



Memorial Sloan Kettering  
Cancer Center

# Acute Myeloid Leukemia: Targets and Curability, so Close But a Journey So Far

Martin S. Tallman, M.D.  
Chief, Leukemia Service  
Memorial Sloan Kettering Cancer Center  
Professor of Medicine  
Weill Cornell Medical College  
New York, NY



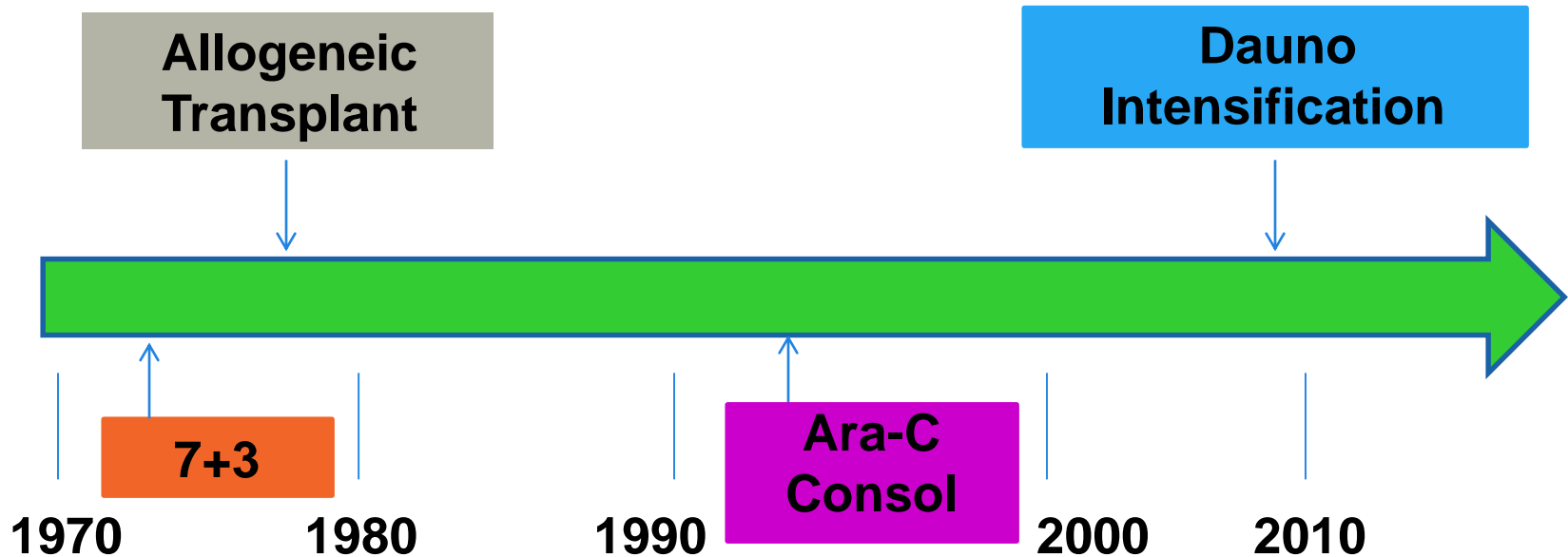
# Disclosure

Principal investigator role	Cellerant, ADC Therapeutics, Orsenix, Arog, Bioline
Employee	None
Consultant	None
Major Stockholder	None
Speakers' Bureau	None
Scientific Advisory Board	Daiichi Sankyo
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Presentation includes the following off-label drug use:  
Gilteritinib, Quizartinib, Crenolanib, Venetoclax, Selinexor, Tamobarotene,  
Entospletinib, Palbociclib, Cobimetinib, Pevonedistat, H3B-8800

