

Acute Myeloid Leukemia and Myelodysplastic Syndromes:

Where are we in 2025?

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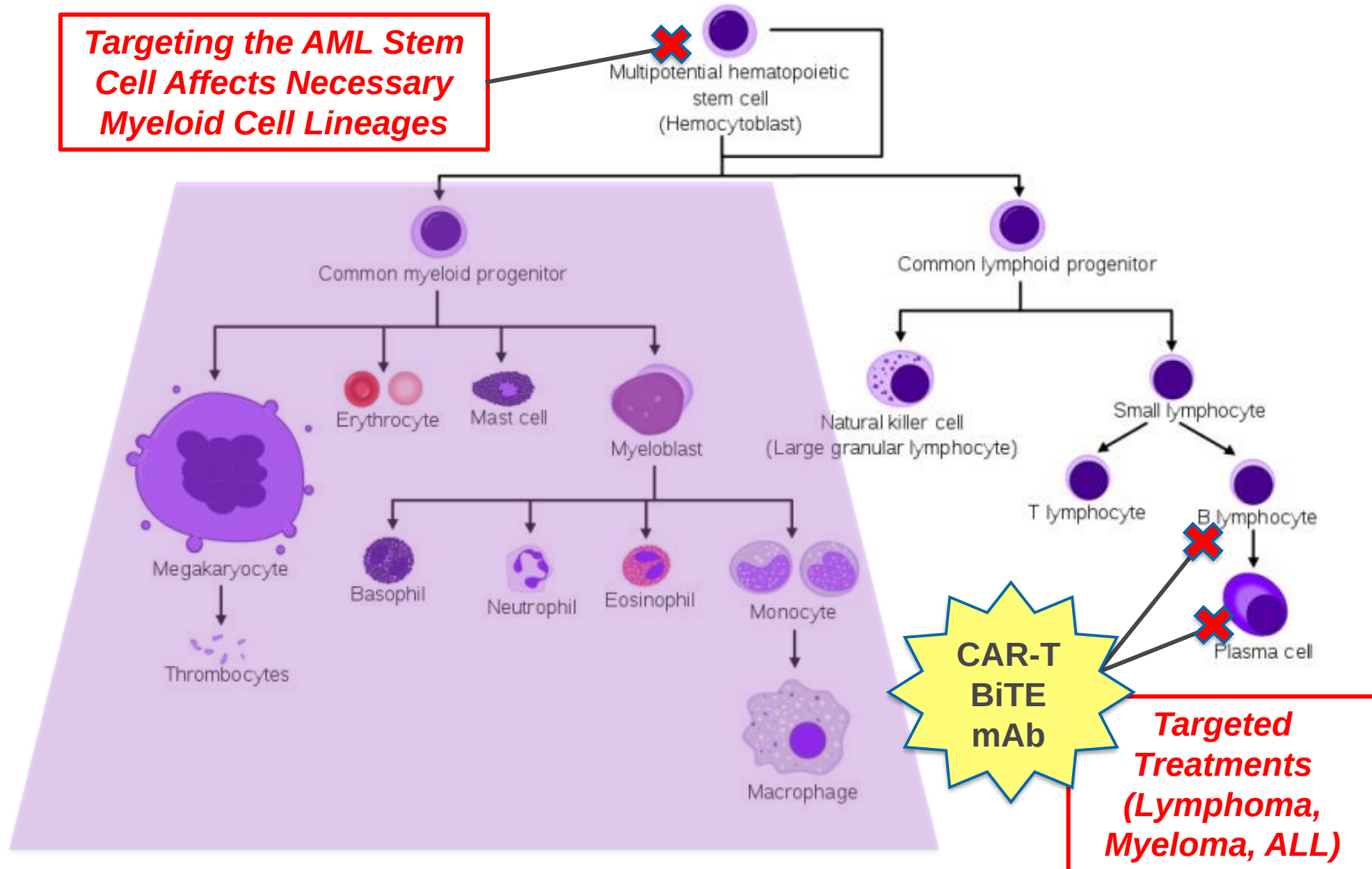
UC San Diego Moores Cancer Center

Disclosures

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- Research Funding: Function Oncology

Normal Hematopoiesis

Targeting the AML Stem Cell Affects Necessary Myeloid Cell Lineages



Overview

- AML
 - Blast Threshold (WHO 5th Edition vs ICC)
 - AML-defining cytogenetic/genetic lesions
 - Risk Stratification (ELN 2022)
 - Treatment challenges
- MDS
 - Risk Stratification (IPSS-R vs IPSS-M)
 - Sequencing therapies for lower risk MDS
 - Challenges in higher risk MDS

AML DIAGNOSIS

AML Diagnosis

- **Previously: $\geq 20\%$ myeloid blasts in bone marrow or blood**
 - ▶ t(15;17), t(8;21), inv(16)/t(16;16) regardless of blast percentage
 - ▶ Myeloid sarcoma/chloroma (a tumor composed of myeloblasts)
- **WHO and ICC 2022 Updates**
 - ▶ Both retain recurrent genetic abnormalities as a primary consideration
 - ▶ For specific genetic abnormalities, ICC mandates $\geq 10\%$ blasts in the bone marrow or blood but WHO does not specify a blast cut-off (next slide)
 - ▶ *BCR::ABL1* requires $\geq 20\%$ blasts for both ICC and WHO
 - ▶ AML with MDS-related changes defined by cytogenetic or gene mutation criteria, not morphologic dysplasia
 - ▶ *ICC created a new category of “MDS/AML” for blasts 10-19% with designated genetic mutations*
 - ▶ *TP53* mutated AML, exclusive to ICC, requires $\geq 20\%$ blasts and VAF $\geq 10\%$

Two New Classification Systems in 2022

AML-Defining Genetic Lesion	AML Blast Threshold	
	WHO	ICC
t(15;17)(q24.1;q21.2)/ <i>PML::RARA</i> and other <i>RARA</i> rearrangements	Any	≥10%
t(8;21)(q22;q22.1)/ <i>RUNX1::RUNX1T1</i>	Any	≥10%
inv(16)(p13.1;q22) or t(16;16)(p13.1;q22)/ <i>CBFB::MYH11</i>	Any	≥10%
t(9;11)(p21.3;q23.3)/ <i>MLLT3::KTM2A</i> and other <i>KMT2A</i> rearrangements	Any	≥10%
t(6;9)(p22.3;q34.1)/ <i>DEK::NUP214</i>	Any	≥10%
inv(3)(q21.3;q26.2) or t(3;3)(q21.3;q26.2); <i>GATA2::MECOM(EVI1)</i>	Any	≥10%
<i>MECOM</i> rearrangements	Any	≥10%
Other recurring translocations including <i>NUP98</i> and <i>RBM15:MRTF1</i>	Any	≥10%
<i>NPM1</i> mutated	Any	≥10%
In-frame bZIP <i>CEBPA</i> mutations	≥20%	≥10%
<i>TP53</i> mutation	-	≥20%
t(9;22)(q34.1;q11.2)/ <i>BCR::ABL1</i>	≥20%	≥20%
MDS-related cytogenetic/genetic abnormalities	≥20%	≥10%

