

בסיס ביולוגי וטיפול קו ראשון ב-AML

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November 2025

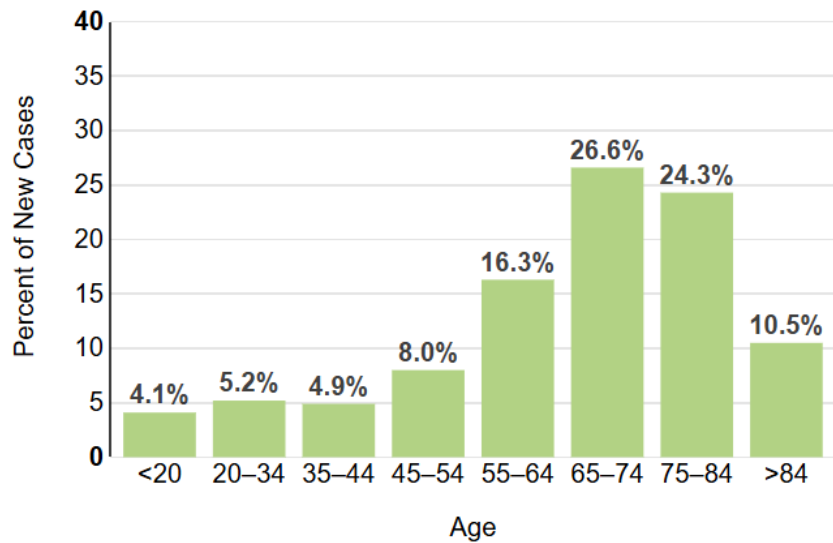
owolach@gmail.com

Acute Myeloid Leukemia (AML)



5-Year
Relative Survival

32.9%



Acute myeloid leukemia is most frequently diagnosed among people aged 65-74.

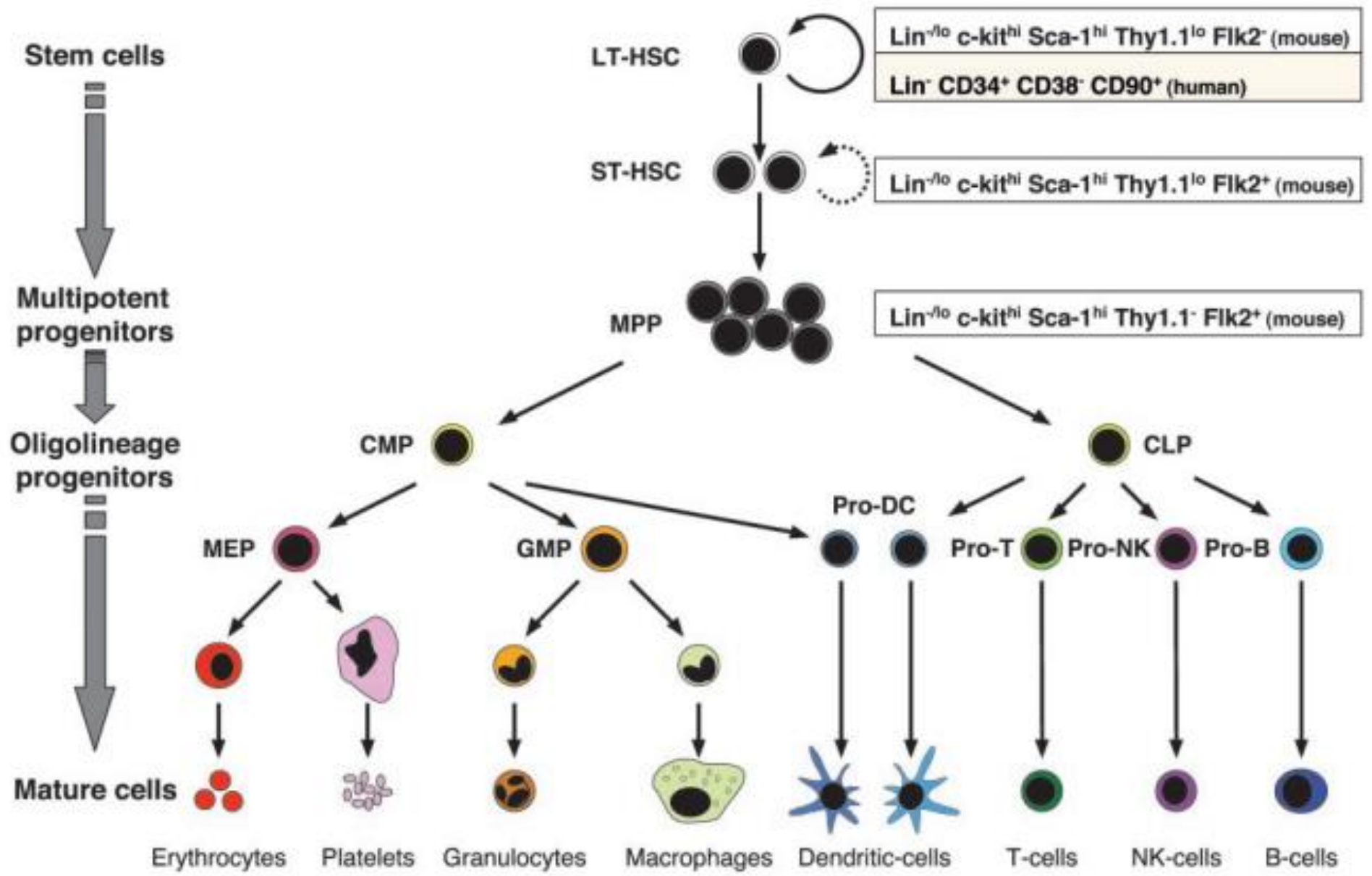
Median Age
At Diagnosis

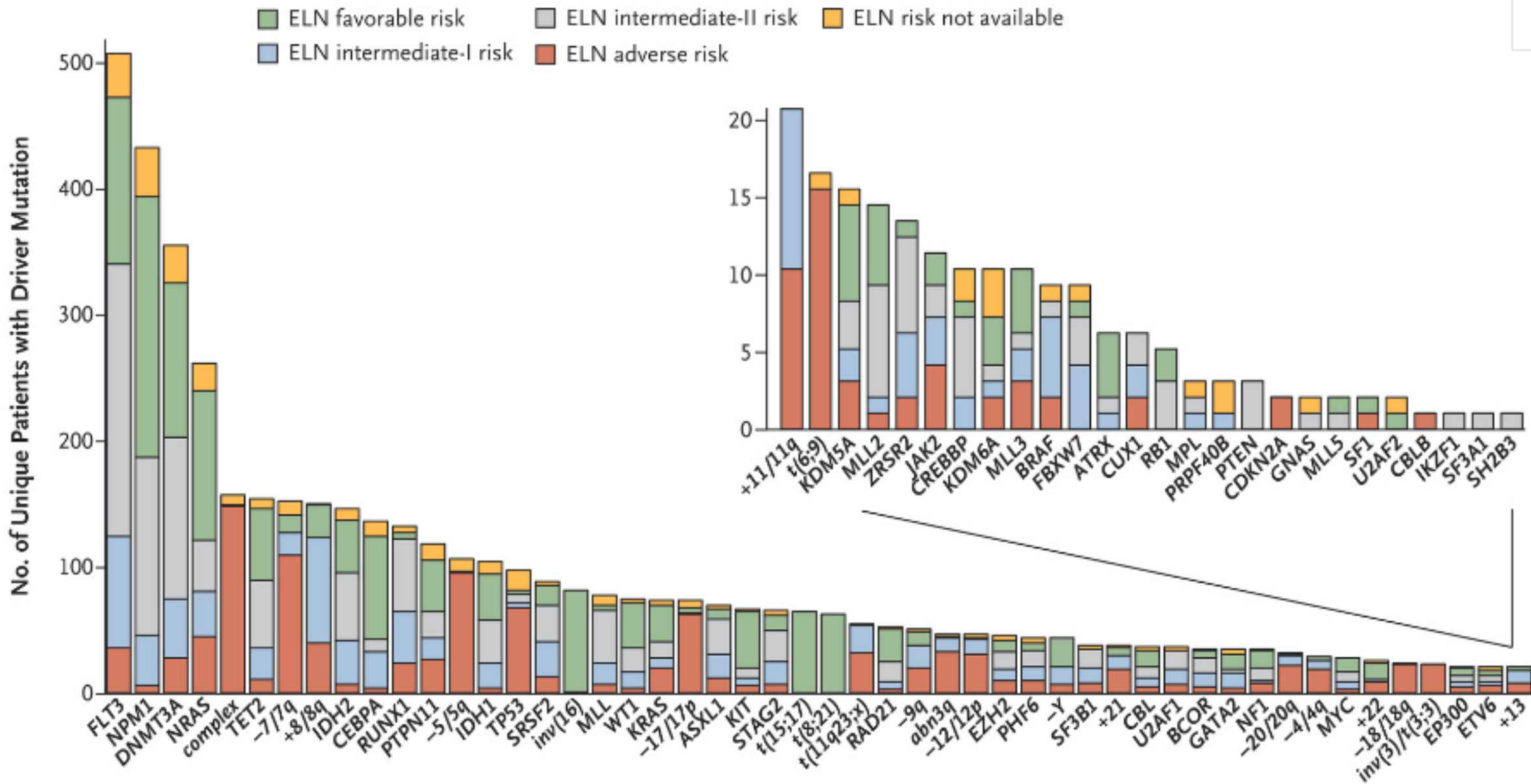
69

Common Types of Cancer	Estimated New Cases 2025	Estimated Deaths 2025
1. Breast Cancer (Female)	316,950	42,170
2. Prostate Cancer	313,780	35,770
3. Lung and Bronchus Cancer	226,650	124,730
4. Colorectal Cancer	154,270	52,900
5. Melanoma of the Skin	104,960	8,430
6. Bladder Cancer	84,870	17,420
7. Kidney and Renal Pelvis Cancer	80,980	14,510
8. Non-Hodgkin Lymphoma	80,350	19,390
9. Uterine Cancer	69,120	13,860
10. Pancreatic Cancer	67,440	51,980
-	-	-
Acute Myeloid Leukemia	22,010	11,090

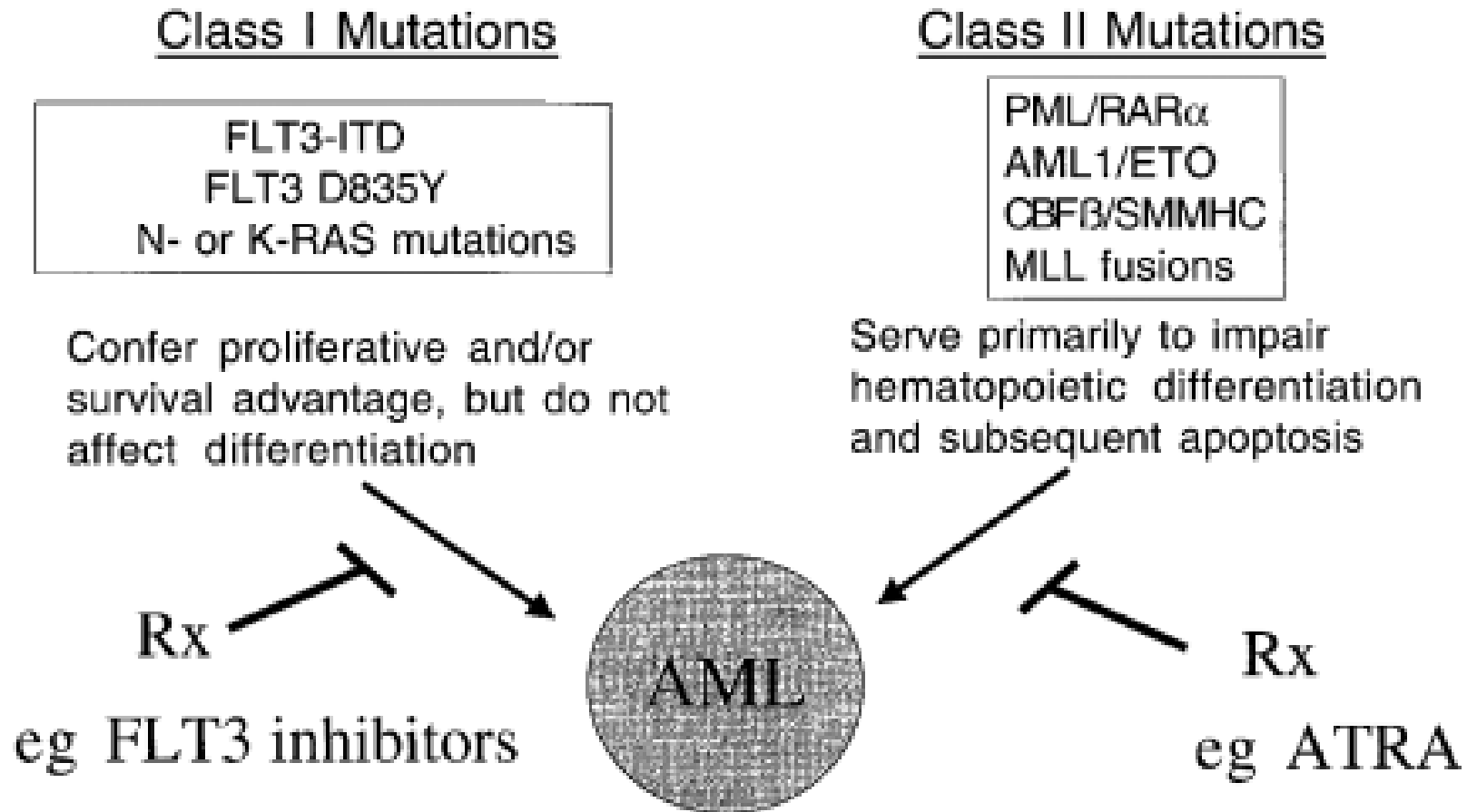
Source SEER data base

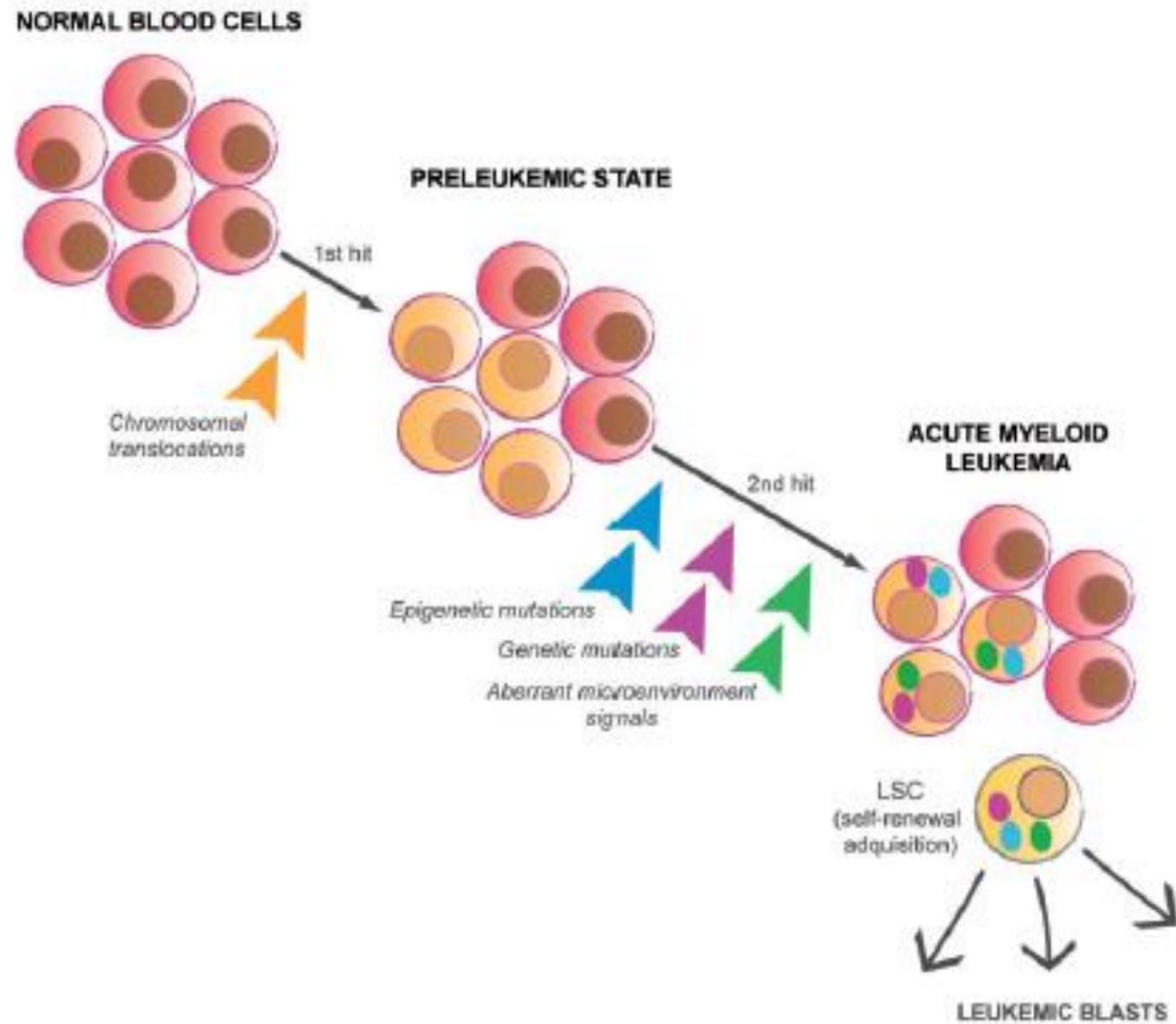
<https://seer.cancer.gov/statfacts/html/aml.html>



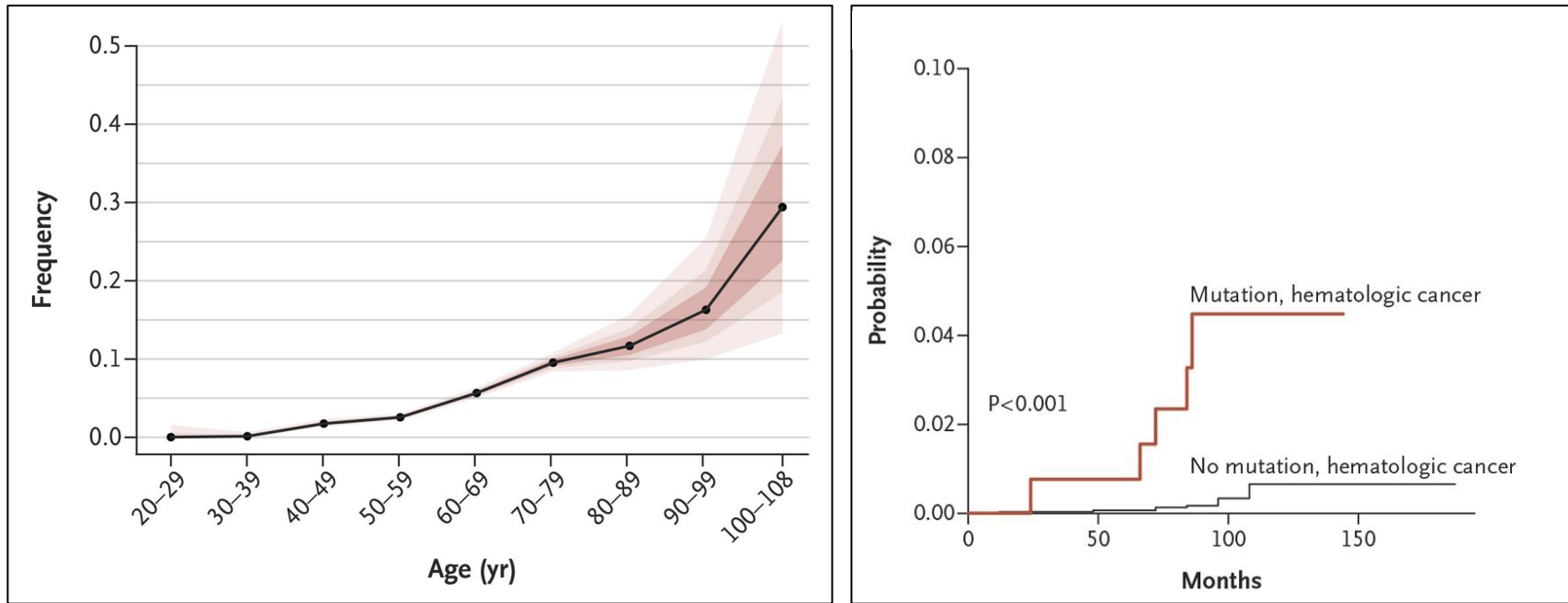


Concepts in the evolution of AML





Clonal hematopoiesis of indeterminate potential (CHIP)

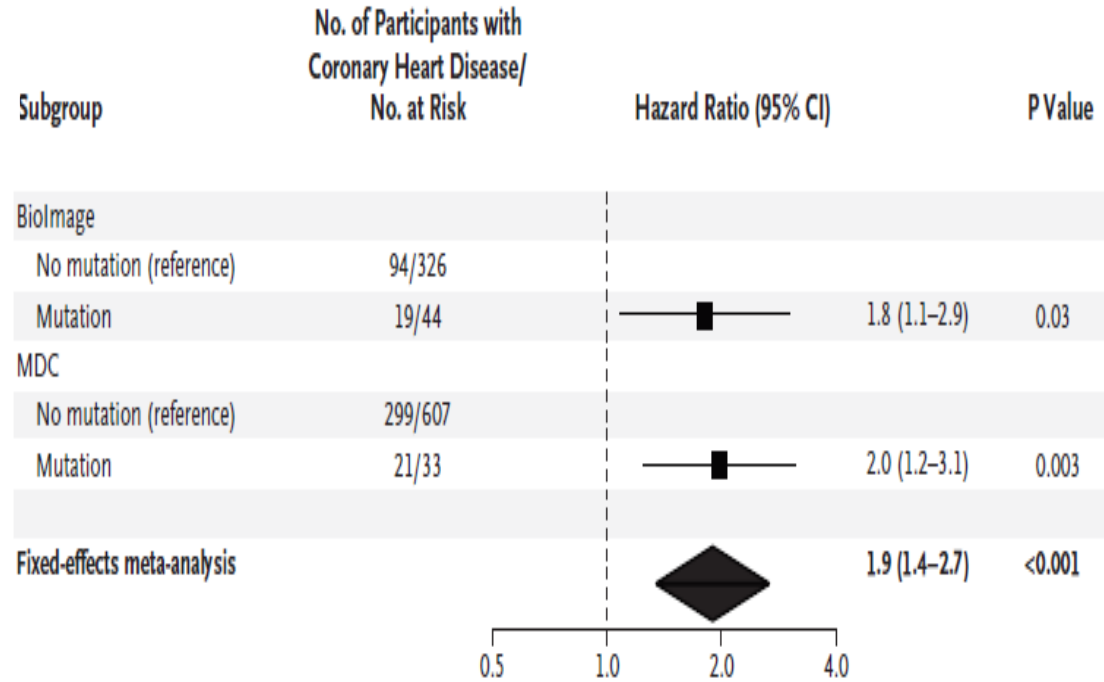


Acquired mutations in peripheral blood of healthy adults.
Associated with increased risk of hematologic malignancy (1%/yr).

Jaiswal et al, NEJM 2014; Genovese et al, NEJM 2014
Xie et al, Nat Med 2014; Young et al, Nat Comm 2016

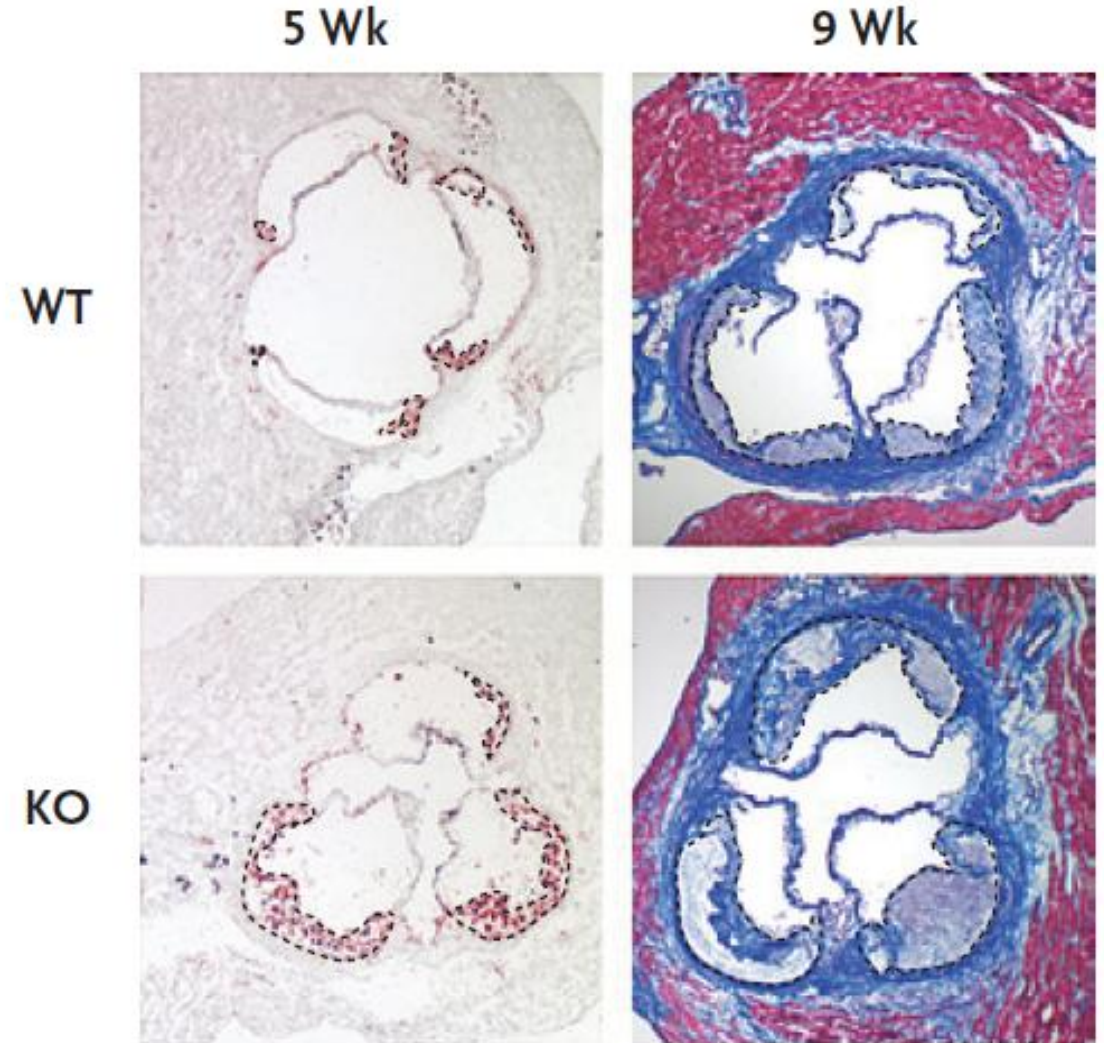
CHIP & Heart disease

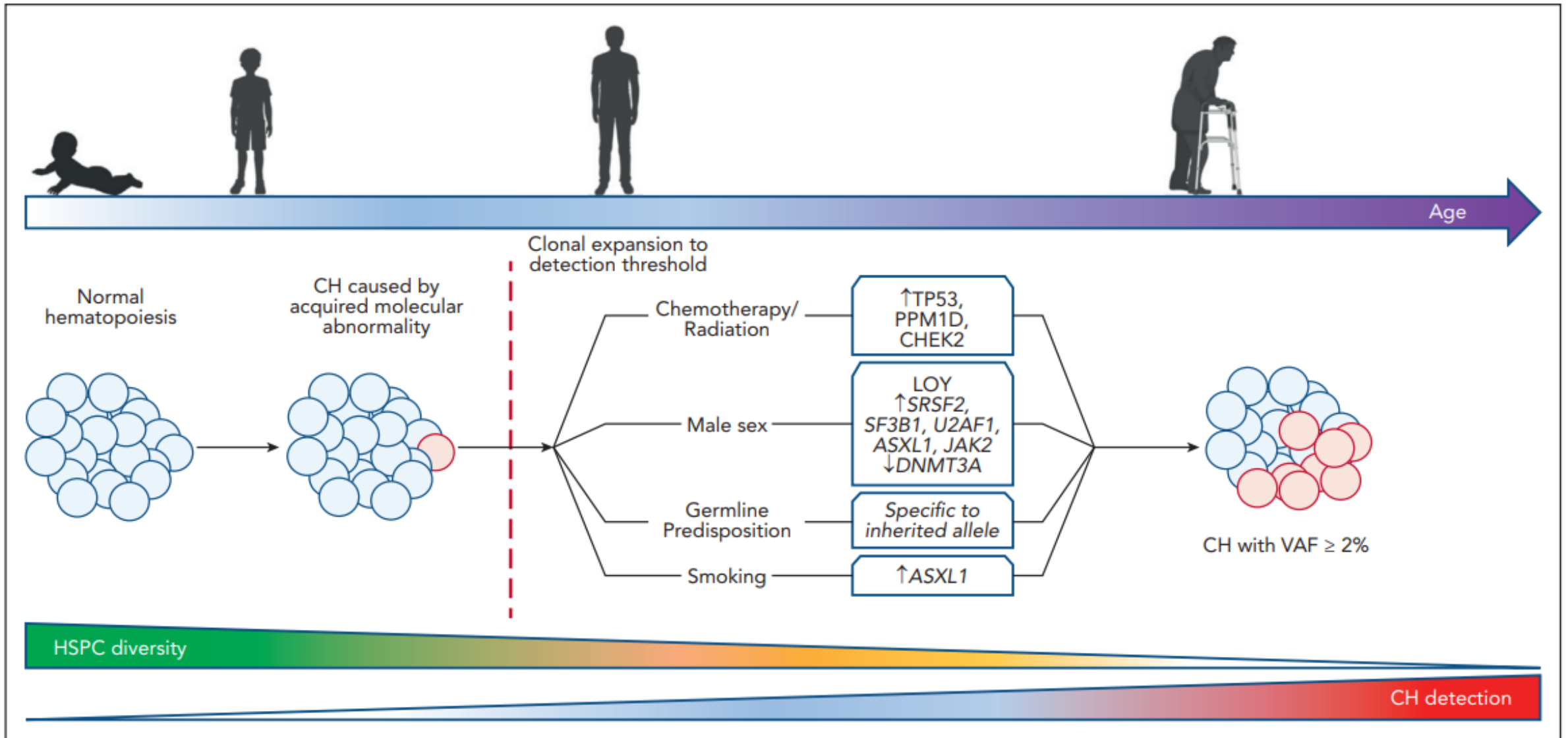
A CHIP and Coronary Heart Disease

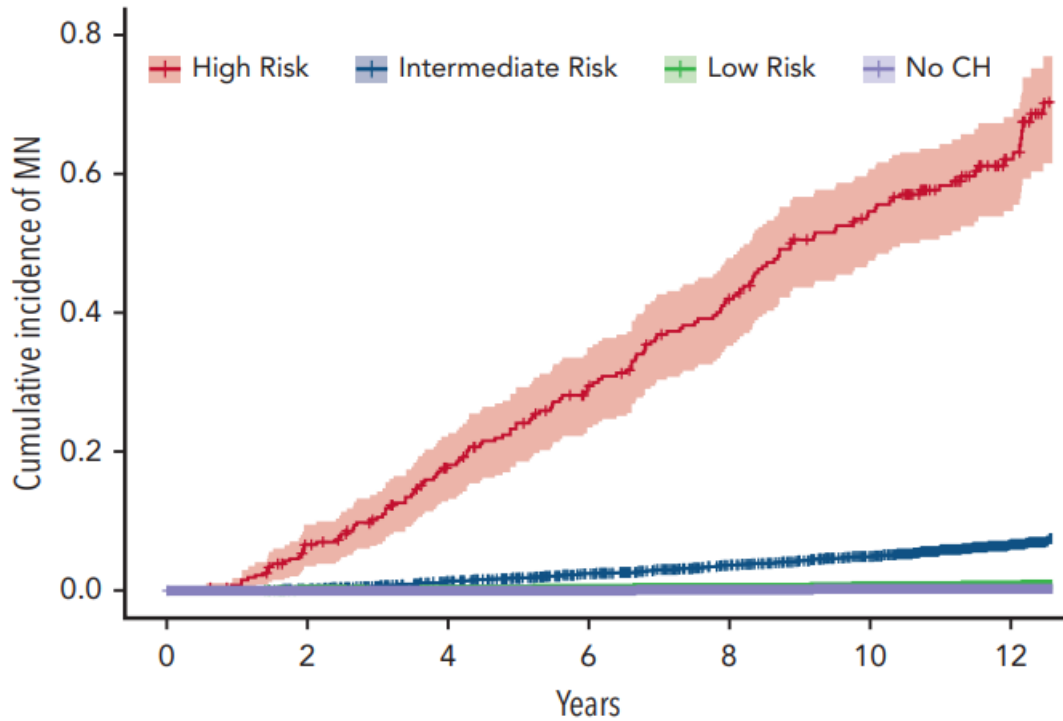


	Events	OR	P value
<i>DNMT3A</i>	31/46	1.4 (0.7–2.8)	0.29
<i>TET2</i>	12/13	8.3 (1.2–357.5)	0.02
<i>ASXL1</i>	8/8	Undefined	0.02
<i>JAK2</i>	16/16	Undefined	<0.001

Aortic-Root Sections, According to *Tet2* Status



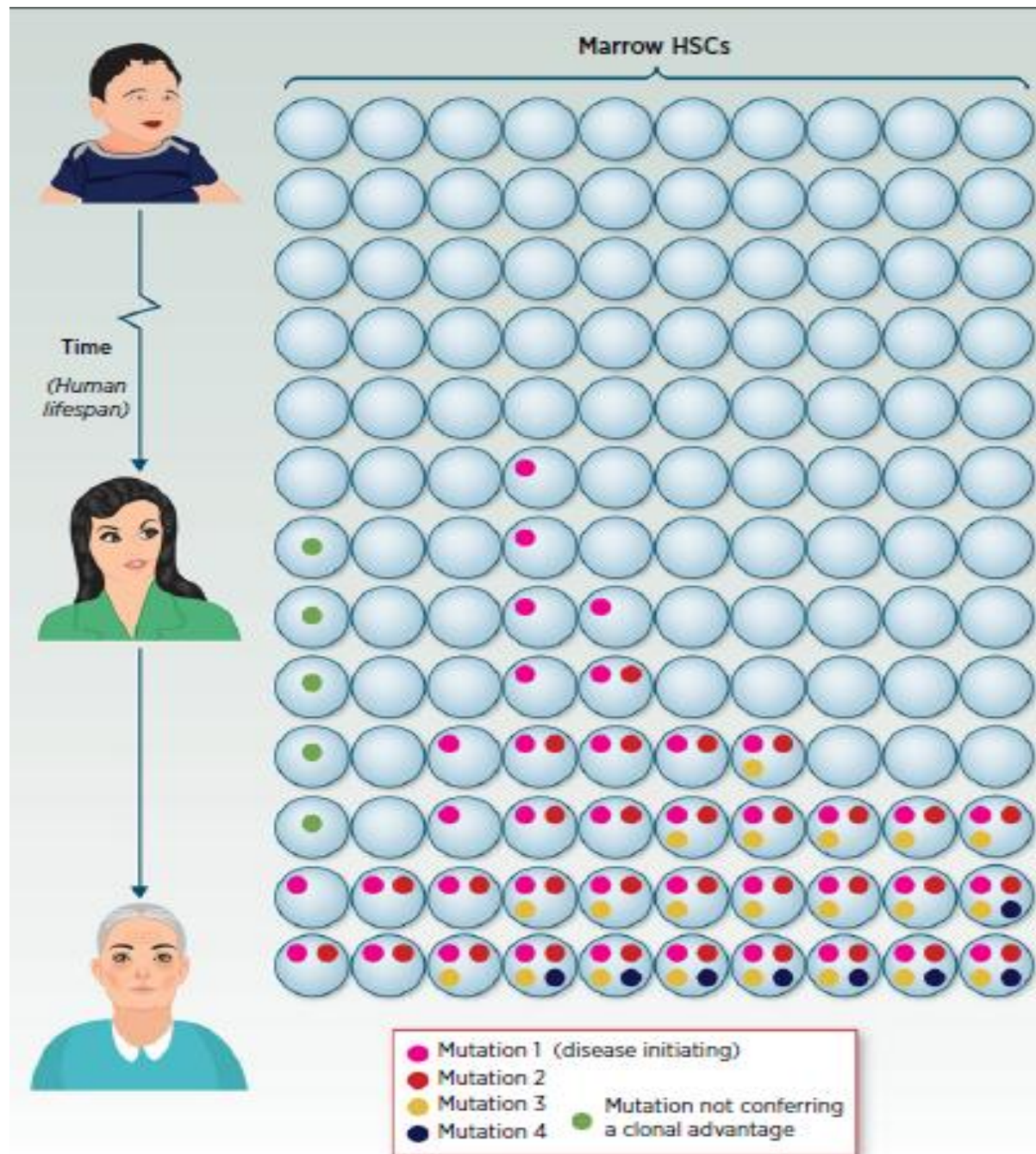




	Number at risk						
	0	2	4	6	8	10	12
High Risk	265	237	193	156	123	90	38
Intermediate Risk	4855	4763	4614	4449	4266	4062	1758
Low Risk	24715	24588	24325	24013	23613	23161	10721
No CH	441121	438984	435665	431461	426319	420058	197093

Years

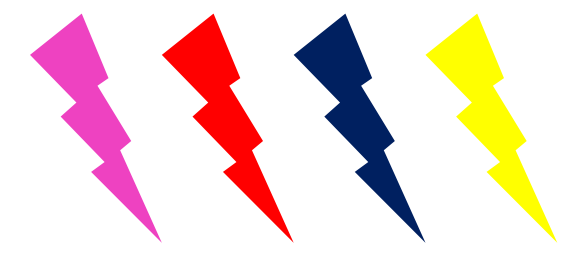
CHRS Prognostic Variable Scores					
Prognostic Variable	0.5	1	1.5	2	2.5
Single <i>DNMT3A</i>	present	absent	-	-	-
High Risk Mutation	-	absent	-	-	present
Mutation Number	-	1	-	≥ 2	-
Variant Allele Fraction	-	< 0.2	-	> 0.2	-
Red Cell Distribution Width	-	< 15	-	-	≥ 15
Mean Corpuscular Volume	-	< 100	-	-	> 100
Cytopenia	-	CHIP	CCUS	-	-
Age	-	< 65y	≥ 65y	-	-



