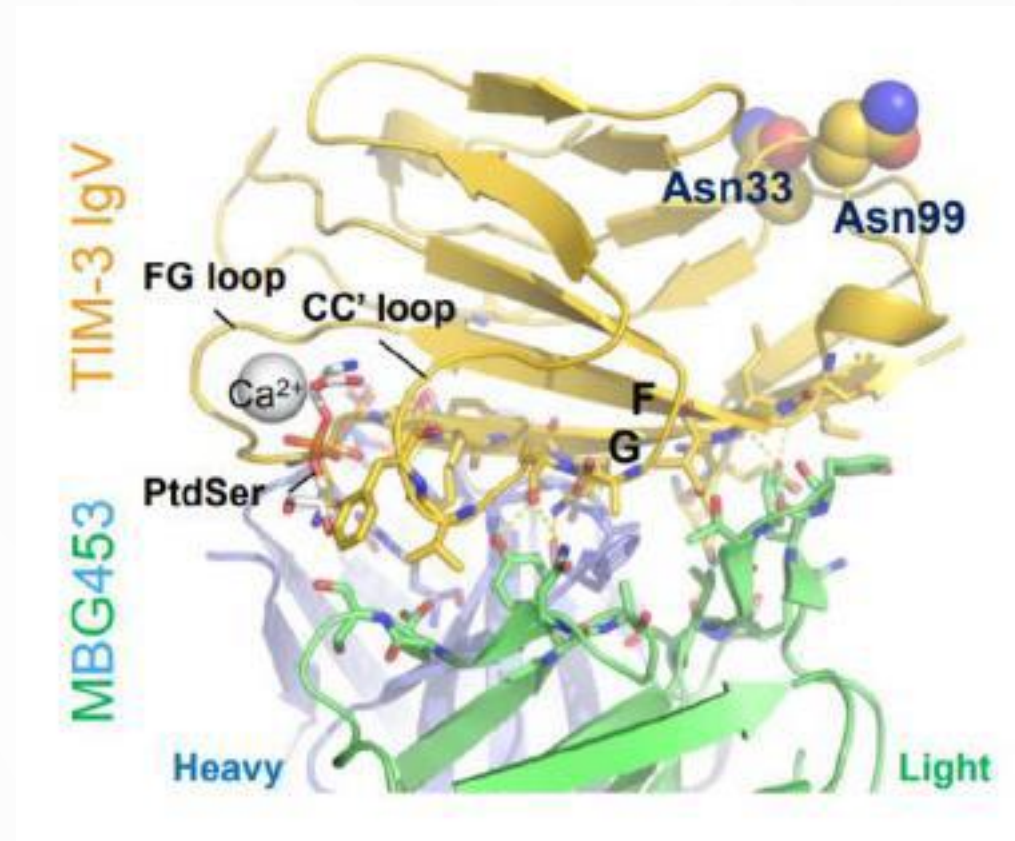
Clinical Blockade of PD-1 and LAG3: Potential Mechanisms of Action



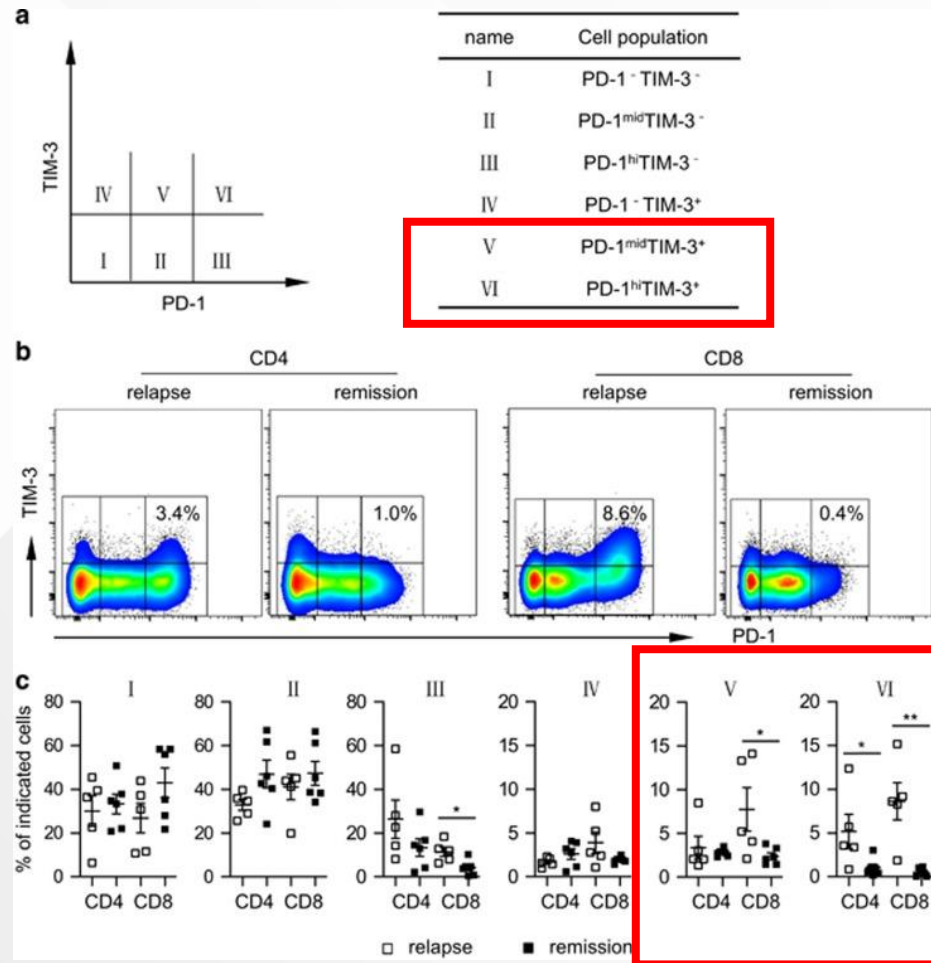
T-cell Immunoglobulin Domain and Mucin Domain 3



Sabatolimab (MBG453)