Rapid NGS Myeloid Panel

- Available at UCSD since May 2024 (Dr. Wei Song)
- Interrogates 45 DNA target genes and 35 RNA fusion driver genes
- Needed for diagnosis (AML-defining lesions), risk stratification, treatment selection

DNA panel: hotspot genes (28)		DNA panel: full genes (17)		RNA panel: fusion driver genes (35)			RNA panel: expression genes (5)
ANKRD26	KRAS	ASXL1	PRPF8	ABL1	HMGA2	NUP98	BAALC
ABL1	MPL	BCOR	RB1	ABL2	JAK2	NUP214	MECOM
BRAF	MYD88	CALR	RUNX1	BCL2	KAT6A (MOZ)	PAX5	MYC
CBL	NPM1	CEBPA	SH2B3	BRAF	KAT6B	PDGFRA	SMC1A
CSF3R	NRAS	ETV6	STAG2	CCND1	KMT2A	PDGFRB	WT1
DDX41	PPM1D	EZH2	TET2	CREBBP	KMT2A PTDs	RARA	
DNMT3A	PTPN11	IKZF1	TP53	EGFR	MECOM	RUNX1	
FLT3 (ITD, TKD)	SMC1A	NF1	ZRSR2	ETV6	MET	TCF3	
GATA2	SMC3	PHF6		FGFR1	MLLT10	TFE3	
HRAS	SETBP1			FGFR2	MRTFA (MKL1)	ZNF384	
IDH1	SF3B1			FUS	MYBL1		
IDH2	SRSF2				MYH11		
JAK2	U2AF1				NTRK2		
KIT	WT1				NTRK3		

AML Defining Lesions

AML RISK STRATIFICATION

AML Risk (Staging) Is Determined by Genetic Abnormalities

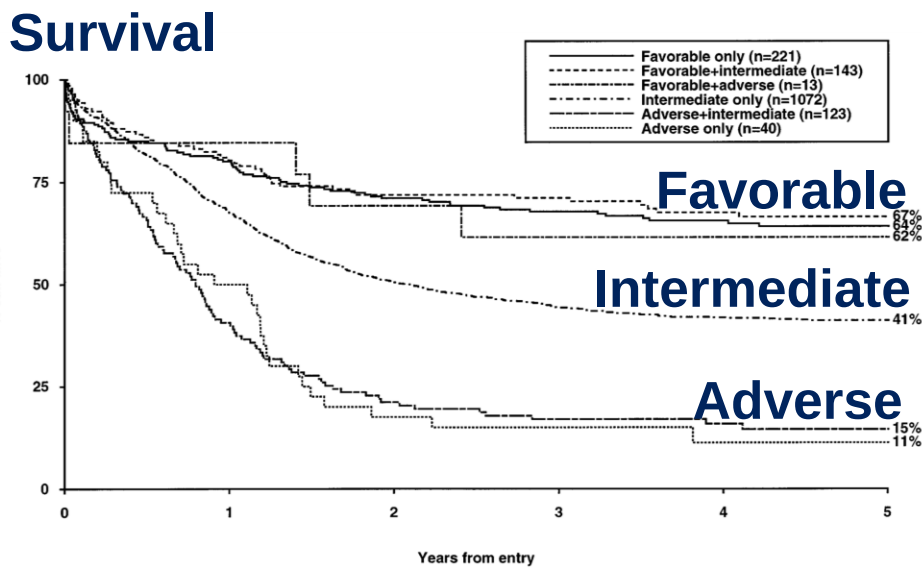
Risk Category	Genetic Abnormality
Favorable (15%)	t(8;21)(q22;22.1)/RUNX1::RUNX1T1 inv(16)(p13.1q22) or t(16;16)(p13.1;q22)/CBFB::MYH11 Mutated NPM1 without FLT3-ITD bZIP in-frame mutated CEBPA
Intermediate (50%)	Mutated NPM1 with FLT3-ITD Wild-type NPM1 with FLT3-ITD (without adverse-risk genetic lesions) t(9;11)(p21.3;q23.3)/MLLT3::KMT2A Cytogenetic and/or molecular abnormalities not classified as favorable or adverse
Poor/Adverse (35%)	t(6;9)(p23.3;q34.1)/DEK::NUP214 t(v;11q23.3)/KMT2A-rearranged t(9;22)(q34.1;q11.2)/BCR::ABL1 t(8;16)(p11.2;p13.3)/KAT6A::CREBBP inv(3)(q21.3q26.2) or t(3;3)(q21.3;q26.2)/GATA2, MECOM(EVI1) t(3q26.2;v)/MECOM(EVI1)-rearranged -5 or del(5q); -7; -17/abn(17p) Complex karyotype, monosomal karyotype Mutated ASXL1, BCOR, EZH2, RUNX1, SF3B1, SRSF2, STAG2, U2AF1 and/or ZRSR2 Mutated TP53