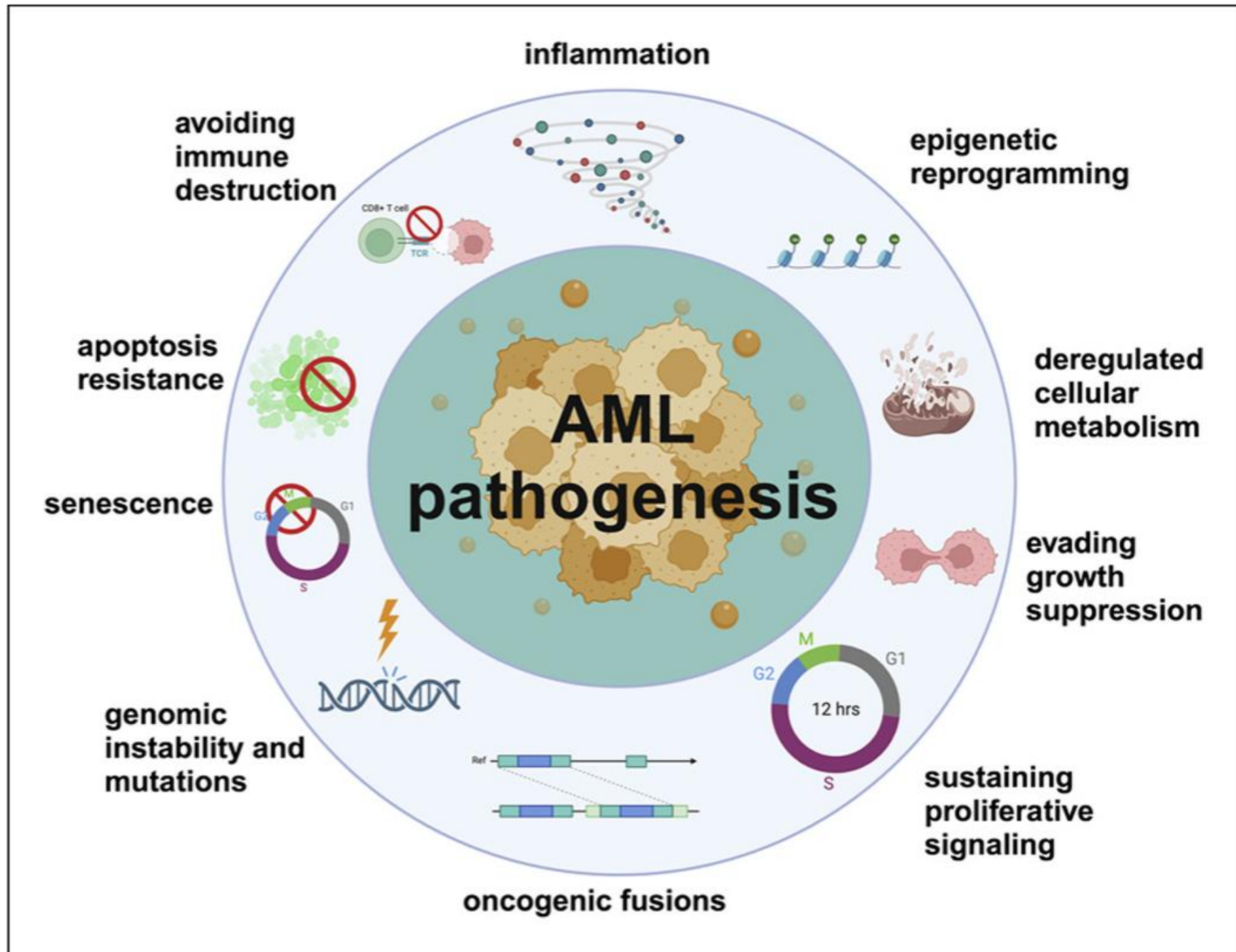
תאי אב תקינים
Normal hematopoiesis



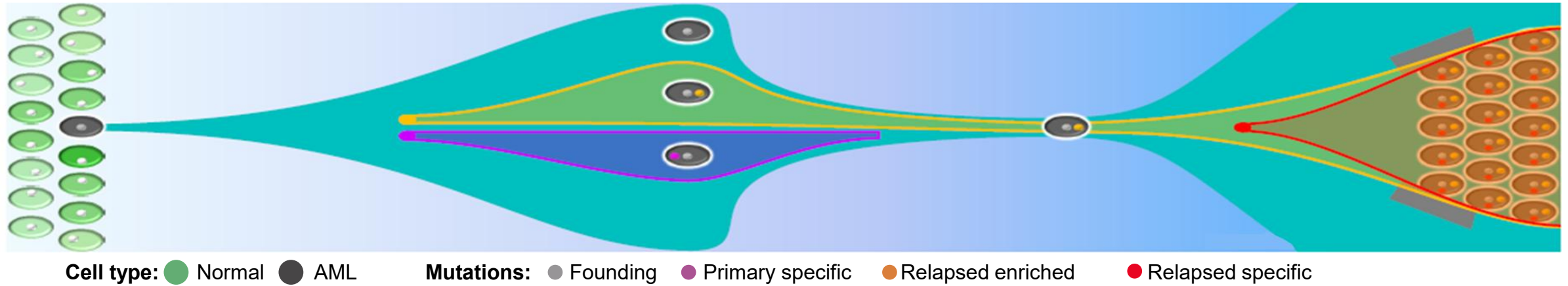
שינוי גנטי נרכש (סומטי)
Clonal hematopoiesis



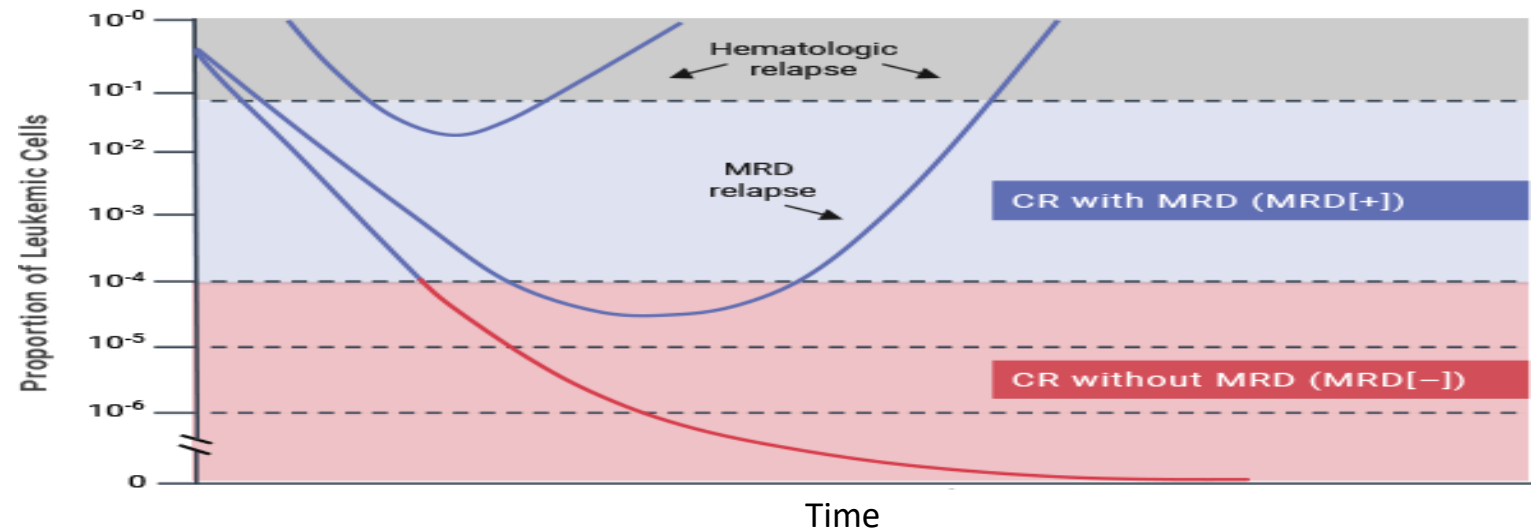
מחלה
MDS/AML



AML is a heterogenous, clonally complex disease¹



Genetic, epigenetic and metabolic insights inform therapeutic goals²

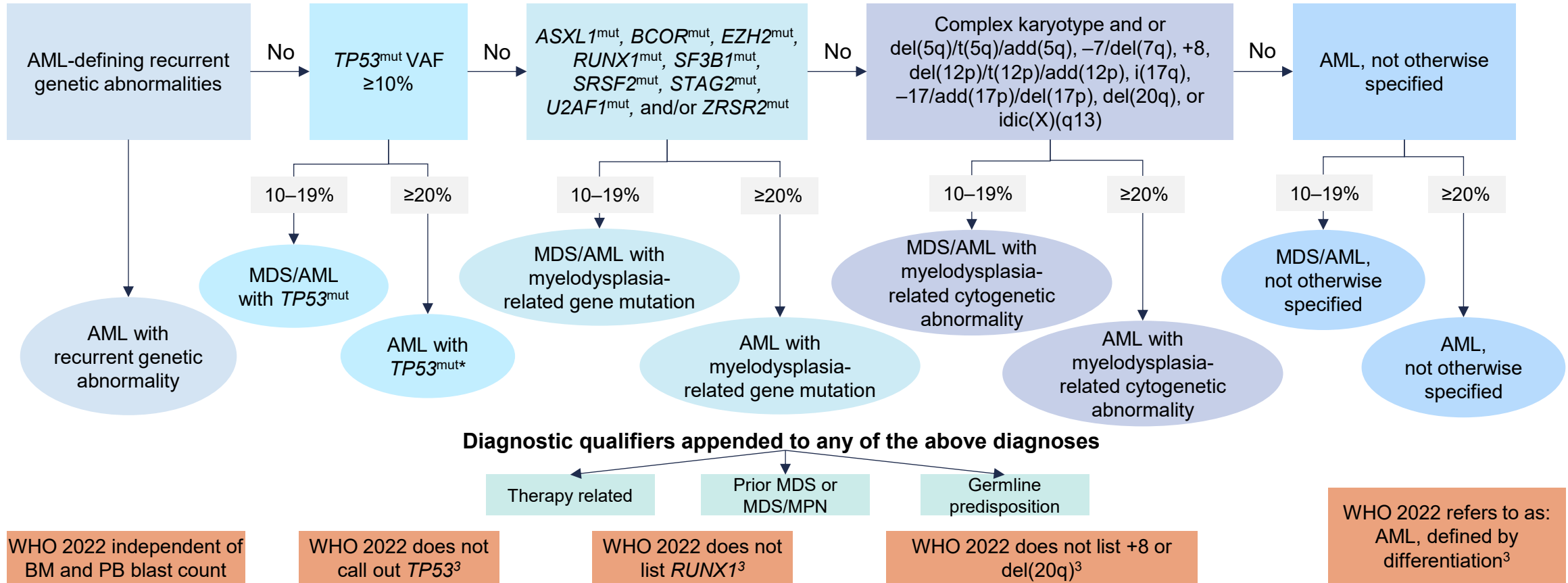


1. Ding L, et al. *Nature* 2012; **481**:506–510.

2. Short NJ, et al. *Am J Hematol* 2019; **94**:257–265.

ICC 2022 and WHO 2022 classification

International Consensus Classification of AML – Hierarchy ≥10% myeloid blasts or blast equivalents in bone marrow or blood



* The presence of TP53 mutation (VAF ≥10% with/without loss of TP53^{wt}) defines the entity AML with TP53^{mut}; mut, mutated; VAF, variant allelic frequency; wt, wild type; 1. Adapted from Döhner H, et al. Blood 2022; 140:1345–1377; 2. Arber DA, et al. Blood 2022; 140:1200–1228; 3. Khoury JD, et al. Leukemia 2022; 36:1703–1719

ICC vs. WHO

Diagnosis of AML

ICC 2022¹

WHO 5th Edition²

NPM1 mutation

≥10% blasts

Regardless of blast count

TP53 mutation

≥20% blasts

Not a separate category

CEBPA mutation

≥10% blasts
bZIP mutation only

≥20% blasts
Biallelic or *bZIP* mutation

AML-MR

0-19% blasts: MDS/AML
≥20% blasts: AML

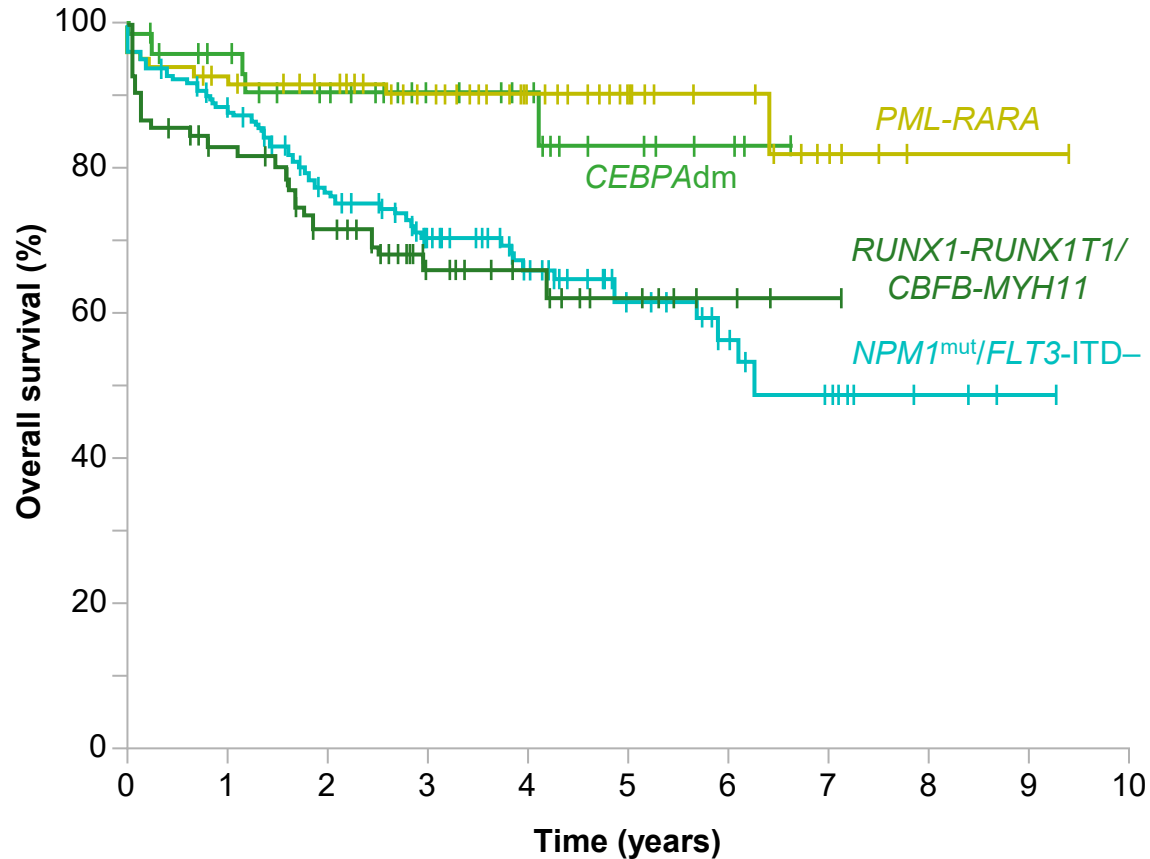
≥20% blasts
RUNX1^{mut}, +8 not included

AML, acute myeloid leukemia; AML-MR, AML-myelodysplasia-related; HMA, hypomethylating agent; IC, intensive chemotherapy; ICC, International Consensus Classification; MDS, myelodysplastic syndrome; Tx, treatment; Ven, venetoclax; WHO, World Health Organization.

1. Arber DA, et al. *Blood*. 2022;140(11):1200-1228. 2. Khoury JD, et al. *Leukemia*. 2022;36(7):1703-1719. 3. Chandra DJ, et al. *Blood Rev*. 2024 Mar;64:101156. doi: 10.1016/j.blre.2023.101156. Epub 2023 Nov 25.

Risk stratification can inform outcomes

OS in patients aged <60 years with *de novo* AML* (N=867)¹



ELN 2017²

ELN 2022³

Favorable

- *t(8;21); RUNX1-RUNX1T1*
- *inv(16) or t(16;16); CBFB-MYH11*
- Mutated *NPM1* without *FLT3-ITD* or *FLT3-ITD*^{low†}
- **Biallelic-mutated *CEBPA***

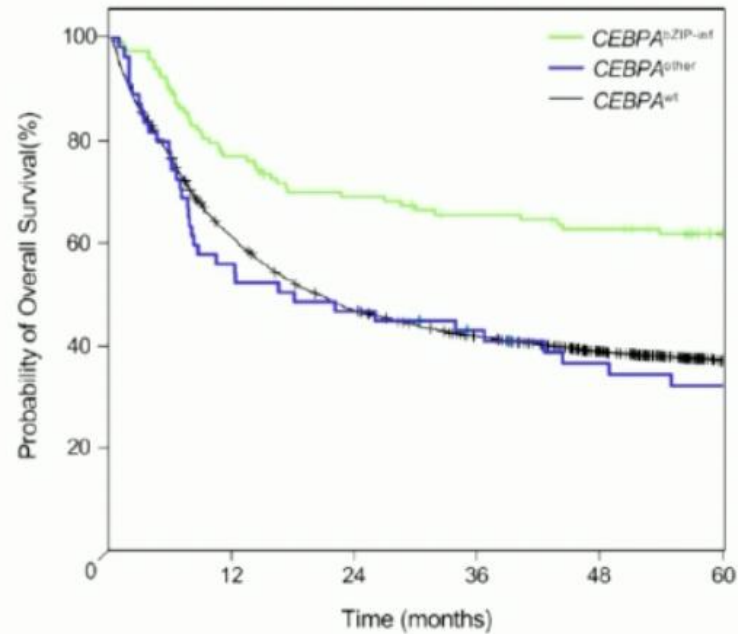
- bZIP in-frame mutated *CEBPA*

* This population of patients had been previously treated with intensive chemotherapy; † Low allelic ratio (<0.5) vs high allelic ratio (≥0.5).

1. Haferlach C, et al. *Blood* 2016; **128**:286; 2. Döhner H, et al. *Blood* 2017; **129**:424–447; 3. Döhner H, et al. *Blood* 2022; doi: 10.1182/blood.2022016867.

Survival analysis by *CEBPA* mutation status

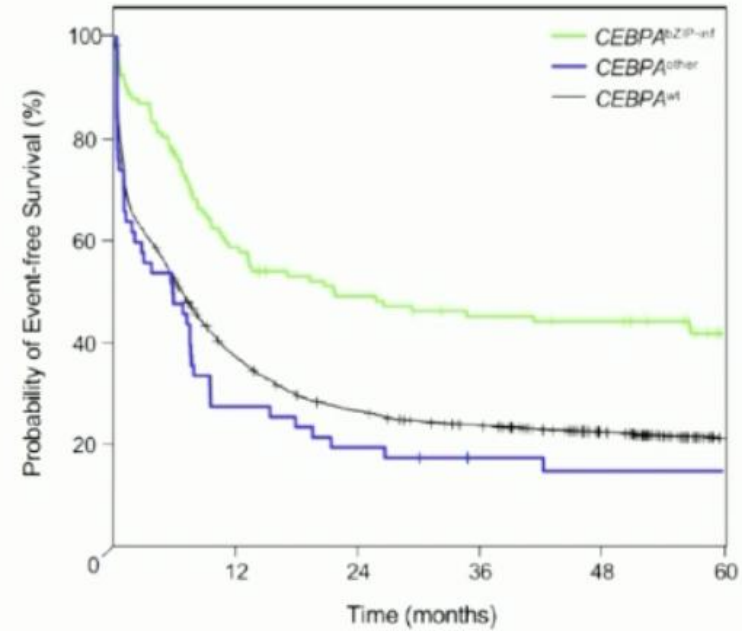
Overall survival



<i>CEBPA</i> ^{wt}	4225	2472	1777	1461	1169	921
<i>CEBPA</i> ^{bZIP-inf}	154	117	102	93	77	66
<i>CEBPA</i> ^{other}	82	42	30	25	20	15

	<i>CEBPA</i> ^{wt}	<i>CEBPA</i> ^{other}
<i>CEBPA</i> ^{other}	0.480	-
<i>CEBPA</i> ^{bZIP-inf}	<0.001	<0.001

Event-free survival



<i>CEBPA</i> ^{wt}	4225	1593	1100	926	768	625
<i>CEBPA</i> ^{bZIP-inf}	154	92	75	68	60	49
<i>CEBPA</i> ^{other}	82	22	15	11	9	8

	<i>CEBPA</i> ^{wt}	<i>CEBPA</i> ^{other}
<i>CEBPA</i> ^{other}	0.160	-
<i>CEBPA</i> ^{bZIP-inf}	<0.001	<0.001

➤ Favorable outcome is confined to AML with bZIP in-frame *CEBPA* mutations, irrespective of mono- or biallelic

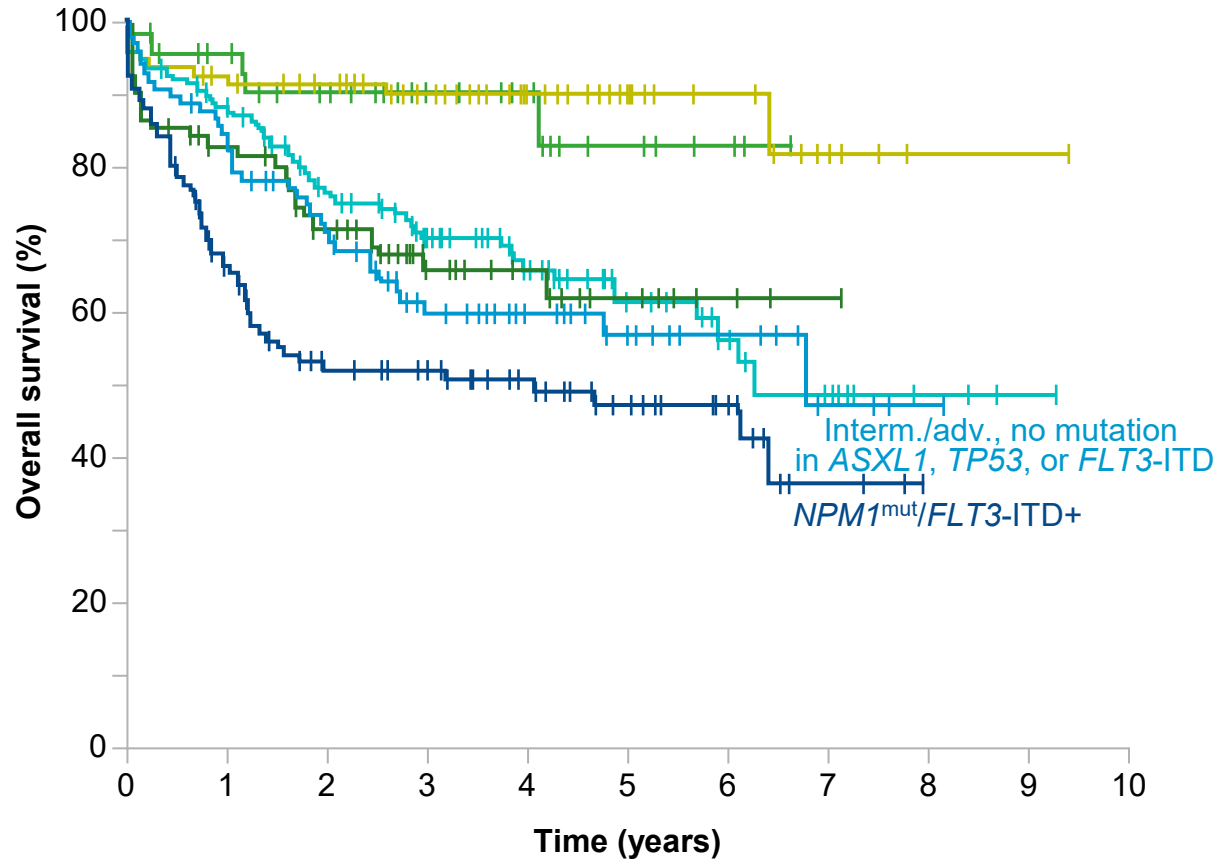
Similar data:

Tarlock K, et al. Blood 2021;138(13):1137; Wakita S, et al. Blood Advances 2021;6(1):238; Rucker F, et al. EHA 2022

Taube F et al. Blood. 2022;139(1):87-103.

Risk stratification can inform outcomes

OS in patients aged <60 years with *de novo* AML* (N=867)¹



ELN 2017²

ELN 2022³

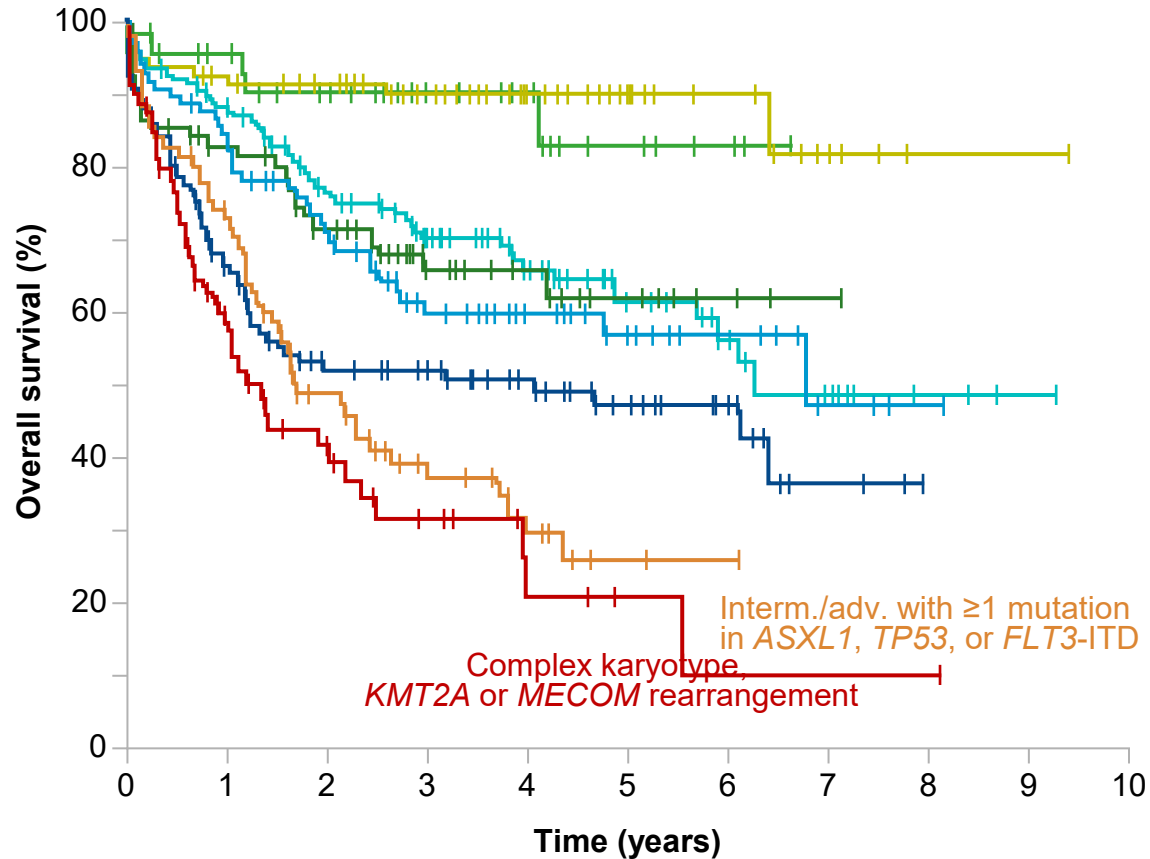
Favorable	<ul style="list-style-type: none"> • t(8;21); <i>RUNX1-RUNX1T1</i> • inv(16) or t(16;16); <i>CBFB-MYH11</i> • Mutated <i>NPM1</i> without <i>FLT3-ITD</i> or <i>FLT3-ITD</i>^{low†} • Biallelic-mutated <i>CEBPA</i> 	<ul style="list-style-type: none"> • bZIP in-frame mutated <i>CEBPA</i>
Intermediate	<ul style="list-style-type: none"> • Mutated <i>NPM1</i> and <i>FLT3-ITD</i>^{high†} • Wild-type <i>NPM1</i> without <i>FLT3-ITD</i> or <i>FLT3-ITD</i>^{low*,†} • t(9;11); <i>MLL3-KMT2A</i> • Abnormalities not classified as favorable/adverse 	<ul style="list-style-type: none"> • Any <i>FLT3-ITD</i>

* This population of patients had been previously treated with intensive chemotherapy; † Low allelic ratio (<0.5) vs high allelic ratio (≥0.5).

1. Haferlach C, et al. *Blood* 2016; **128**:286; 2. Döhner H, et al. *Blood* 2017; **129**:424–447; 3. Döhner H, et al. *Blood* 2022; doi: 10.1182/blood.2022016867.

Risk stratification can inform outcomes

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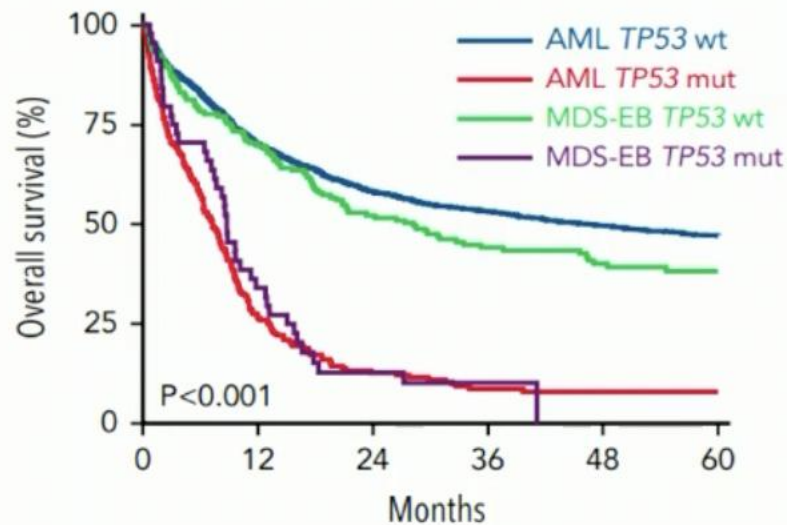
Favorable	<ul style="list-style-type: none"> t(8;21); <i>RUNX1-RUNX1T1</i> inv(16) or t(16;16); <i>CBFB-MYH11</i> Mutated <i>NPM1</i> without <i>FLT3-ITD</i> or <i>FLT3-ITD</i>^{low†} Biallelic-mutated <i>CEBPA</i> 	<ul style="list-style-type: none"> bZIP in-frame mutated <i>CEBPA</i>
Intermediate	<ul style="list-style-type: none"> Mutated <i>NPM1</i> and <i>FLT3-ITD</i>^{high†} Wild-type <i>NPM1</i> without <i>FLT3-ITD</i> or <i>FLT3-ITD</i>^{low*,†} t(9;11); <i>MLL3-KMT2A</i> Abnormalities not classified as favorable/adverse 	<ul style="list-style-type: none"> Any <i>FLT3-ITD</i>
Adverse	<ul style="list-style-type: none"> t(6;9)(p23;q34.1); <i>DEK-NUP214</i> t(v;11q23.3); <i>KMT2A</i> rearranged t(9;22)(q34.1;q11.2); <i>BCR-ABL1</i> inv(3)(q21.3q26.2) or t(3;3)(q21.3;q26.2); <i>GATA2</i>, <i>MECOM(EVI1)</i> -5 or del(5q); -7; -17/abn(17p) Complex or monosomal karyotype Wild-type <i>NPM1</i> and <i>FLT3-ITD</i>^{high†} Mutated <i>RUNX1</i>, <i>ASXL1</i>, or <i>TP53</i> 	<ul style="list-style-type: none"> Mutated <i>ASXL1</i>, <i>BCOR</i>, <i>EZH2</i>, <i>RUNX1</i>, <i>SF3B1</i>, <i>SRSF2</i>, <i>STAG2</i>, <i>U2AF1</i>, or <i>ZRSR2</i> Mutated <i>TP53</i>

* This population of patients had been previously treated with intensive chemotherapy; † Low allelic ratio (<0.5) vs high allelic ratio (≥0.5).

1. Haferlach C, et al. *Blood* 2016; **128**:286; 2. Döhner H, et al. *Blood* 2017; **129**:424–447; 3. Döhner H, et al. *Blood* 2022; doi: 10.1182/blood.2022016867.

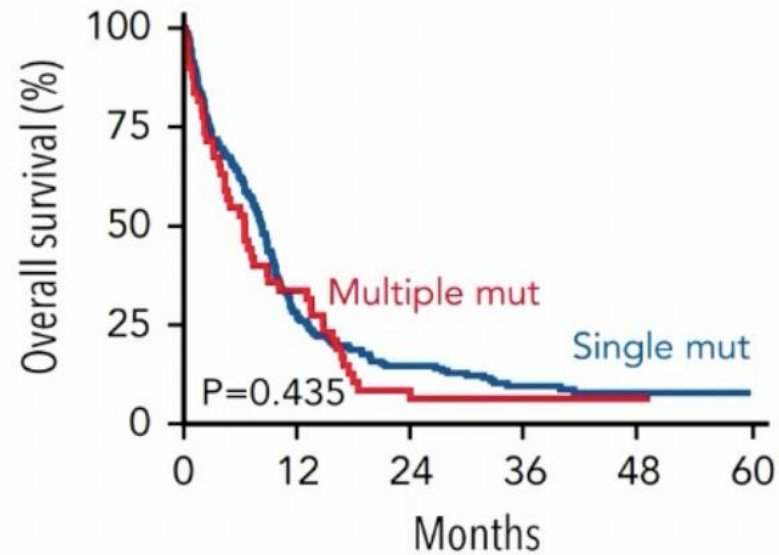
TP53 mutation in AML and MDS with excess blasts

- Accumulating evidence from both a clinical and molecular perspective that AML and MDS with mutated *TP53* represent a distinct molecular disease entity



No. at risk:

AML <i>TP53</i> wt	1805	1255	1014	807	572	397
AML <i>TP53</i> mut	186	50	23	14	9	5
MDS-EB <i>TP53</i> wt	165	114	81	62	45	35
MDS-EB <i>TP53</i> mut	44	15	5	1	0	0

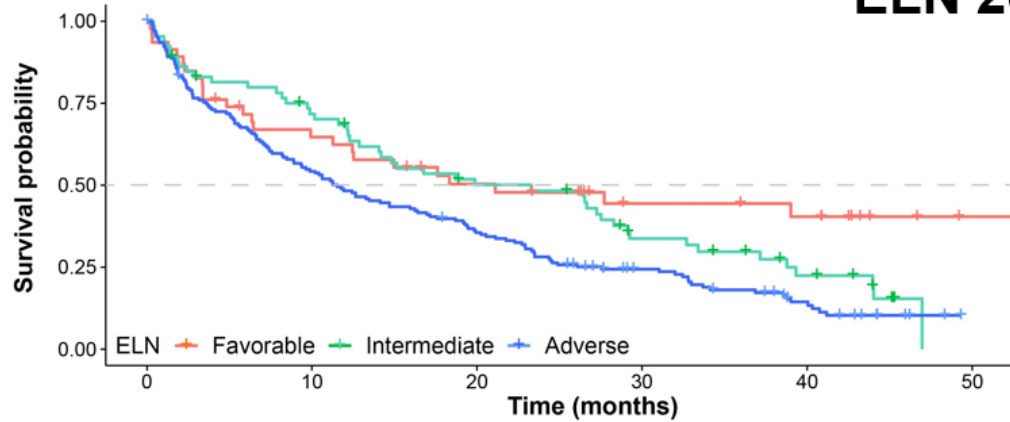


No. at risk:

Single mut	181	49	25	12	6	3
Multiple mut	49	16	3	3	3	2

ELN risk stratification perform poorly in lower-intensity treated patients

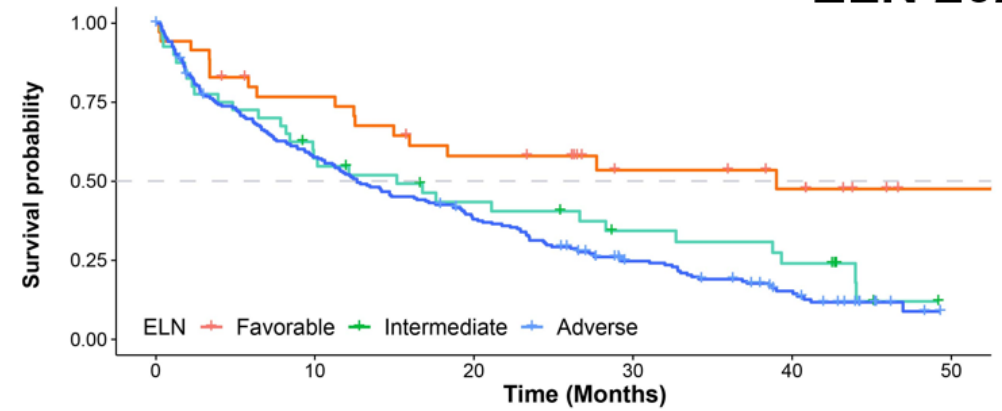
ELN 2017



		Patients at risk					
		0	10	20	30	40	50
ELN	Favorable	46	28	20	12	10	2
	Intermediate	65	44	29	17	9	0
	Adverse	168	90	58	31	14	0

ELN 2017	n	Events	Median OS, mo (95% CI)
Favorable	46	25	21.09 (9.92 – NE)
Intermediate	65	48	23.26 (12.85 – 28.29)
Adverse	168	141	11.53 (8.87 – 16.23)