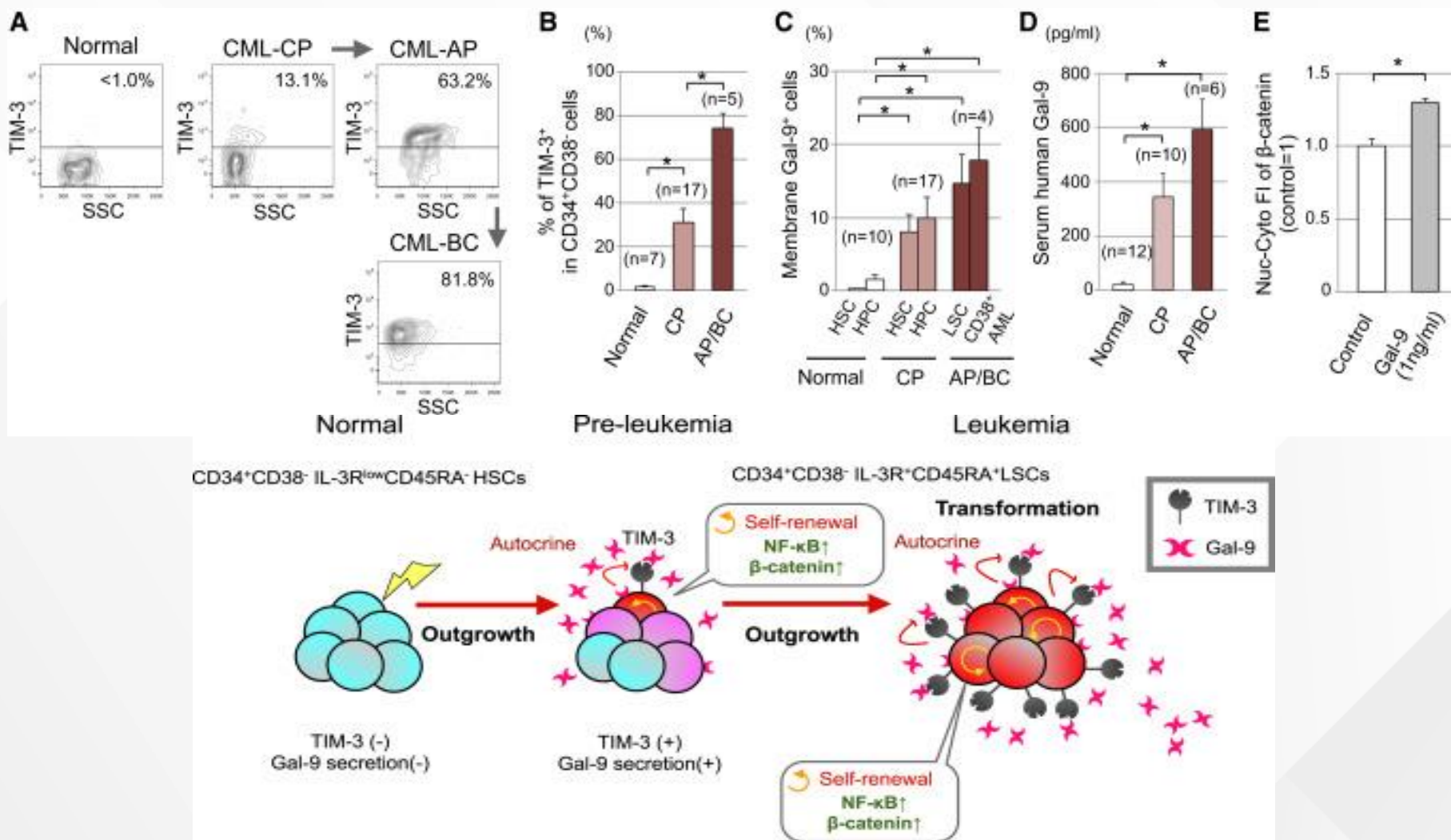


TIM3/PD-1 Expression and Post-HCT Relapse



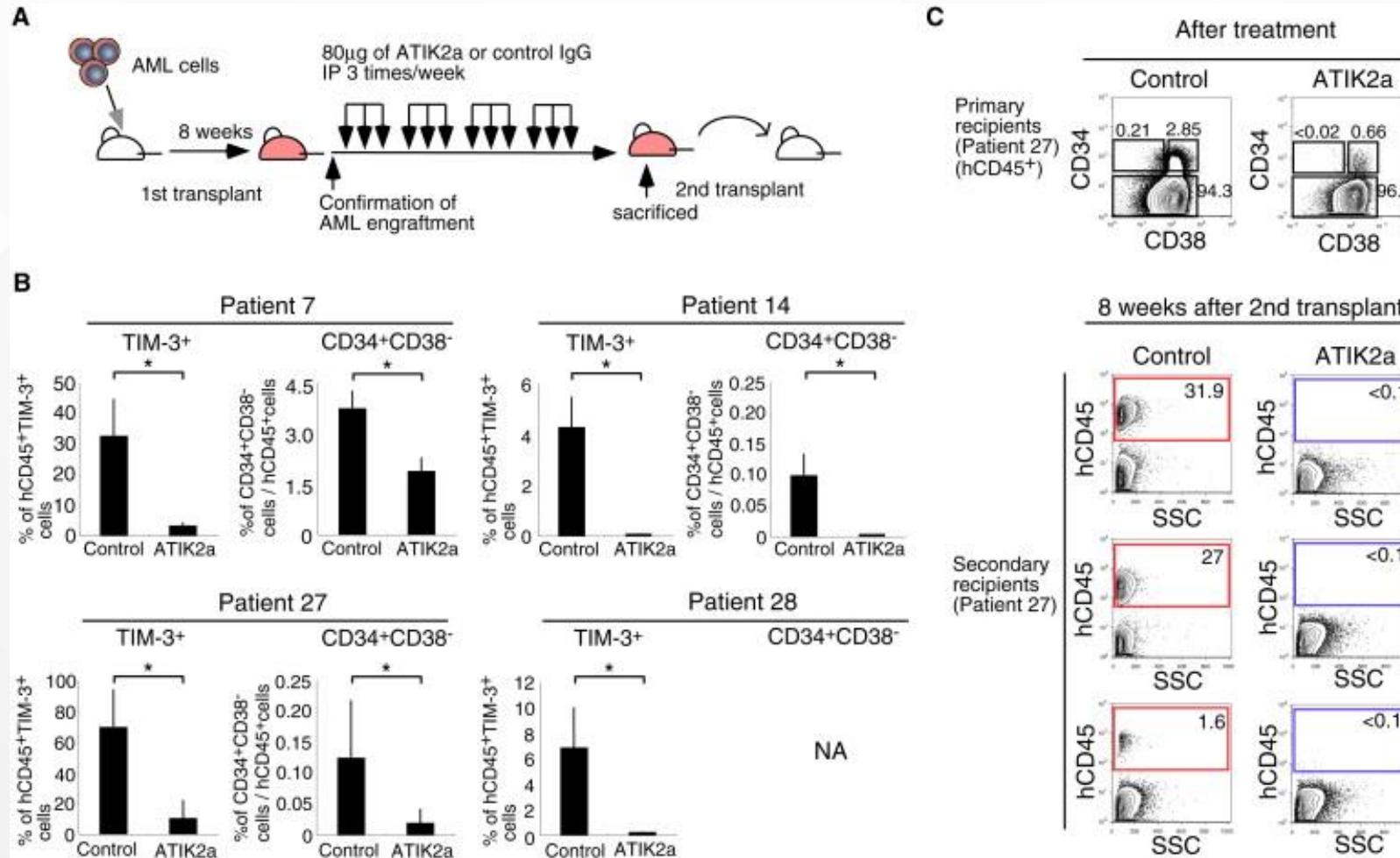
- Increase in PD-1⁺/TIM3⁺ CD8 T-cells at relapse
- Consistent with immune evasion and TIM3 upregulation playing a potential role

TIM3 Expression on Leukemic Progenitors



CML-CP, chronic myelogenous leukemia-chronic phase; CML- AP, chronic myelogenous leukemia-accelerated phase; CML-BC, chronic myelogenous leukemia-blast crisis; HSCs, hematopoietic stem cells; LCSs, leukemic stem cells; TIM3, T-cell immunoglobulin mucin-3. Kikushige et al. *Cell Stem Cell* 2015;17(3):341-352.

TIM3 Inhibition in AML



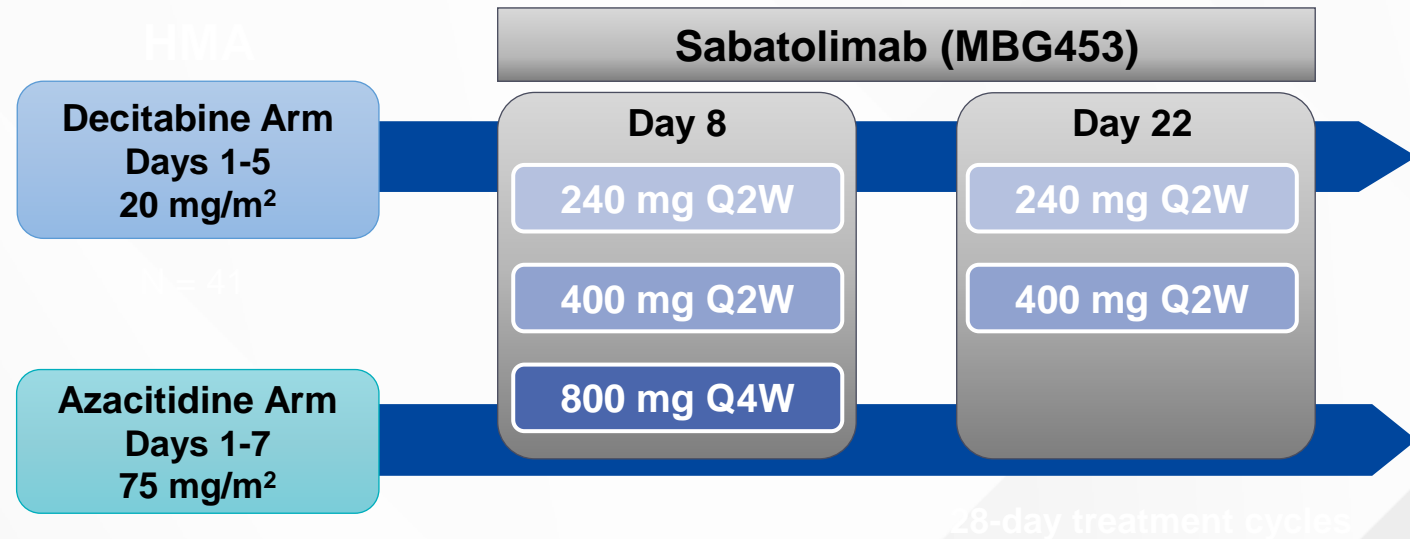
Phase 1b Study of Sabatolimab + HMA in MDS and AML

vHR/HR-MDS: IPSS-R
high- or very high-risk MDS

ND-AML: Unfit, newly
diagnosed AML, ineligible
for standard chemotherapy

No prior HMA treatment

ClinicalTrials.gov Identifier: **NCT03066648^a**



Primary Endpoints:

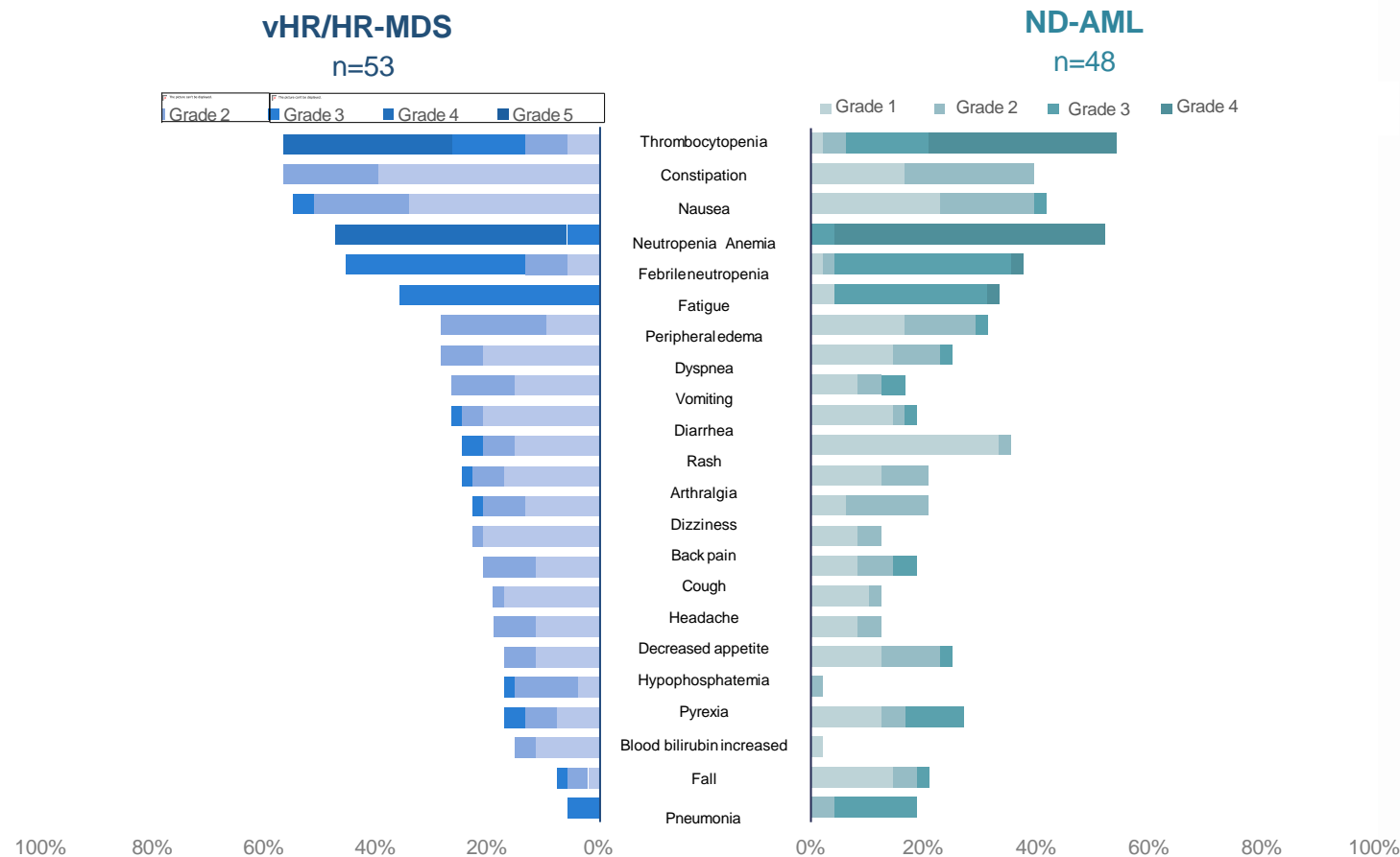
Maximum tolerated dose/recommended dose, safety, and tolerability

Secondary Endpoints:

Preliminary efficacy: Response rates and duration of response

Most Commonly Occurring Adverse Events

(≥15% in either population, regardless of relationship to treatment)