Add
gemtuzumab

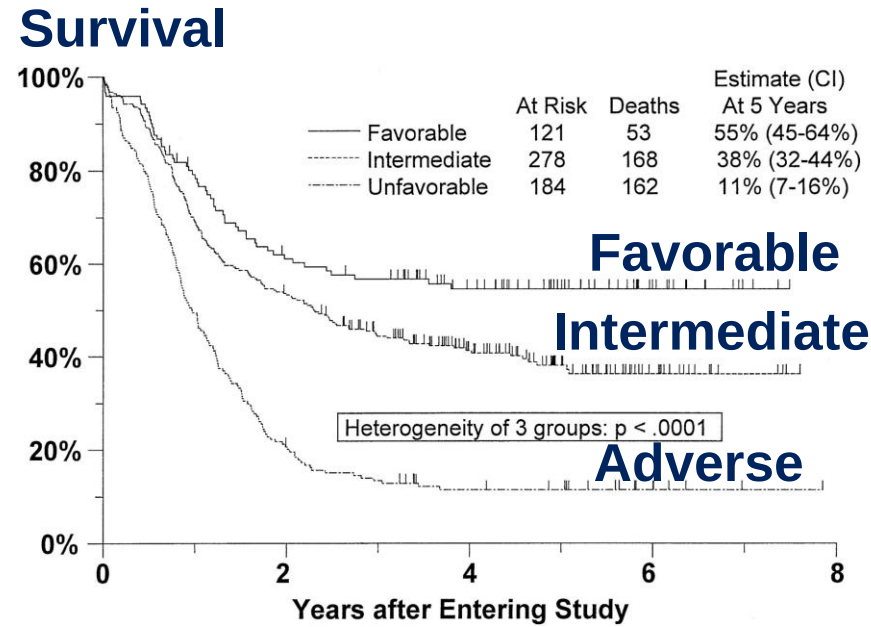
Add
midostaurin,
quizartinib

7+3 vs. CPX-351
vs. HMA/Ven

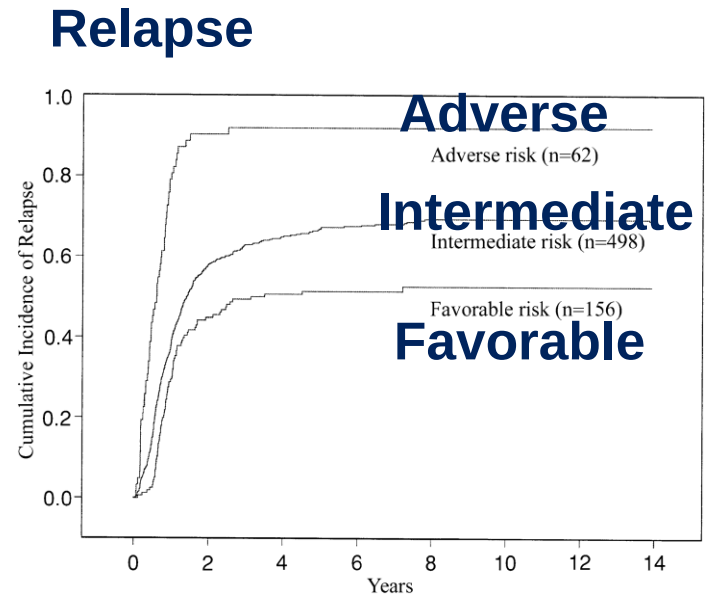
AML 5-Year Survival by Risk Group



UK MRC AML 10
n=1612



SWOG/ECOG
n=609



CALGB
n=1213

Grimwade *Blood* 1998; Slovak *Blood* 2000; Byrd *Blood* 2002

AML 5-Year Survival by Risk Group

	Favorable	Intermediate	Poor
UK MRC AML 10 n=1612	65%	41%	14%
SWOG/ECOG n=609	55%	38%	11%
CALGB n=1213	55%	24%	5%

**Allogeneic HSCT for most
fit/younger (<75 years)
patients who have a donor**

AML TREATMENT

AML Therapies

- **7+3 induction chemotherapy (1973), and other intensive chemo**
**7+3 = 7 days of cytarabine, 3 days of anthracycline (daunorubicin, idarubicin)*
- **Hypomethylating Agents (HMAs): Azacitidine (2004)*, Decitabine (2006)***

**Approved for MDS, used off-label for AML*

- Gemtuzumab (2010 → FDA hold, then approved 2017)
- CPX-351/liposomal cytarabine + daunorubicin (2017)
- Midostaurin (2017)
- Enasidenib (2017)
- **Venetoclax (2018), with HMA**
- Glasdegib (2018)
- Ivosidenib (2018), +/- HMA
- Gilteritinib (2018)
- Azacitidine, oral tablets (2020)
- Olutasidenib (2022)
- Quizartinib (2023)
- Revumenib (2024)

***AML drugs
approved in the
last decade!***

AML Therapies 1970-2010

	FIT PATIENT	UNFIT PATIENT
Newly Diagnosed	7+3 Other intensive chemo	HMA LDAC Best supportive care
Relapsed/Refractory	Intensive chemo HMA	HMA LDAC Best supportive care

AML Therapies **2010 and beyond**

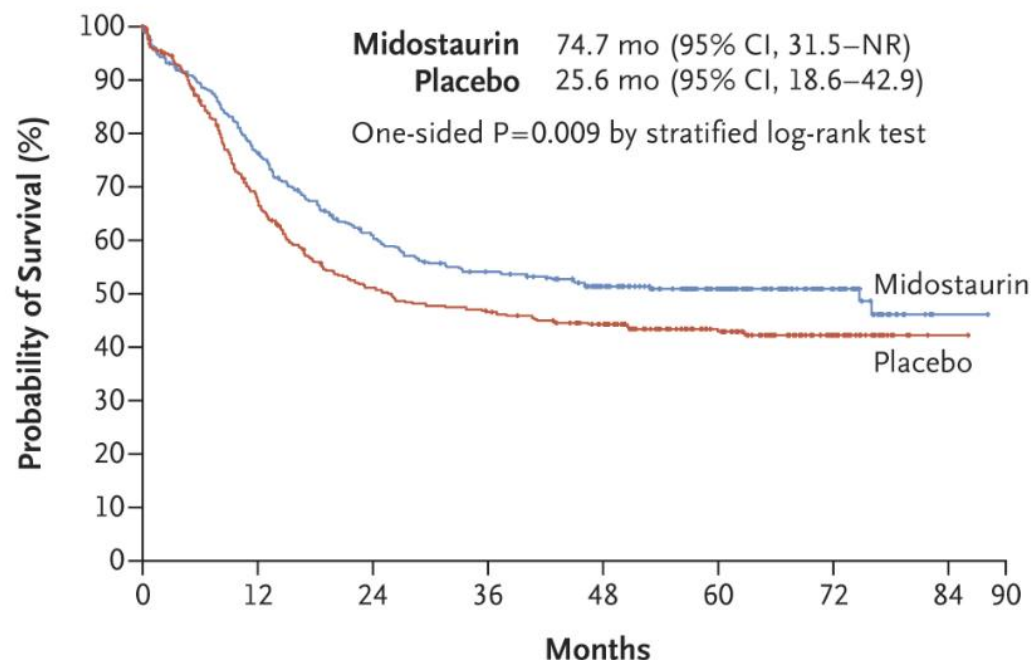
	FIT PATIENT	UNFIT PATIENT
Newly Diagnosed	7+3 7+3 + Gemtuzumab (favorable risk) 7+3 + Mido/Quiz (FLT3-mutated) CPX-351 (t-AML, AML-MRC) ?HMA + Venetoclax	HMA + Venetoclax HMA LDAC + Venetoclax Best supportive care HMA + Ivosidenib (IDH1-mutated) Enasidenib (IDH2-mutated) LDAC + Glasdegib
Relapsed/Refractory	Intensive chemo HMA + Venetoclax Ivosidenib (IDH1-mutated) Olutasidenib (IDH1-mutated) Enasidenib (IDH2-mutated) Gilteritinib (FLT3-mutated) Quizartinib (FLT3-mutated) Gemtuzumab (CD33-positive) Revumenib (KMT2A-r)	HMA + Venetoclax HMA Ivosidenib (IDH1-mutated) Olutasidenib (IDH1-mutated) Enasidenib (IDH2-mutated) Gilteritinib (FLT3-mutated) Quizartinib (FLT3-mutated) Gemtuzumab (CD33-positive) Revumenib (KMT2A-r)