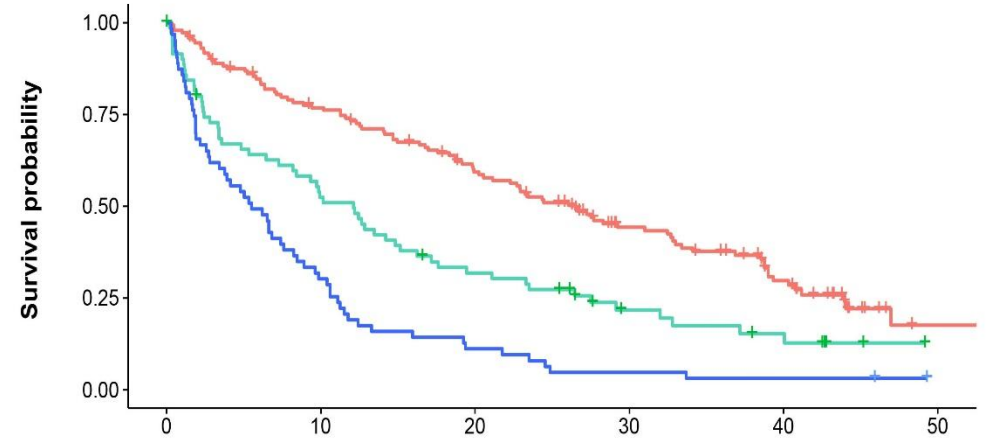
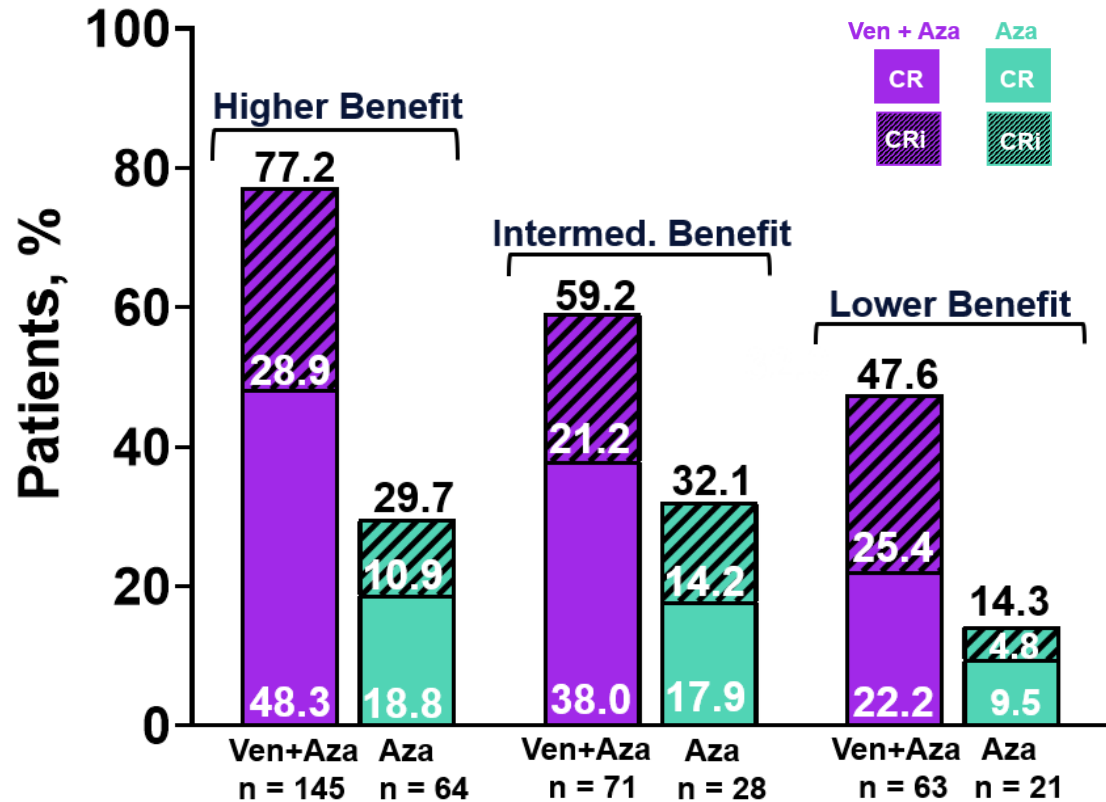
ELN 2022



		Patients at Risk					
		0	10	20	30	40	50
ELN	Favorable	35	25	18	11	8	2
	Intermediate	40	22	15	10	7	0
	Adverse	204	115	74	39	18	0

ELN 2022	n	Events	Median OS, mo (95% CI)
Favorable	35	16	39.0 (12.52 – NE)
Intermediate	40	30	15.15 (8.18 – 28.29)
Adverse	204	168	12.65 (10.41 – 17.15)

Genetic Risk Stratification and Outcomes for Ven-HMA Treated patients



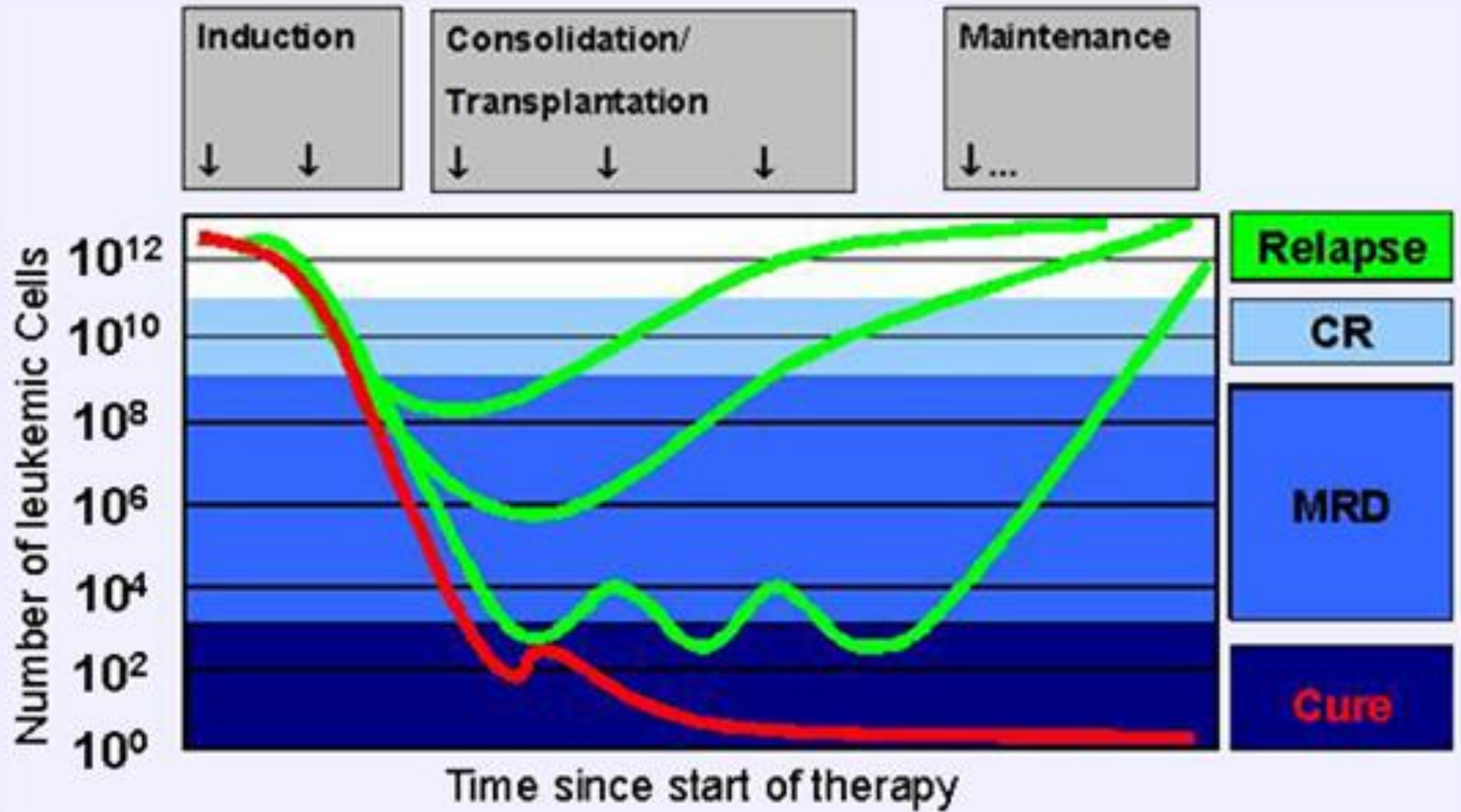
Benefit Group

Benefit Group	Patients at Risk	0	10	20	30	40	50
Higher Benefit	145	107	79	47	25	2	0
Interm. Benefit	71	36	21	10	6	0	0
Lower Benefit	63	19	7	3	2	0	0

Ven + Aza (N = 279)	n	Events	Median OS, months (95% CI)
Higher Benefit	145	96	26.51 (20.24, 32.69)
Intermediate Benefit	71	57	12.12 (7.26 - 15.15)
Lower Benefit	63	61	5.52 (2.79 - 7.59)

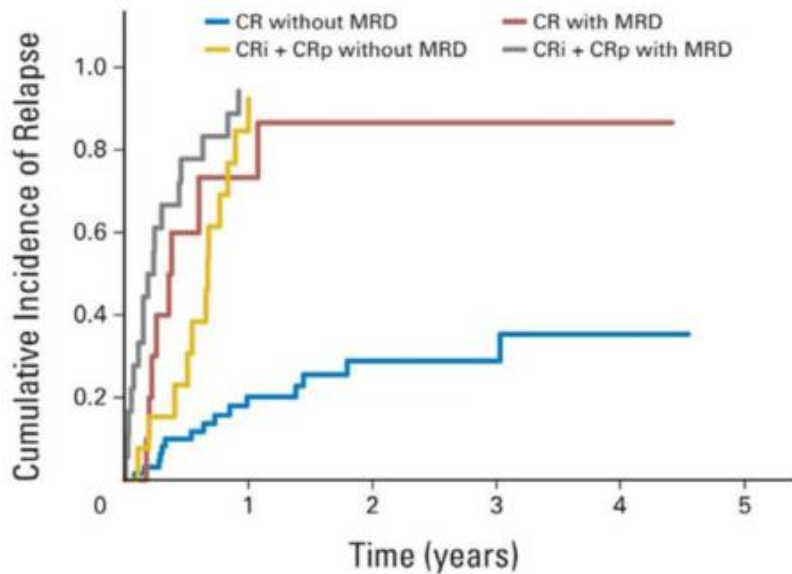
ELN risk classification for patients receiving less-intensive therapies

Risk category	Genetic abnormality
Favorable	<ul style="list-style-type: none">• Mutated <i>NPM1</i> (<i>FLT3-ITD</i>^{neg}, <i>NRAS</i>^{wt}, <i>KRAS</i>^{wt}, <i>TP53</i>^{wt})• Mutated <i>IDH2</i> (<i>FLT3-ITD</i>^{neg}, <i>NRAS</i>^{wt}, <i>KRAS</i>^{wt}, <i>TP53</i>^{wt})• Mutated <i>IDH1</i>^b (<i>TP53</i>^{wt})• Mutated <i>DDX41</i>^c• Other cytogenetic and/or molecular abnormalities^d (<i>FLT3-ITD</i>^{neg}, <i>NRAS</i>^{wt}, <i>KRAS</i>^{wt}, <i>TP53</i>^{wt})
Intermediate	<ul style="list-style-type: none">• Other cytogenetic and molecular abnormalities^d (<i>FLT3-ITD</i>^{pos} and/or <i>NRAS</i>^{mut} and/or <i>KRAS</i>^{mut}; <i>TP53</i>^{wt})
Adverse	<ul style="list-style-type: none">• Mutated <i>TP53</i>



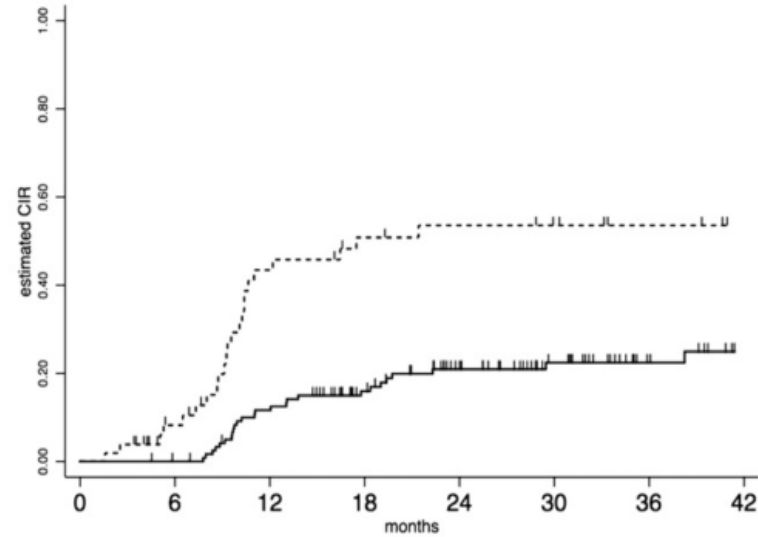
Minimal Residual Disease

Flow-based



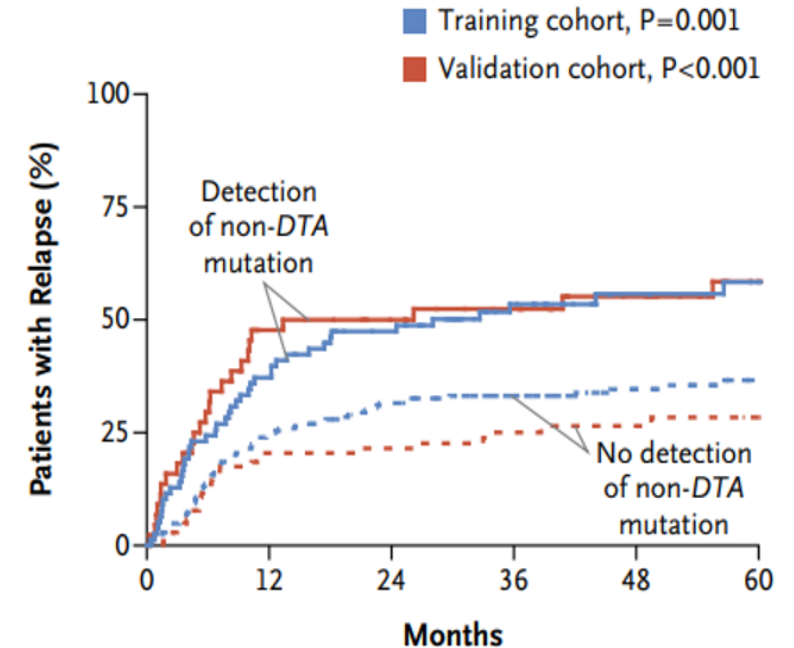
Chen et al. JCO 2015

RT-PCR



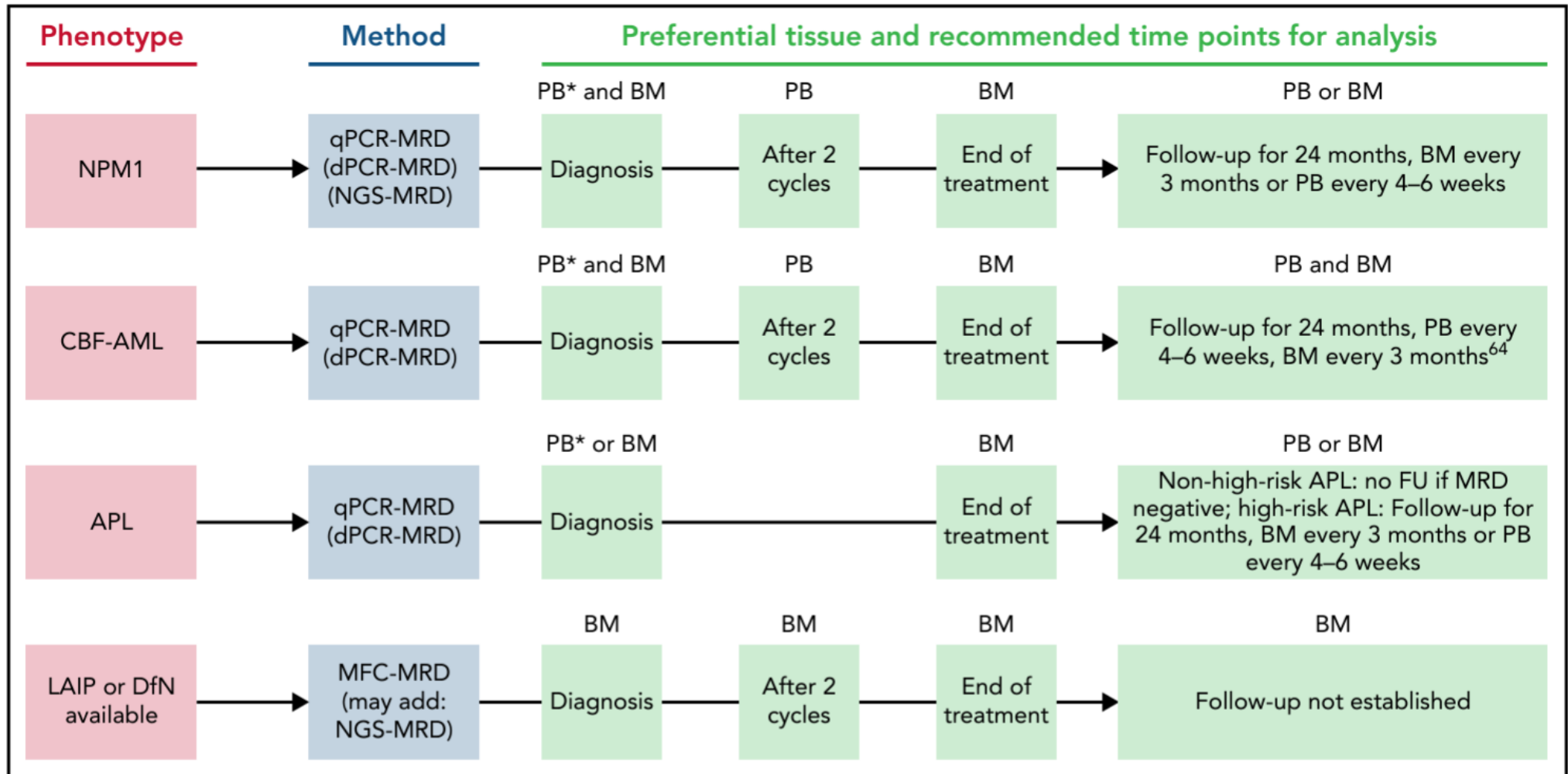
Jourdan Blood 2013; Yin Blood 2012

NGS

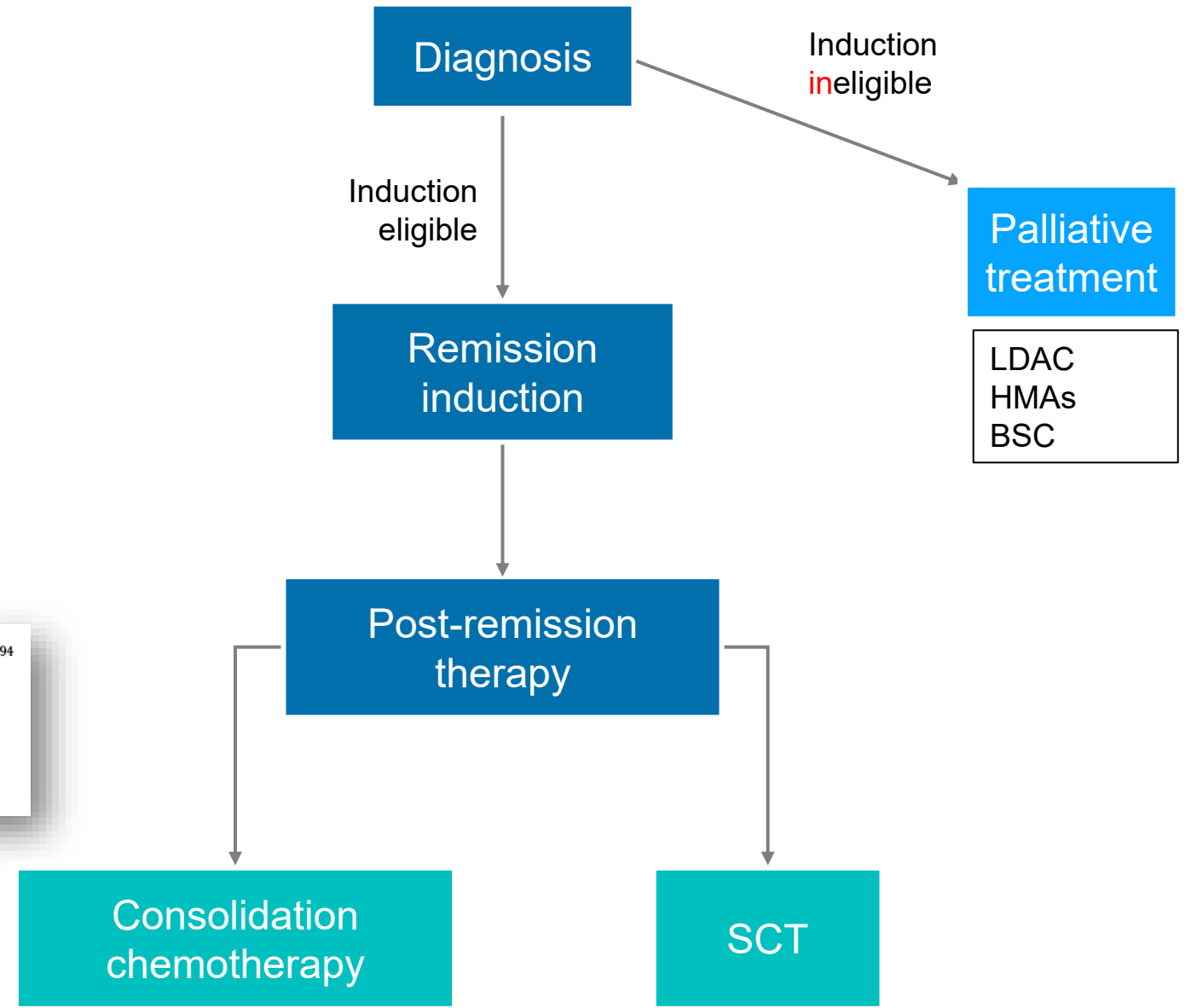
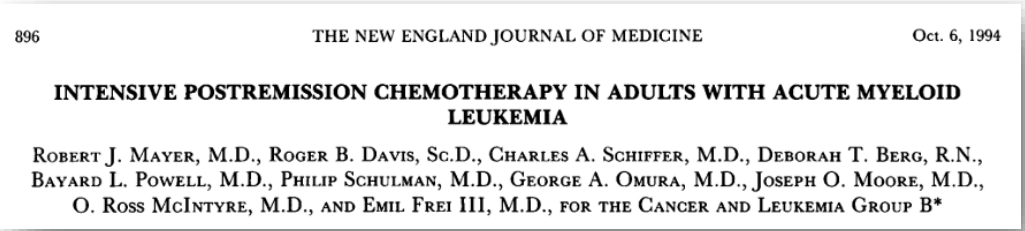
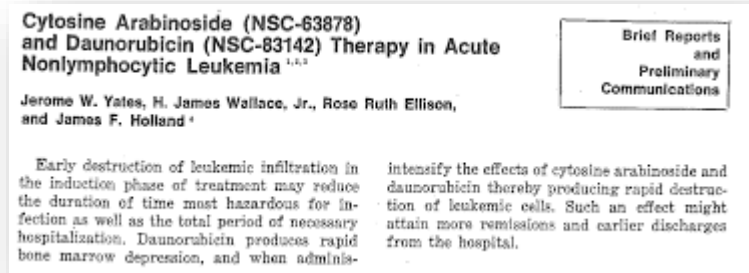


Jongen-Lavrencic et al, NEJM 2018

2021 Update on MRD in acute myeloid leukemia: a consensus document from the European LeukemiaNet MRD Working Party



The historical AML “dogma”

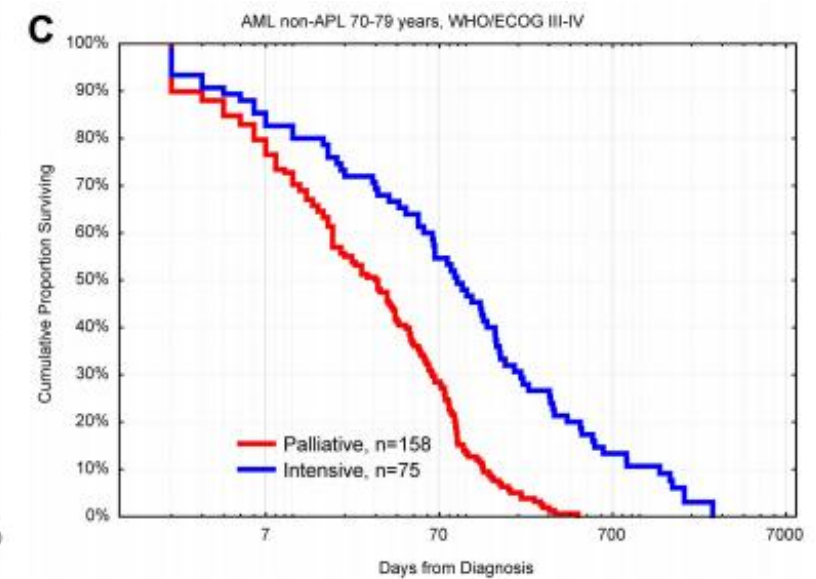
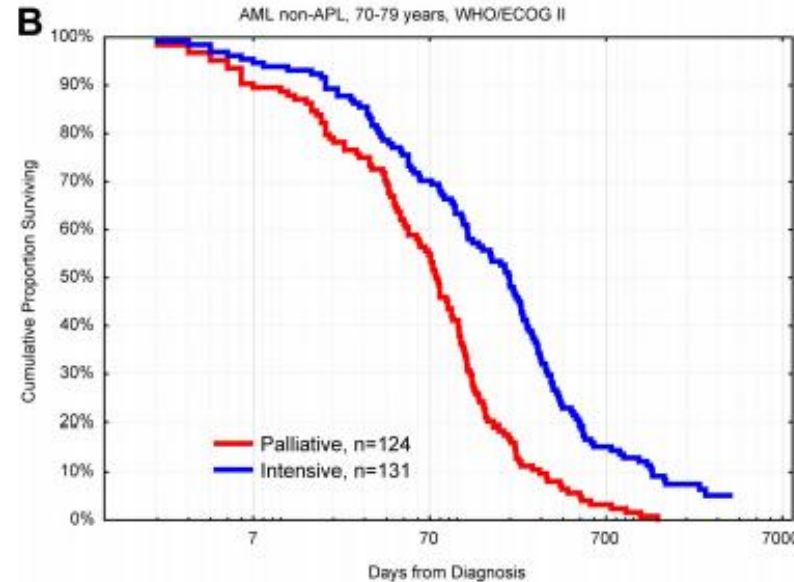
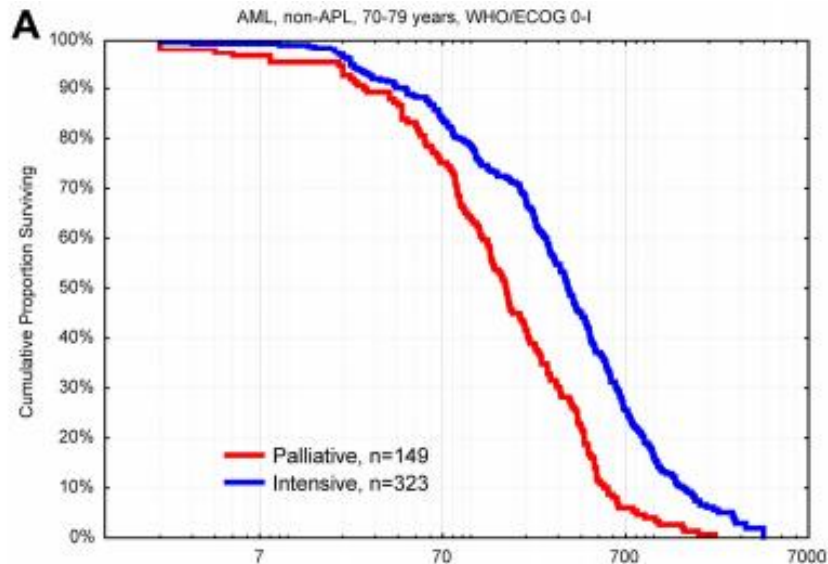


Resistance to therapy, **not** TRM – the leading cause of death in older AML

Resistance vs treatment-related mortality (TRM)^a

Age	Patients (N)	Complete remission	TRM	Resistant
<56 years	368	64%	9%	27%
56–65 years	246	46%	17%	37%
66–75 years	274	39%	24%	37%
>75 years	80	33%	31%	36%

Based on Appelbaum et al. Blood 2006; 107:3481–3485. [10].

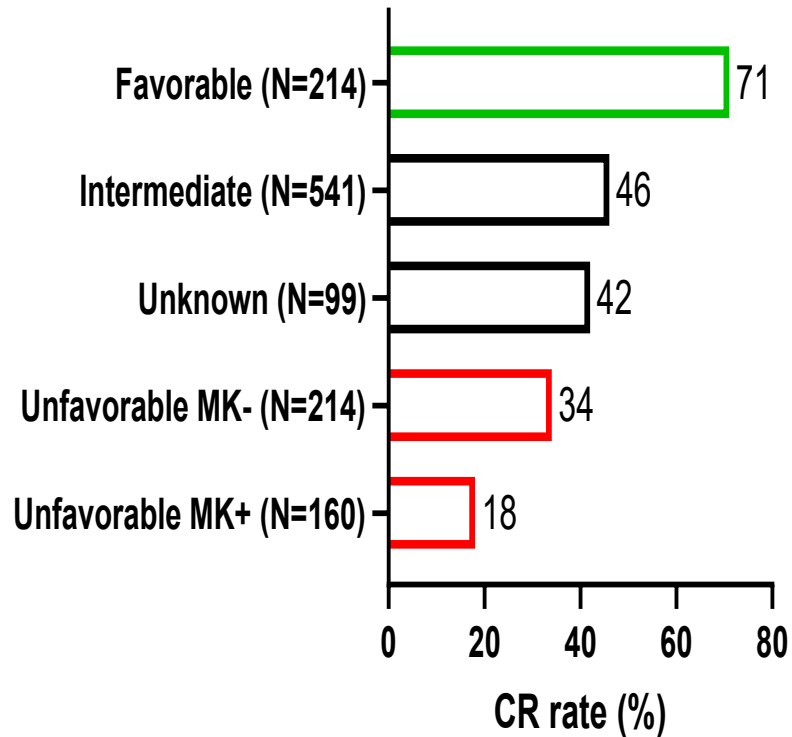


Response

versus

Early death

Remission and cytogenetics in multiple SWOG trials (N=1344)



Adapted from
Medeiros et al. Blood 2010, Sep 30;116(13):2224-8

analysis by number of adverse prognostic factors

No. of adverse factors	No. of patients (%)	8-wk mortality, %	CR, %	Survival		
				Median, mo	2-y, %	3-y, %
0	122 (28)	16	57	11.3	30	22
1	170 (40)	31	52	5.3	15	7
2	100 (23)	55	29	1.5	7	6
≥ 3	38 (9)	71	16	0.5	0	0

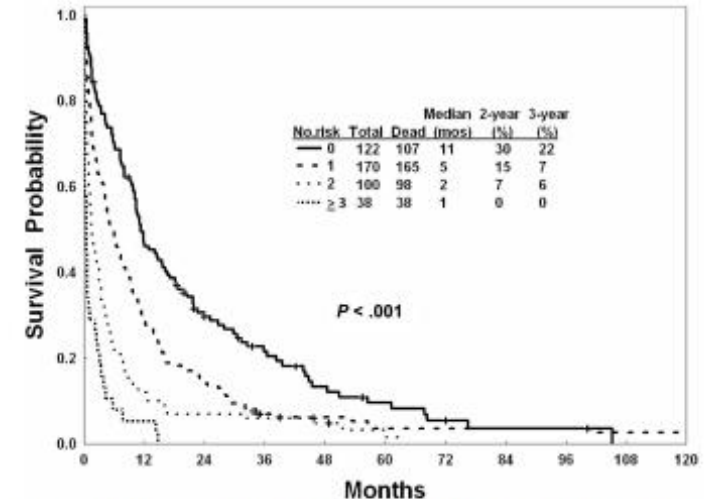
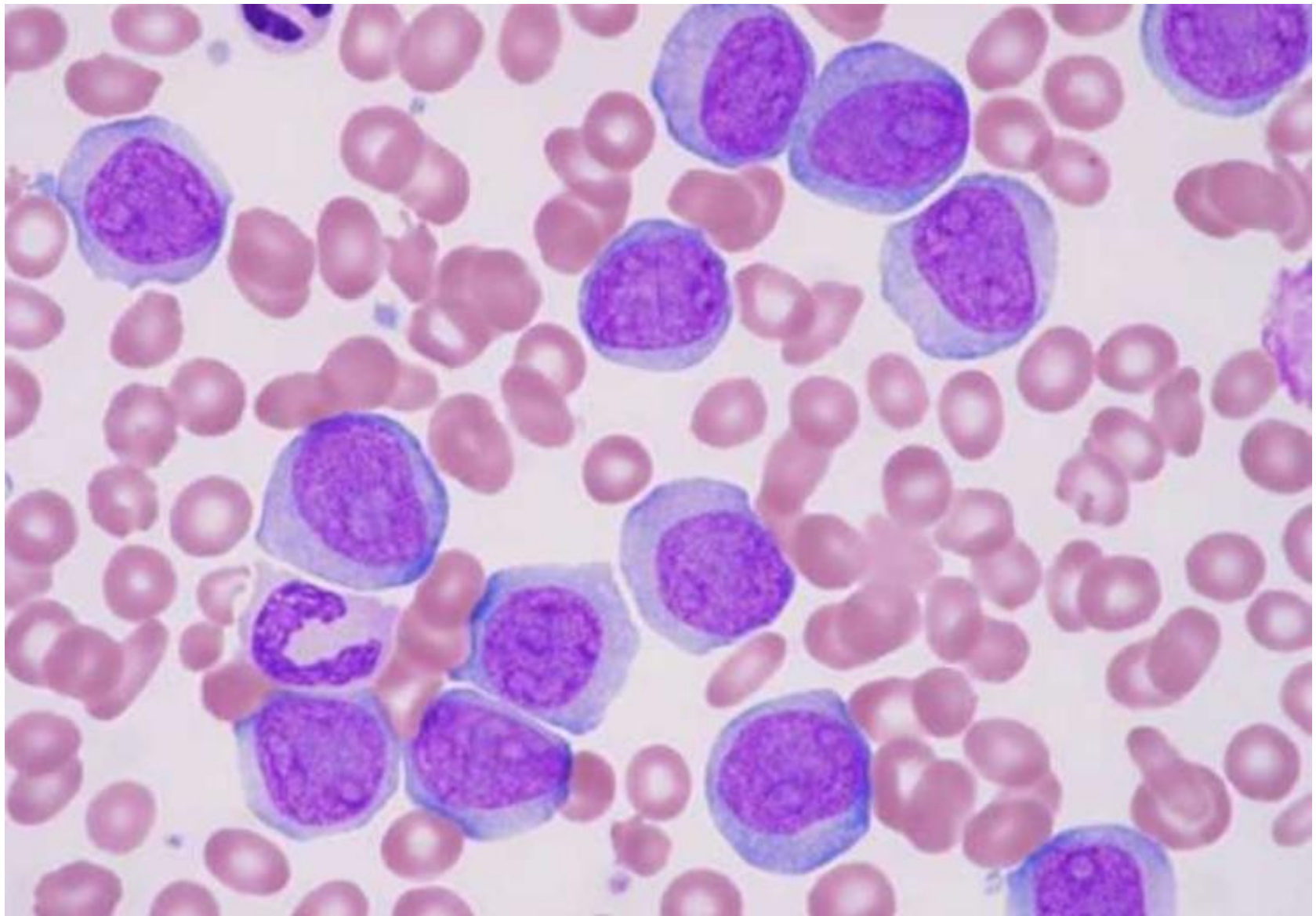
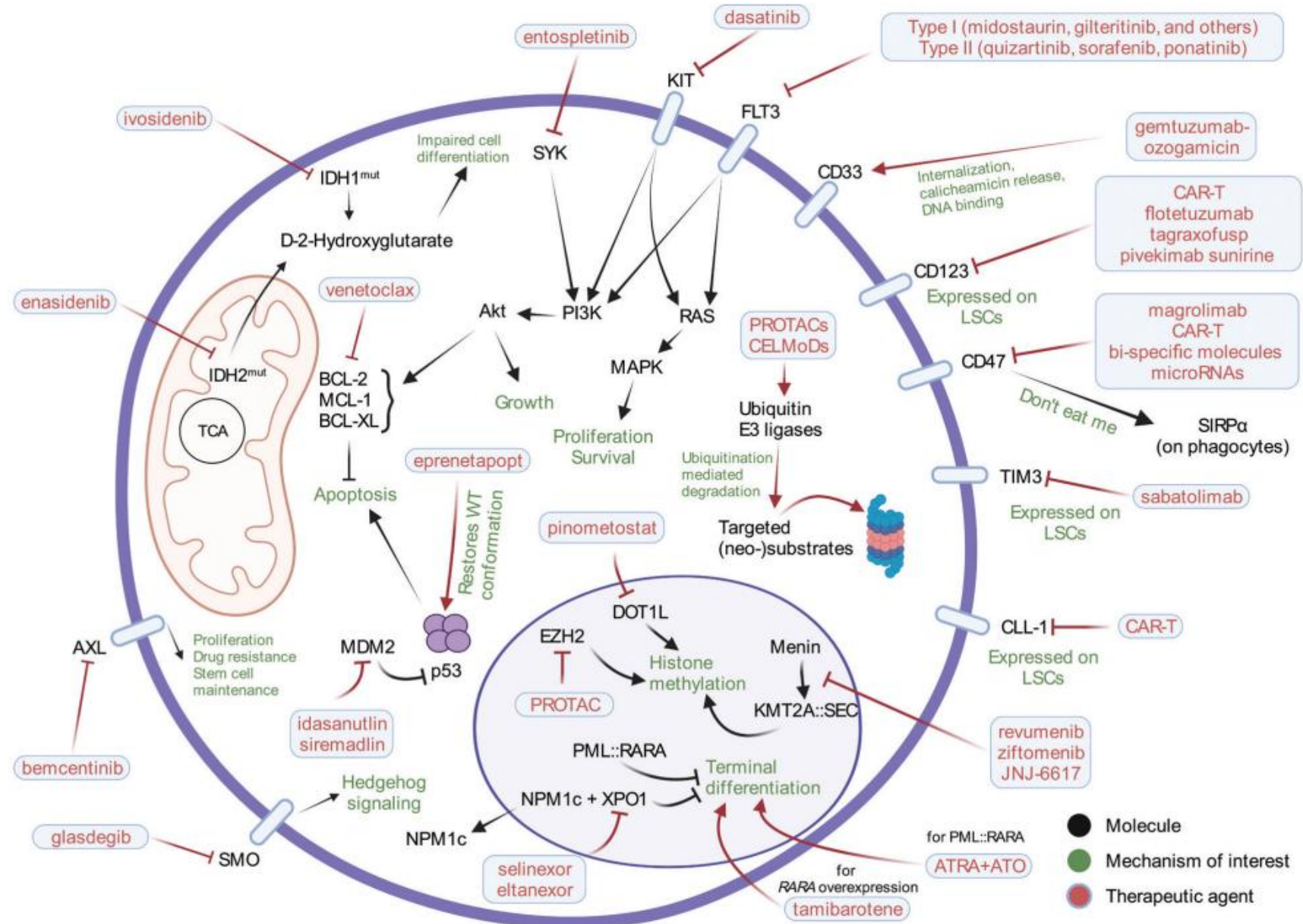


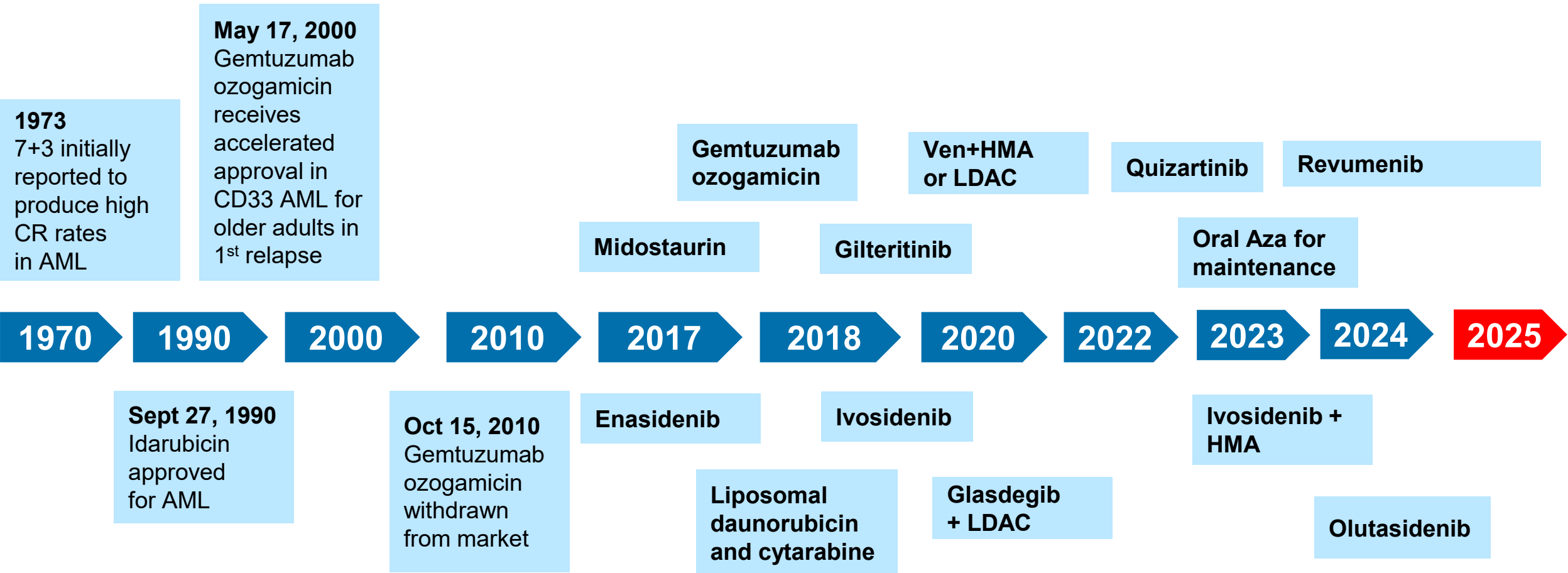
Figure 4. Survival of 430 elderly patients with AML by number of independent risk factors for 8-week mortality.