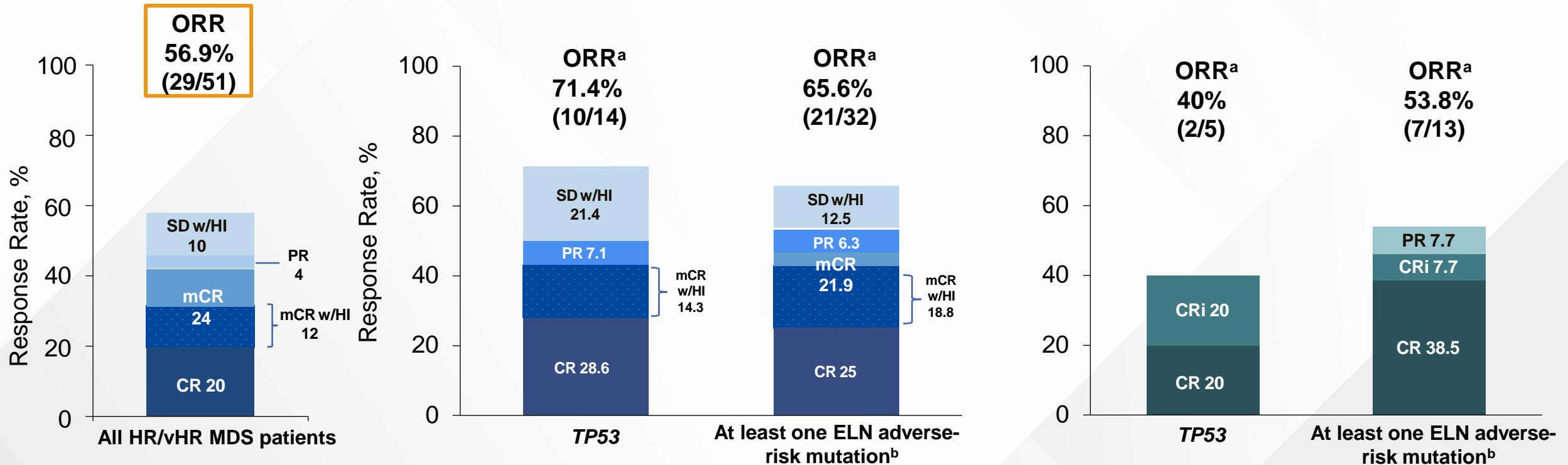
vHR/HR-MDS and ND-AML AEs

- Most common reported AEs were cytopenias, GI symptoms, fatigue
- Low rate of sabatolimab dose modification:
 - 1/101 (1%) patients had dose reduction
- 38/101 (38%) patients had dose interruption (cycle delay >7d) due to AE
 - No patient with vHR/HR-MDS and only 3 with ND-AML discontinued treatment due to an AE
- One patient with neutropenic colitis reported as suspected to be related to study treatment died of septic shock. No other treatment-related deaths were reported
- No DLTs in vHR/HR-MDS and only 1 in ND-AML

Sabatolimab + HMA Response Rates

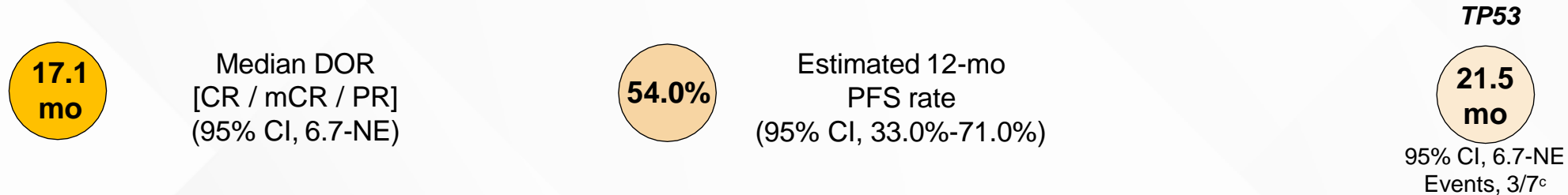
vHR/HR-MDS

ND-AML



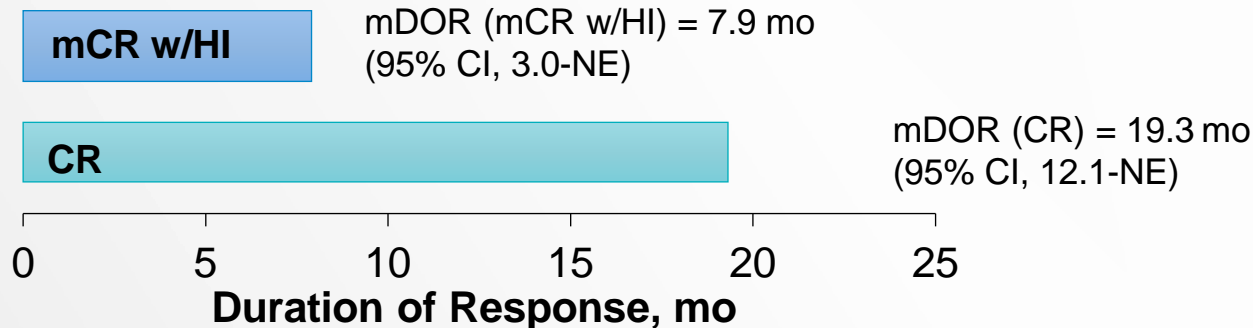
^aEvaluable patients included patients with a valid baseline and at least 1 postbaseline bone marrow assessment or if they had disease progression or disease-related death prior to the first marrow assessment.
 CR, complete response; CRi, CR with incomplete blood recovery; ELN, European LeukemiaNet; HI, hematologic improvement; HMA, hypomethylating agent; mCR, marrow complete response;
 ND-AML, newly diagnosed acute myeloid leukemia; ORR, overall response rate; PR, partial response; SD, stable disease; HR, high risk; vHR, very high risk. Brunner et al. *Blood* 2021;138(suppl 1):244.

Duration of Responses to Sabatolimab + HMA in MDS

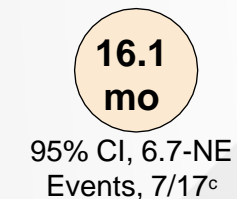


Median Duration of Response by response category

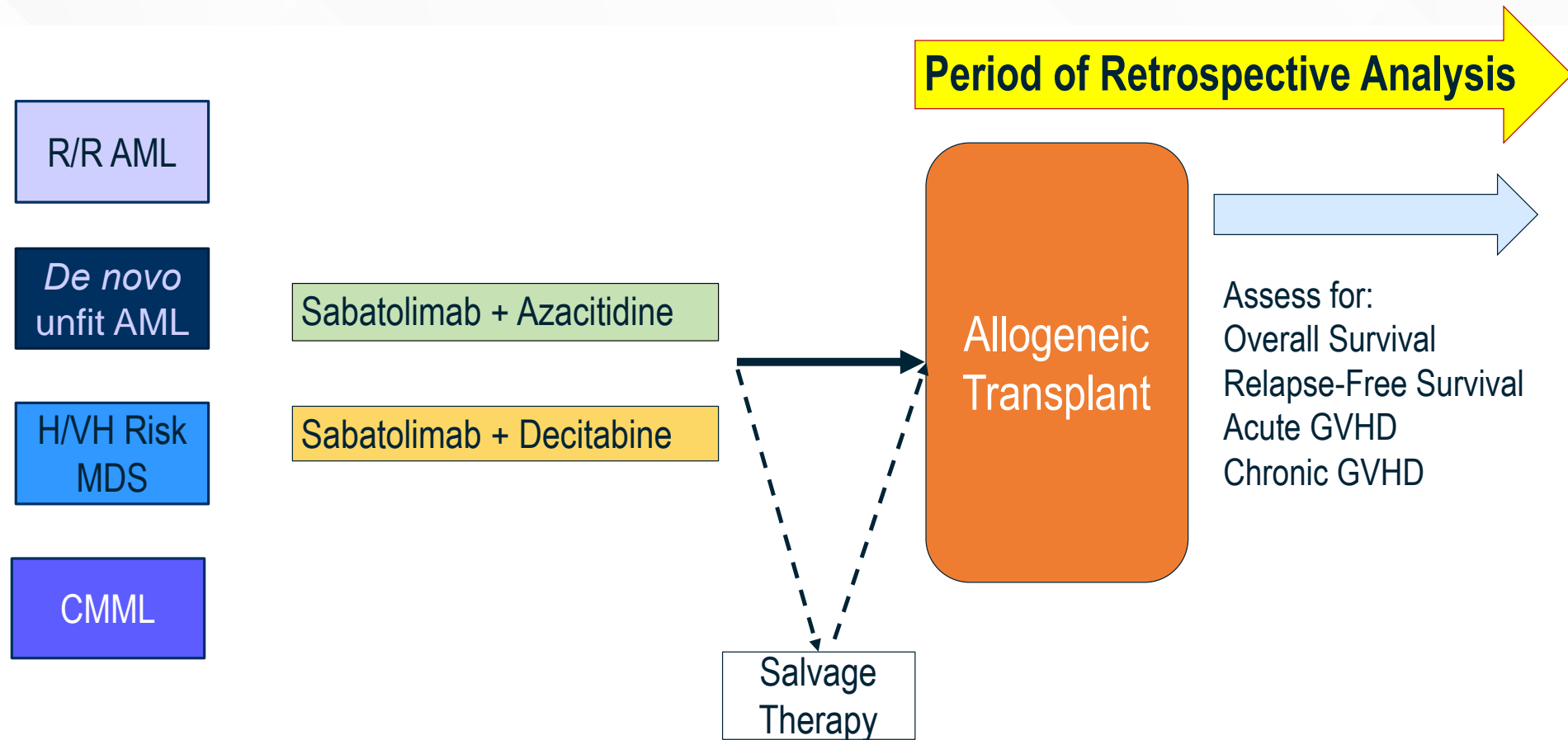
mDOR for mCR and PR could not be estimated