**AML Treatment Question 1:
Which FLT3 inhibitor for a chemo-eligible patient
with newly diagnosed AML?**

C10603/RATIFY

7+3/Midostaurin vs. 7+3/Placebo

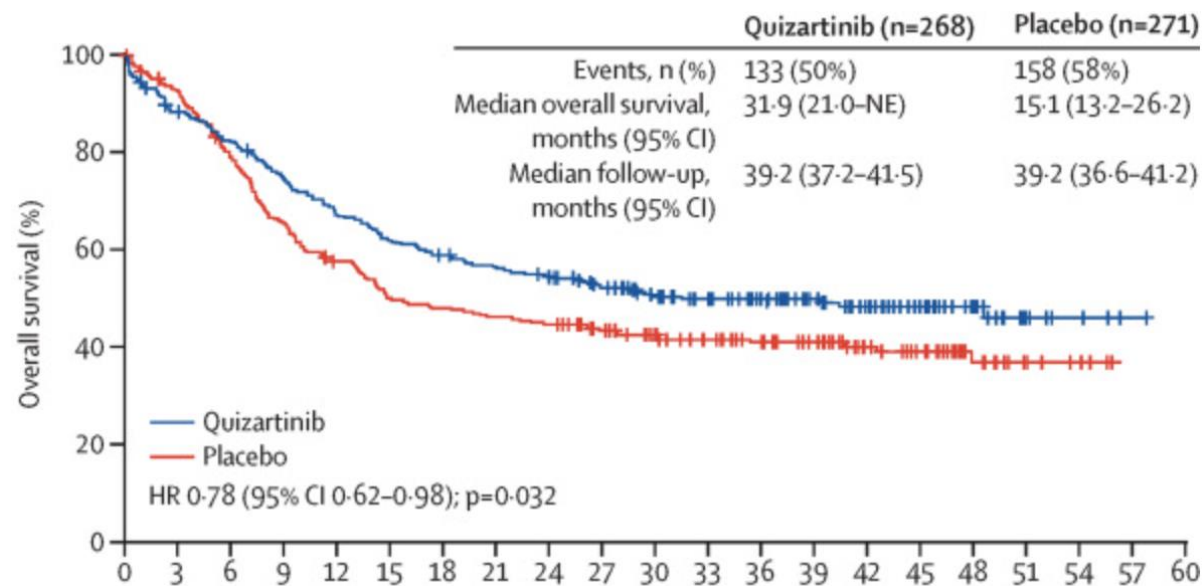
Median Overall Survival



OS 74.7 vs 25.6 mo, $p=0.009$
 OS benefit in ITD only
 OS in CR1 allo: 28.1 vs 22.7 mo, $p=0.07$
 GI toxicity, rash

QUANTUM-FIRST

7+3/Quizartinib vs. 7+3/Placebo

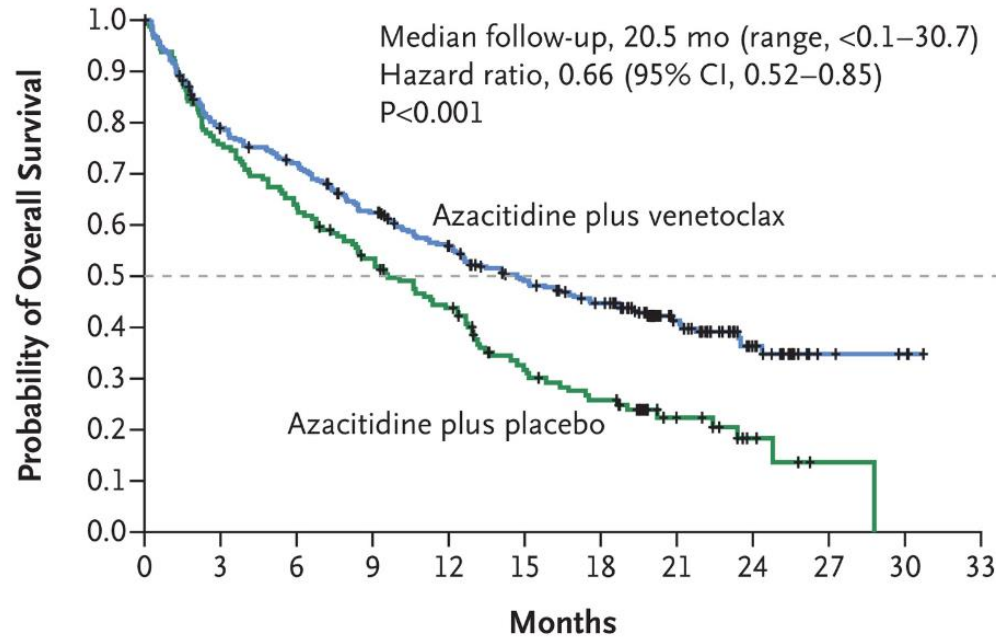


OS 31.9 vs. 15.1 mo, $p=0.032$
 Activity against ITD only
 Enrolled up to age 75 years
 QTc prolongation

**AML Treatment Question 2:
Which treatment for the non chemo-eligible
patient with IDH1-mutated AML?**

VIALE-A

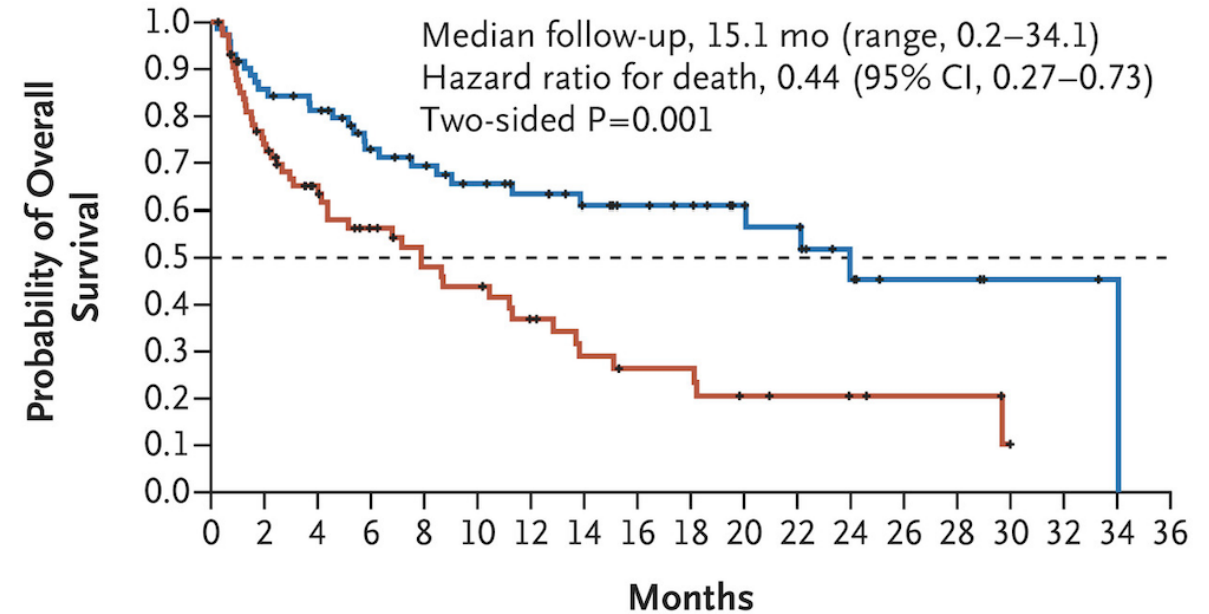
Azacitidine/Venetoclax vs. Azacitidine/Placebo