Kantarjian Blood 2010, Nov 25;116(22):4422-9

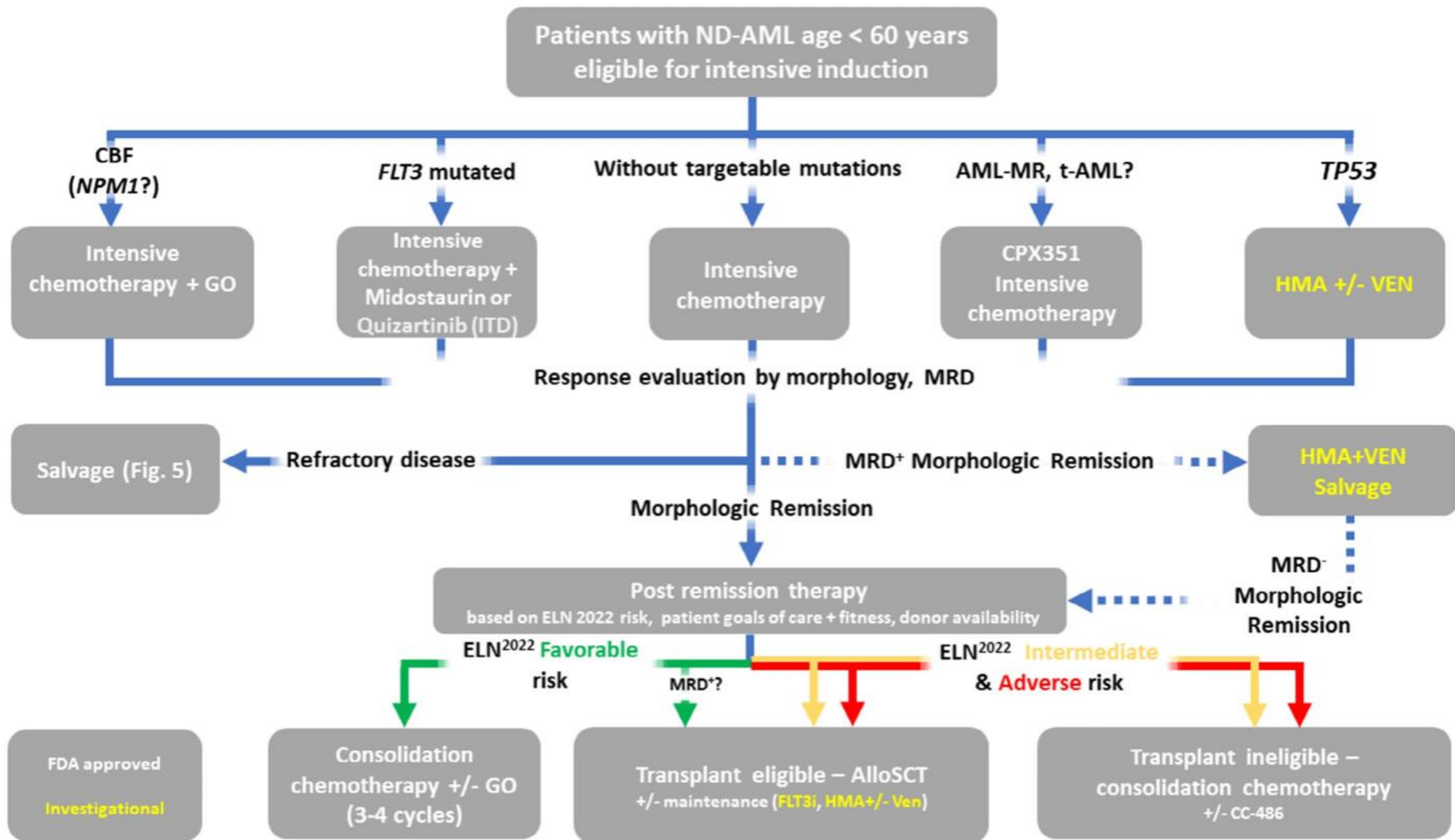


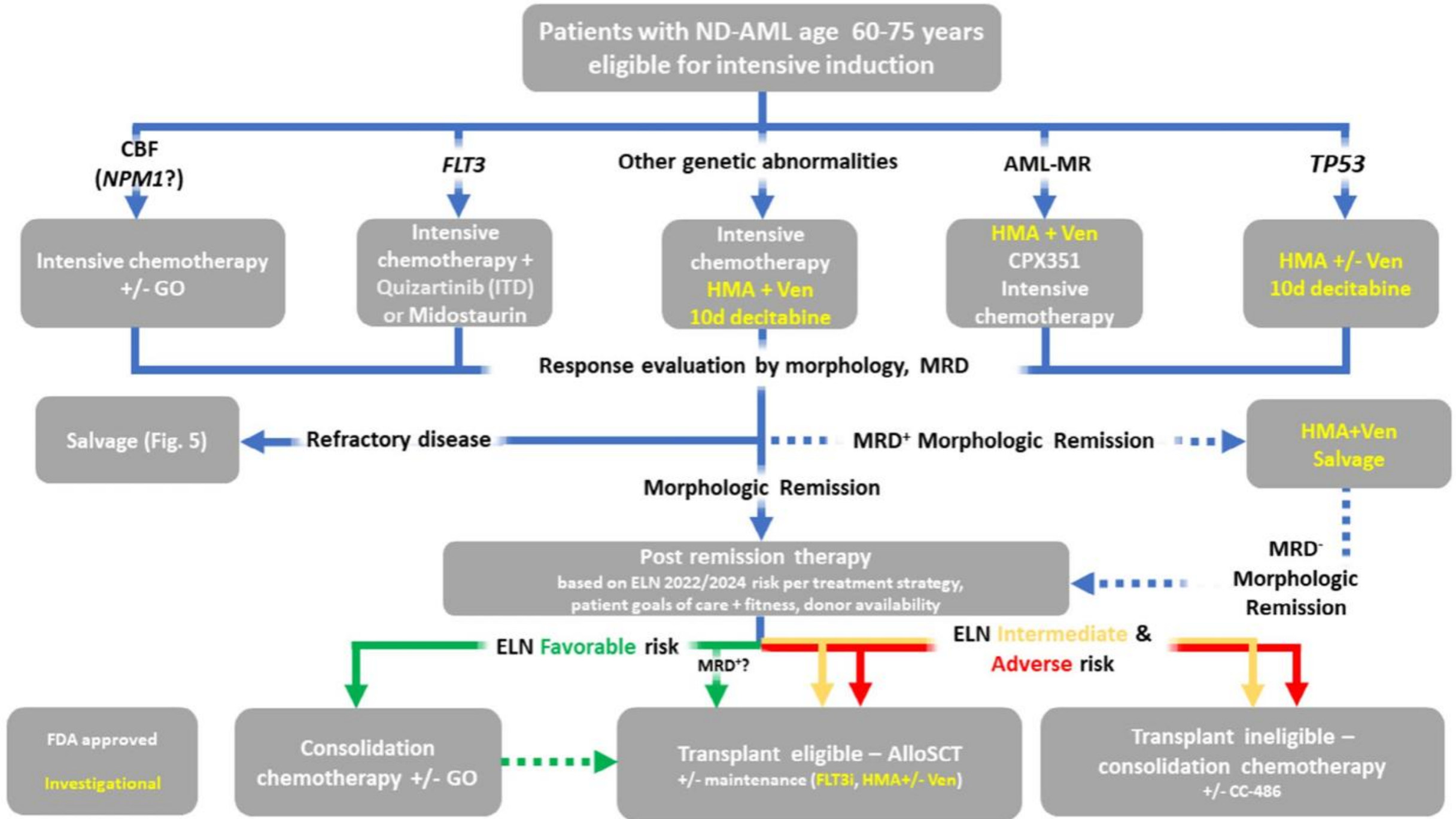


The 'era' of targeted therapy – recent FDA approvals



Differentiation and Combination therapies...





Goals of Care

Intensive
induction

Lower-intensity
approach



Fitness



Efficacy



Toxicity



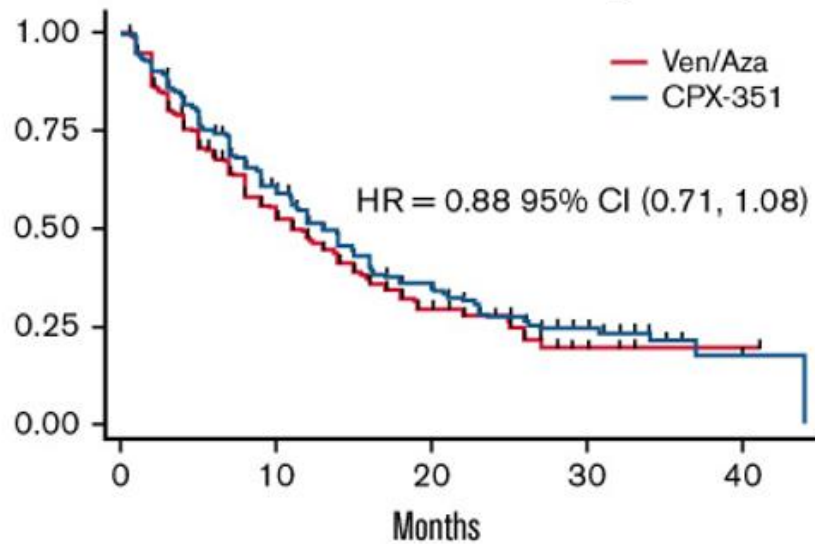
Quality of Life



CPX-351 vs. HMA-Ven

Retrospective comparative study (FLATIRON & University of Pennsylvania)

Overall survival from diagnosis



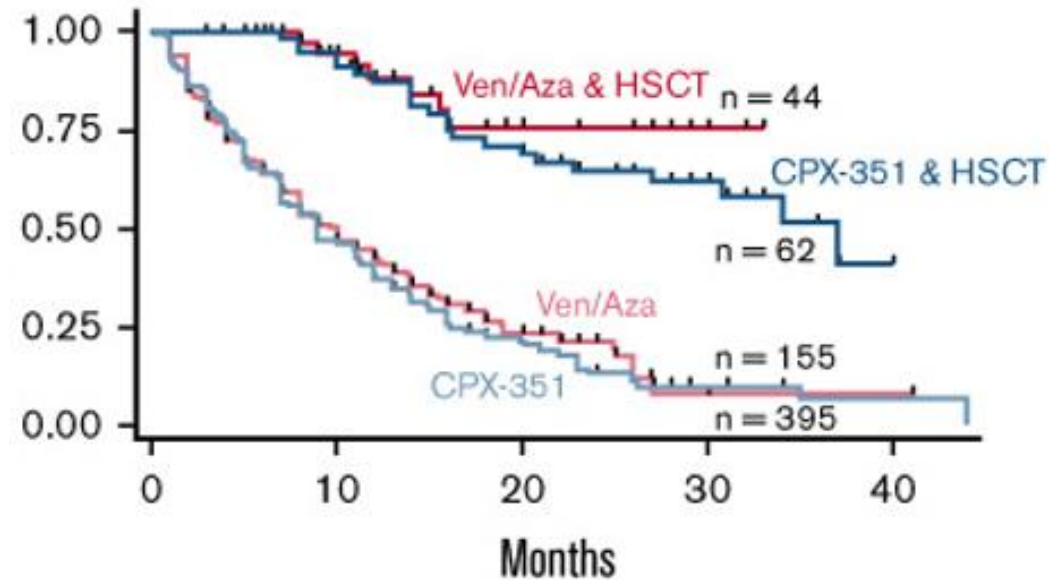
Number at risk

	0	10	20	30	40
Ven/Aza	439	168	38	5	1
CPX-351	217	111	59	23	4



Median OS	13 months	11 months
Median OS w/ allogeneic transplant	Not reached	37 months
Median OS w/o allogeneic transplant	10 months	9 months

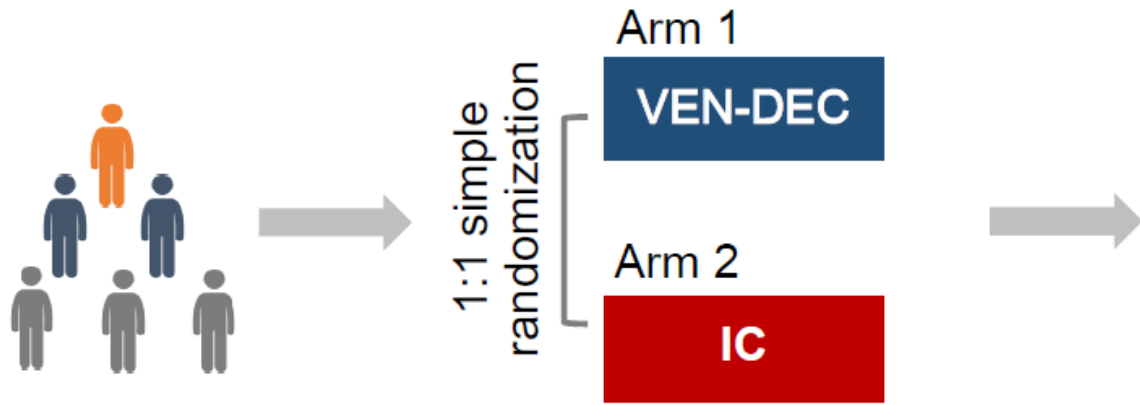
Transplant critical for long-term benefit



Flatiron & UPHS	CPX-351 n = 217	Venetoclax & azacitidine n = 439
Median cycles (range)	2 (1-5)	4 (1-28)
30 day mortality % (95% CI)	5% (2%-8%)	5% (3%-7%)
60 day mortality % (95% CI)	10% (6%-14%)	13% (10%-16%)
Diagnosis of infection ¹ % (95% CI)	51% (42%-61%)	20% (15%-25%)

IC vs. HMA-Ven

Prospective randomized phase II study



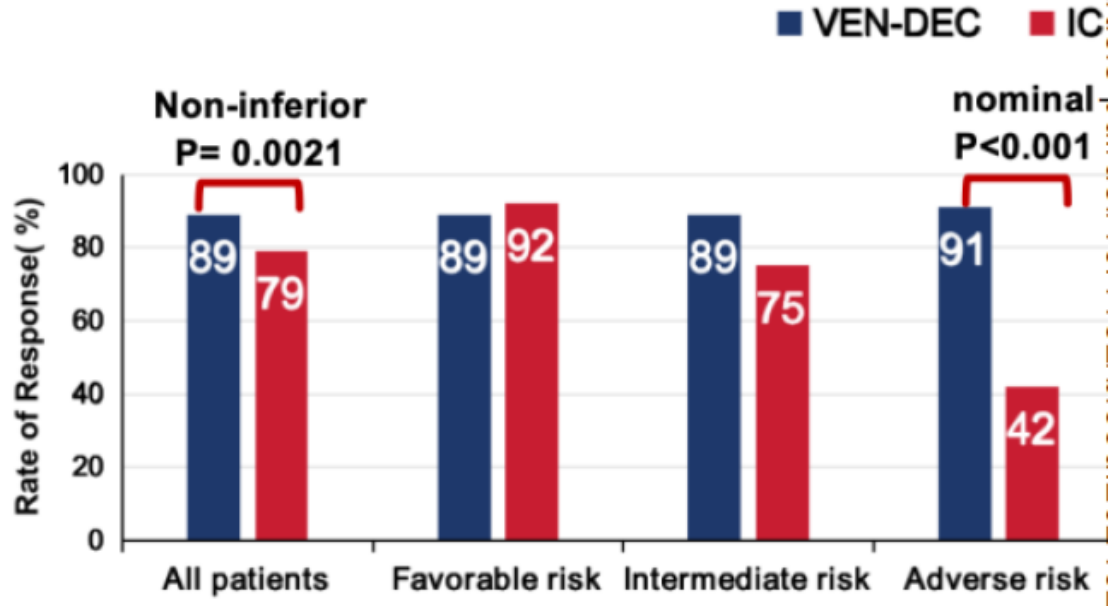
188 Patients

Induction/Re-induction

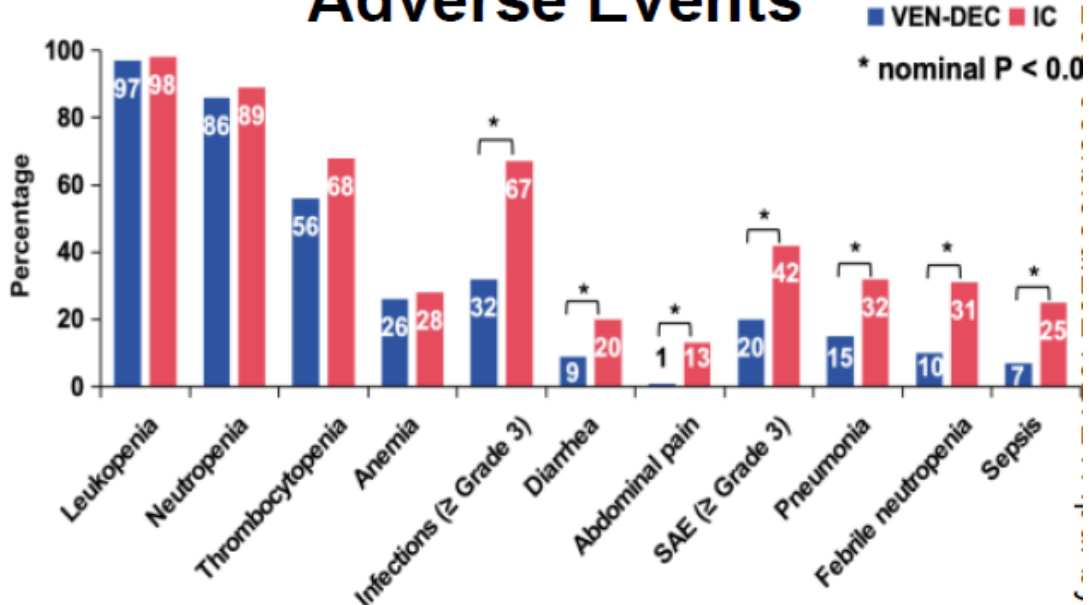
- Untreated AML
- 18-59 years old
- No extramedullary disease
- No prior Venetoclax / HMA use

- VEN-DEC:
Venetoclax (target 400 mg/d) × 28d
Decitabine (20 mg/m²/d) × 5d
- IC:
Idarubicin (12 mg/m²/d) × 3d
Cytarabine (100 mg/m²/d) × 7d

CR/CRi for Induction



Adverse Events



Jing Lu et al Blood 2025.

Decitabine is not approved in Japan and is used in combination with Azacitidine. Please refer to the Venclexta package insert for further details.

Abstract Number : abs25-8236

Abstract Title : Results from paradigm - a phase 2 randomized multi-center study comparing azacitidine and venetoclax to conventional induction chemotherapy for newly diagnosed fit adults with acute myeloid leukemia

Category: 600s - Hematologic Malignancy

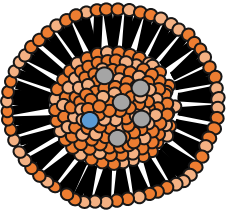
Review Category: 617. Acute Myeloid Leukemias: Commercially Available Therapies

Authors

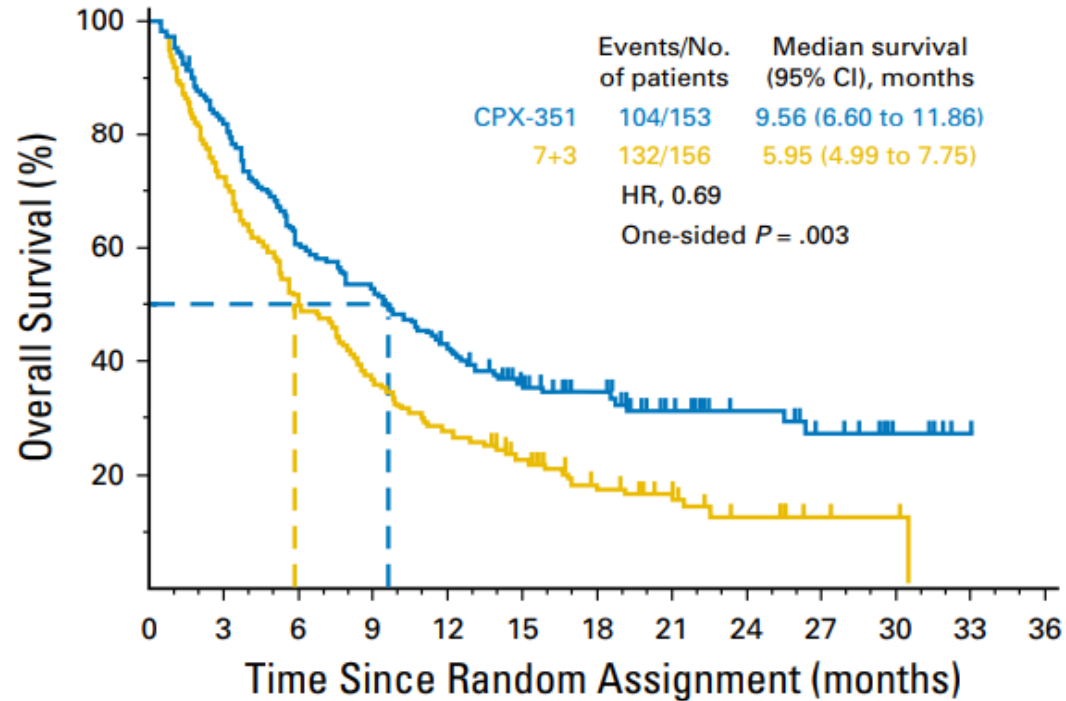
Amir Fathi¹, Alexander Perl², Geoffrey Fell³, Brian Jonas⁴, Brittany Ragon⁵, Alice Mims⁶, Uma Borate⁶, Gabriel Mannis⁷, Karen Quillen⁸, Max Stahl^{3, 9}, Paul Koller¹⁰, Andrew Artz¹⁰, Monzr M. Al Malki¹⁰, Guido Marcucci¹⁰, Mary Linton Peters⁸, Timothy Graubert¹, Peter Westervelt^{11, 12}, Philip Amrein¹, Hanno Hock¹, Andrew Brunner¹, Gabriela Hobbs¹, Rupa Narayan¹, Michelle Lee¹, Brandon Aubrey¹, Alyssa Watson¹¹, Richard Hao¹¹, Shilton Dhaver¹¹, Michael Grunwald⁵, Yi-Bin Chen¹, Andrew Matthews², Christopher Hourigan¹³, Brent Wood¹⁴, Donna Neuberg³, Areej El-Jawahri¹, Ibrahim Aldoss¹⁰

¹ Massachusetts General Hospital / Harvard Medical School, Boston, MA, United States, ² Abramson Cancer Center, University of Pennsylvania, Philadelphia, PA, United States, ³ Dana Farber Cancer Institute, Boston, MA, United States, ⁴ University of California Davis Comprehensive Cancer Center, Sacramento, CA, United States, ⁵ Levine Cancer Institute, Atrium Health, Charlotte, NC, United States, ⁶ The Ohio State University, Columbus, OH, United States, ⁷ Stanford University School of Medicine, Stanford, CA, United States, ⁸ Beth Israel Deaconess Medical Center, Boston, MA, United States, ⁹ Yale School of Medicine, New Haven, CT, United States, ¹⁰ City of Hope, Duarte, CA, United States, ¹¹ Massachusetts General Hospital, Boston, MA, United States, ¹² MaineHealth, Brunswick, ME, United States, ¹³ Fralin biomedical research institute, Virginia Tech University, Roanoke, VA, United States, ¹⁴ Children's Hospital Los Angeles, University of Southern California, Los Angeles, MA, United States



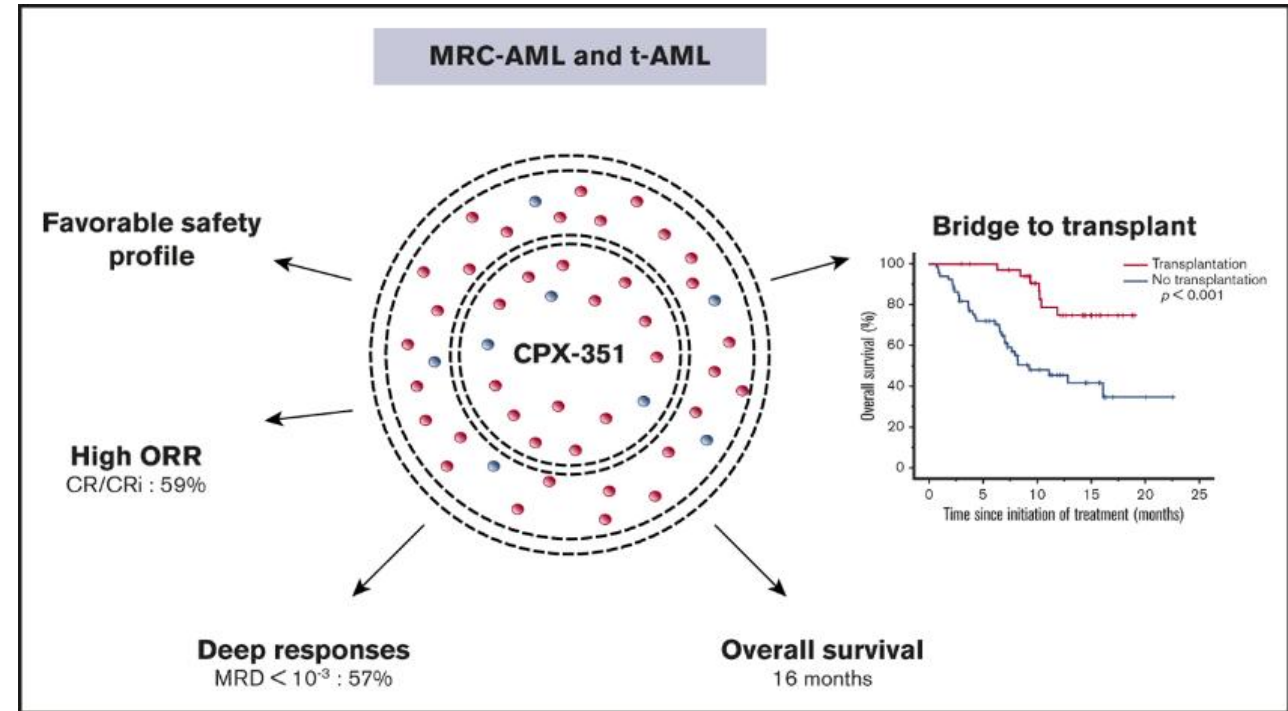


CPX-351 for Secondary AML

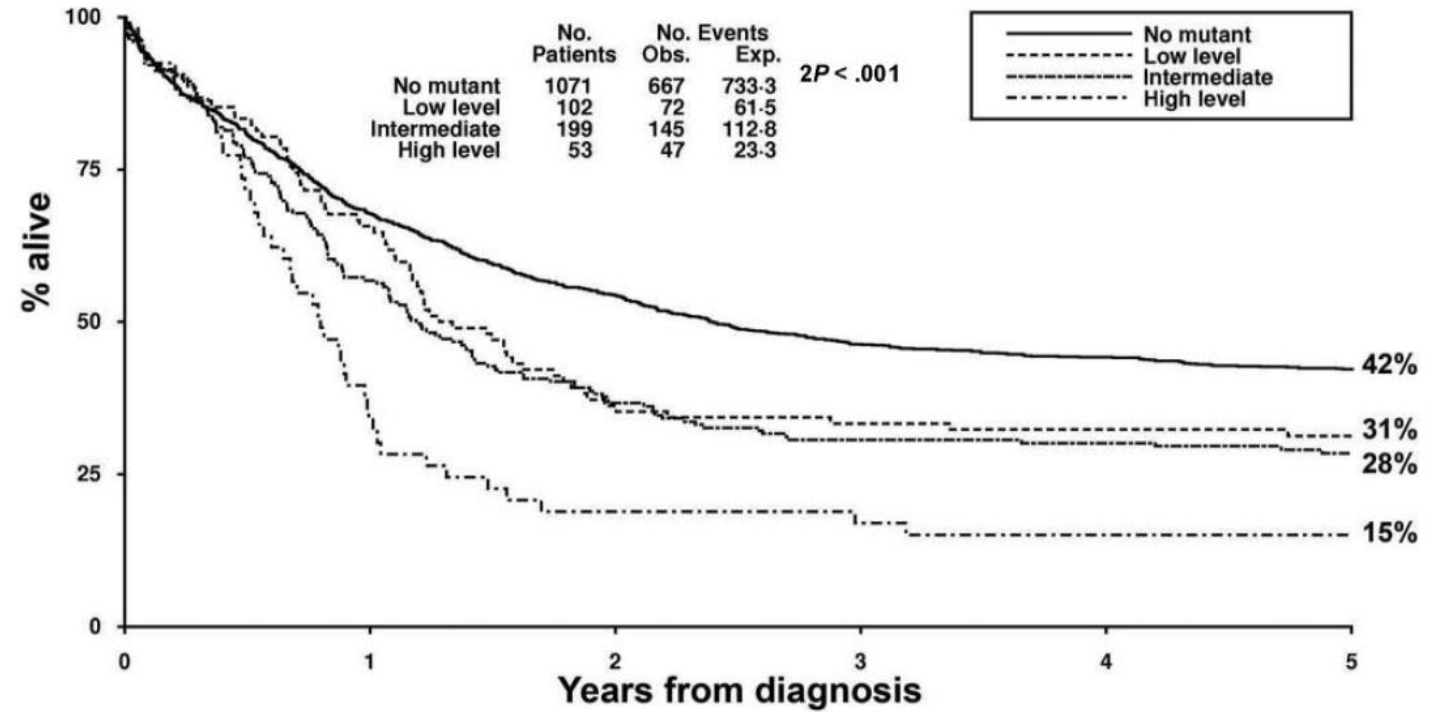
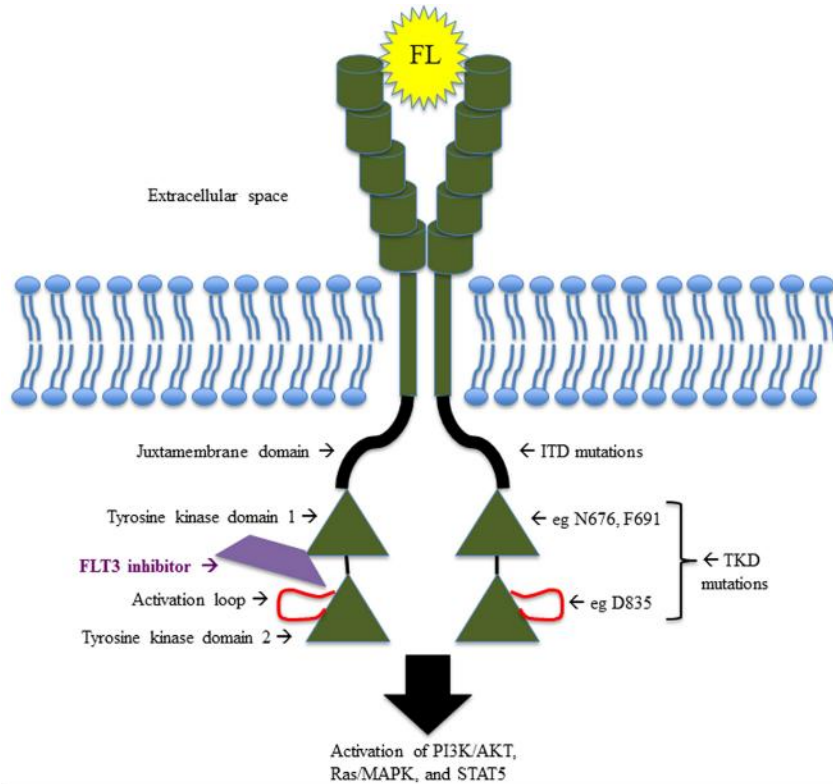


No. at risk

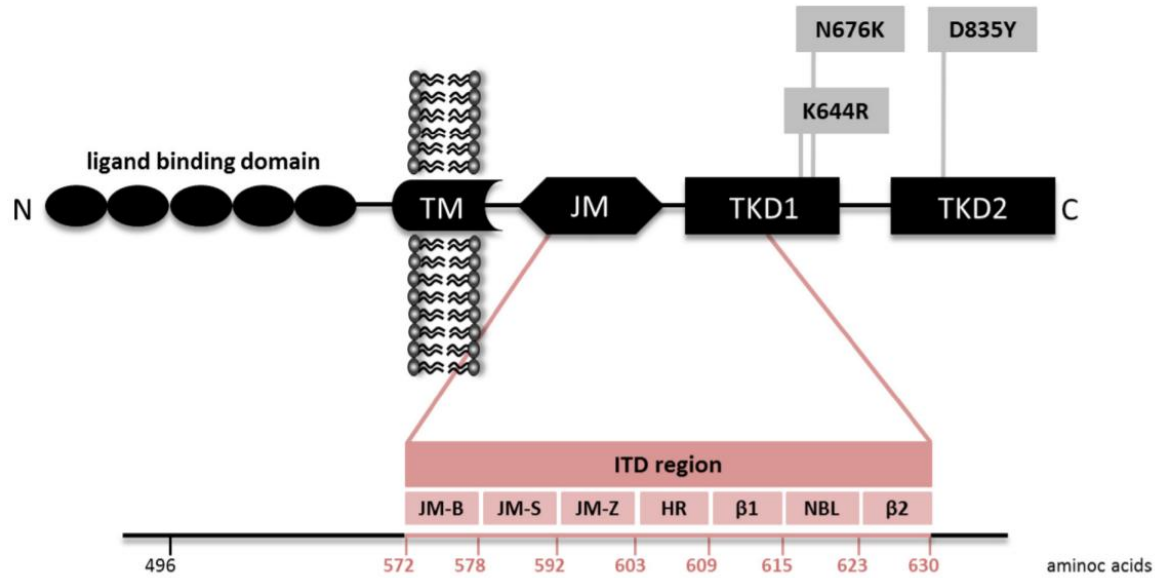
CPX-351	153	122	92	79	62	46	34	21	16	11	5	1
7+3	156	110	77	56	43	31	20	12	7	3	2	0



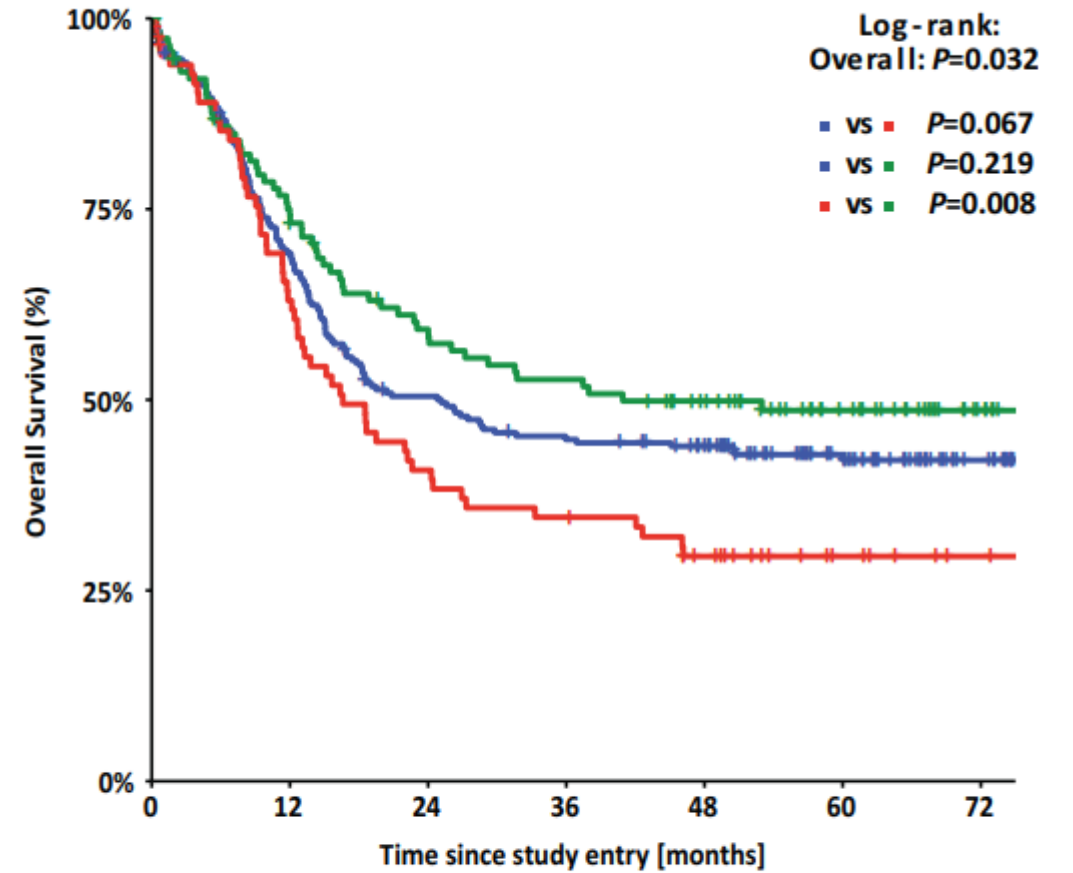
Fms-like Tyrosine Kinase 3



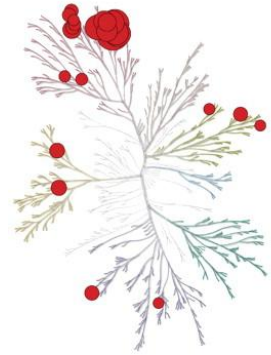
ITD insertion site



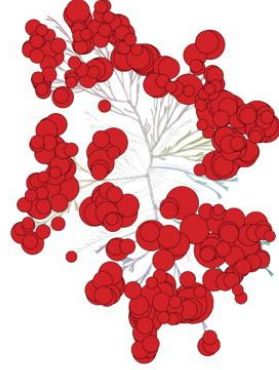
A



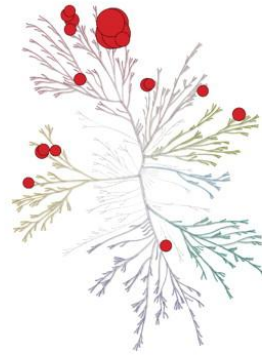
JMDsole	251	163	116	102	92	59	23
JMD/TKD1	117	82	63	56	46	28	10
TKD1sole	84	51	33	28	20	10	5