At least one ELN adverse-risk mutation^b



Retrospective Study: Transplant After Sabatolimab Exposure

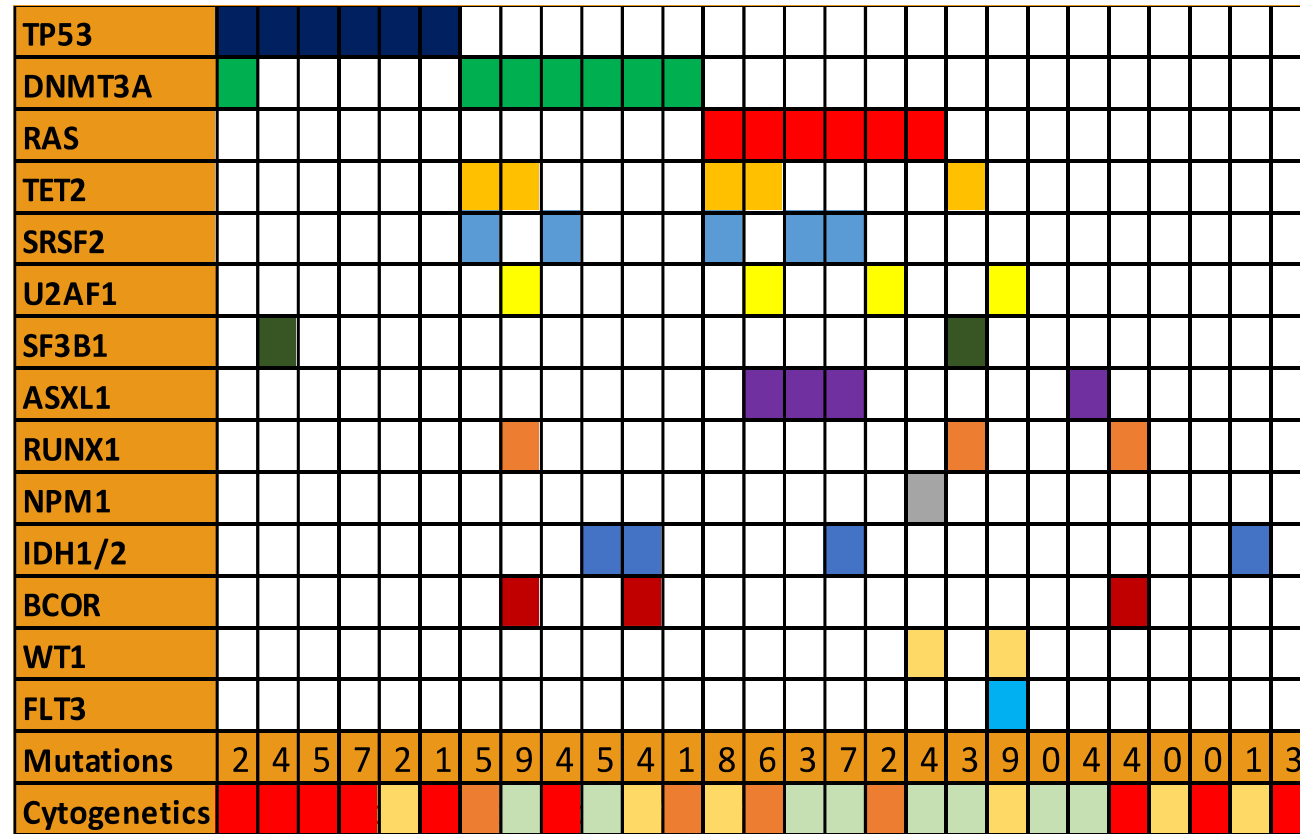


Patient Characteristics

Age (median, range)	67 (23-77)
Male Sex	18 (64%)
WHO Category	
AML	6 (21%)
MDS	19 (68%)
CMML	3 (11%)
Cytogenetic Risk	
Intermediate	14 (52%)
Normal	8 (30%)
Adverse	13 (48%)
Complex	9 (33%)
IPSS-R (median, range)	5.5 (3.5-9.0)
ELN High Risk Mutation	14 (50%)

HMA Therapy	
Azacitidine	16 (57%)
Decitabine	12 (43%)
Best Overall Response Prior to HCT	
CR	10 (36%)
mCR/CRI	9 (32%)
PR/Hi	2 (8%)
NR/SD	7 (25%)
Conditioning Intensity	
MAC	4 (17%)
RIC	20 (83%)
Donor Source	
MRD	6 (21%)
MUD	18 (64%)
MMUD/Haplo	4 (14%)

Molecular Profiling Identified Several Very High-Risk Molecular Features

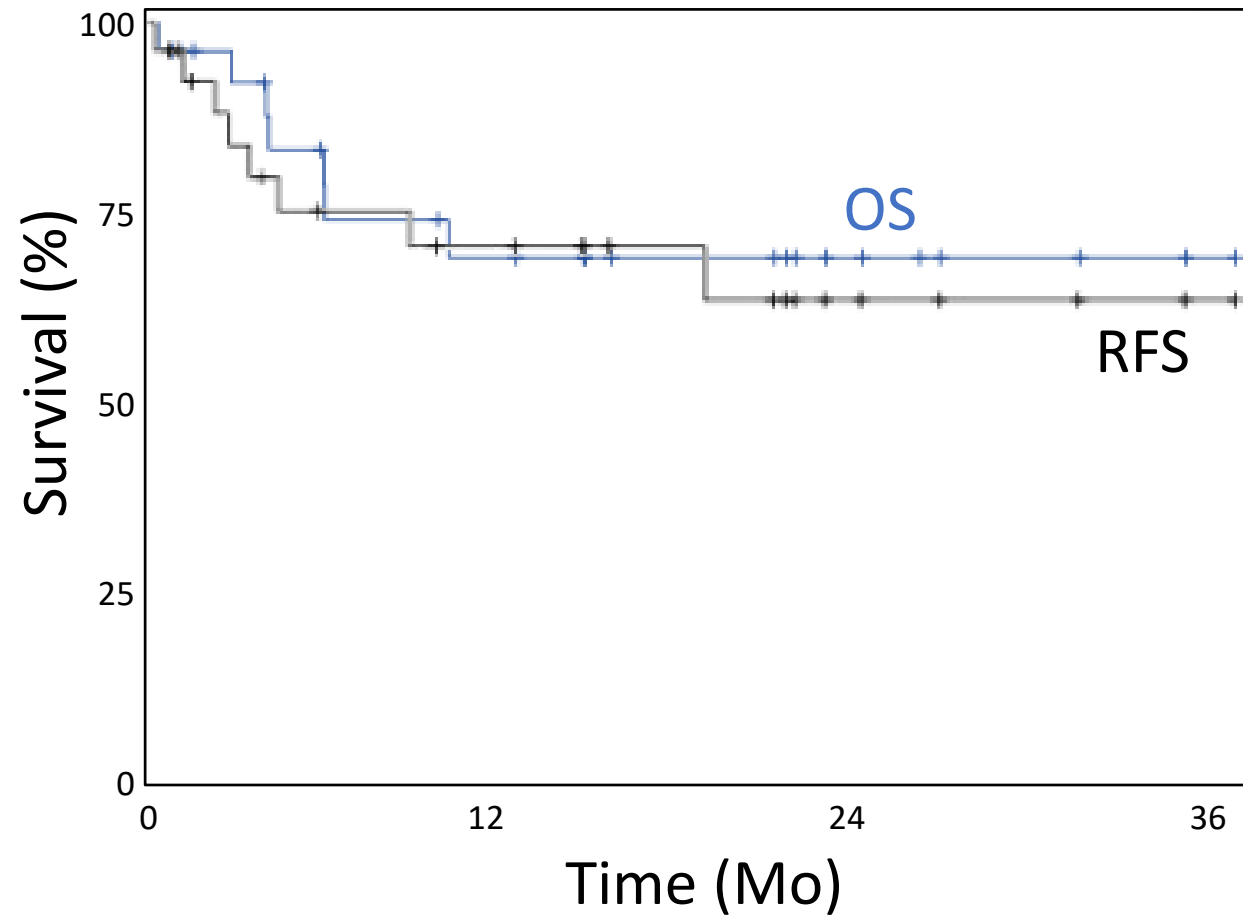


Cytogenetics: ■ Complex ■ Adverse ■ Intermediate ■ Normal

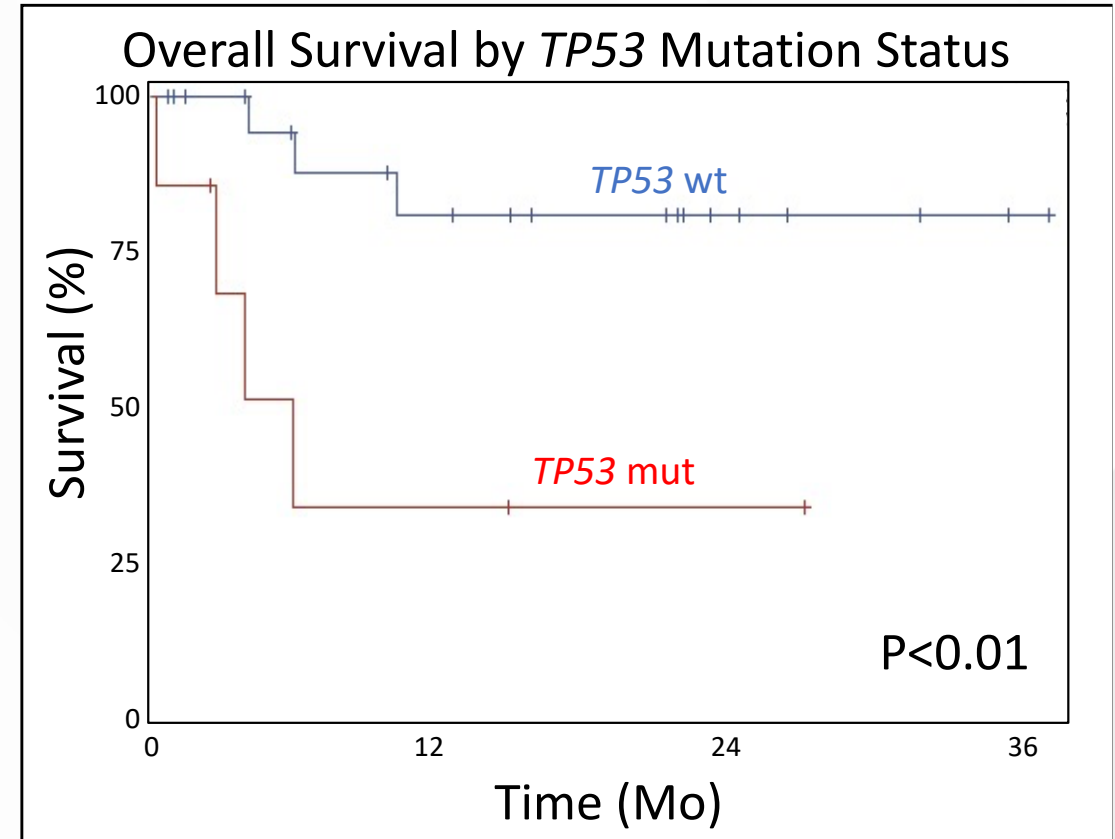
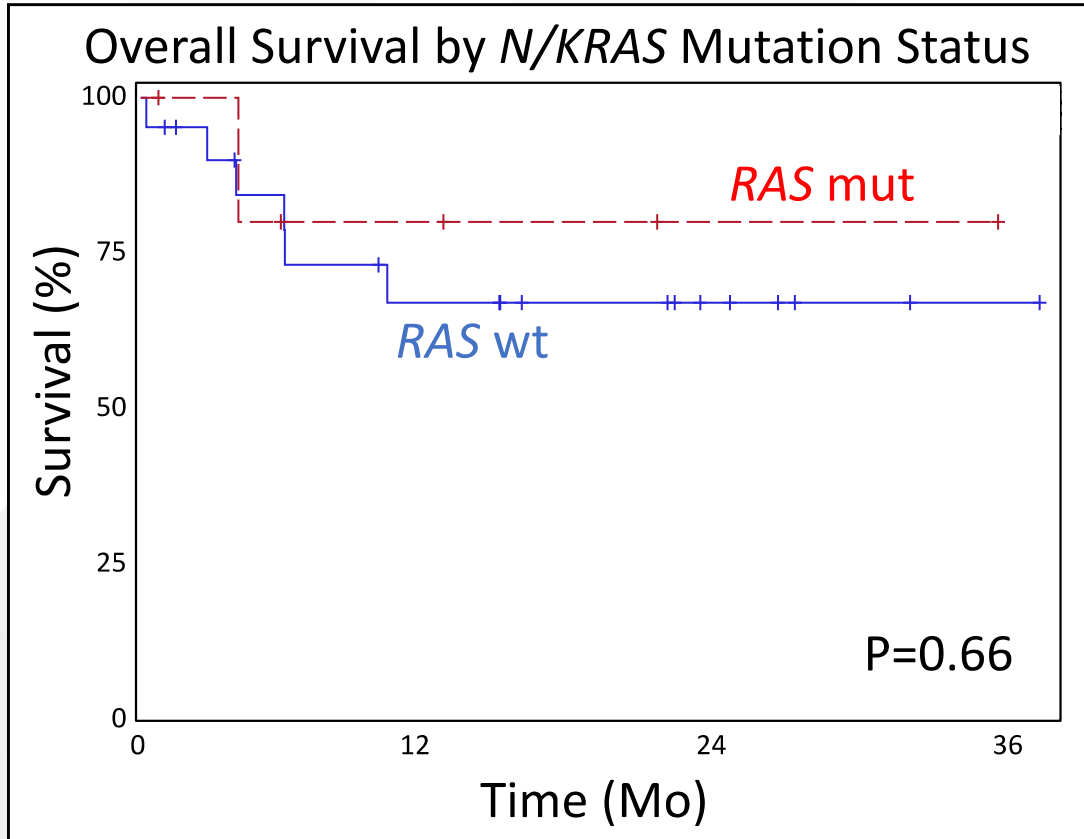
Investigator Reported GVHD Events