OS 14.7 vs 9.6 mo, $p<0.001$
 ORR 66.4 vs 28.3%, $p<0.001$
**IDHm, ORR 75.4% vs 10.7%*
TTR 1 vs 2.6 mo, DOR 17.8 vs 13.9 mo
 Gr 3/4 febrile neutropenia 30 vs 10%
 Gr 3/4 thrombocytopenia 45 vs 38%

AGILE

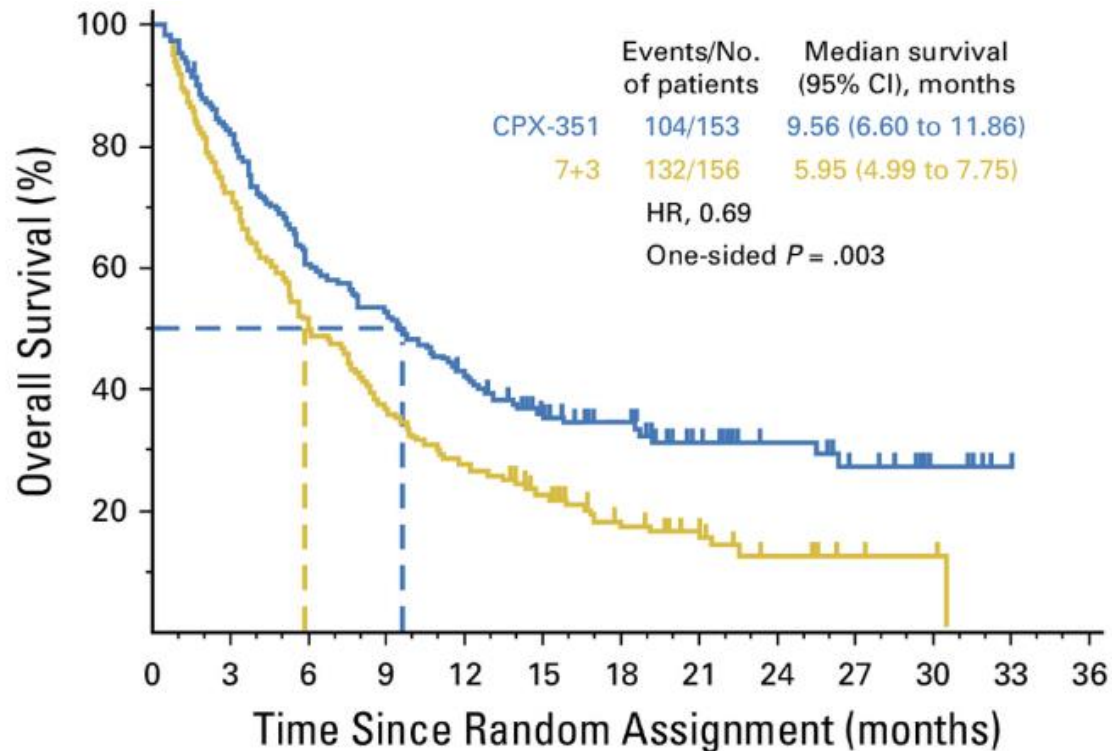
Azacitidine/Ivosidenib vs. Azacitidine/Placebo



OS 24 vs 7.9 mo, $p=0.001$
 ORR 63 vs 19%, $p<0.001$
 TTR 2.1 vs 3.7 mo, DOR 22.1 vs 9.2 mo
DS 14 vs 8%
 Gr 3/4 febrile neutropenia 28 vs 34%
 Gr 3/4 thrombocytopenia 24 vs 21%

**AML Treatment Question 3:
For adverse risk AML,
intensive chemo vs HMA/Ven?**

CPX-351 vs. 7+3



sAML (MDS, therapy-related), 60-75yr
OS 9.56 vs 5.95 mo, $p=0.009$

CR/CRi 47.7 vs 33.3%

60-day mortality 13.7 vs 21.2%

Gr 5 infection 7.2 vs 2.6%

Gr 5 bleeding 2.6 vs 2.6%

Subgroup	CPX-351	7+3
tAML, n=63	46.7%	36.4%
sAML, prior HMA, n=105	36%	32.7%
sAML, no prior HMA n=40	66.7%	36.8%
Poor Risk cytogenetics, n=155	43.1%	21.7%

Subgroup	Azacitidine + Venetoclax	Azacitidine + Placebo
AML-MRC, n=141	60.9%	22.4%
sAML, n=107	66.7%	22.9%
Poor Risk cytogenetics, n=160	52.9%	23.2%
TP53, n=52	55.3%	0%

Targeted Therapies

Differentiation Syndrome Risk

IDH1 inhibitors

- **Ivosidenib**

- ▶ Newly diagnosed AML (age ≥ 75) and R/R AML, R/R MDS
- ▶ Ph 1 study, untreated AML age ≥ 75 , n=34, CR/CRh 42%, median OS 12.6mo → R/R AML, n=179, CR/CRh 30%, median OS 8.8mo
- ▶ Ph 1 study, R/R MDS, n=18, CR 38.9%, med CR duration NR (1.9-80.8+ mo)

- **Olutasidenib**

- ▶ Ph 1/2 study, R/R AML, n=147, CR/CRh 35%, median OS 11.6mo

IDH2 inhibitor

- **Enasidenib**

- ▶ Ph 1/2 study, R/R AML, n=239, ORR 40%, CR 19%, median OS 9.3mo

IDH2 inhibitor

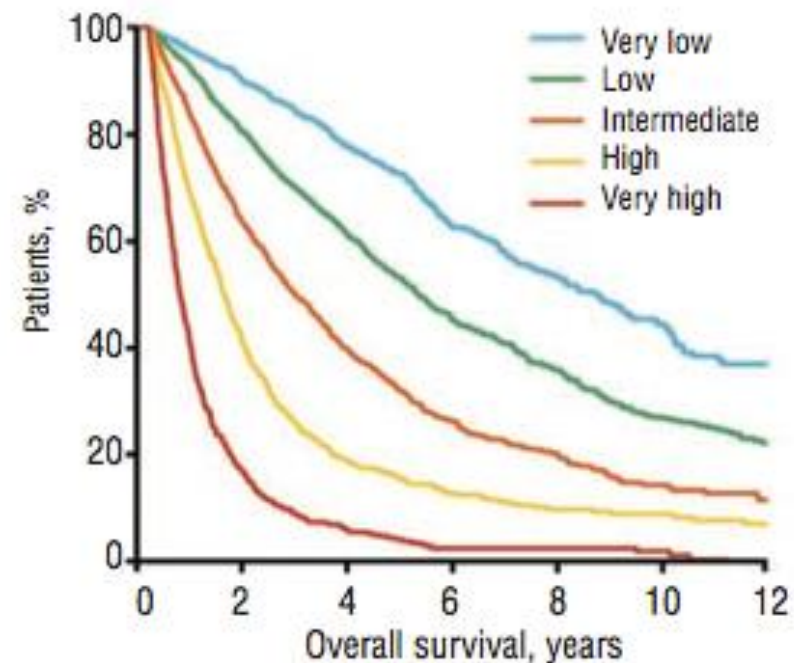
- **Revumenib**

- ▶ Ph 1/2 study, R/R acute leukemia (adult+pediatric) with *KMT2A*-r, n=104, CR+CRh 21.2%, ORR 63.2%, median duration 6.4mo