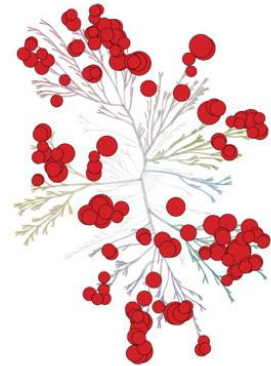
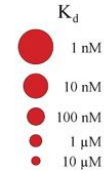
AC220



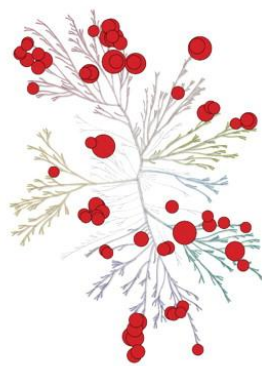
CEP-701



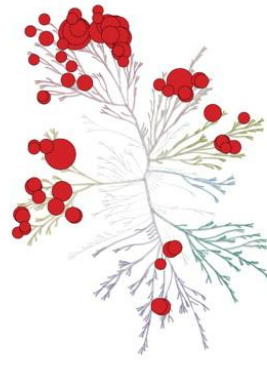
MLN-518



PKC-412



CGP-52421



Sorafenib



Sunitinib

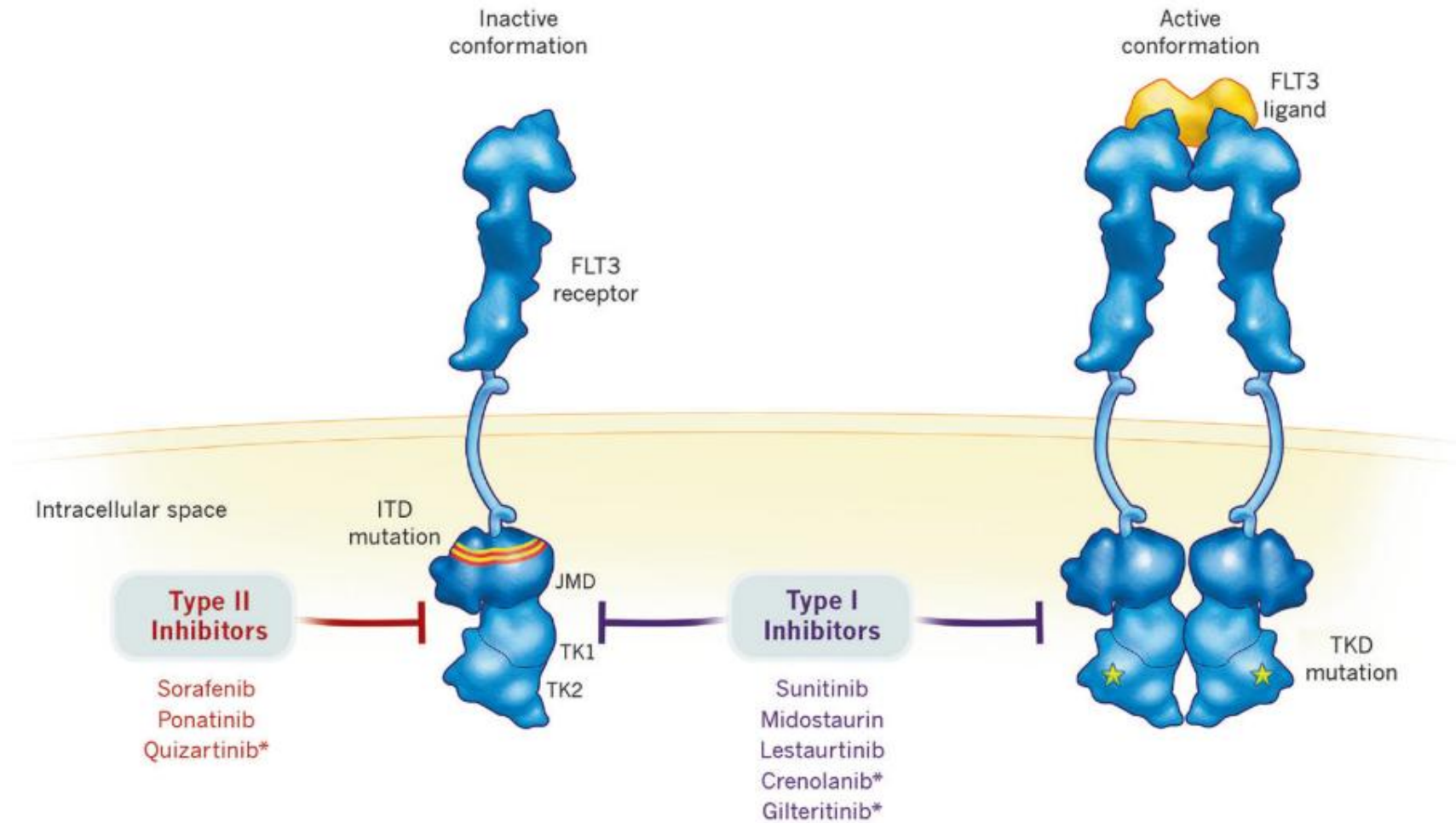
Potency (IC_{50})

Lestaurtinib (CEP-701)	700nM
Midostaurin (PKC-412)	1000nM

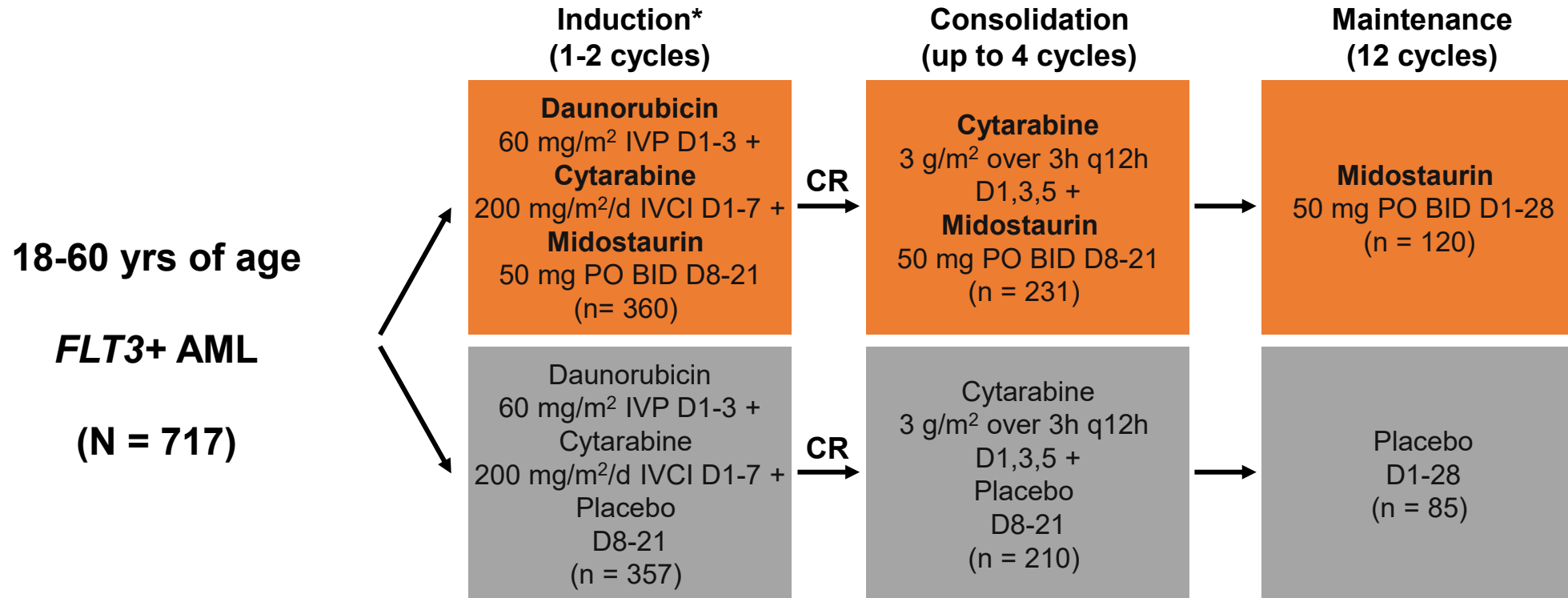
Potency (IC_{50})

Sorafenib	265nM
Quizartinib (AC220)	18nM

Summary of first- and next-generation *FLT3* inhibitors

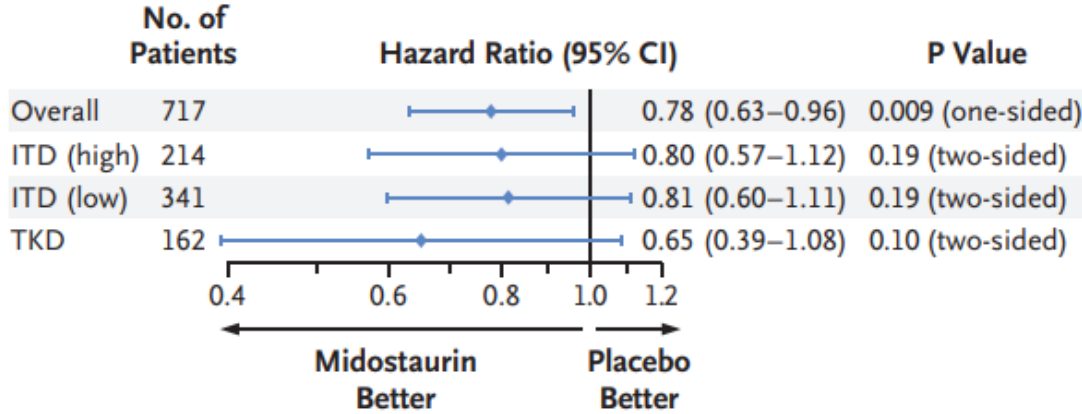
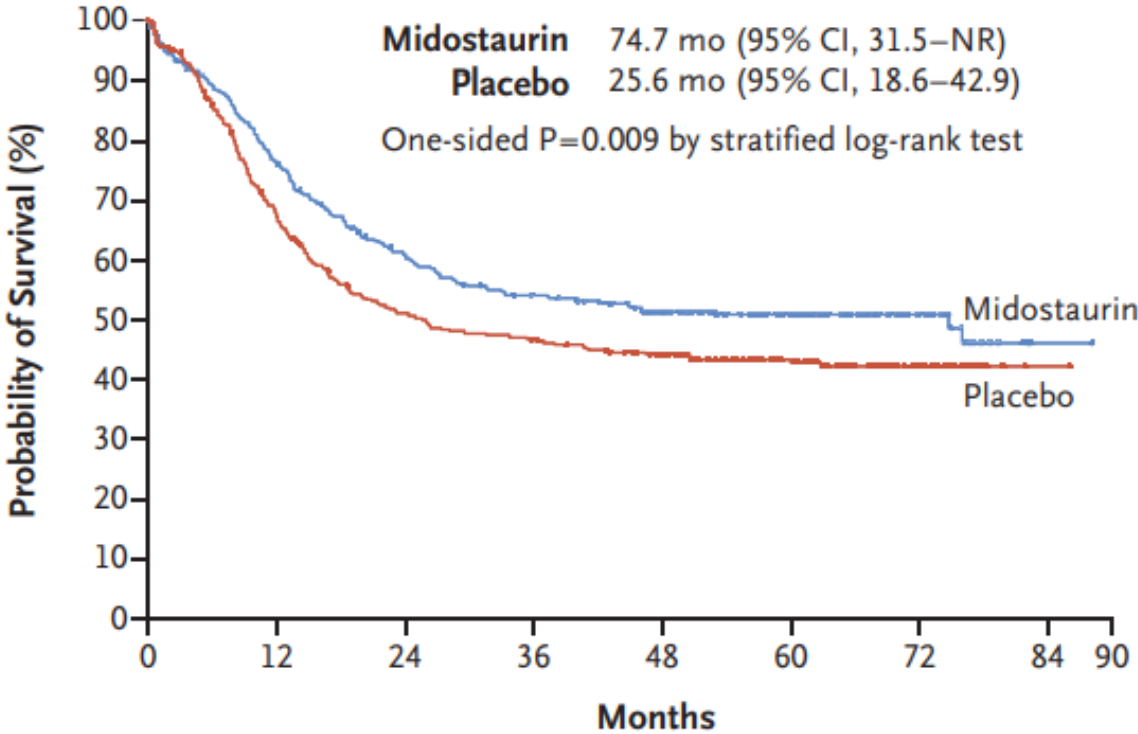


RATIFY: Midostaurin in FLT3-Positive AML



*Hydroxyurea allowed for ≤ 5 days prior to induction therapy.

RATIFY: Midostaurin in FLT3-Positive AML

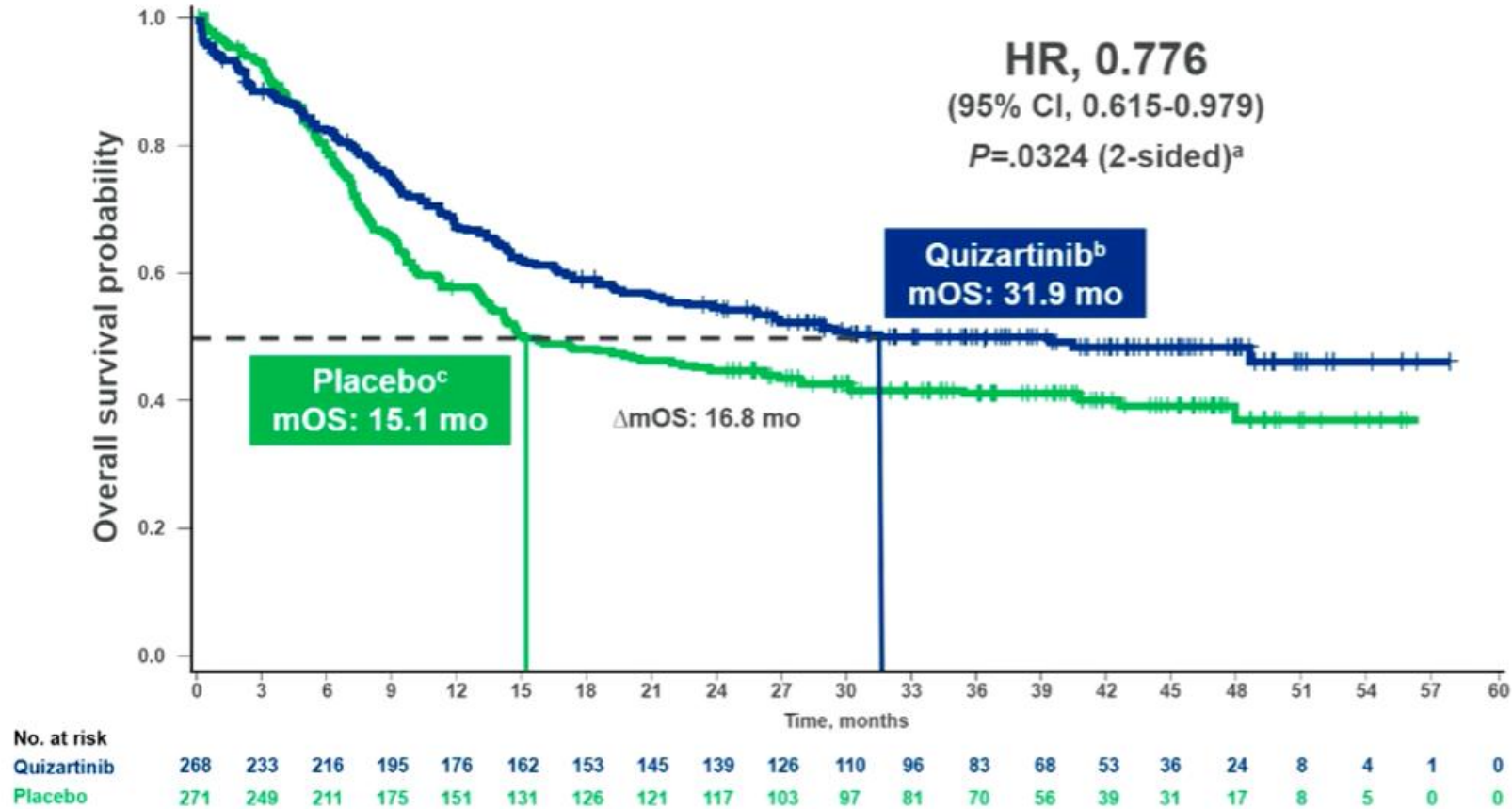


No. at Risk	0	12	24	36	48	60	72	84	90
Midostaurin	360	269	208	181	151	97	37	1	
Placebo	357	221	163	147	129	80	30	1	

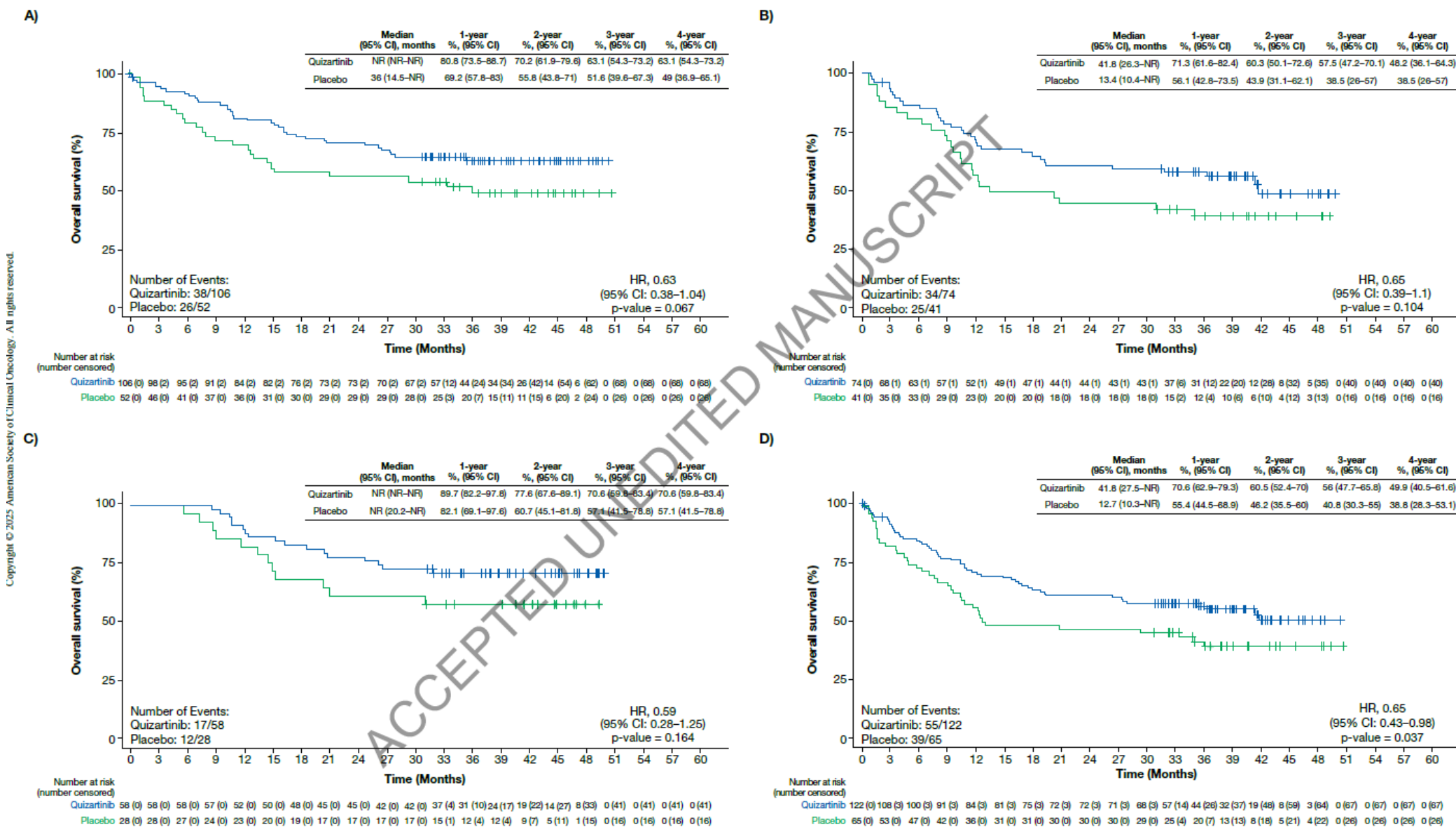
Quantum-First study

Newly Diagnosed *FLT3*-ITD+ AML; Ph3 Quizartinib + Chemotherapy

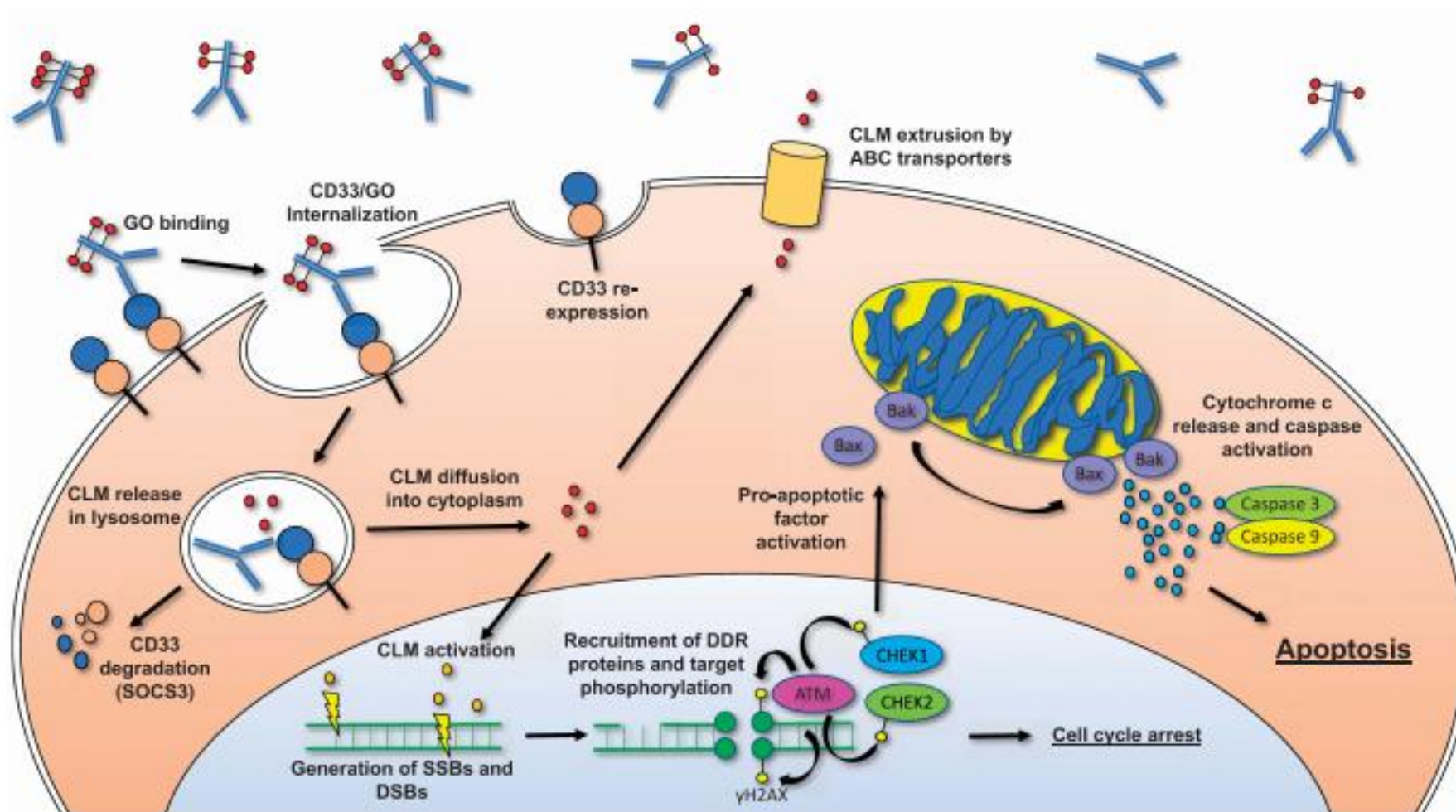
Primary Endpoint: Overall Survival



Quizartinib for FLT3—ITD negative AML – Quiwi trial



Gemtuzumab Ozogamicin (GO, Mylotarg[®])

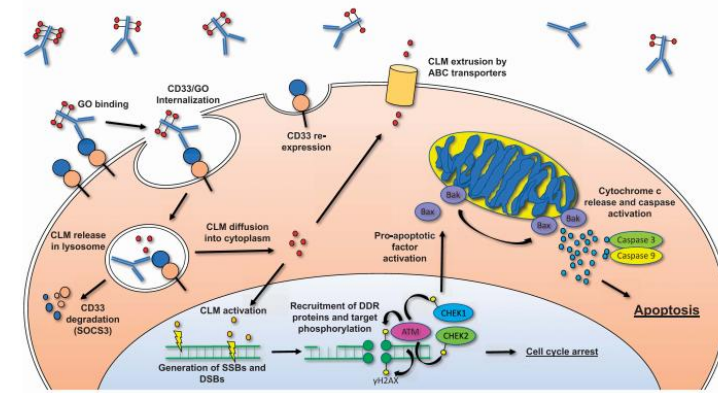


Gemtuzumab ozogamicin in acute myeloid leukemia: a remarkable saga about an active drug

Jacob M. Rowe^{1,2} and Bob Löwenberg³

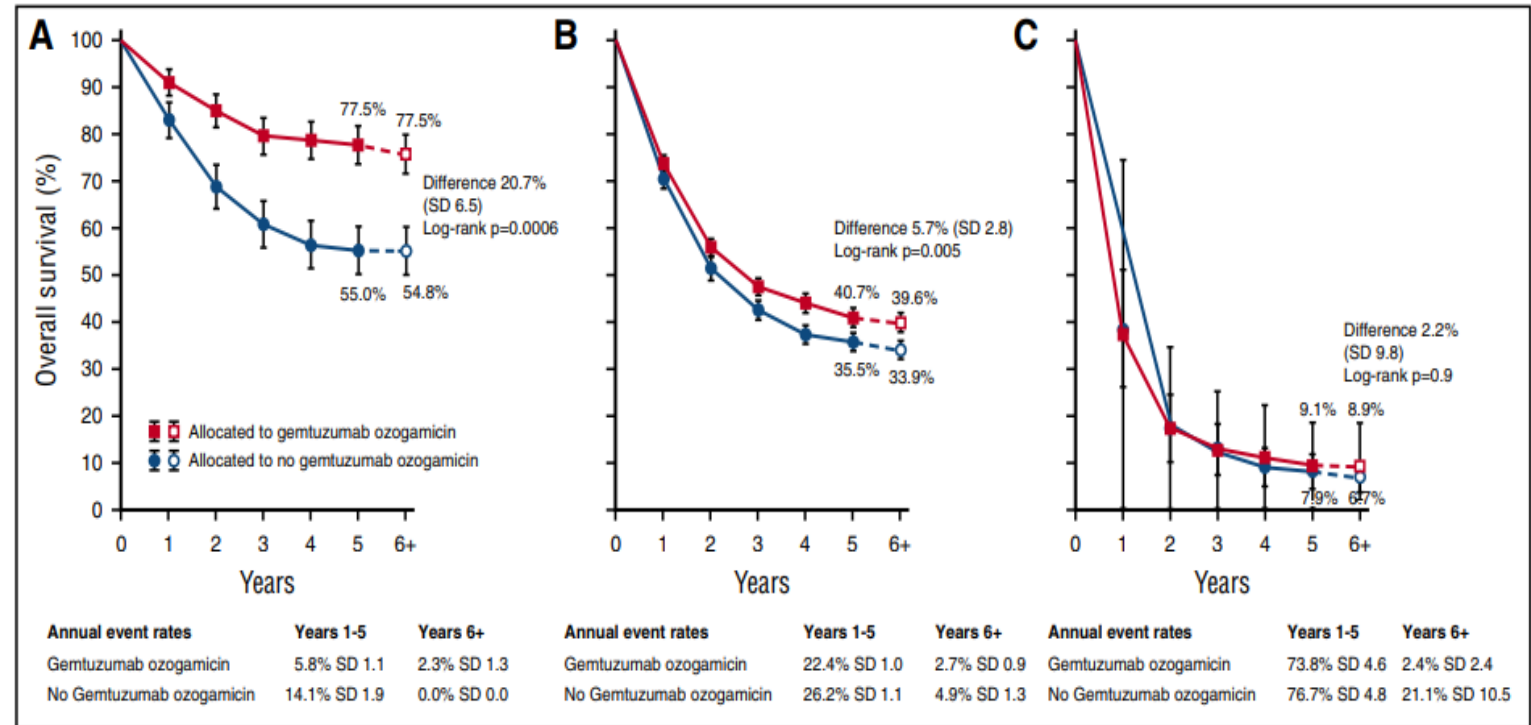
¹Shaare Zedek Medical Center, Jerusalem, Israel; ²Technion, Israel Institute of Technology, Haifa, Israel; and ³Department of Hematology, Erasmus University Medical Center, Rotterdam, The Netherlands

Improving patient outcome gemtuzumab ozogamicin

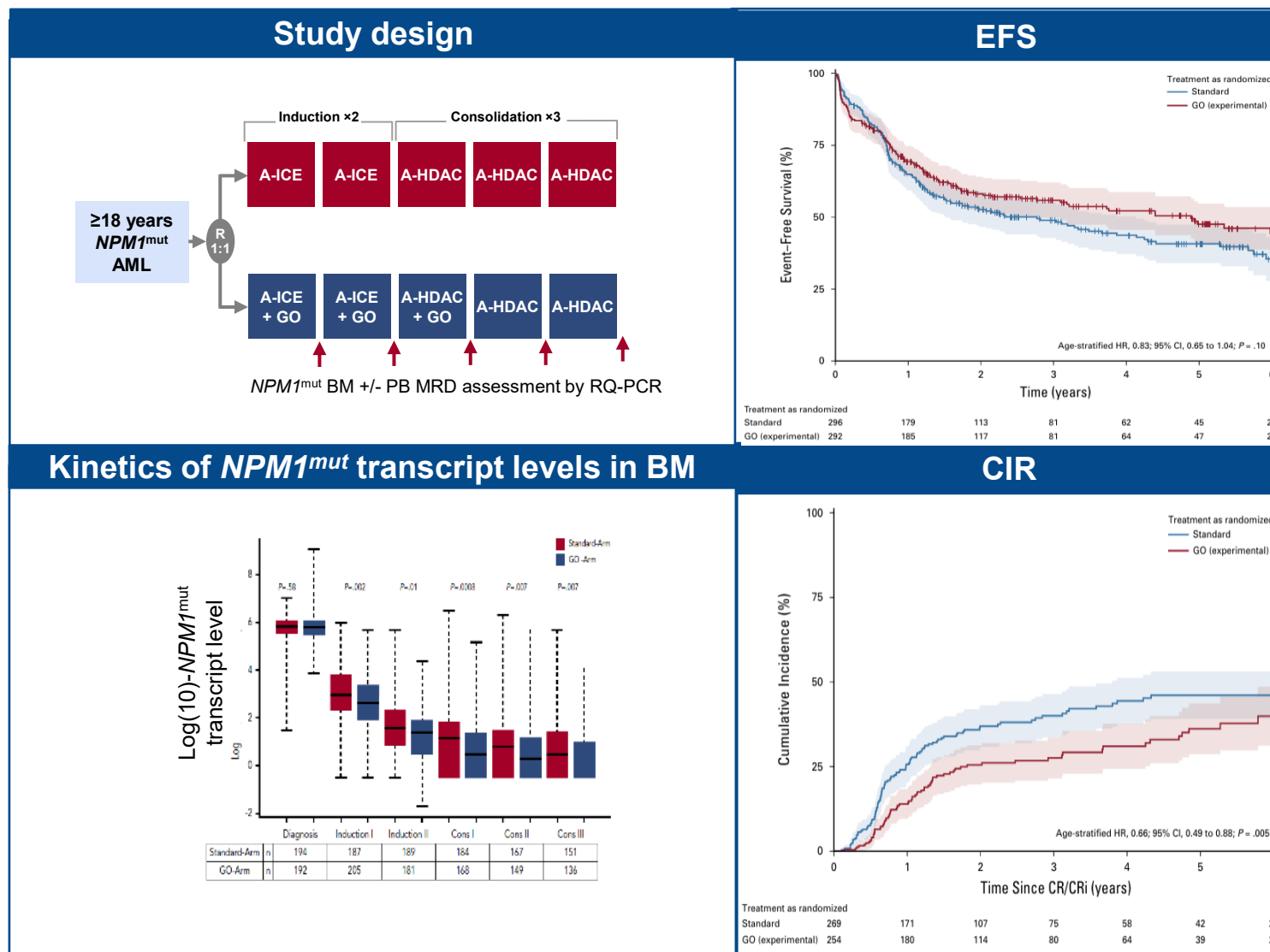


<p>MRC AML 15 1,099 patients 5-year OS of all patients: GO 43%, no GO 41%</p>
<p>SWOG-0106 595 patients 5-year OS of all patients: GO 46%, no GO 50%</p>
<p>NCRI AML 16 1,115 patients 4-year OS of all patients: GO 20%, no GO 15%</p>
<p>GOELAMS AML2006IR 238 patients 5-year OS of all patients: GO 53%, no GO 46%</p>
<p>ALFA-0701 278 patients 2-year OS of all patients: GO 53%, no GO 41%</p>

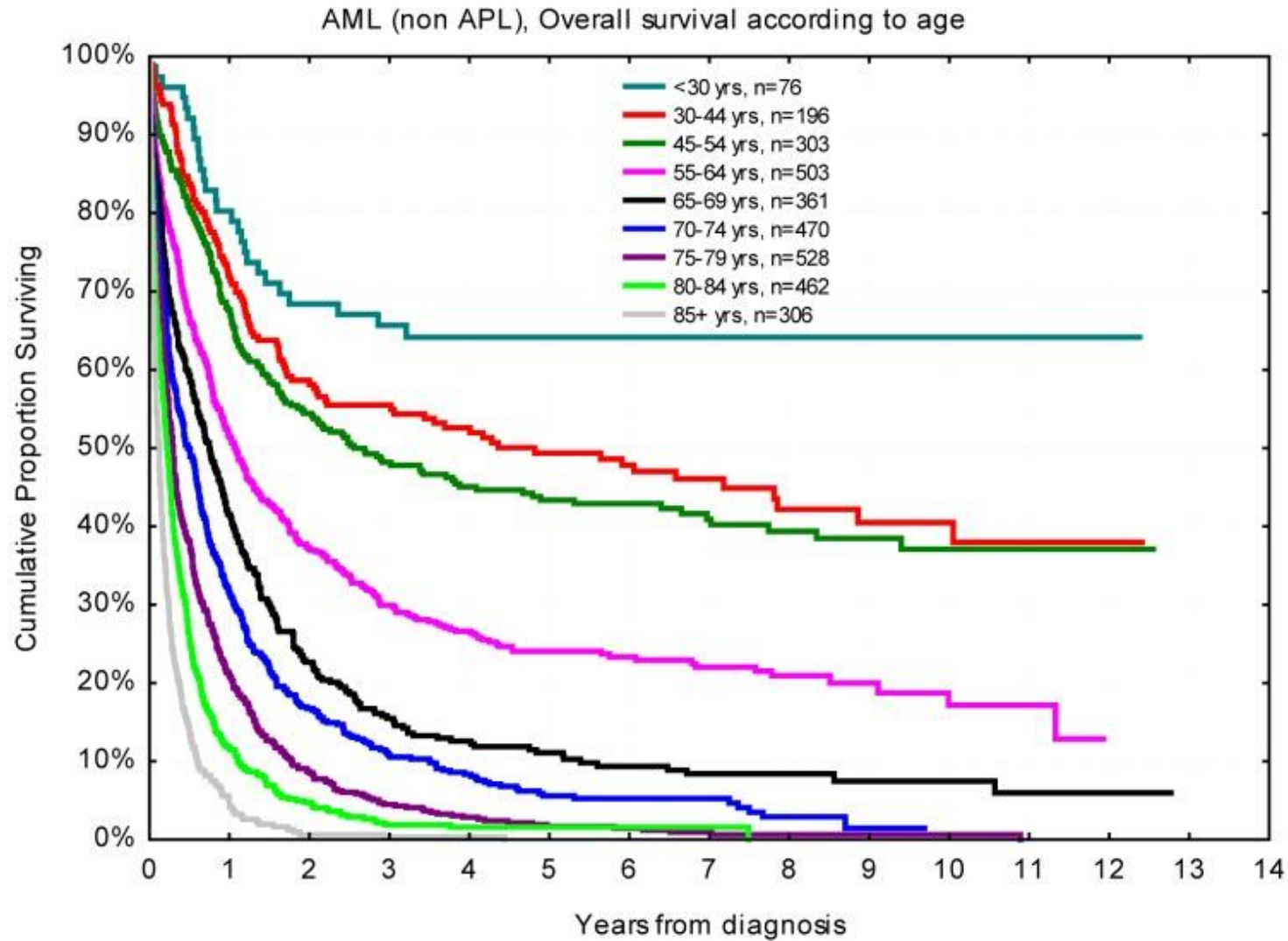
<p>Meta-analysis</p> <p>5-year OS of all patients GO 34.6% No GO 30.7%</p> <p>5-year OS favorable risk GO 77.5% No GO 55%</p> <p>5-year OS intermediate risk GO 40.7% No GO 35.5%</p> <p>5-year OS unfavorable risk GO 9.1% No GO 7.9%</p>