- Acute GVHD was seen in 16 patients; maximum grade 3-4 aGVHD occurred in 4 patients:
 - 2 patients with stage 4 GI disease, 1 with stage 3 GI disease, and 1 patient with stage 4 skin GVHD
 - One patient died on hospice after G4 aGVHD
- Chronic GVHD requiring systemic immunosuppression was seen in 8 patients, none of which have died or relapsed
- One patient also received spartalizumab (PD-1) and had grade 2 skin aGVHD and no cGVHD

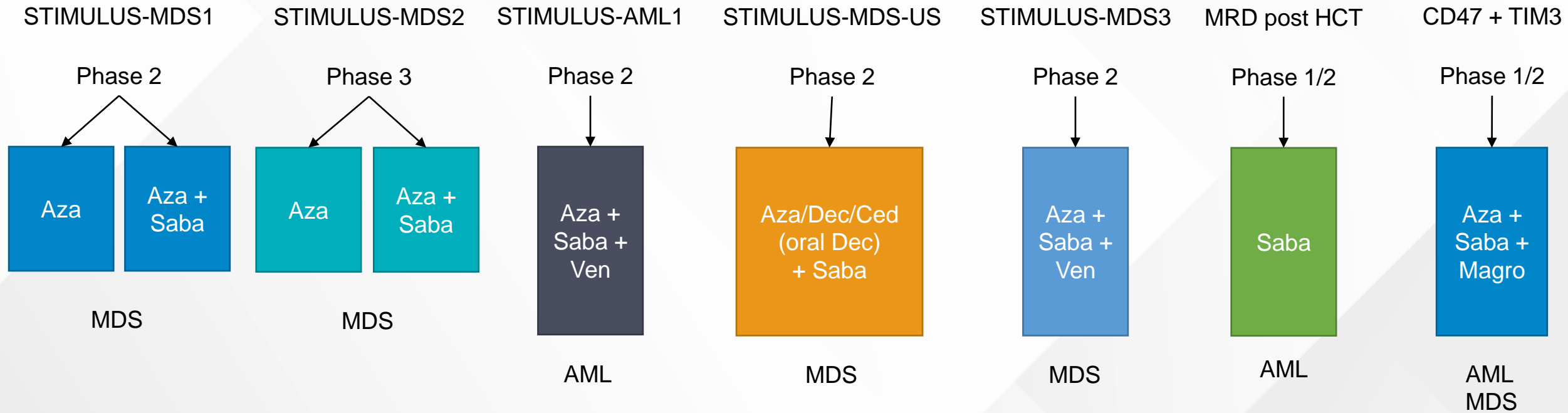
Overall and Relapse-Free Survival Post-HCT