MDS Updates

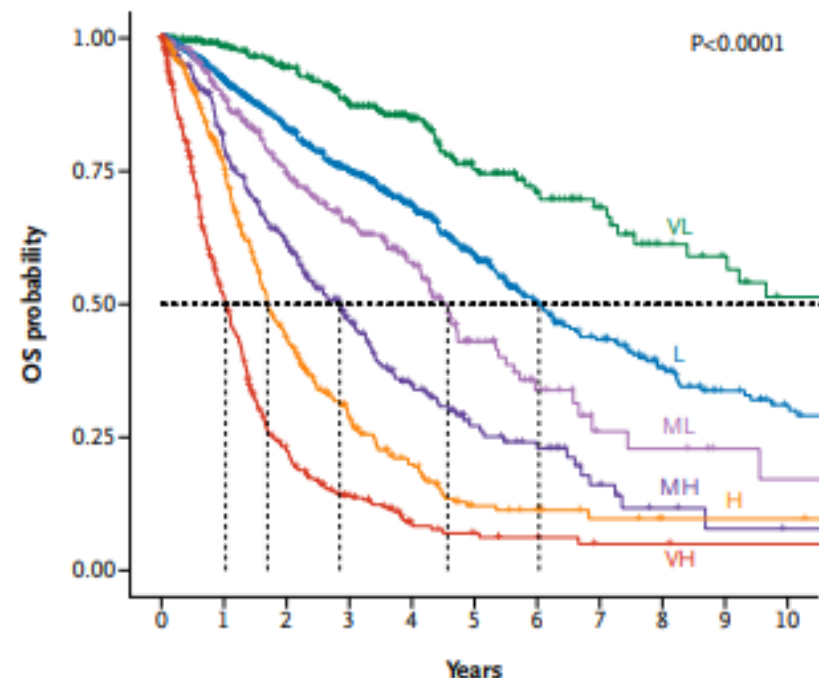
MDS Risk Stratification (Staging)

IPSS-R



Cytogenetic Risk
Bone marrow blast %
Hemoglobin
Platelet count
Absolute Neutrophil Count

IPSS-M



Cytogenetic Risk
Bone marrow blast %
Hemoglobin
Platelet count
(Absolute Neutrophil Count)

Mutation data

ASXL1, CBL, DNMT3A, ETV6, EZH2, FLT3, IDH2, KRAS, MLL^{PTD}, NPM1, NRAS, RUNX1, SF3B1, SRSF2, TP53^{multihit}, U2AF1

Approved MDS Therapies

Lower Risk MDS

- Erythropoiesis Stimulating Agents (ESA)
- Lenalidomide (2005)
- Luspatercept (2020)
- Ivosidenib (2023)
- Imetelstat (2024)
- Hypoplastic MDS: Immunosuppressive Therapy (IST)*

Higher Risk MDS

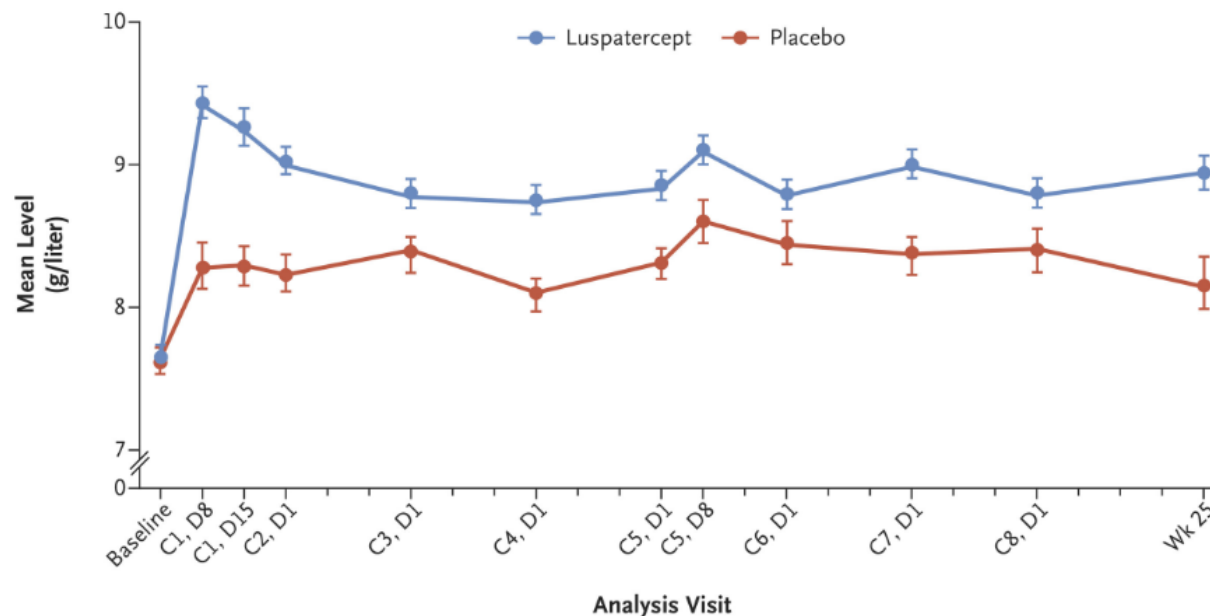
- Hypomethylating Agents aka HMAs
 - Azacitidine, IV or SQ (2004)
 - Decitabine, IV (2006)
 - Decitabine/Cedazuridine, PO (2020)
- Ivosidenib (2023)
- Hematopoietic Stem Cell Transplantation

MDS Treatment Question 1: Luspatercept vs ESA for lower-risk MDS?