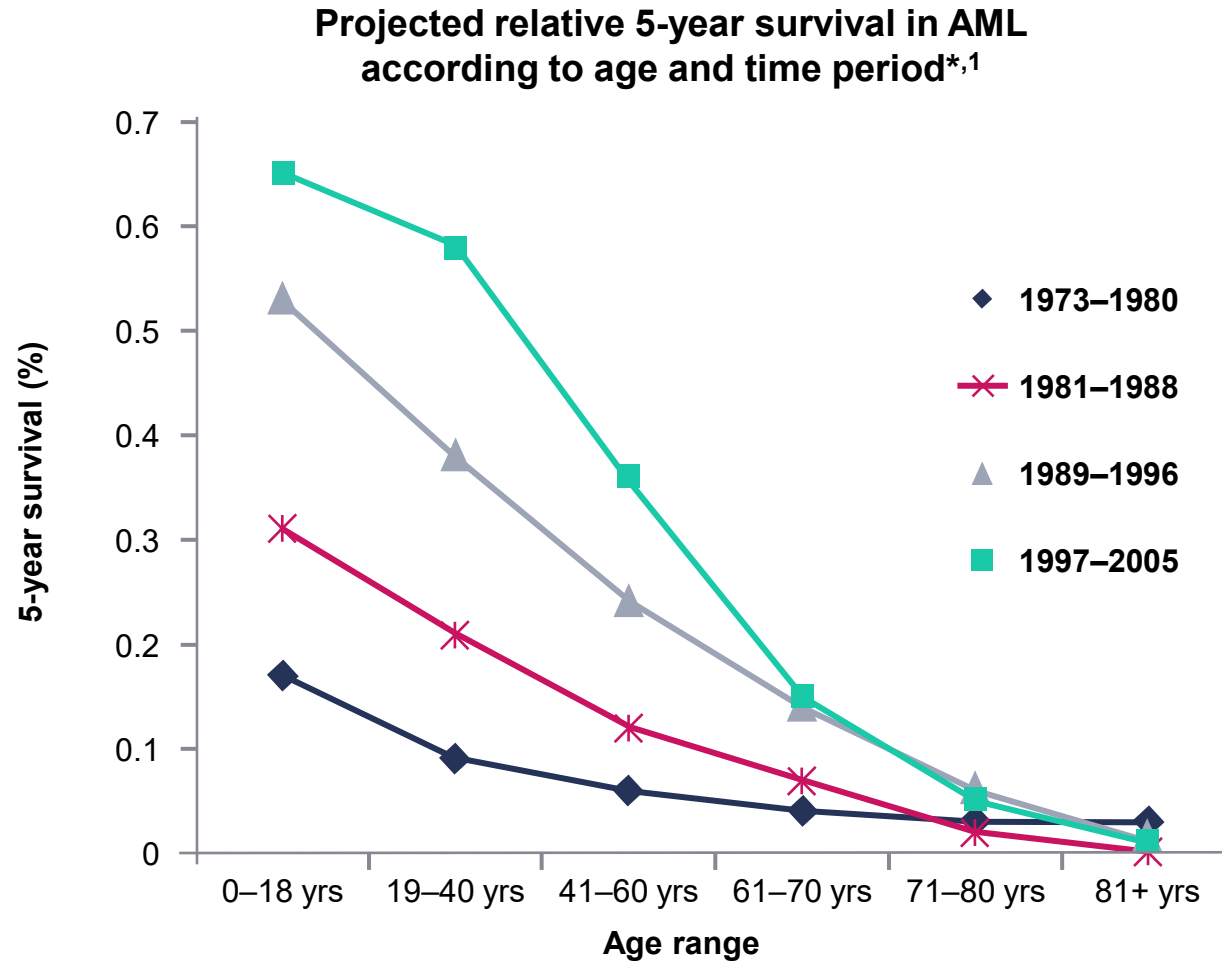
Gemtuzumab ozogamicin in *NPM1*-mutated AML: Results from prospective AMLSG 09-09 phase 3 study



Age and outcome in AML



The challenge of treating AML



* Point estimates are from Derolf, *et al.*²

1. Juliusson G, *et al. Blood* 2012; **119**:3890-3899;
2. Derolf AR, *et al. Blood* 2009; **113**:3666-3672.

Treatment decision-making remains complex

Lack of consensus on the definition of eligibility for intensive chemotherapy

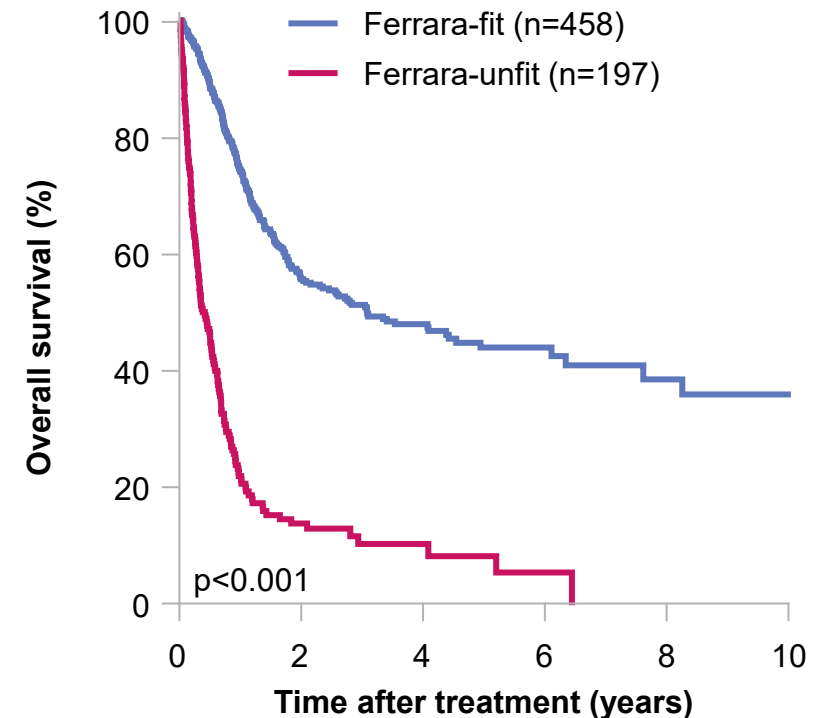
Ferrara criteria¹

Consensus criteria of SIE-SIES-GITMO for defining fit/unfit for IC^{1,3}

- Advanced age (>75 years)
- Severe organ comorbidities*
- Active infection resistant to anti-infective therapy
- Cognitive impairment
- Low performance score (ECOG PS ≥ 3)

* And any other comorbidity that the physician judges to be incompatible with IC.
GITMO, Italian Group for Bone Marrow Transplantation;
IC, intensive chemotherapy; ND, newly diagnosed; SIE, Italian Society of Hematology;
SIES, Italian Society of Experimental Hematology; TRM, treatment-related mortality.

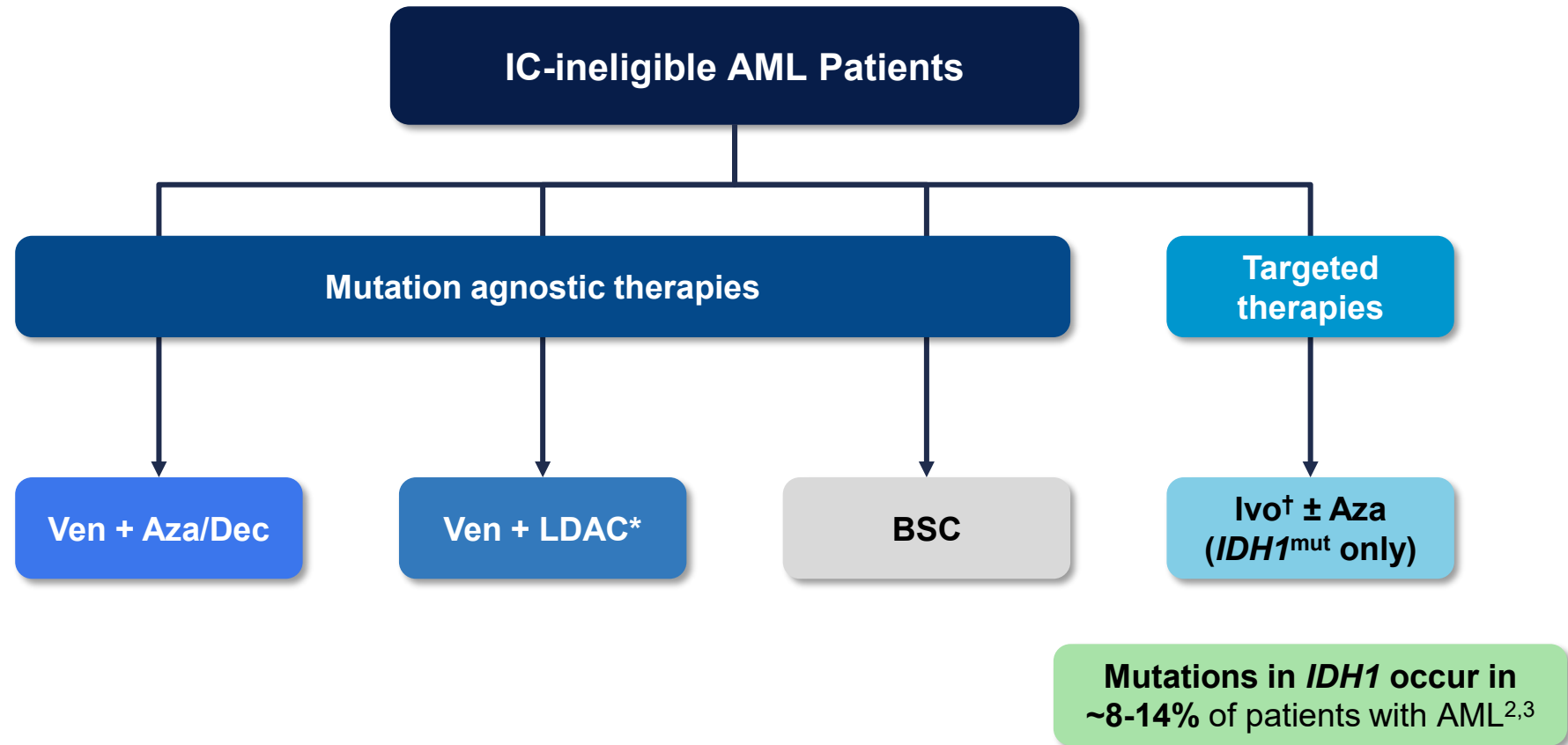
OS by Ferrara category



1. Ferrara F, et al. *Leukemia* 2013; **27**:997–999;
2. Walter RB, et al. *J Clin Oncol* 2011; **29**:4417–4423;
3. Chen EC & Garcia JS. *Hematology Am Soc Hematol Educ Program* 2020; **2020**:41–50.

Selected treatment options according to ELN:

IC-ineligible patients with AML



*LDAC + venetoclax is not approved by the EMA for use in patients with AML. †Ivosidenib monotherapy is currently not approved by the EMA.

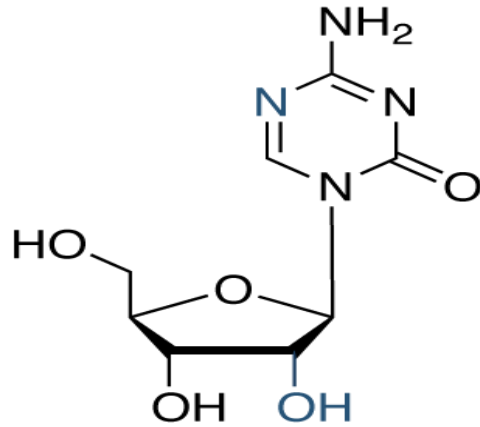
AML, acute myeloid leukemia; Aza, azacitidine; BSC, best supportive care; Dec, decitabine; ELN, European LeukemiaNet;

IC, intensive chemotherapy; Ivo, ivosidenib; LDAC, low-dose cytarabine; Ven, venetoclax.

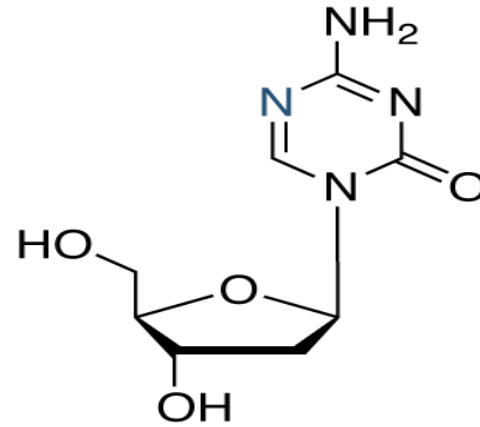
1. Döhner H et al. *Blood*. 2022;140(12):1345-1377. 2. Medeiros BC et al. *Leukemia*. 2017;31:272-281. 3. Issa GC et al. *Blood Cancer J*. 2021;11:107.

Hypomethylating Agents

Inhibitors of DNA methyl transferases:



5-azacytidine
(azacitidine)



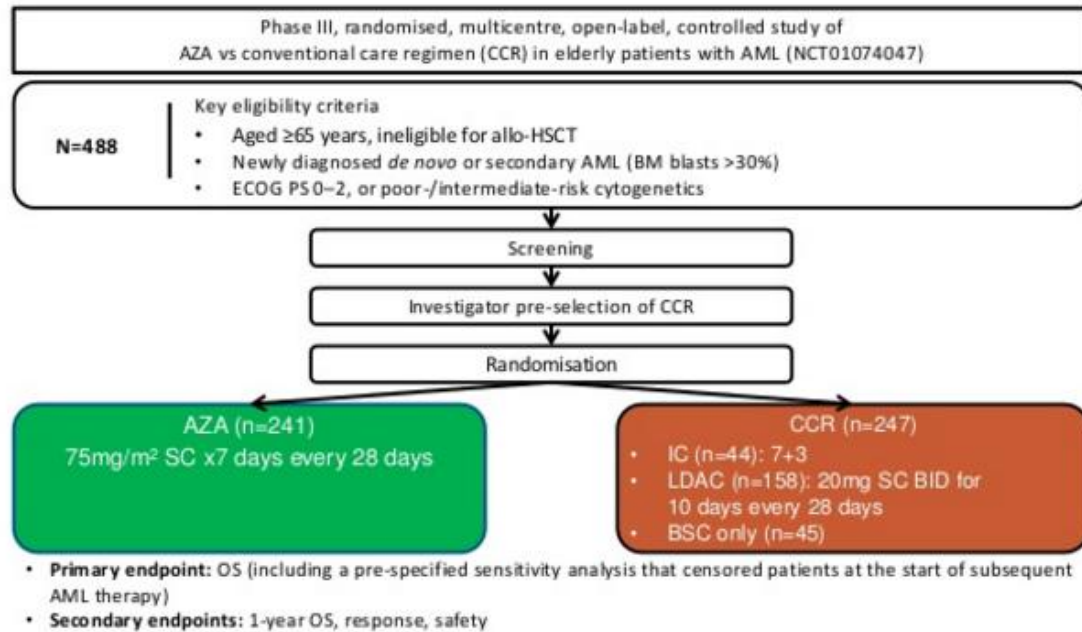
5-aza-2'-deoxycytidine
(decitabine)

Both incorporate into DNA and cause hypomethylation (DEC > AZA)

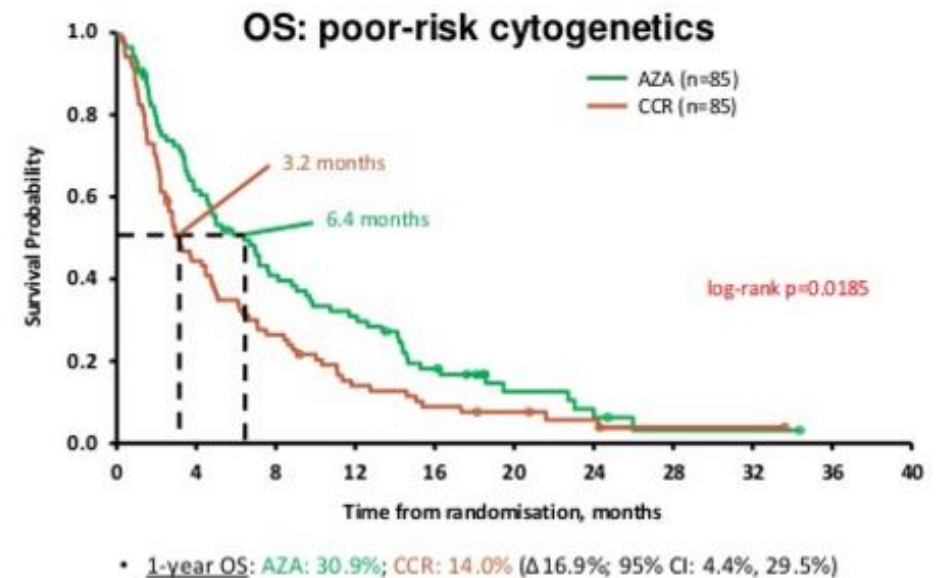
AZA preferentially causes DNA damage and induces apoptosis

Monotherapy with AZA

Study design of the phase III randomised AZA-AML-001 trial



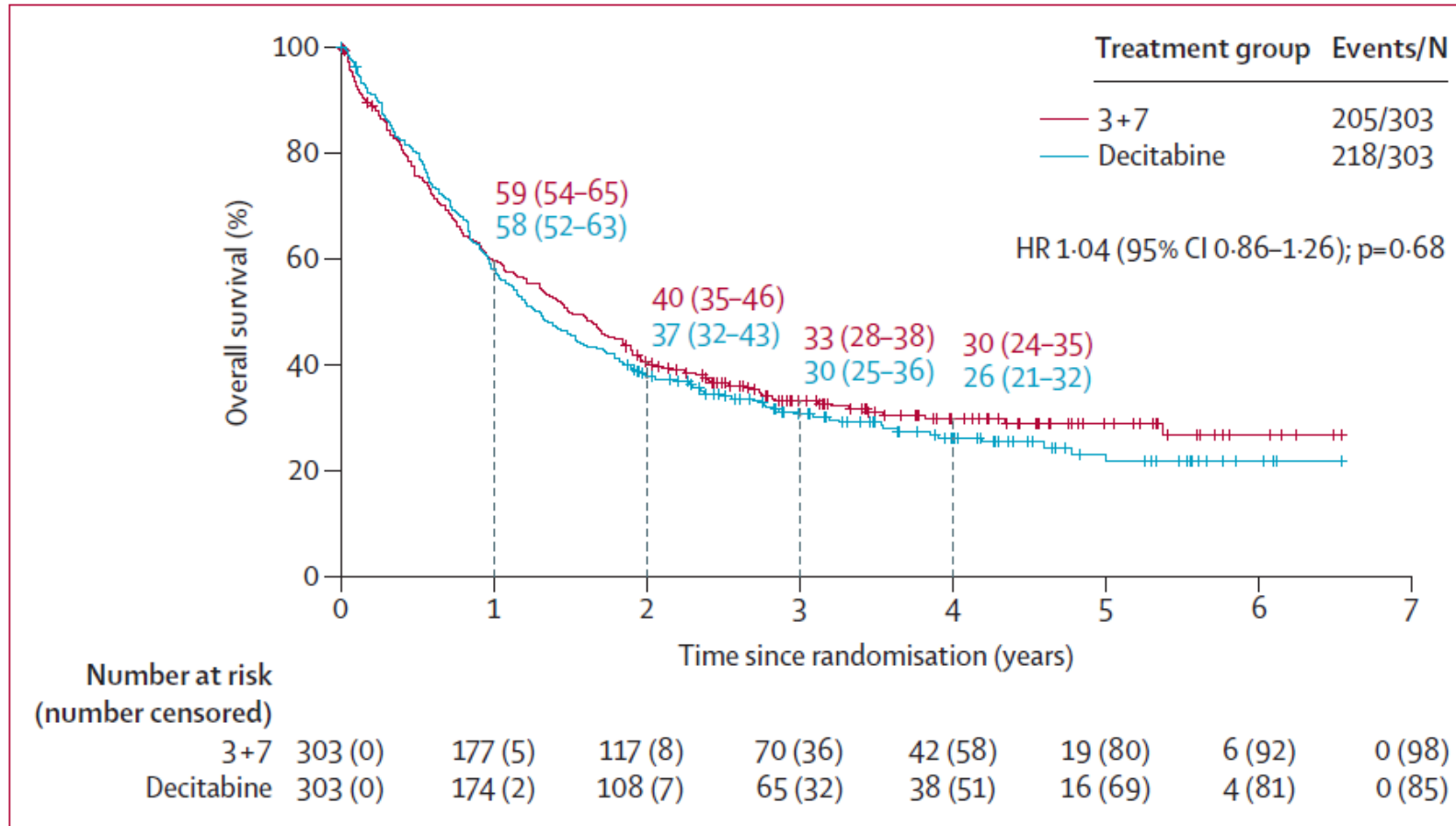
Subanalysis of AZA-AML-001 trial Outcomes according to cytogenetic risk



Patients with poor-risk cytogenetics treated with AZA had significantly longer median OS than those treated with CCR

Decitabine vs. '7+3'

Open-label, randomized, controlled, phase 3 trial (EORTC, GIMEMA, German MDS Study Group)



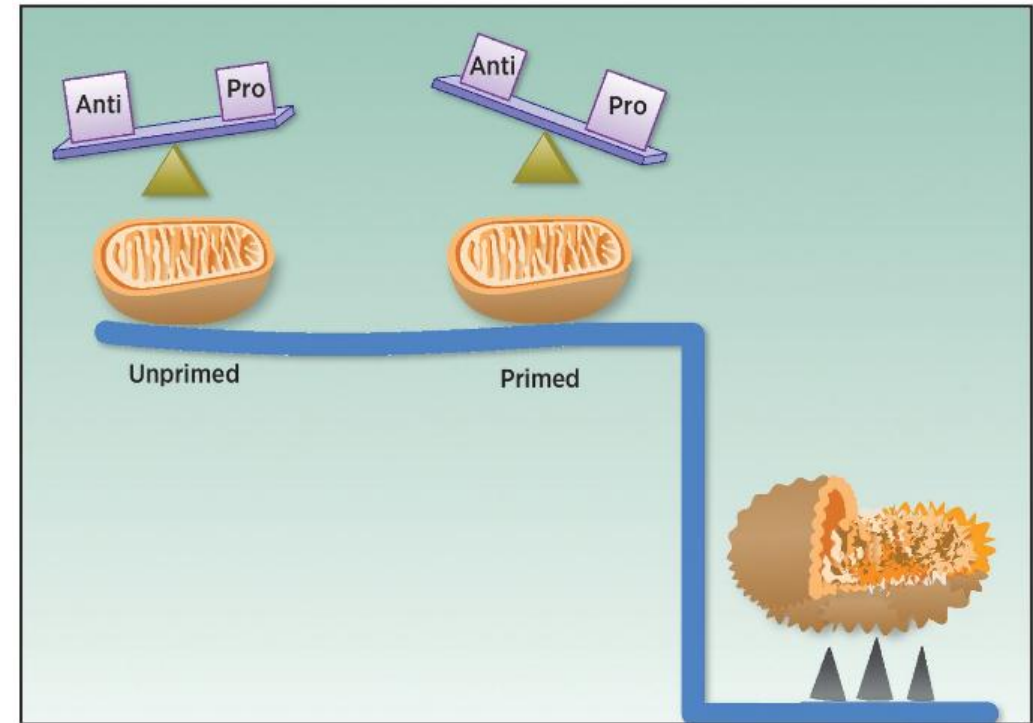
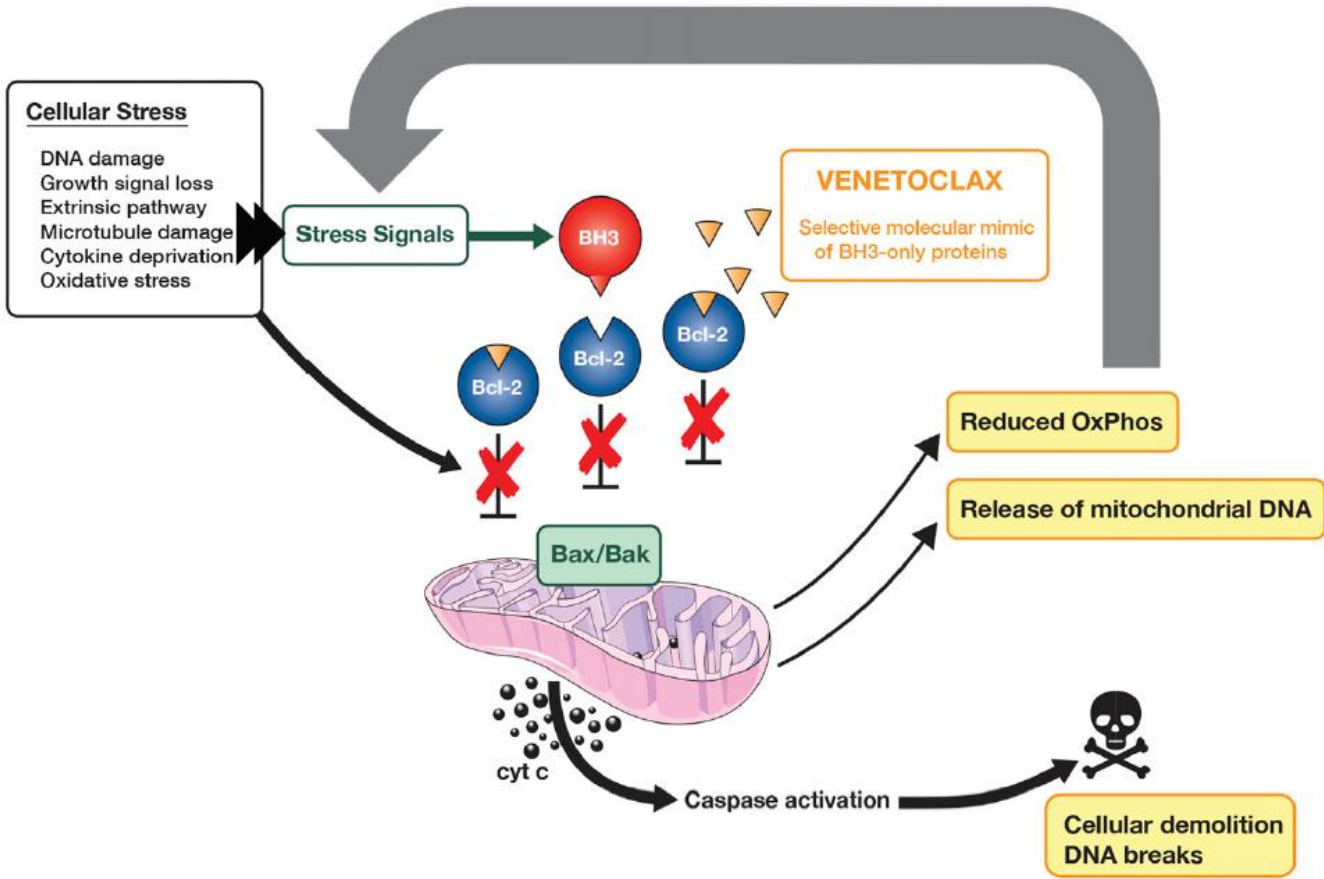
HSCT

- Decitabine, 122 patients (40%)
- 7+3, 118 patients (39%)

Median time to HSCT

- Decitabine, 4.2 mos. (IQR 3.0-5.3)
- 7+3, 3.5 mos. (2.9-5.0)

606 pts, >60 years, Newly-Diagnosed AML



VIALE A

AZA-PBO vs. AZA-VEN

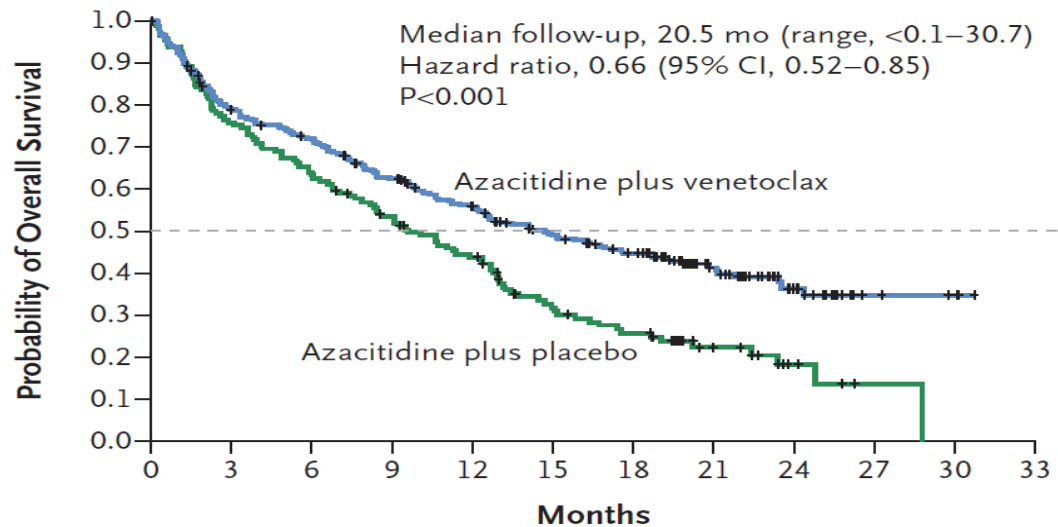
ARM A

Venetoclax: 400 mg PO, daily, days 1–28
+ Azacitidine 75 mg/m² SC/IV days 1–7

2:1 Randomization
N = 433*

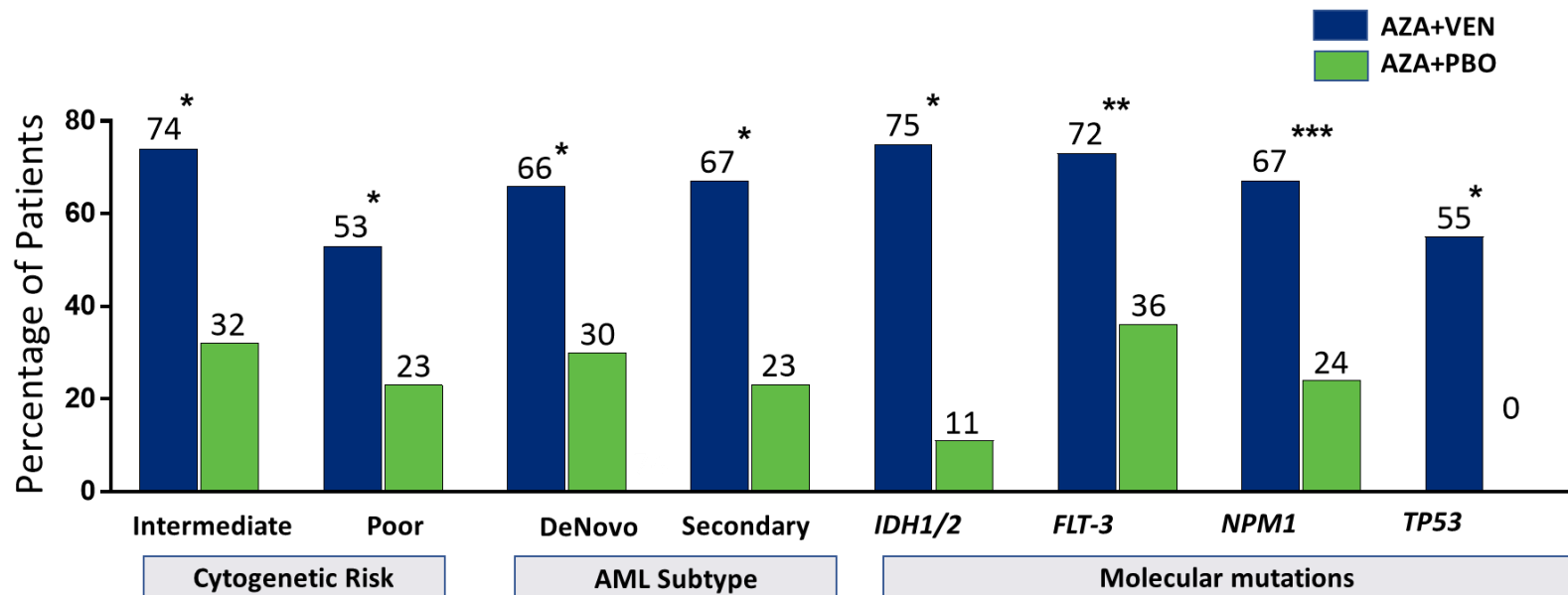
Placebo daily, days 1–28
+ Azacitidine 75 mg/m² SC/IV days 1–7

ARM B



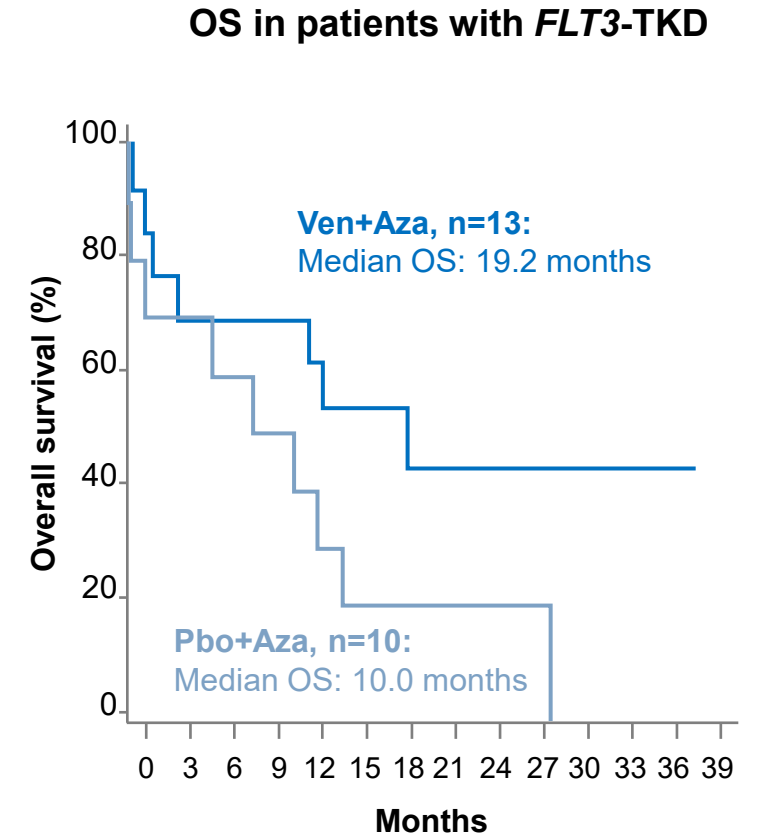
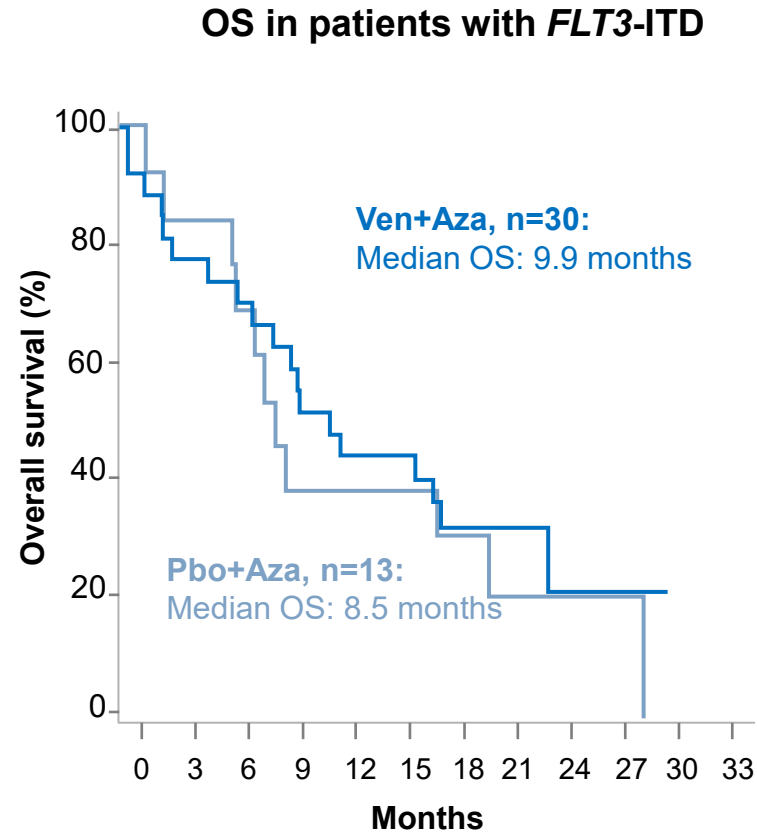
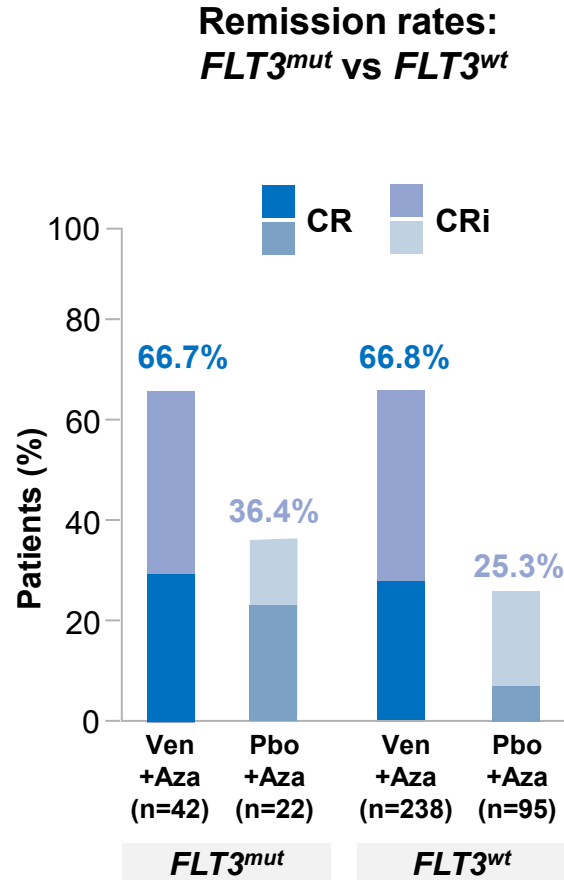
No. at Risk

Azacitidine plus venetoclax	286	219	198	168	143	117	101	54	23	5	3	0
Azacitidine plus placebo	145	109	92	74	59	38	30	14	5	1	0	0



Impact of *FLT3* mutation on outcomes with doublet regimen Ven + Aza

Pooled analysis of patients with ND AML in VIALE-A and phase 1b study (N=498)

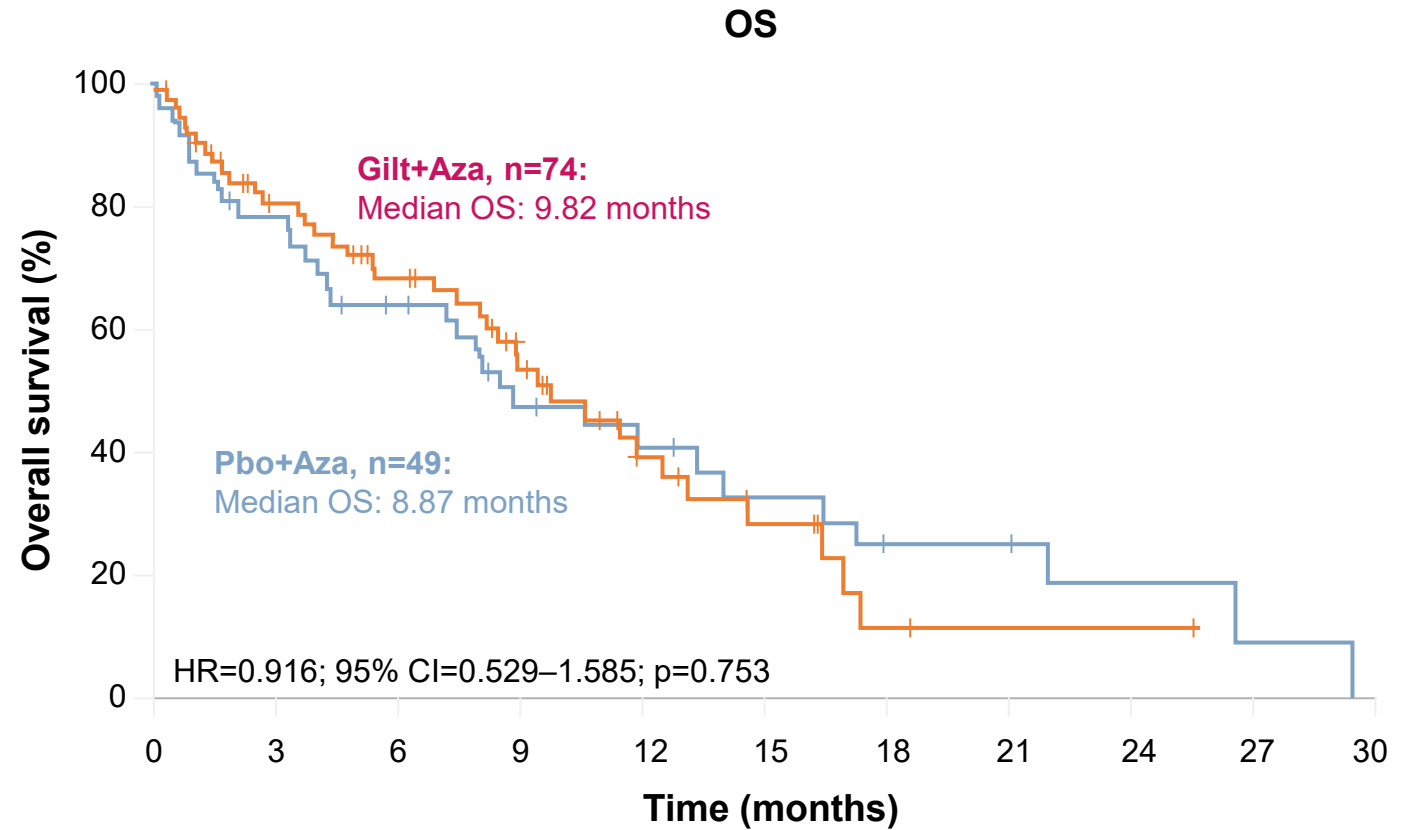
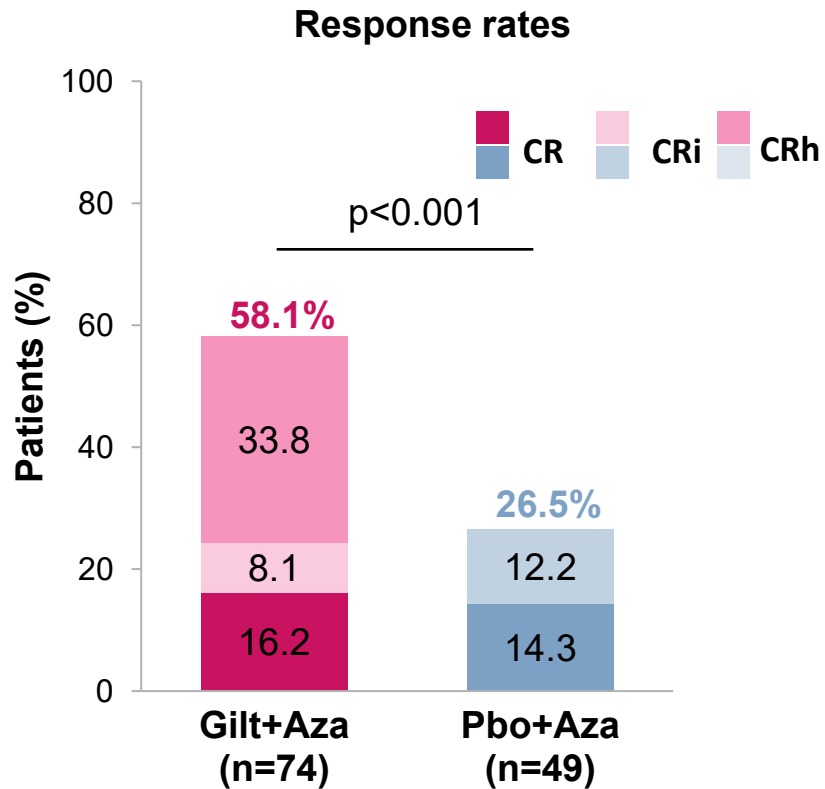


The safety and tolerability of the Ven+Aza in patients with *FLT3*^{mut} were similar to patients with *FLT3*^{wt} mutation status.