Overall Survival by Mutation Status

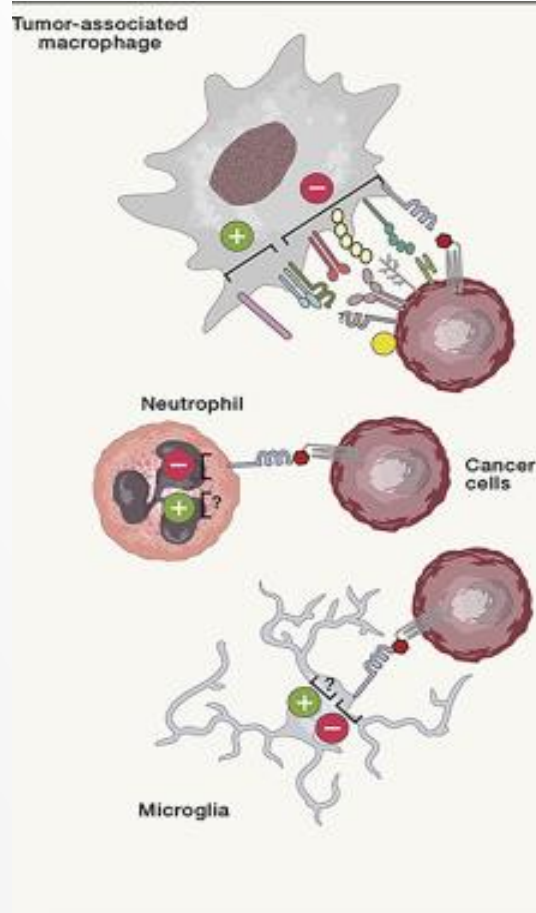


Sabatolimab in MDS and AML

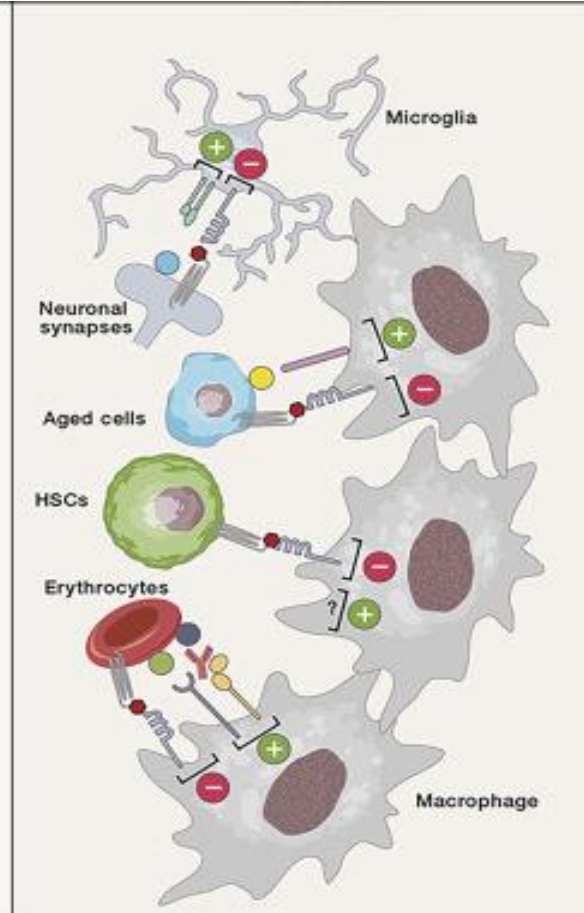


Targeting CD47 in MDS

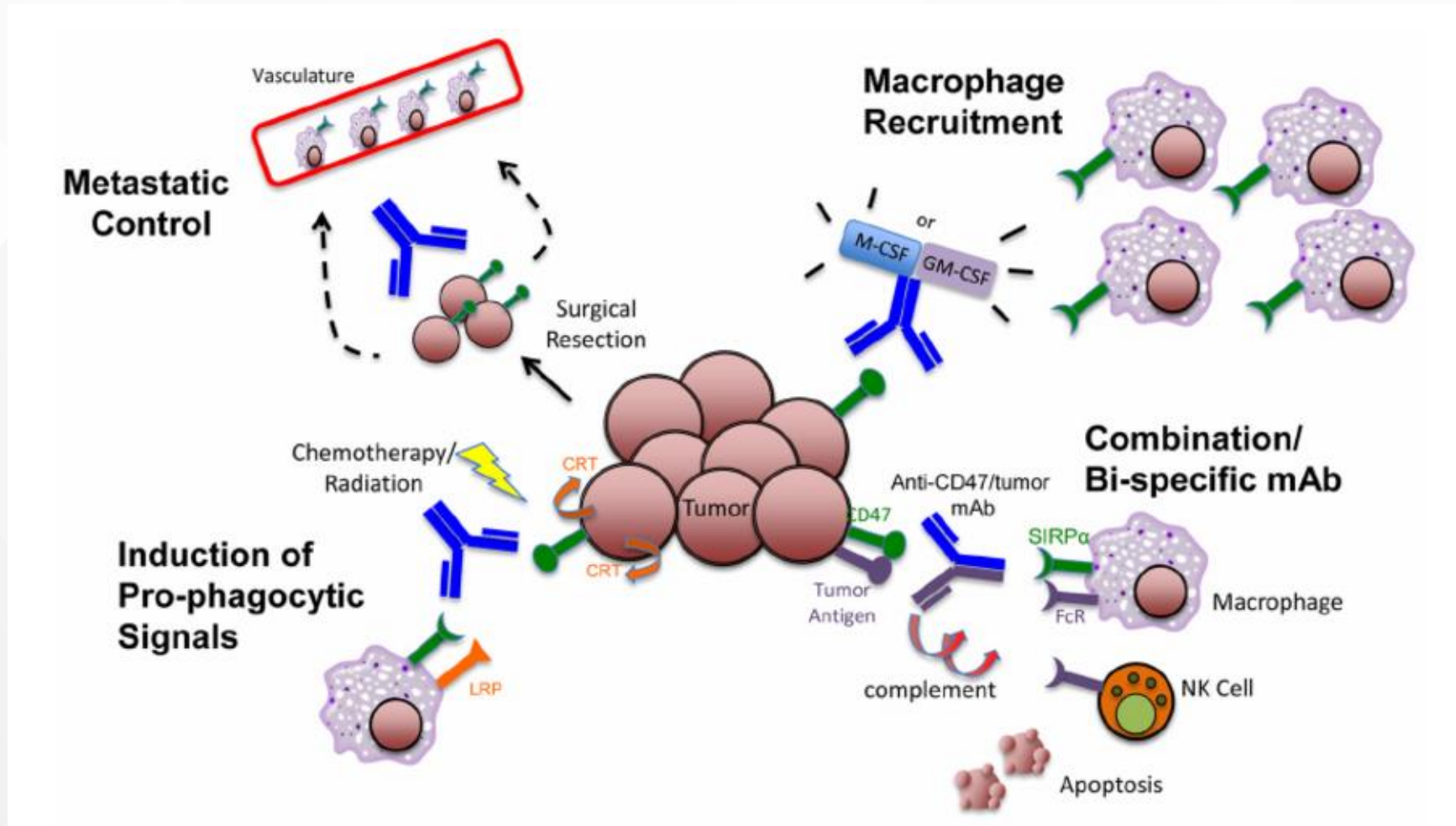
Cancer



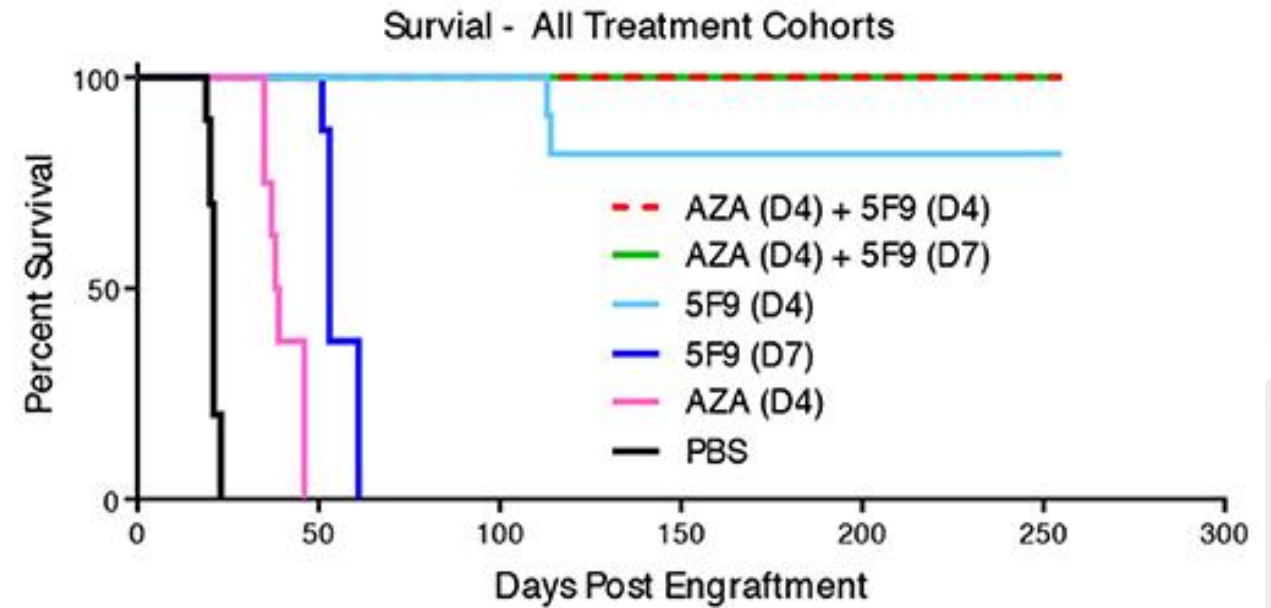
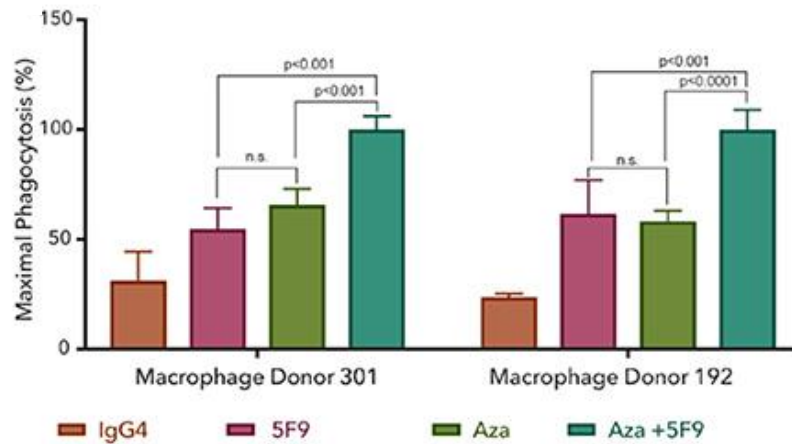
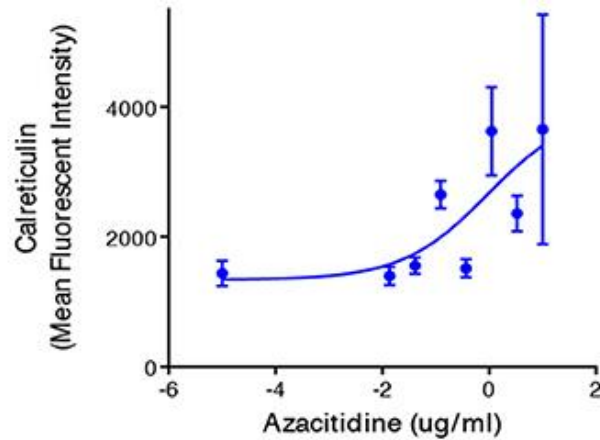
Tissue Hemostasis



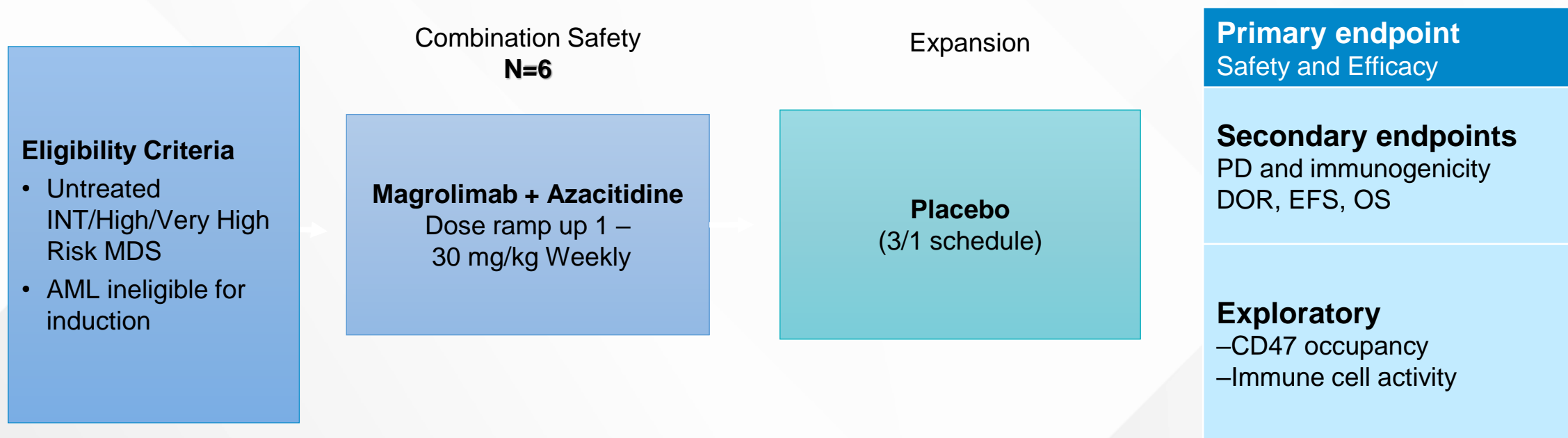
Targeting CD47 in MDS



CD47 and Azacitidine



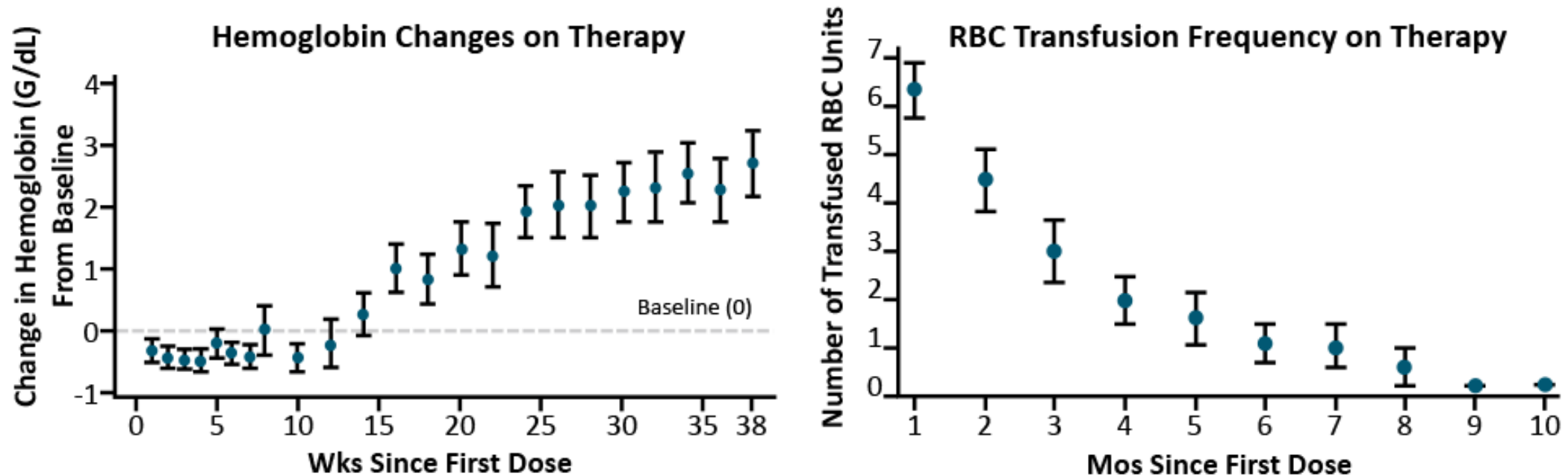
Magrolimab and Azacitidine for HR-MDS



Patient Characteristics

Characteristic	1L MDS 5F9+AZA (N=35)	1L AML 5F9+AZA (N=27)
Median age (range)	70 (47-80)	74 (60-89)
ECOG Performance Status: 0 1 2	13 (37%) 21 (60%) 1 (3%)	9 (33%) 16 (59%) 2 (7%)
Cytogenetic Risk: Favorable Intermediate Poor Unknown/missing	0 10 (29%) 23 (66%) 2 (6%)	0 2 (7%) 18 (67%) 7 (26%)
WHO AML classification: MRC Recurrent abnormalities Therapy-related NOS	—	19 (70%) 2 (7%) 1 (4%) 5 (19%)
WHO MDS classification: RS and single/multi-lineage dysplasia Multilineage dysplasia Excess blasts Unclassifiable/unknown/missing	3 (9%) 6 (17%) 19 (54%) 7 (20%)	—
IPSS-R (MDS): Intermediate High Very High Unknown/missing	11 (31%) 18 (51%) 5 (14%) 1 (3%)	—
Therapy-related MDS Unknown/missing	11 (31%) 1 (3%)	—
Harboring a <i>TP53</i> mutation	4 (11%)	11 (41%)