MEDALIST

Luspatercept vs. Placebo Lower-risk MDS-RS, R/R ESA

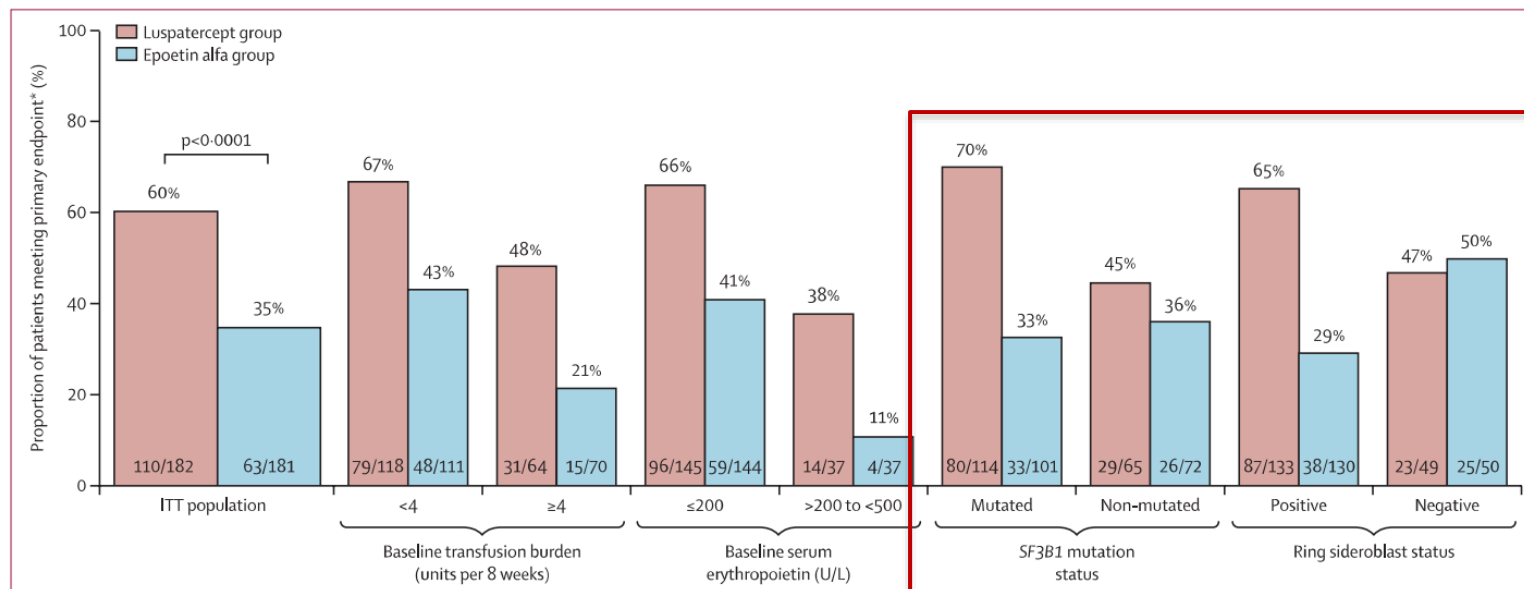
- RBC-TI (8wk) 38 vs 13%, $p < 0.001$
- RBC-TI (48wk) 33 vs 12%, $p < 0.001$
- OS similar in both groups (46+ mo)



COMMANDS

Luspatercept vs. ESA Lower-risk MDS, untreated

- RBC-TI (12wk, 24 wk) 59 vs 31%, 48 vs 29%
- RBC-TI (≥ 1 yr, ≥ 1.5 yr) 45 vs 28%, 30 vs 14%
- **Significant benefit to *most* subgroups (*SF3B1*, not RS-neg)**



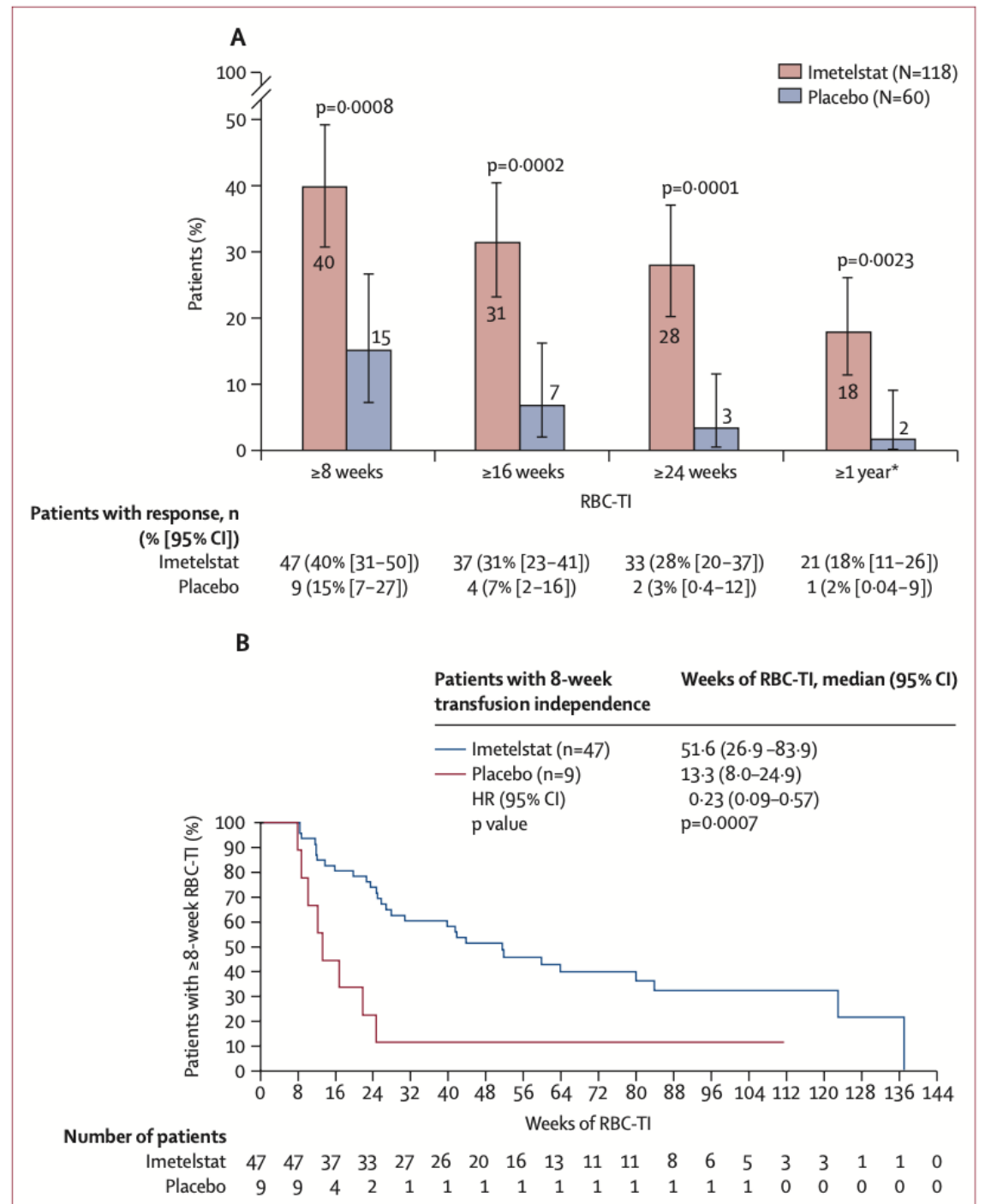
**MDS Treatment Question 2:
Luspatercept vs Imetelstat vs ESA
for lower-risk MDS, *SF3B1*-WT, RS-neg?**

IMerge

Imetelstat vs. Placebo

Lower-risk MDS-RS, R/R ESA

- RBC-TI (8wk) 40 vs 9%, $p=0.0008$
- RBC-TI (24wk) 28 vs 3%, $p=0.0001$
- RBC-TI (8wk, 24wk) with prior therapy:
 - ESA: 40%, 28%
 - ESA ineligible: 36%, 14%
 - Luspatercept: 29%, 20%
 - Lenalidomide: 23%, 12%
- **Toxicity (Gr 3/4)**
 - **Neutropenia 68 vs 3%**
 - **Thrombocytopenia 62 vs 8%**
 - **Median duration 1.9wk, 1.4wk**



MDS Treatment Question 3: HMA monotherapy vs HMA/Venetoclax for higher-risk MDS?