AZA, azacitidine; ITD, internal tandem duplication; mut, mutated; ND, newly diagnosed; Pbo, placebo; TKD, tyrosine kinase domain; Ven, venetoclax; wt, wild type.

CR/CRi rates and OS with Gilt + Aza vs Aza in *FLT3^{mut}* AML

LACEWING: Randomized, phase 3 trial in patients ineligible for intensive chemotherapy

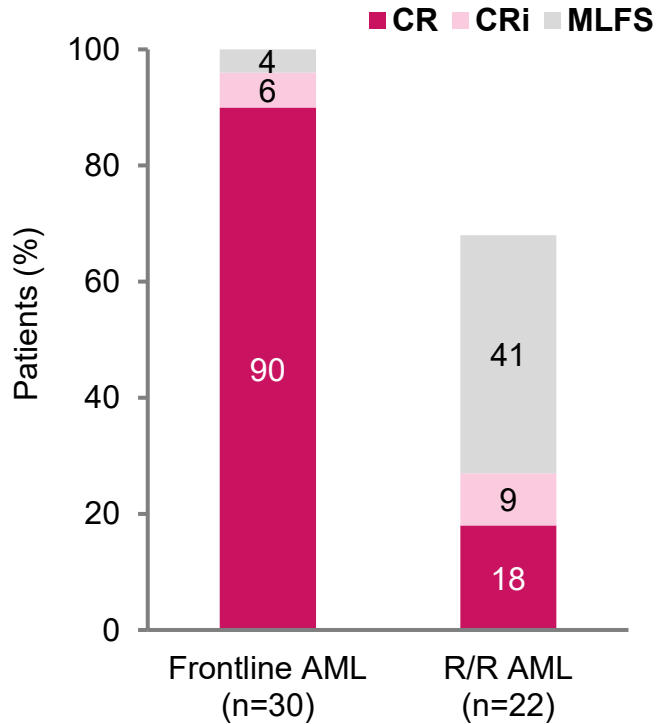


AEs rates (any grade and Gr≥3) were similar in both arms
No new safety signals were reported

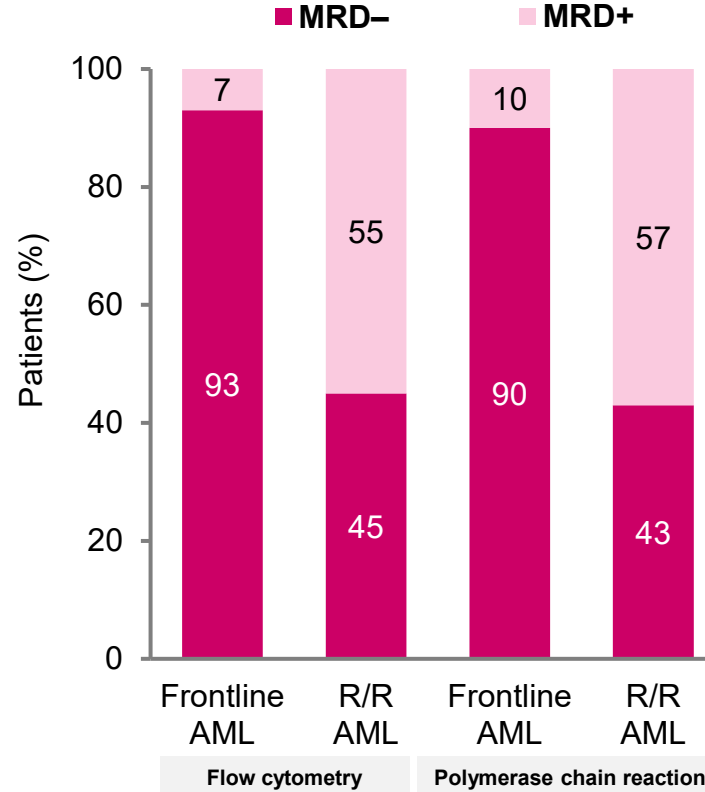
Ven+Aza+Gilt in IC-ineligible *FLT3*^{mut} AML

Phase 1/2 study in untreated IC-ineligible patients with *FLT3*^{mut} AML (N=52)

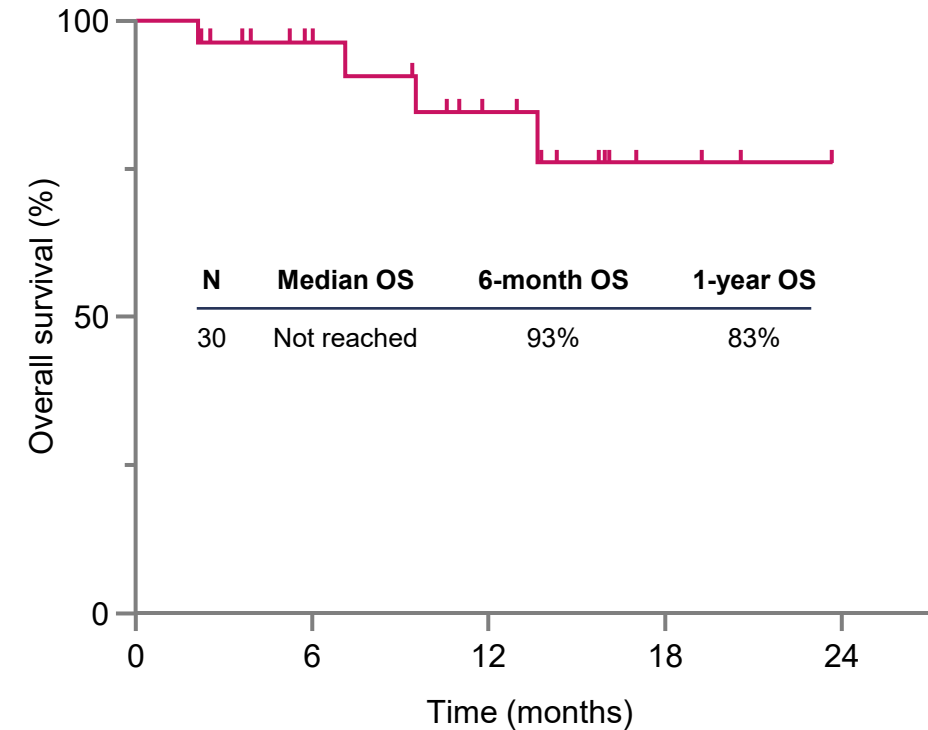
Response



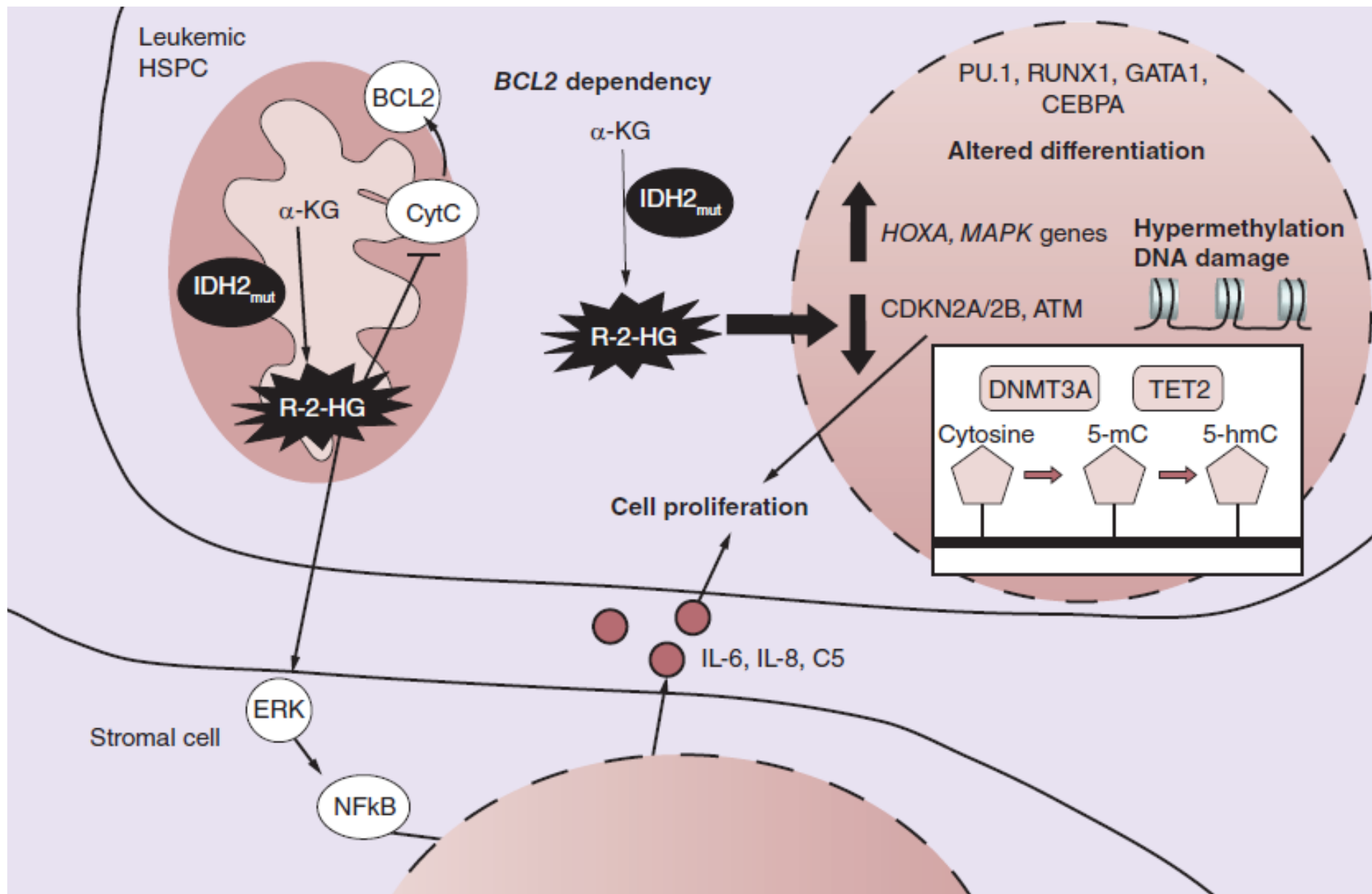
Best minimal residual disease response



Overall survival (ND)

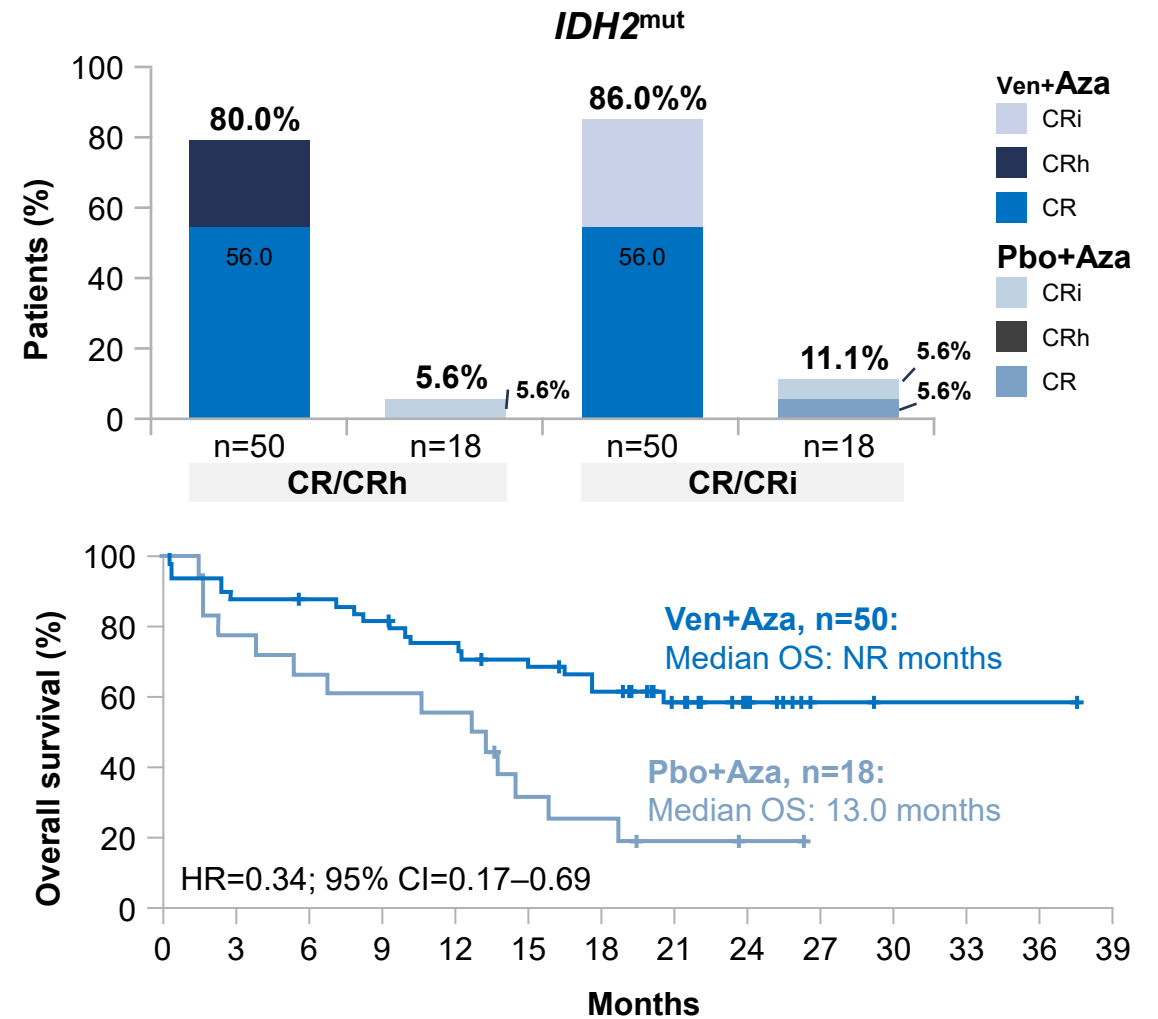
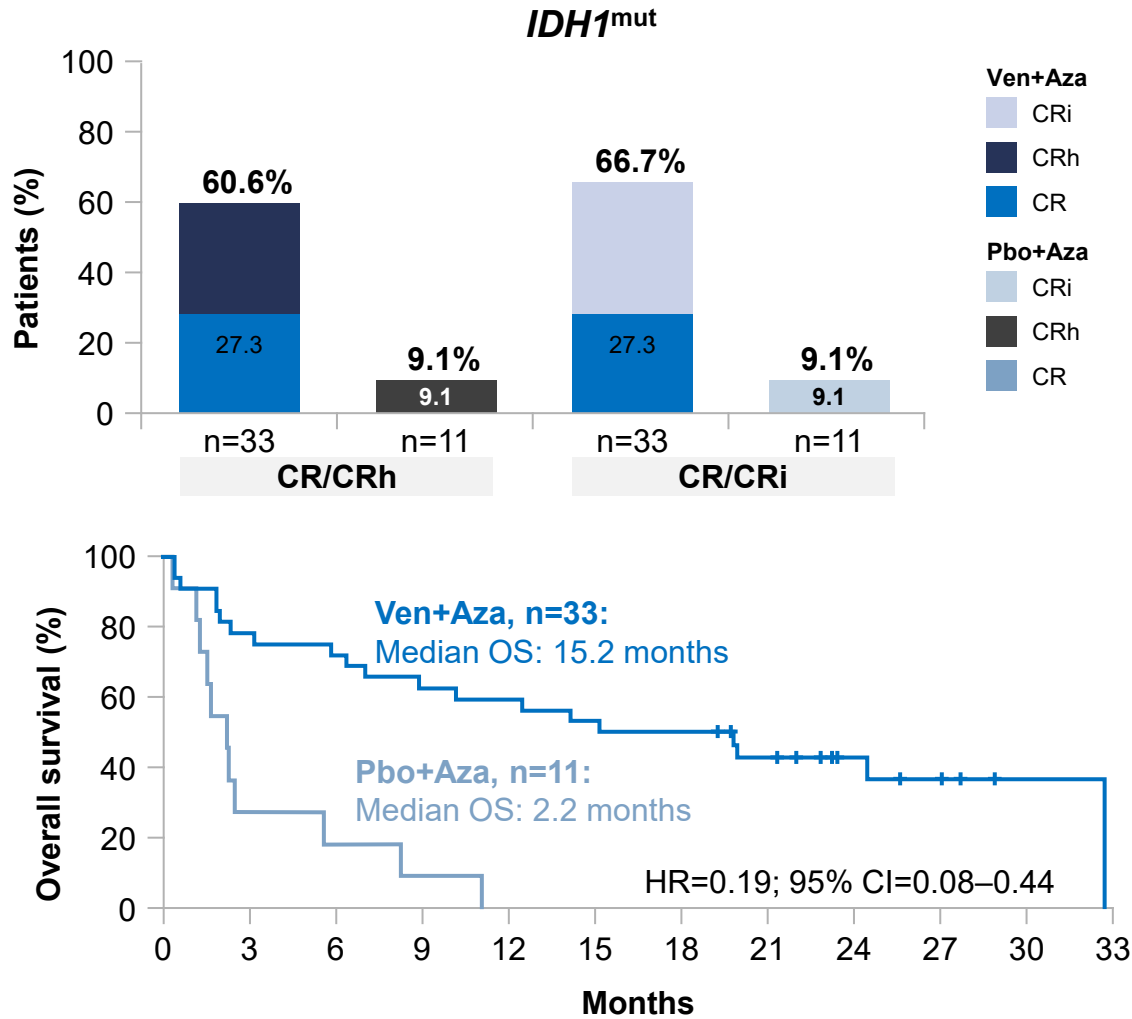


Most common Grade ≥ 3 hematologic adverse events ($>10\%$ of patients) were infections, febrile neutropenia, sepsis, GI bleeding, and hypotension. Myelosuppression was common but manageable with mitigation strategies



Response rates and OS of patients with doublet regimen Ven + Aza

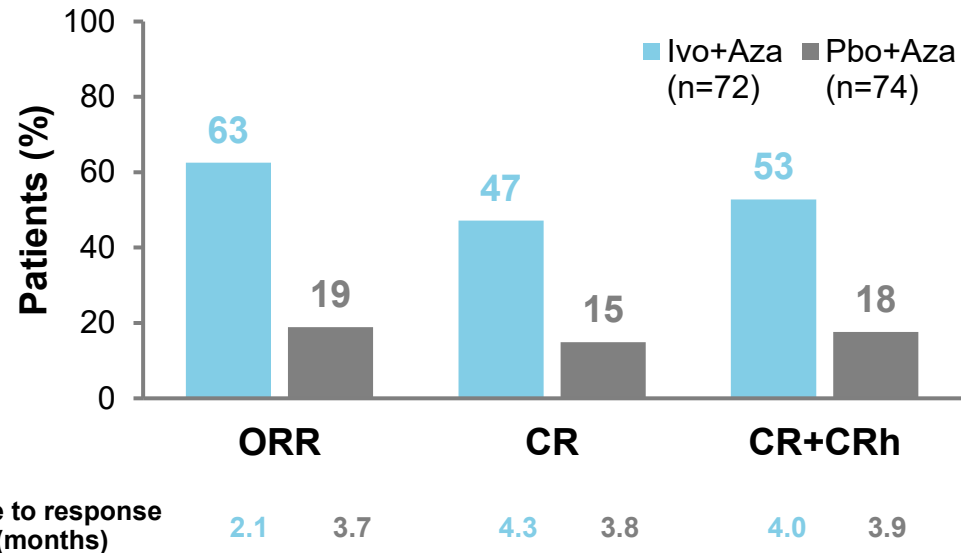
Pooled analysis of ND AML with $IDH1^{mut}$ or $IDH2^{mut}$ in VIALE-A and phase 1b study (N=498)



AGILE: Overall Survival and Safety Summary

Randomized Phase 3 trial evaluating ivosidenib + aza in newly diagnosed IDH1^{mut} AML

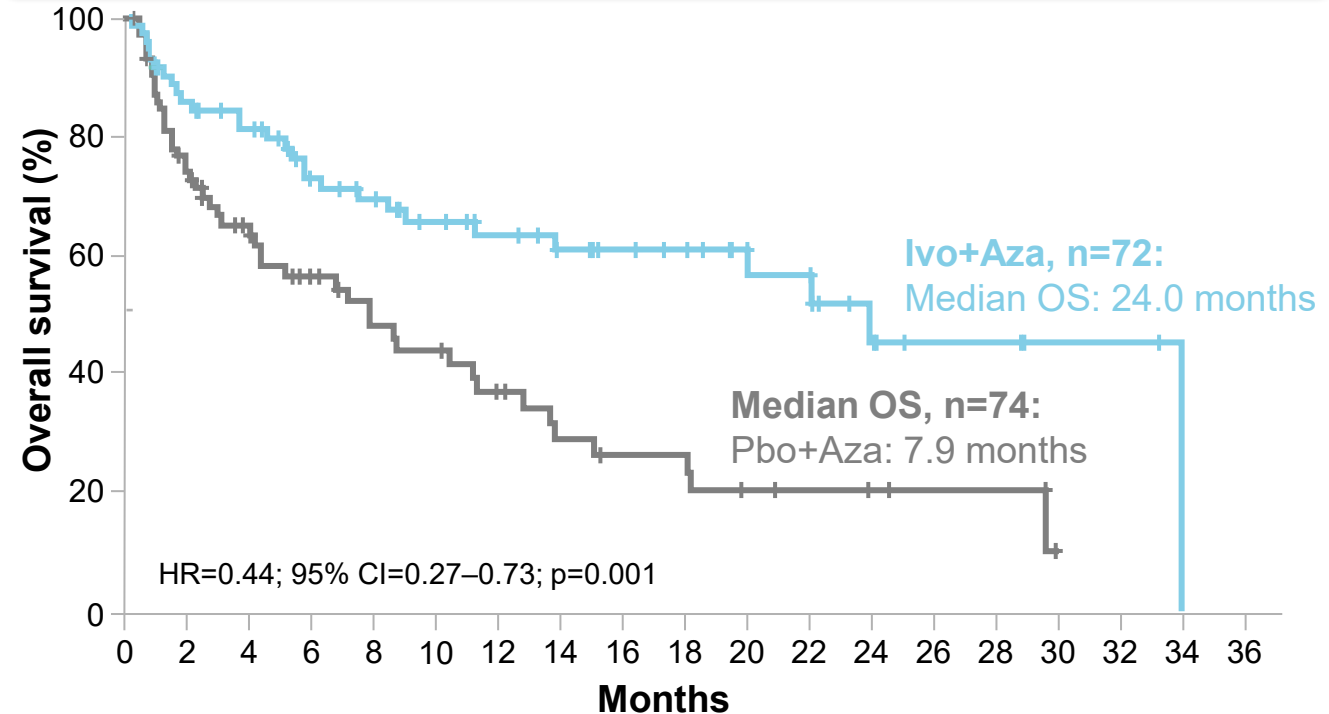
Response



Most common Grade ≥3 hematologic AEs: anemia, febrile neutropenia, neutropenia, and thrombocytopenia

Most common Grade ≥3 non-hematologic AE: pneumonia

AGILE (Ivo + Aza): OS (secondary endpoint)¹



Grade ≥3 differentiation syndrome and QT interval prolongation of ECG occurred in both arms

Discontinuation due to AEs: Ivo+Aza 19 (27%) and Pbo+Aza 19 (26%)

Median follow-up: 15.1 months. AE, adverse event; AML, acute myeloid leukemia; Aza, azacitidine; CI, confidence interval; ECG, electrocardiogram; HR, hazard ratio; Ivo, ivosidenib; mut, mutated; OS, overall survival; Pbo, placebo.

1. Montesinos P, et al. *N Engl J Med*. 2022;386(16):1519–1531.

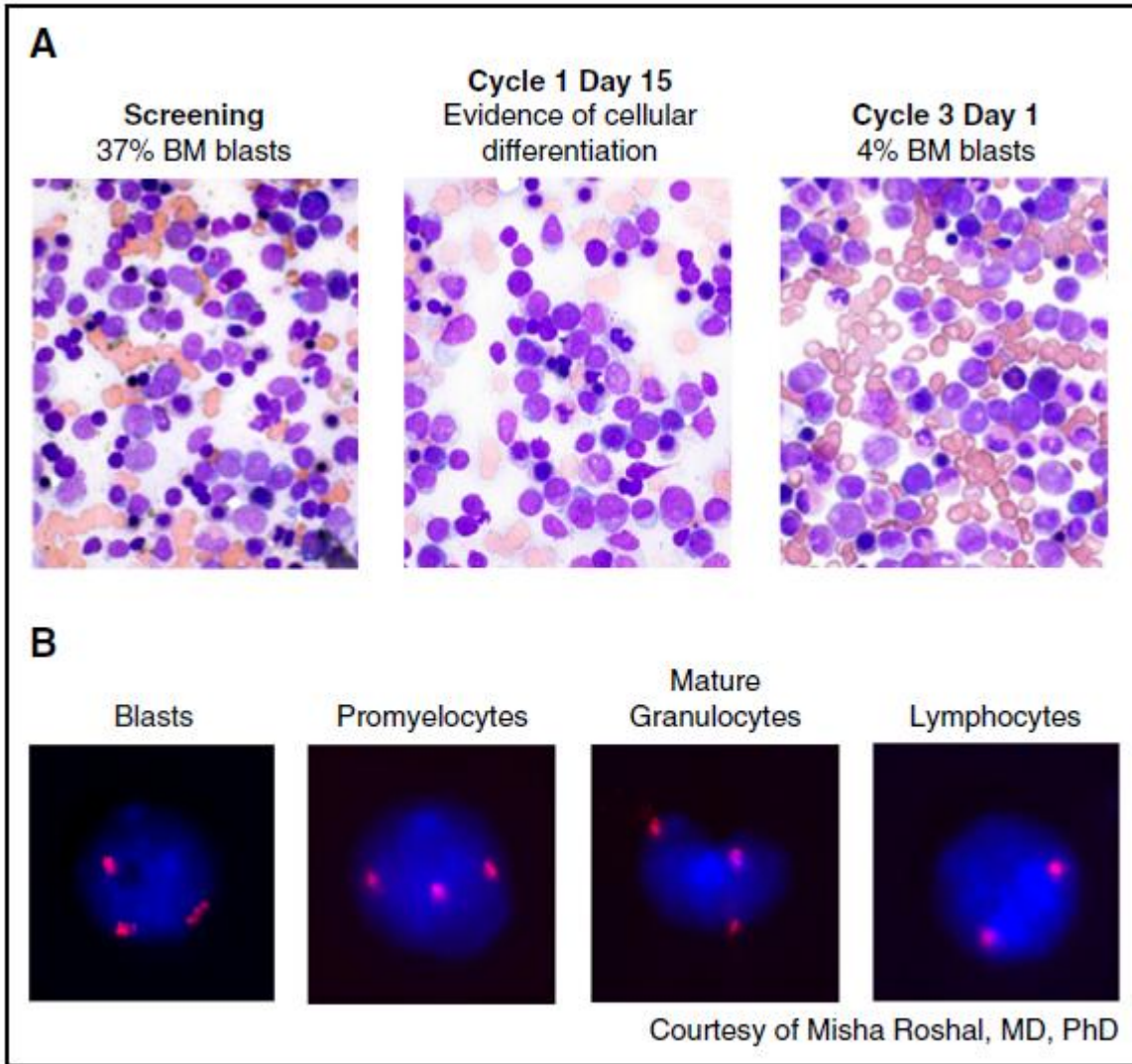
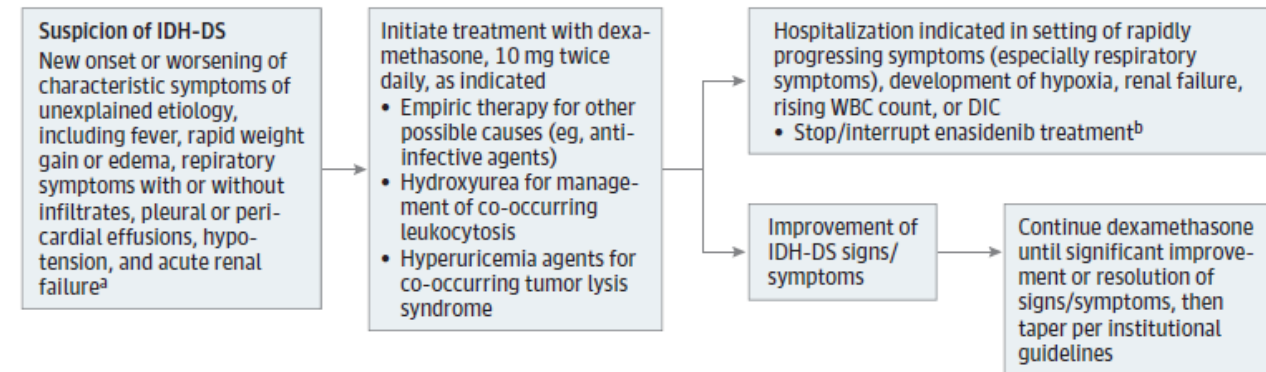


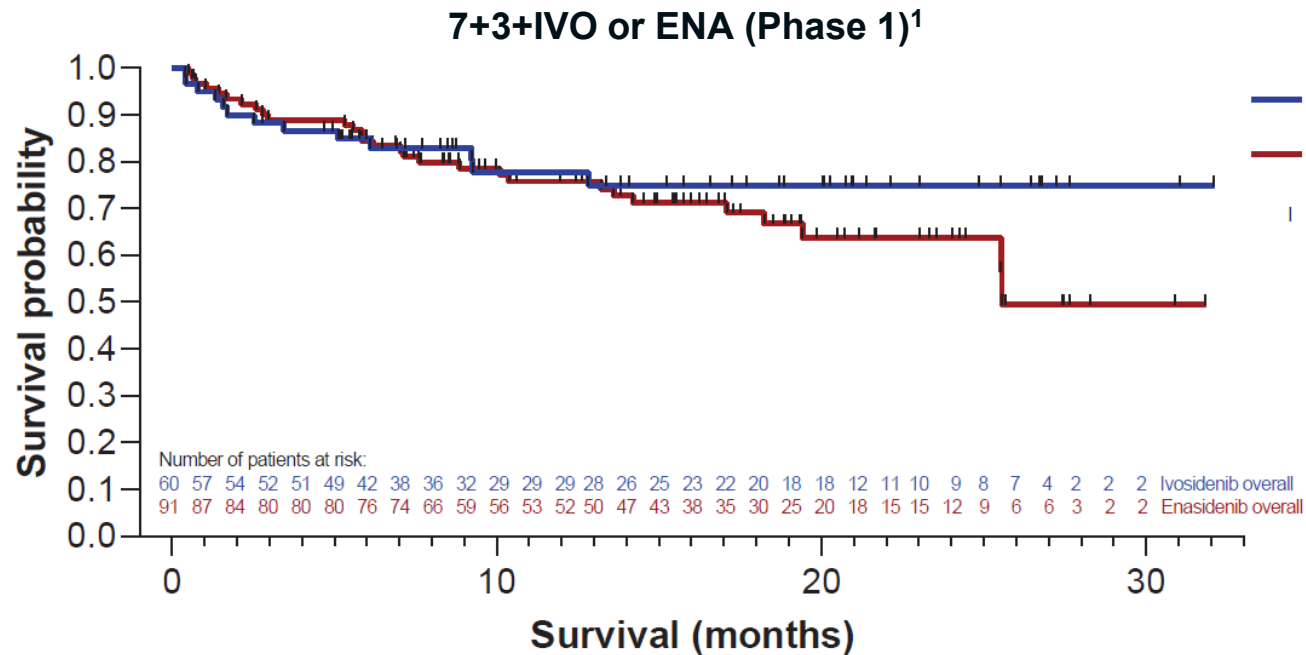
Figure. Differentiation Syndrome Review Committee Amended Protocol for Isocitrate Dehydrogenase Differentiation Syndrome (IDH-DS) Diagnosis and Management