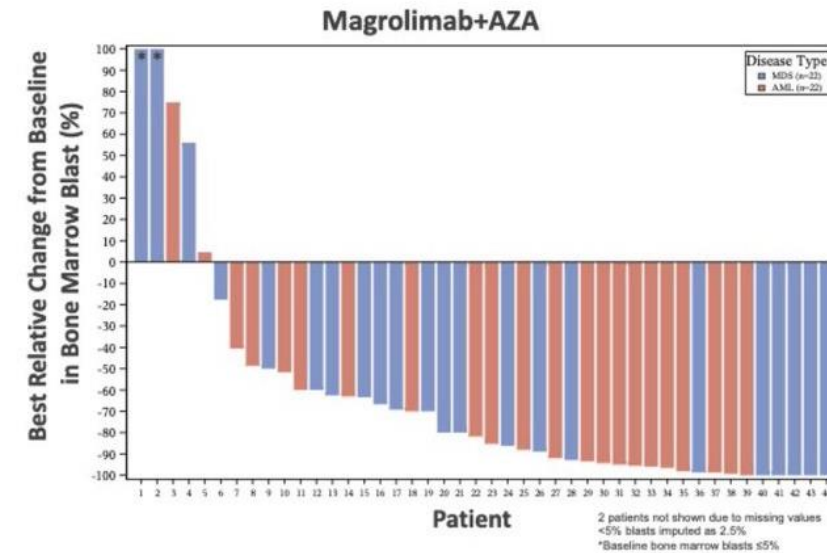
Hemoglobin and Transfusions with Magrolimab and Azacitidine



CD47 “Don’t Eat Me” Checkpoint

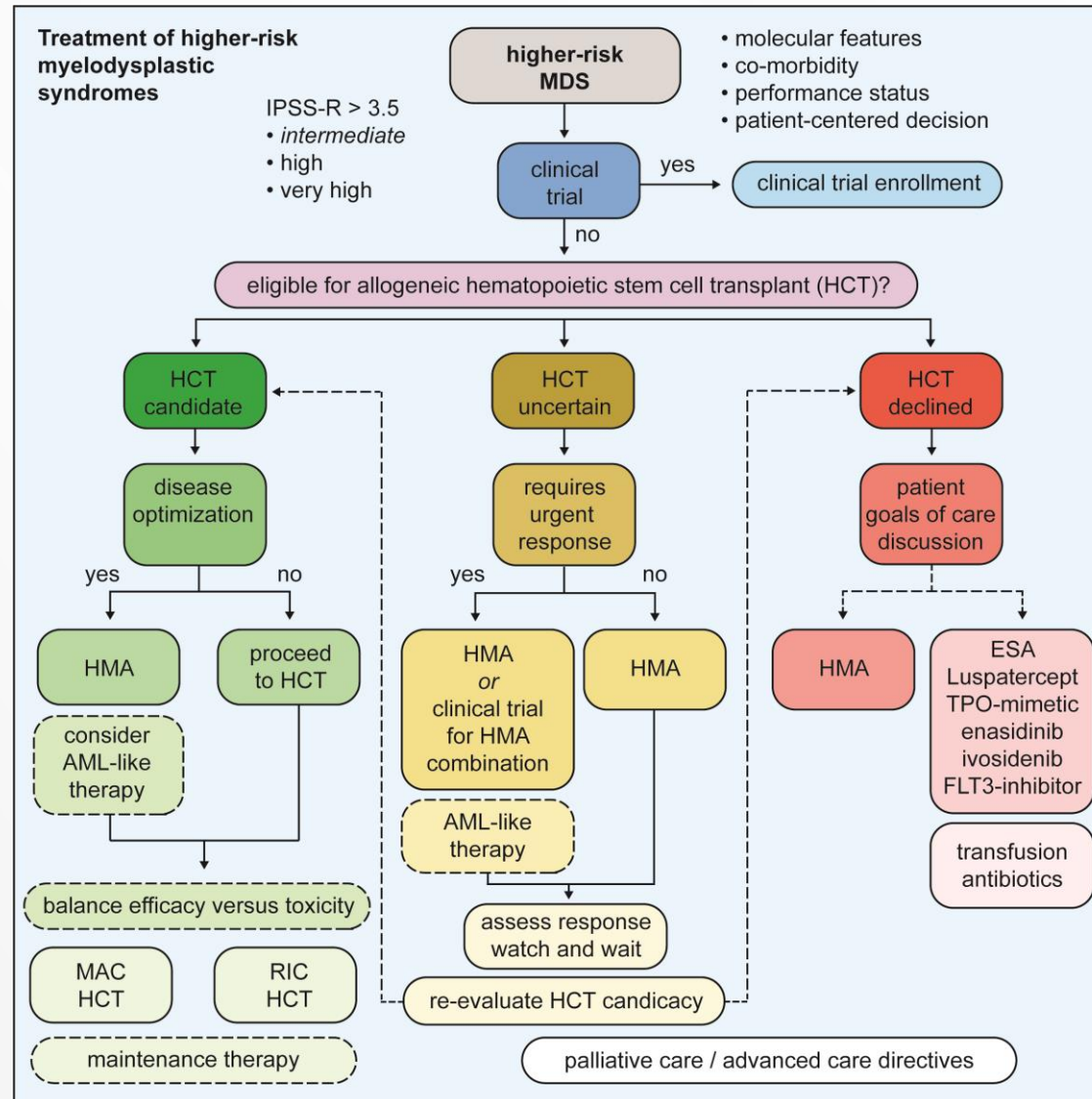
Anti-Leukemic Activity is Observed with Magrolimab + AZA in MDS and AML

Best Overall Response	1L MDS (N = 24)	1L AML (N=22)
ORR	22 (92%)	14 (64%)
CR	12 (50%)	9 (41%)
CRi	-	3 (14%)
PR	0	1 (5%)
MLFS/marrow CR	8 (33%) 4 with marrow CR = HI	1 (5%)
Hematologic improvement (HI)	2 (8%)	-
SD	2 (8%)	7 (32%)
PD	0	1 (5%)



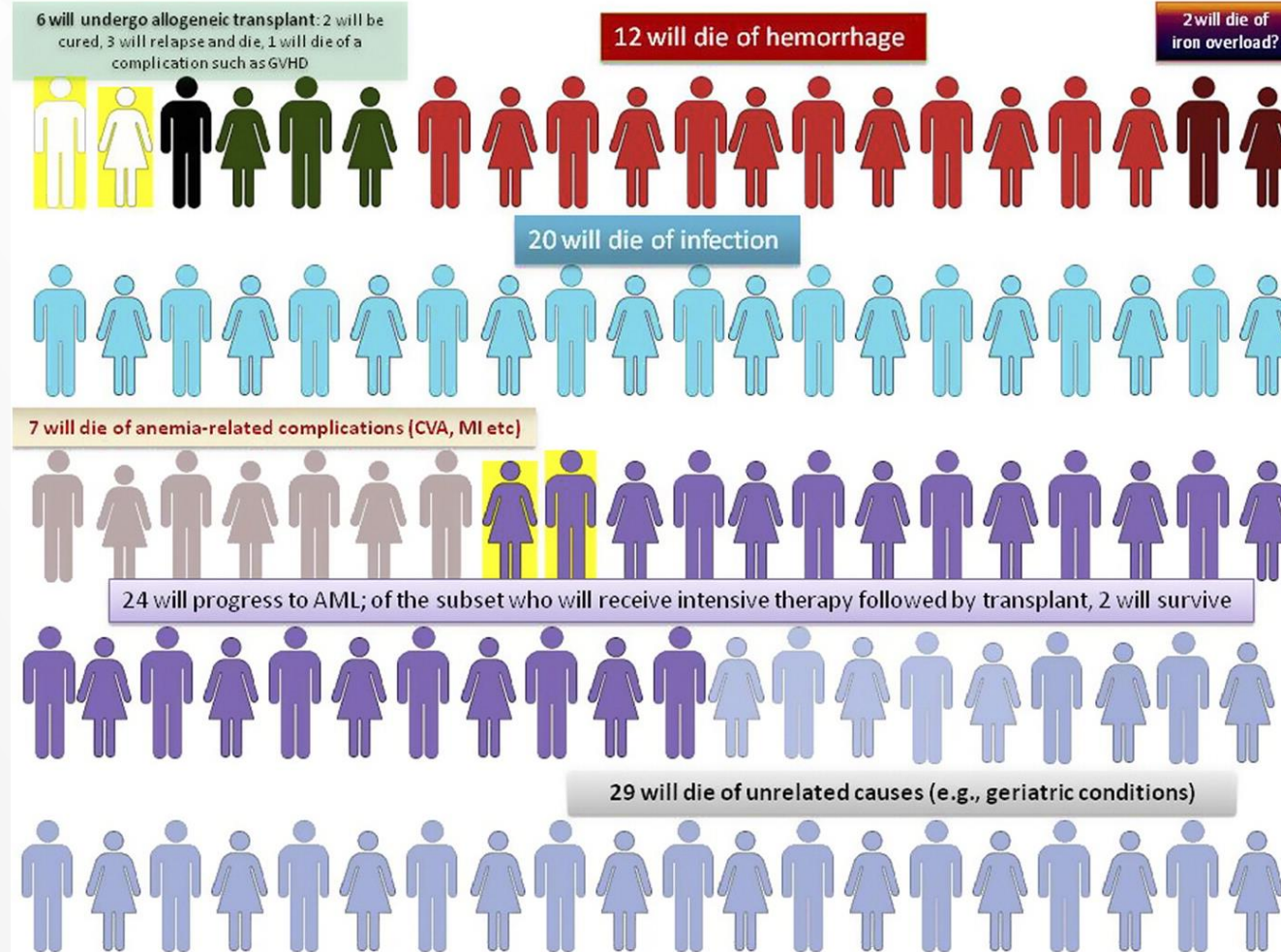
- Magrolimab + AZA induces a 92% ORR (50% CR) in MDS and 64% ORR (55% CR/CRi) in AML
- Median time to response is 1.9 months, more rapid than AZA alone
- Magrolimab + AZA efficacy compares favorably to AZA monotherapy