Phase 1b, Garcia (ASH 2021, 2023)

- Aza/Ven, treatment-naïve, HR MDS
- n=107, median OS 26 mo
- CR 29.9%, duration CR 16.6mo

Phase 1b, Zeidan (Am J Hematol 2023)

- Aza/Ven, R/R MDS
- n=37, median OS 12.6 mo, ORR 39%

VERONA Phase 3 Study

- Results pending
- Planned n=500, treatment-naïve, HR MDS
- 1:1 Aza/Ven vs Aza/Placebo

If MDS/AML category (>10% blasts) and patient is robust/fit, favor HMA/Ven, especially if goal is transplant

Molecular responses in CR or mCR pts at time of 2nd sample

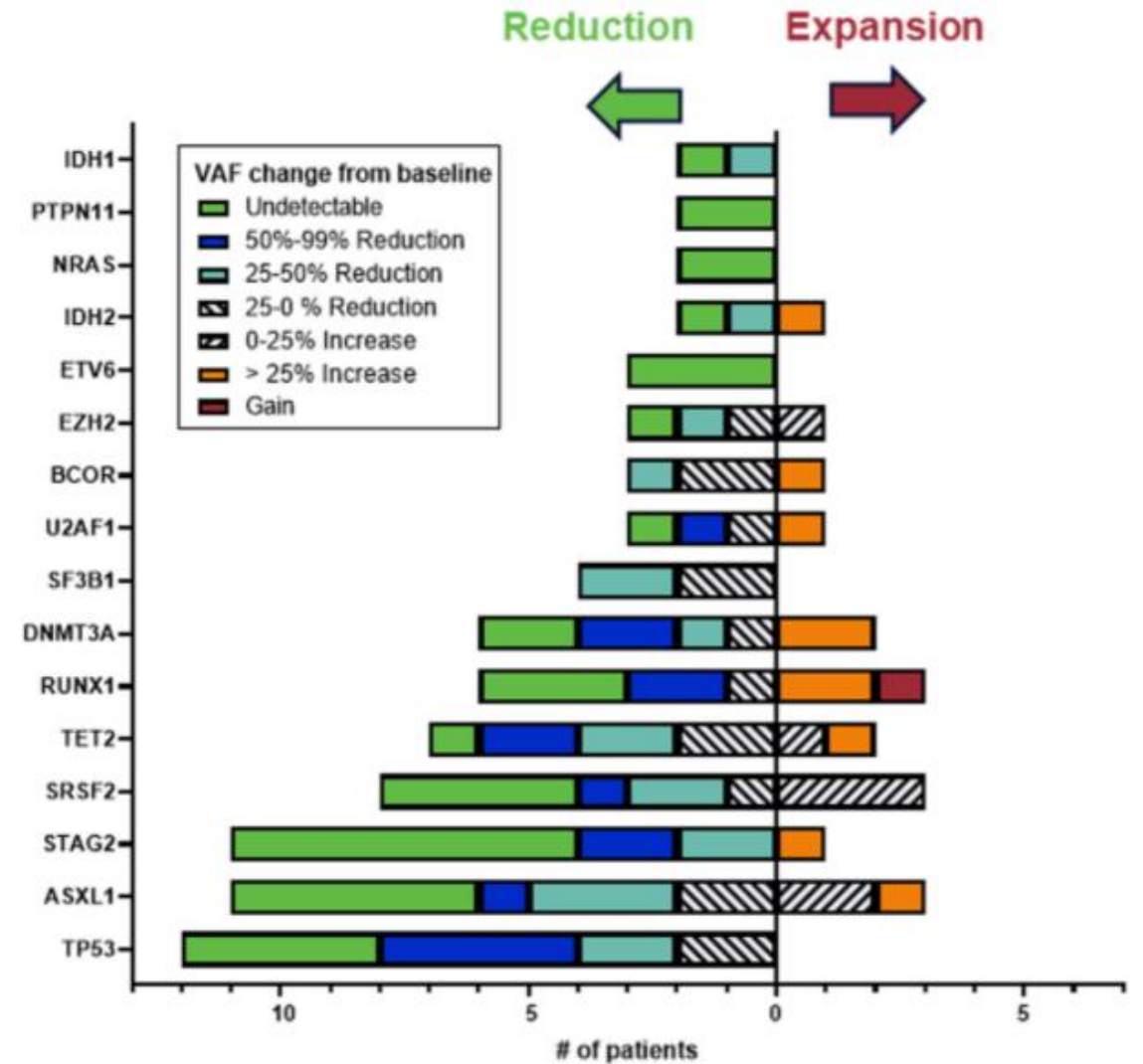


Figure. VAF dynamics in samples collected from patients in CR or mCR at the time of sample acquisition compared to baseline samples collected from bone marrow or the peripheral blood.

“Failed” Drugs in Higher-Risk MDS

- Most experimental treatments are added to an HMA “backbone”
 - Need to balance increased efficacy with more toxicity
- Phase 3 trials that did not find improved outcomes:
 - Pevonedistat, NEDD8 inhibitor, for higher-risk MDS, CMML, AML
 - Magrolimab, CD47 antibody, higher-risk MDS and AML
 - APR-246, restores p53 function, for TP53 mutated higher-risk MDS
 - Sabatolimab, TIM3 inhibitor, higher-risk MDS
 - Tamibarotene, synthetic retinoid, higher-risk MDS

Summary

- **Genetic mutations matter**
 - ▶ Lower/irrelevant blast thresholds if an AML-defining lesion is present
 - ▶ Absolutely needed for AML and MDS staging
- **AML therapies**
 - ▶ Many drug approvals for AML since 2010, after 40+ years of minimal options
 - ▶ 7+3 remains the standard induction chemo for younger/fit patients, with drugs added to this backbone for added efficacy
 - ▶ Hypomethylating Agent (HMA) + Venetoclax is the most transformative regimen for AML since 7+3 – potentially beneficial for most patients with AML (elderly/unfit)
- **MDS therapies**
 - ▶ Luspatercept > ESA for *SF3B1*-mutated or RS+ (unclear for *SF3B1*-WT or RS-neg)
 - ▶ Higher-risk MDS remains challenging to treat

THANK YOU!