Can we Optimize intensive induction outcome

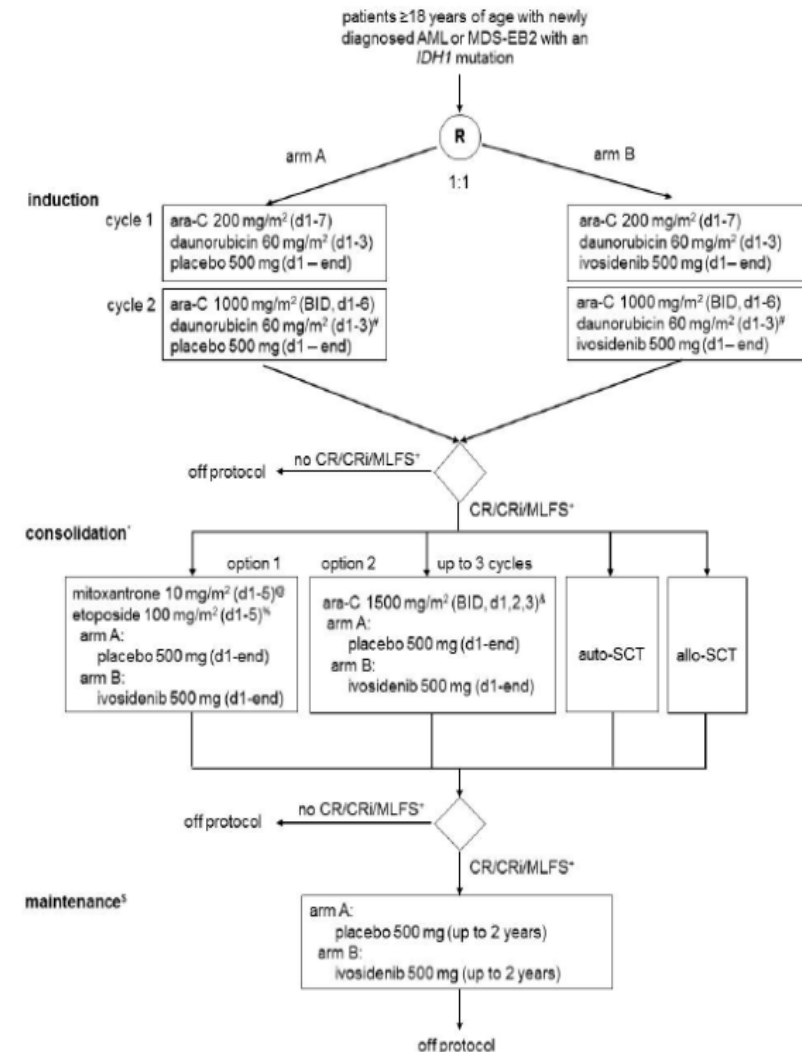
Adding IVO/ENA to intensive induction

HOVON 150 (phase 3)²



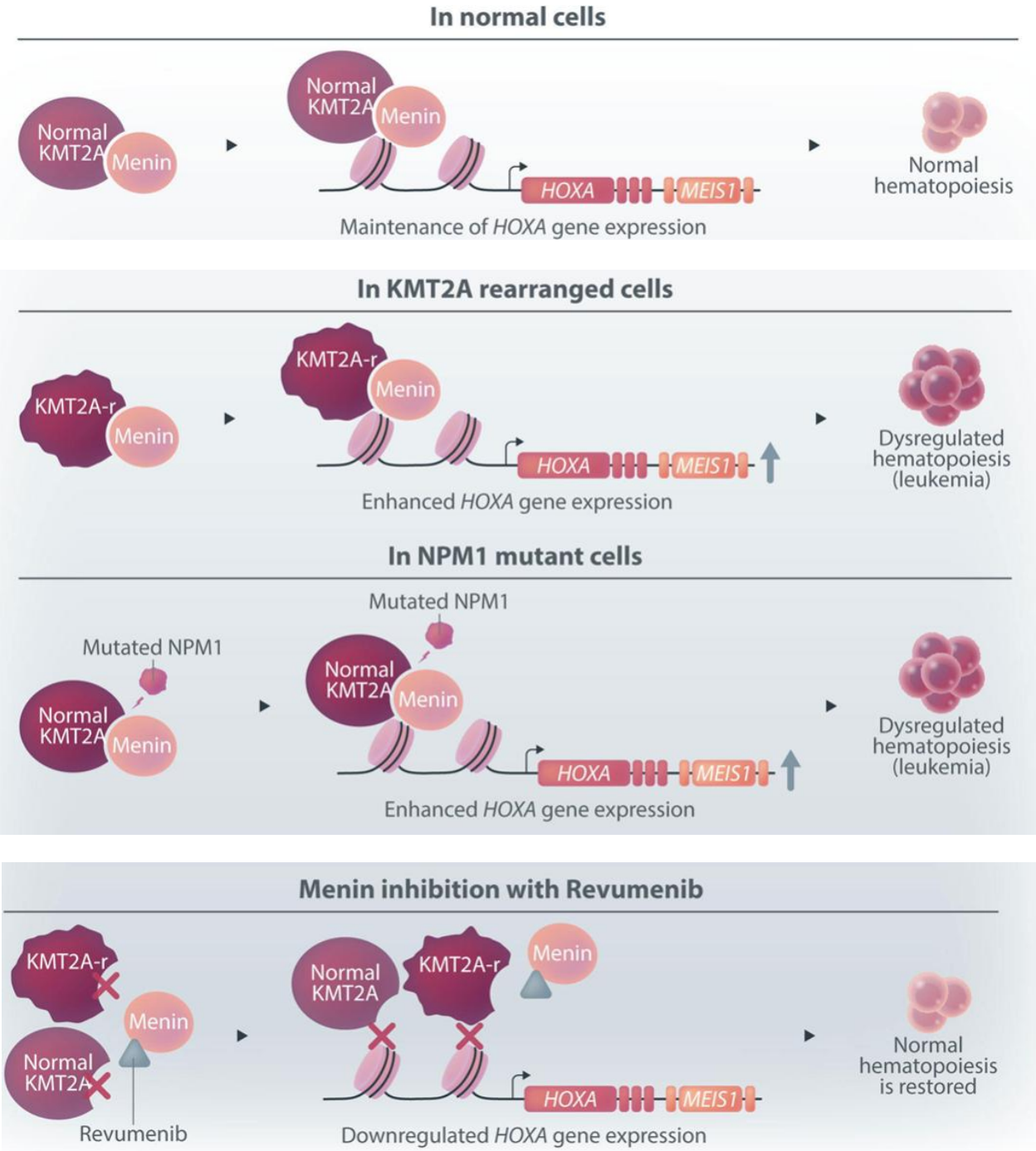
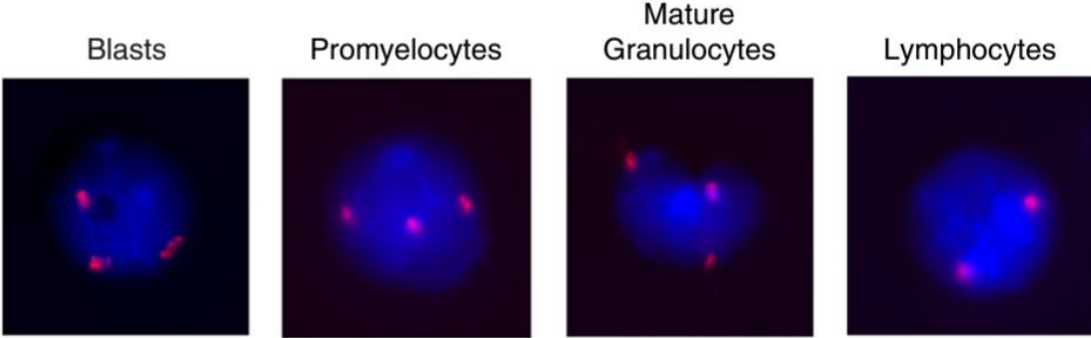
Response category	Ivosidenib 500 mg + chemotherapy, n (%)			Enasidenib 100 mg + chemotherapy, n (%)		
	All, N = 60	De novo AML, n = 42	Secondary AML, n = 18	All, N = 91*	De novo AML, n = 56	Secondary AML, n = 35
CR/CRi/CRp	46 (77)	37 (88)	9 (50)	67 (74)	45 (80)	22 (63)
CR	41 (68)	32 (76)	9 (50)	50 (55)	36 (64)	14 (40)

IDH1 cohort (randomization ivosidenib vs placebo)



Menin inhibitors – An emerging therapeutic class

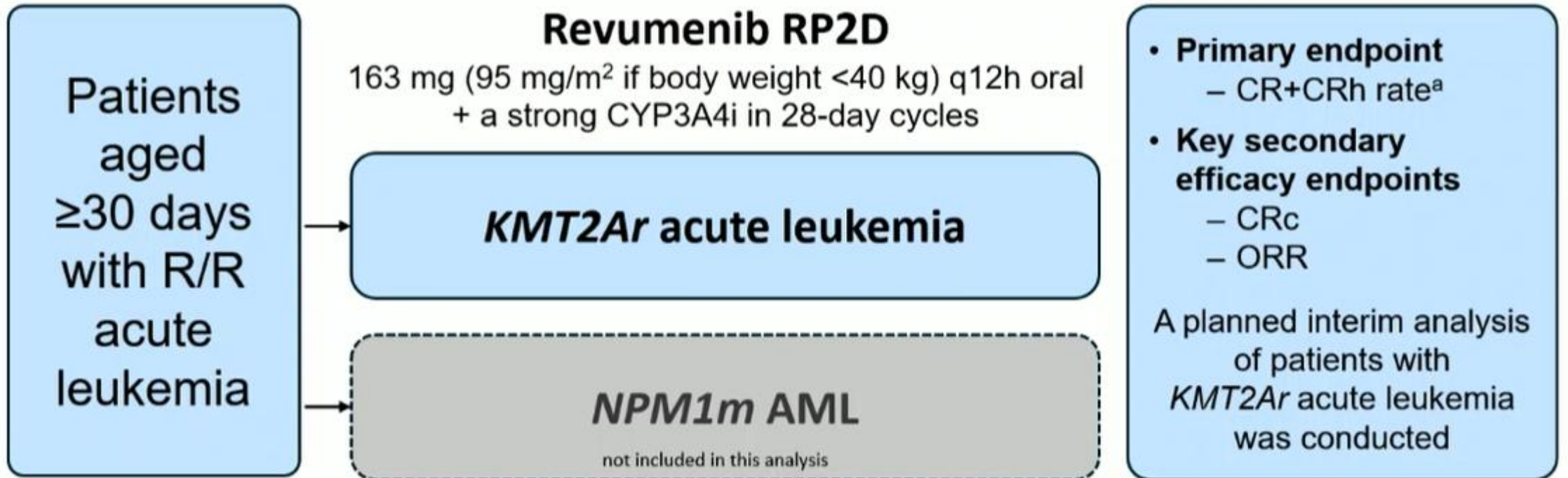
Differentiating Agent
 ATRA/ATO
 IDH Inhibitors
 FLT3 Inhibitors
 Menin Inhibitors
 Other differentiating agents



Adapted from Issa et al. Blood 2024; Stein et al. Blood, 2017; 130 (6): 722-731; Salman & Stein. Haematologica 2024

AUGMENT-101

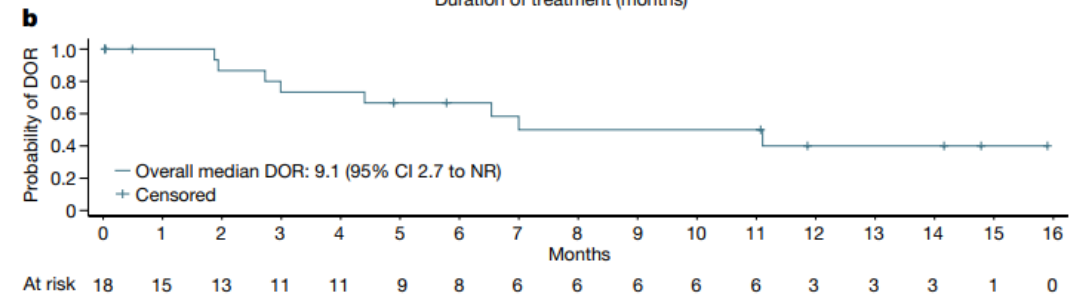
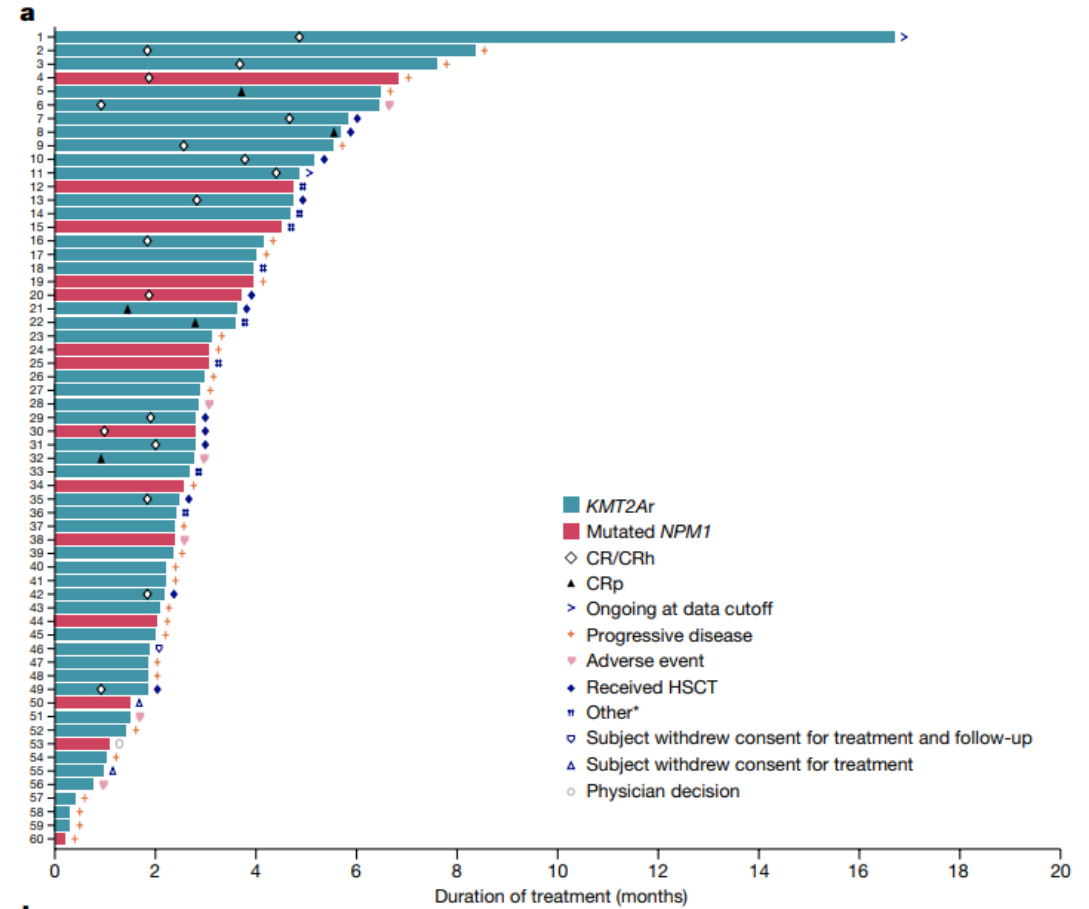
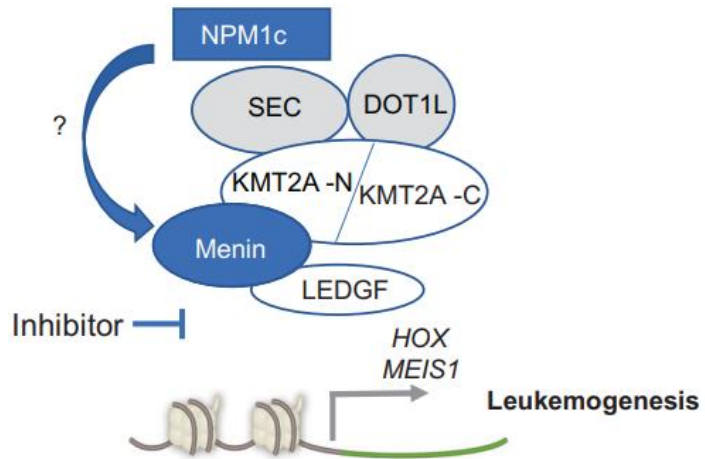
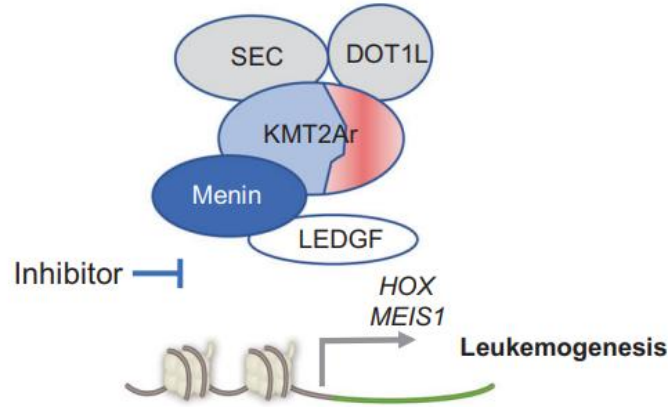
Phase 2 study



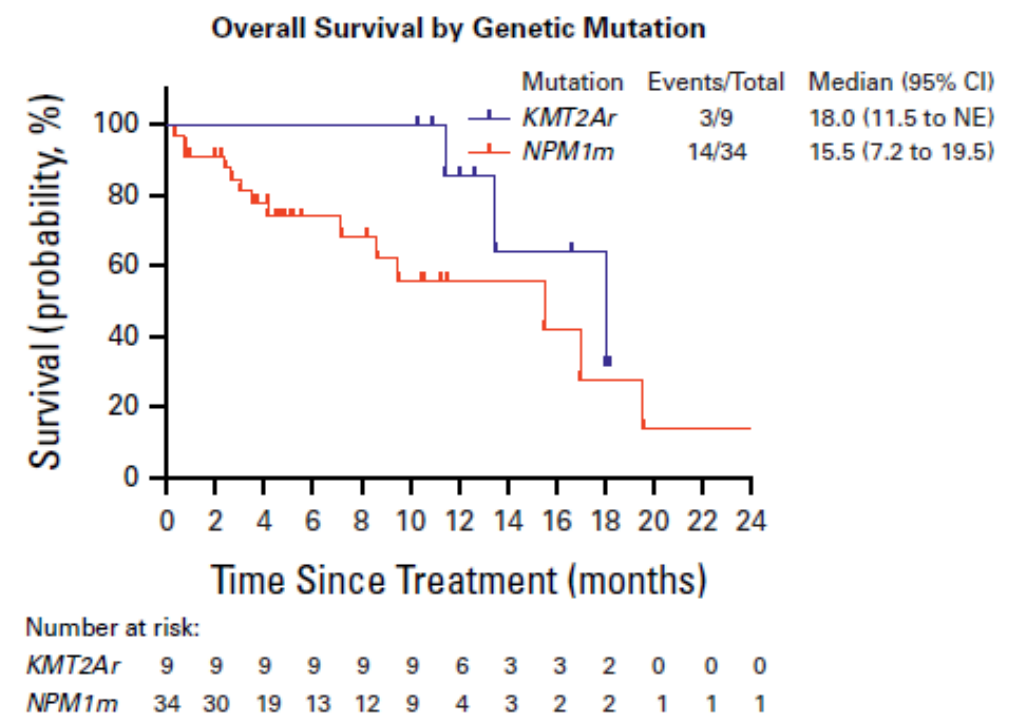
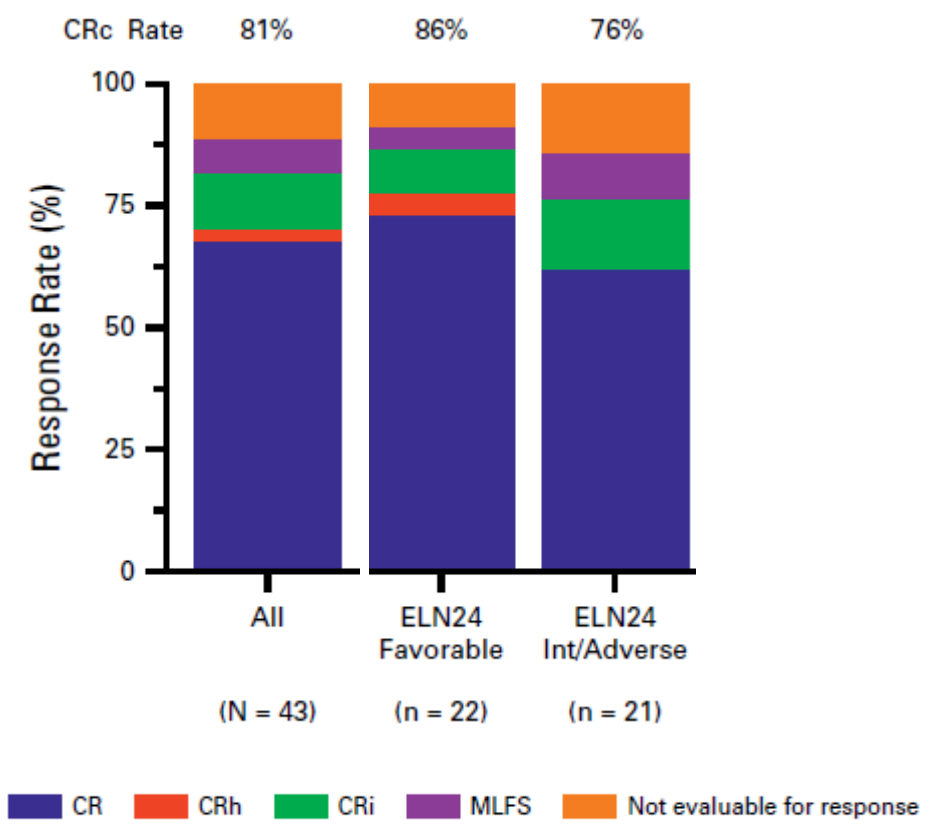
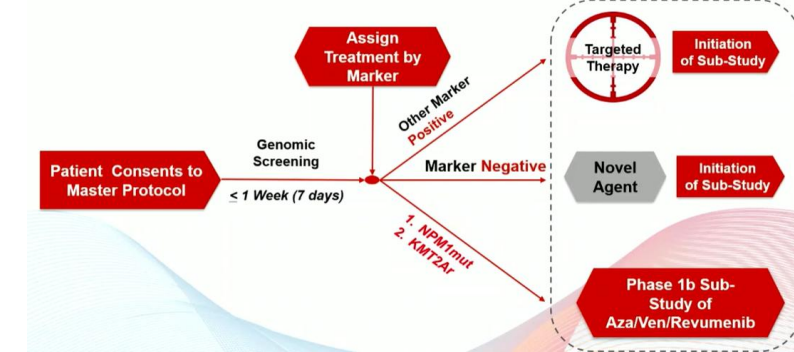
^aCR+CRh rate >10% in adult evaluable population considered lower efficacy bound.

AUGMENT-101 Phase 2 study

Revumenib (SNDX-5613)

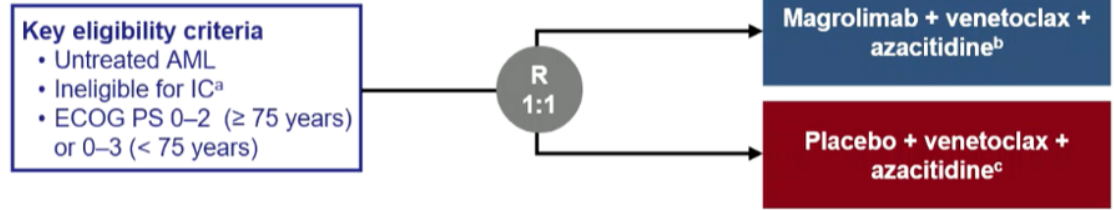


Menin inhibitor triplets, AZA-Ven+revumenib (BEAT AML sub-study)

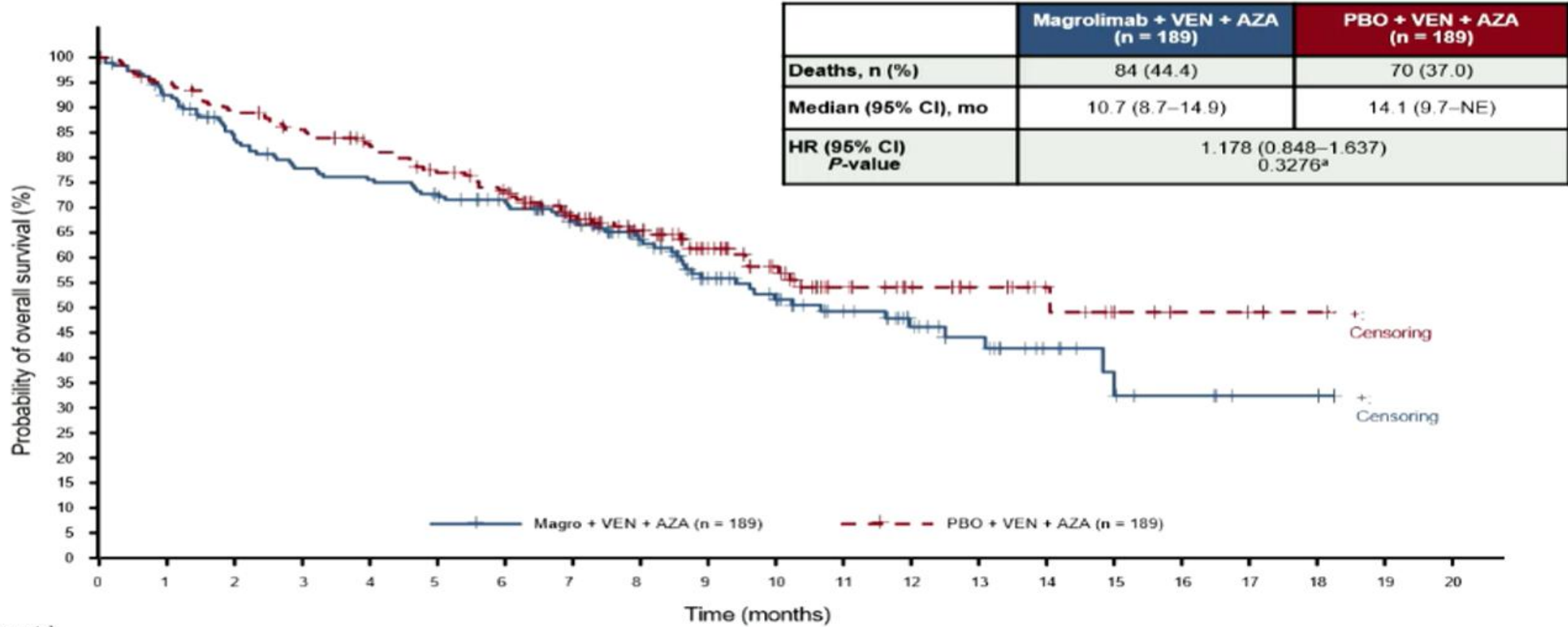


DS in 19% (8 pt), 2> G3
 QTc prolongation in 44% (19 pt), 5> G3

Ven+Aza±Magro in IC-ineligible pts.



Overall Survival at Final Analysis: ENHANCE-3



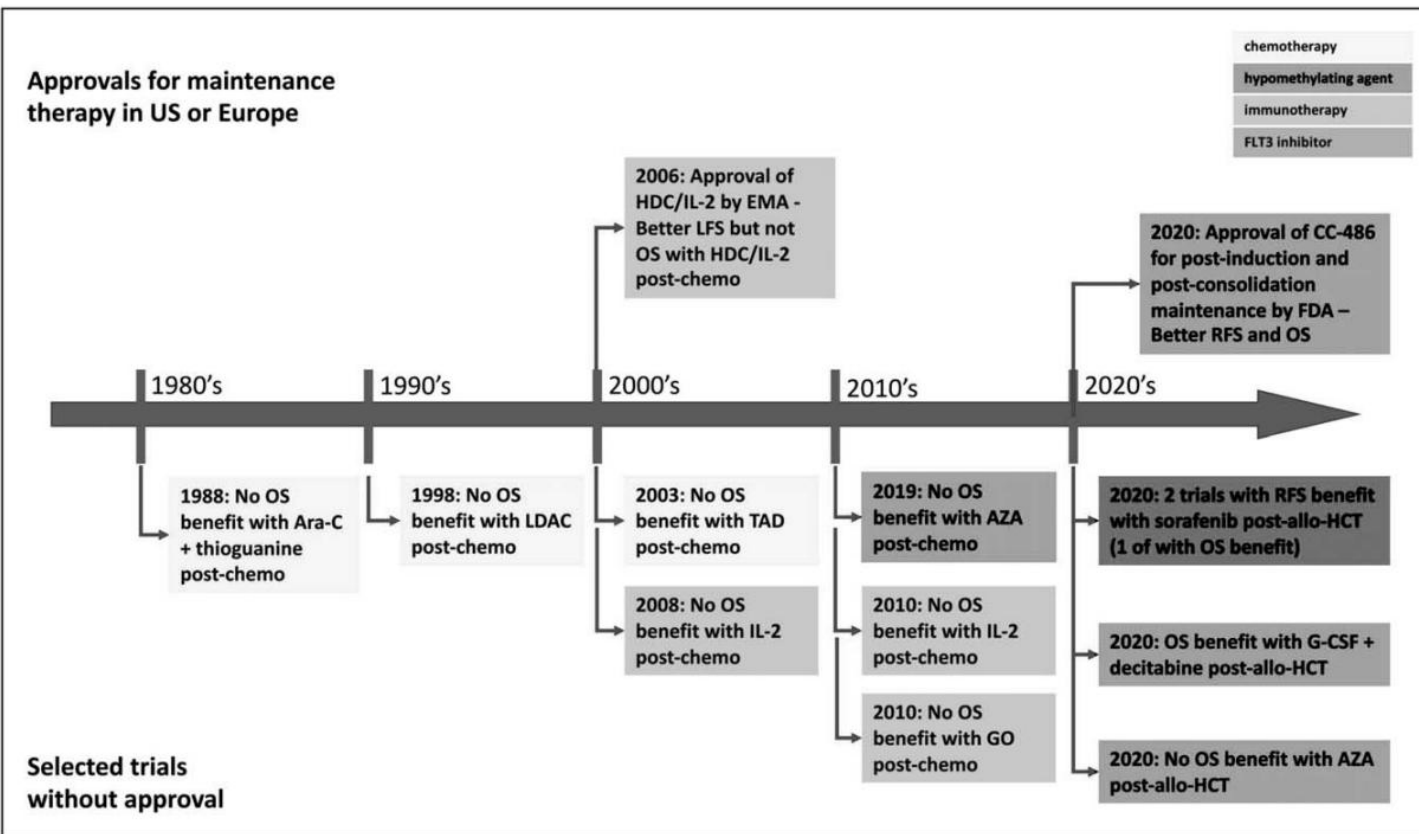
N at risk (events)		0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
Magro + VEN + AZA	(n = 189)	189 (0)	173 (13)	152 (27)	138 (40)	135 (43)	129 (49)	120 (51)	107 (56)	84 (61)	61 (72)	50 (75)	38 (78)	30 (79)	20 (81)	13 (82)	8 (83)	5 (84)	2 (84)	2 (84)	0 (84)	
PBO + VEN + AZA	(n = 189)	189 (0)	174 (9)	163 (19)	154 (26)	146 (30)	134 (40)	125 (47)	104 (54)	84 (59)	63 (63)	46 (66)	29 (69)	22 (69)	16 (69)	12 (69)	8 (70)	3 (70)	3 (70)	1 (70)	0 (70)	

Daver N, et al. *Blood* 2025 .601-611:(5)146

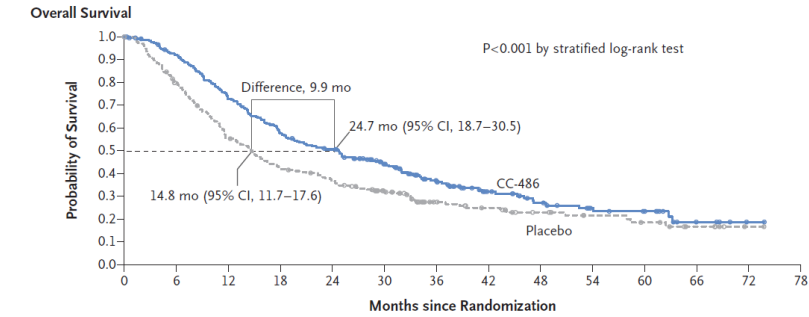
Venetoclax is used in combination with either Azacitidine or low-dose cytarabine in Japan. Please refer to the Venclexta package insert for further details.

Improving patient outcome – maintenance is back...

The long road for effective maintenance¹

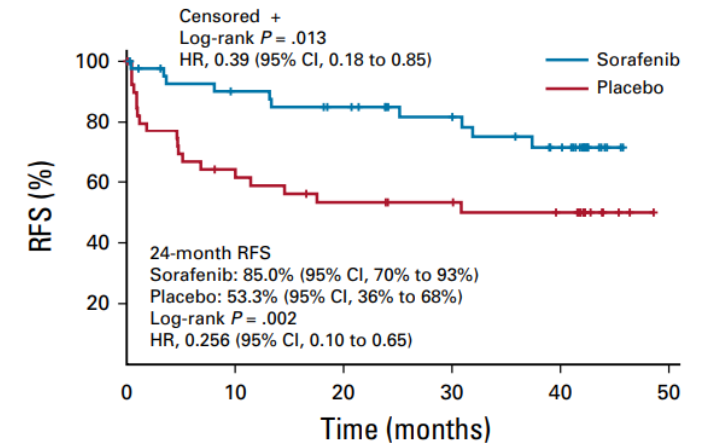


Oral AZA (non-SCT)²



No. at Risk

Months since Randomization	0	6	12	18	24	30	36	42	48	54	60	66	72	78
CC-486	238	213	168	133	115	87	59	37	26	18	15	5	1	0
Placebo	234	183	127	96	82	58	34	27	19	14	11	6	1	0



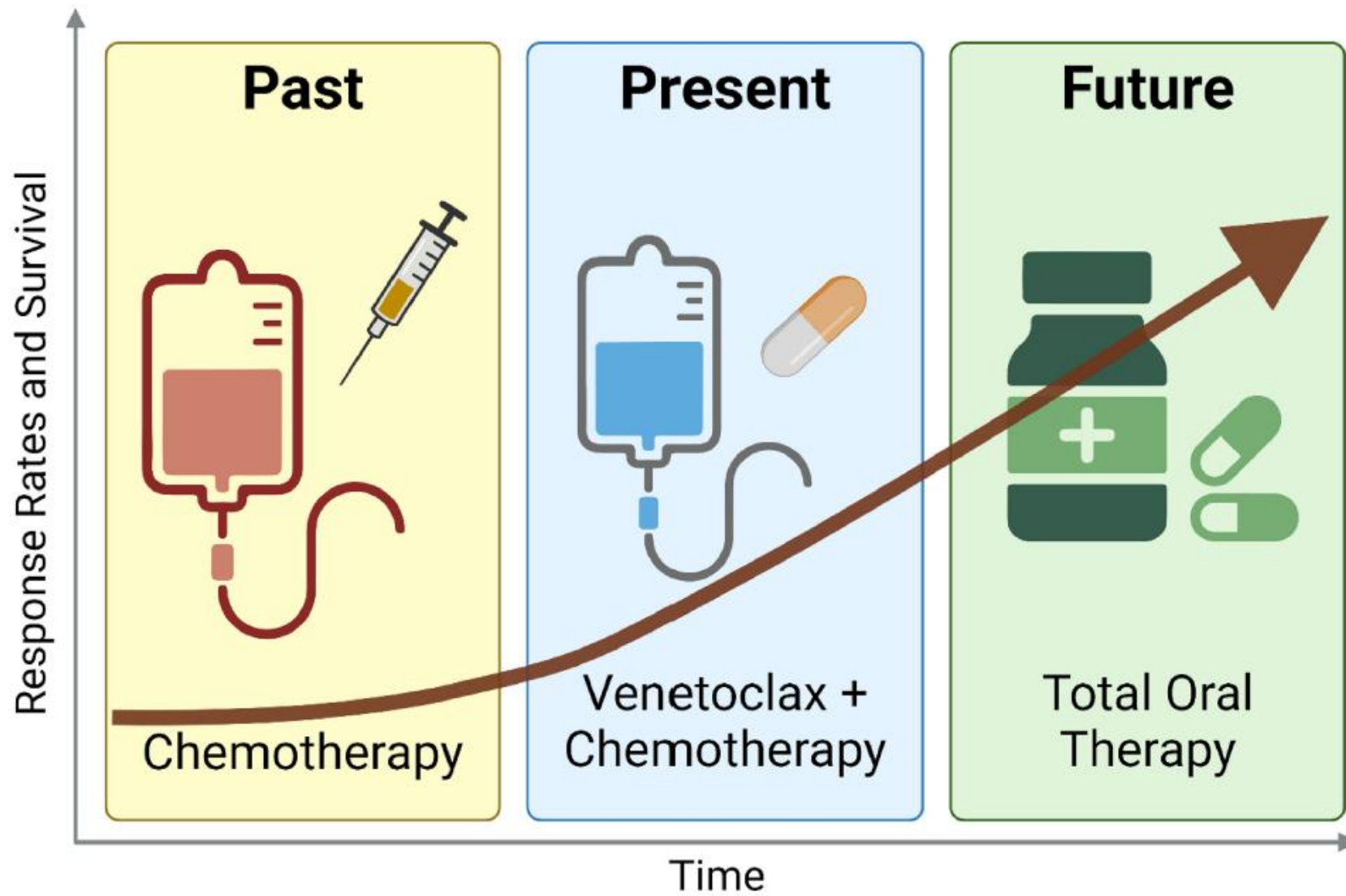
No. at risk:

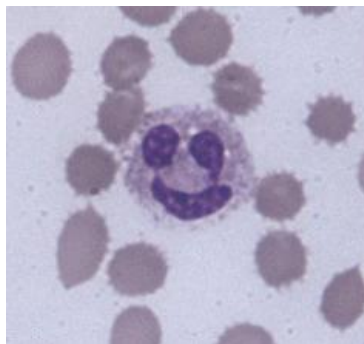
Time (months)	0	10	20	30	40	50
Placebo	40	24	19	17	14	0
Sorafenib	43	35	31	25	18	0

Post allo SCT (FLT3-ITD)³

1. Bewesdorf et al. Curr Opin Oncology 2021 Nov 1;33(6):658-669;

2. Wei et al. N Engl J Med 2020;383:2526-37 3. Burchert et al. JCO 2020 Sep 10;38(26):2993-3002





owolach@gmail.com

Post-remission approach

AML risk group‡	AML risk assessment criteria at diagnosis	MRD after cycle 2	Risk of relapse following consolidation approach		Prognostic scores for NRM that indicate alloHSCT as preferred consolidation		
			Chemotherapy or autoHSCT (%)	AlloHSCT (%)	EBMT score ⁵²	HCT-CI score ⁵³	NRM risk (%)
Good	–t(8;21) or <i>AML1-ETO</i> , WBC <20 –inv16/t(16;16) or <i>CBFB-MYH11</i> –CEBPA-biallelic mutant-positive –FLT3-ITD-negative/NMP1-positive	Positive or negative	35-40	15-20	NA (≤1)	NA (<1)	10-15
Intermediate	–CN –X –Y, WBC <100, CRe –t(8;21) or <i>AML1-ETO</i> plus WBC >20 or mutant KIT	Negative	50-55	20-25	≤2	≤2	<20-25
Poor	–CN –X –Y, WBC <100, CRe –t(8;21) or <i>AML1-ETO</i> , WBC >20 and/or mutant KIT –CN –X –Y, WBC <100, no CRe –CN –X –Y, WBC >100 –CA, but non-CBF, MK-negative, no abn3q26	Positive Positive Negative Negative	70-80	30-40	≤3-4	≤3-4	<30
Very poor	–CN –X –Y, WBC >100 –CA, but non-CBF, MK-negative, no abn3q26, EVI1-negative –MK-positive –abn3q26 –Non-CBF, EVI1-positive –Non-CBF with mutant p53, or –mutant RUNX1, or mutant ASXL1 –or biallelic FLT3-ITD with –FLT3-ITD:FLT3 WT ratio of >0.6	Positive Positive Positive or negative	>90	40-50	≤5	≤5	<40