Treatment of Higher-Risk MDS

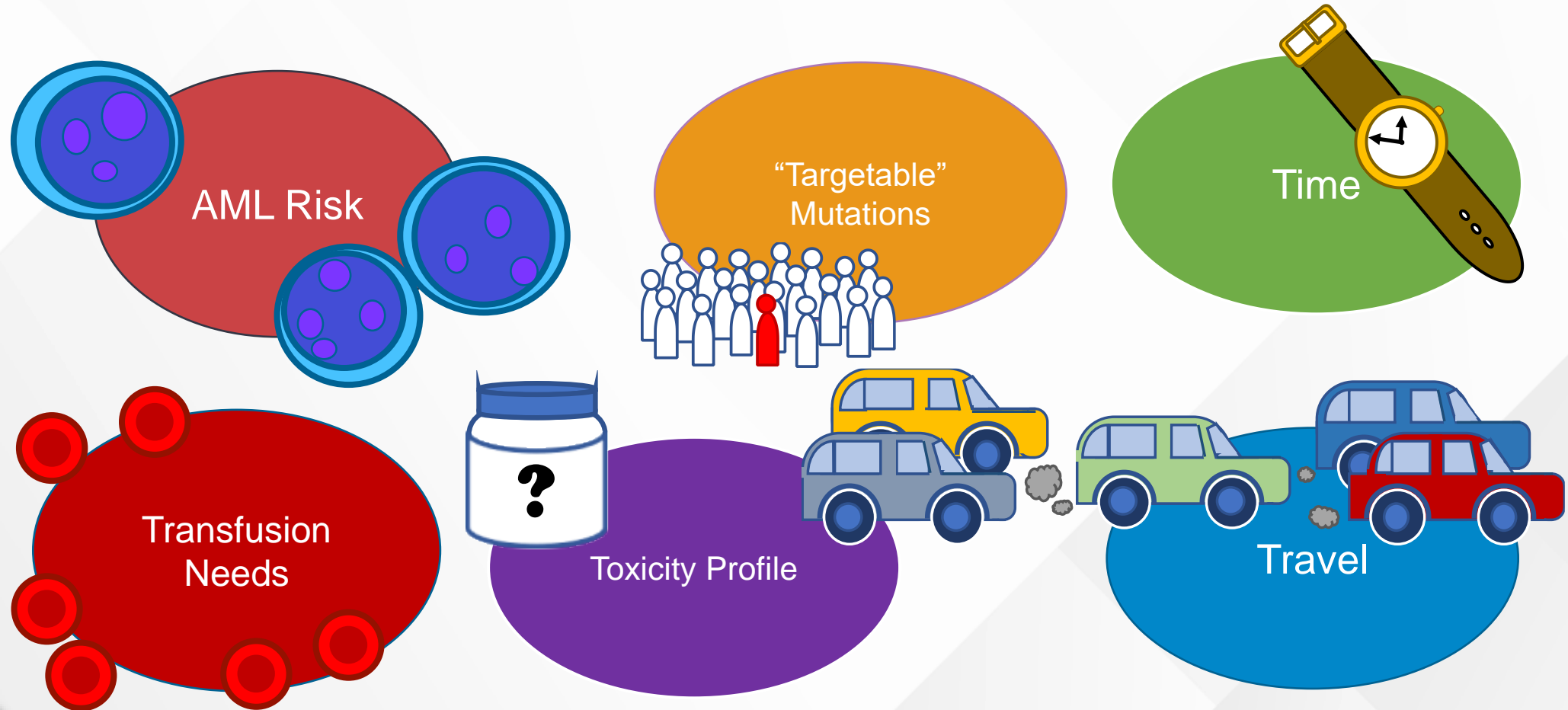


Clinical Outcomes for Patients with MDS

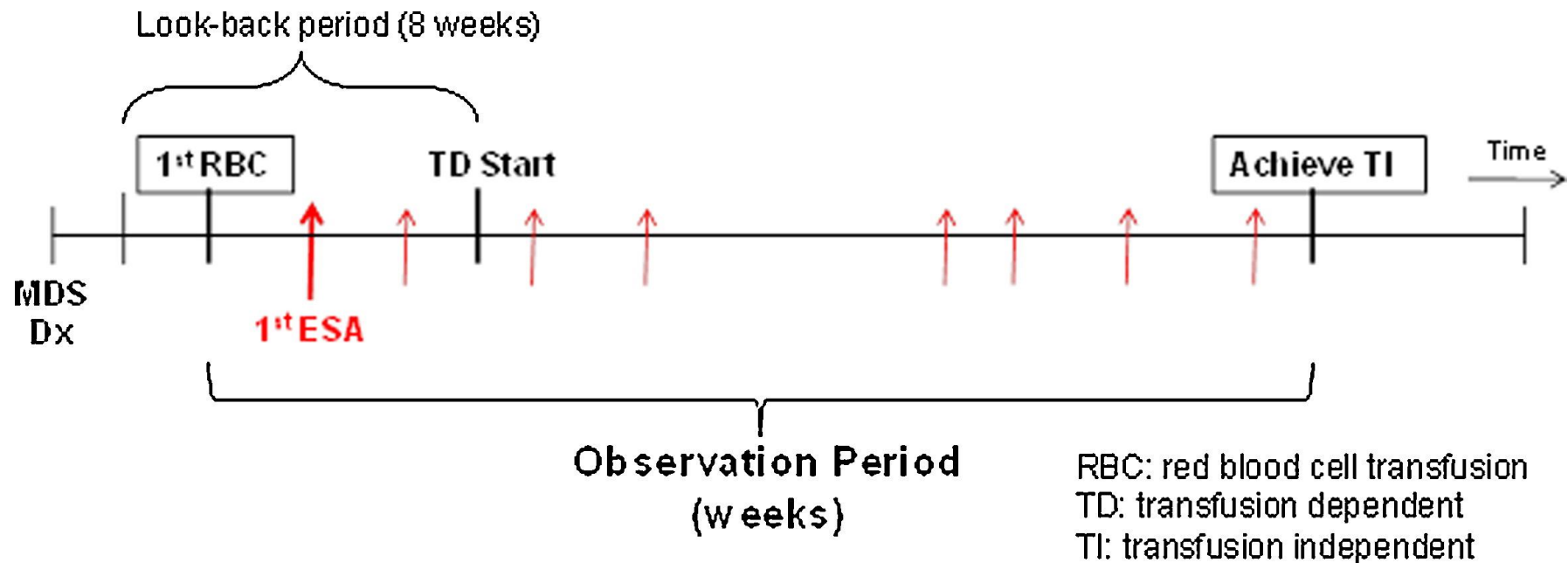
If all of the MDS patients diagnosed in the U.S. this year were represented as 100 people...



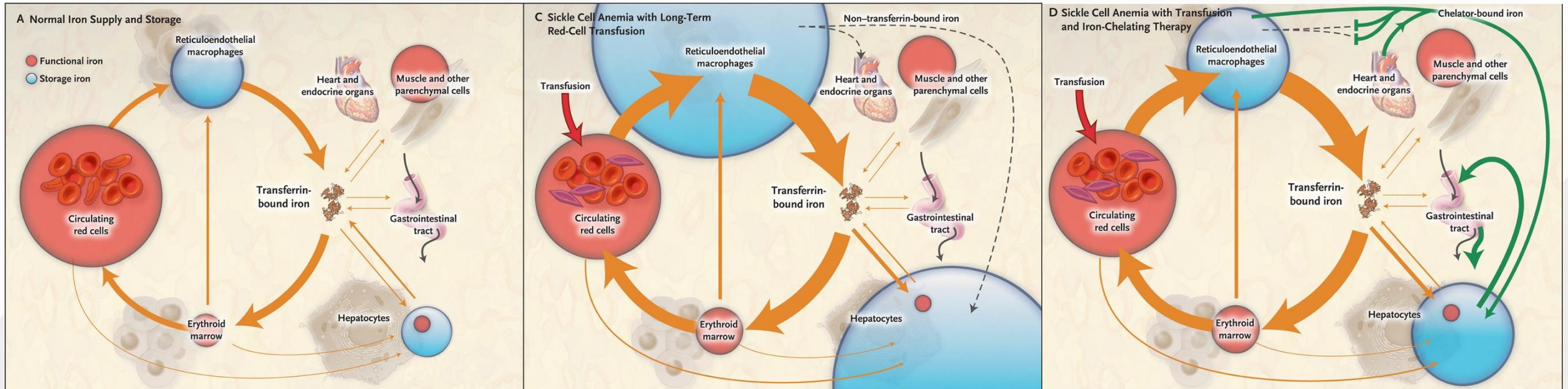
MDS Management: Integrating Many Factors



Transfusion Burden in MDS



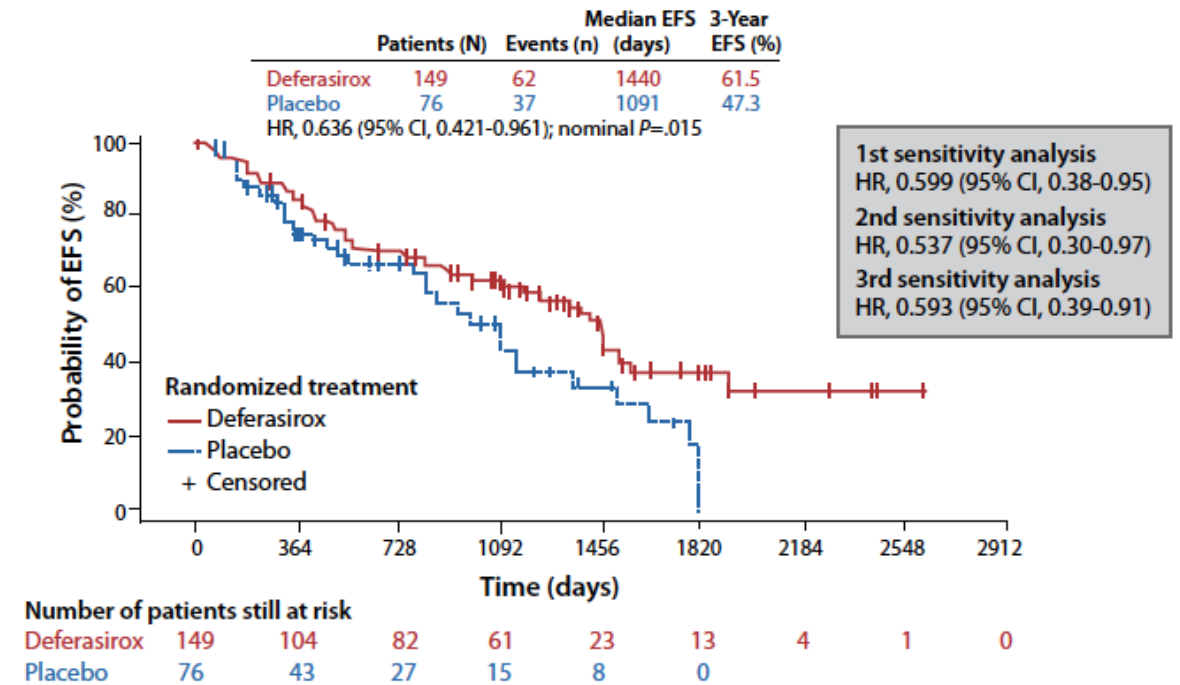
Iron Overload From Transfusions



Iron Chelation: TOLESTO

- Patients with Low/Intermediate-1 MDS randomized to deferasirox or placebo

Event	Deferasirox	Placebo
Death	32.2%	32.9%
AML	6.7%	7.9%
CHF Hospitalization	0.7%	3.9%
Liver Impairment	0.7%	1.3%
Cardiac function decline	2.3%	2.6%



Key Takeaways

- The initial evaluation of MDS requires specialized histopathologic, cytogenetic, and molecular analysis
- Risk stratification is key to determining the treatment goals in MDS
- Patient treatment goals inform treatment selection
- There are numerous alterations in the immune system in MDS that are potential targets to enhance disease control and the duration of responses

Clinical Trial Summary

Therapy	Target	Combination	Trial	Phase	Status (July 2022)
Magrolimab	CD47	+ azacitidine	ENHANCE NCT04313881	3	Recruiting
Sabatolimab	TIM3	+ HMA	STIMULUS-MDS1 NCT03946670	2	Active, not recruiting
		+ azacitidine	STIMULUS-MDS2 NCT04266301	3	Active, not recruiting
		+ azacitidine and venetoclax	STIMULUS-MDS3 NCT04812548	2	Recruiting
		+ HMA	STIMULUS-MDS-US NCT04878432	2	Recruiting
		+ siremadlin (HDM201)	NCT03940352	1b	Recruiting

Harnessing the Power of the Immune System to Manage Higher-Risk MDS