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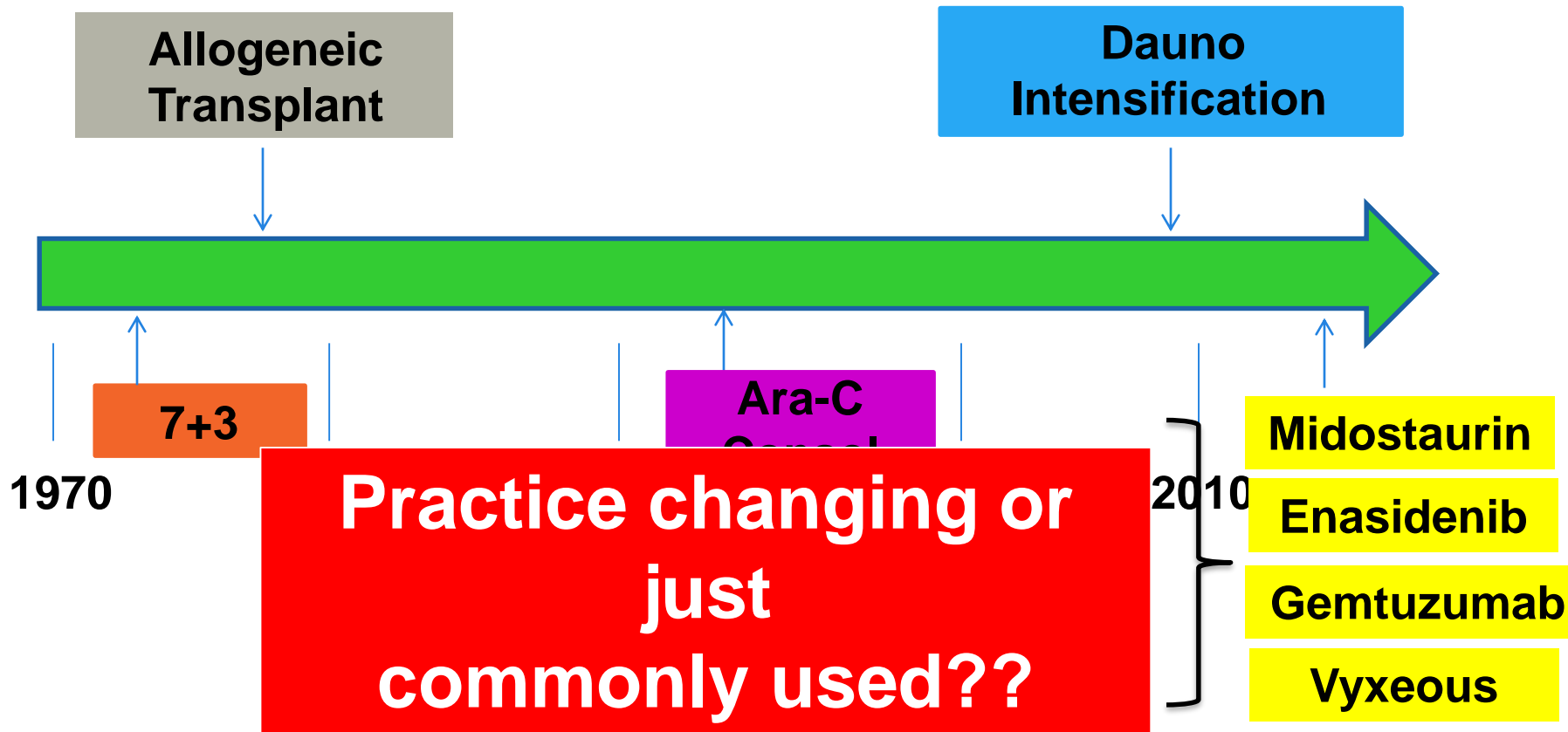
# Practice Changing Treatments in AML



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*Thomas et al. NEJM, 1979; Mayer et al. NEJM, 1994; Fernandez et al. NEJM, 2009; Stone et al. NEJM, 2017*

# Practice Changing Treatments in AML



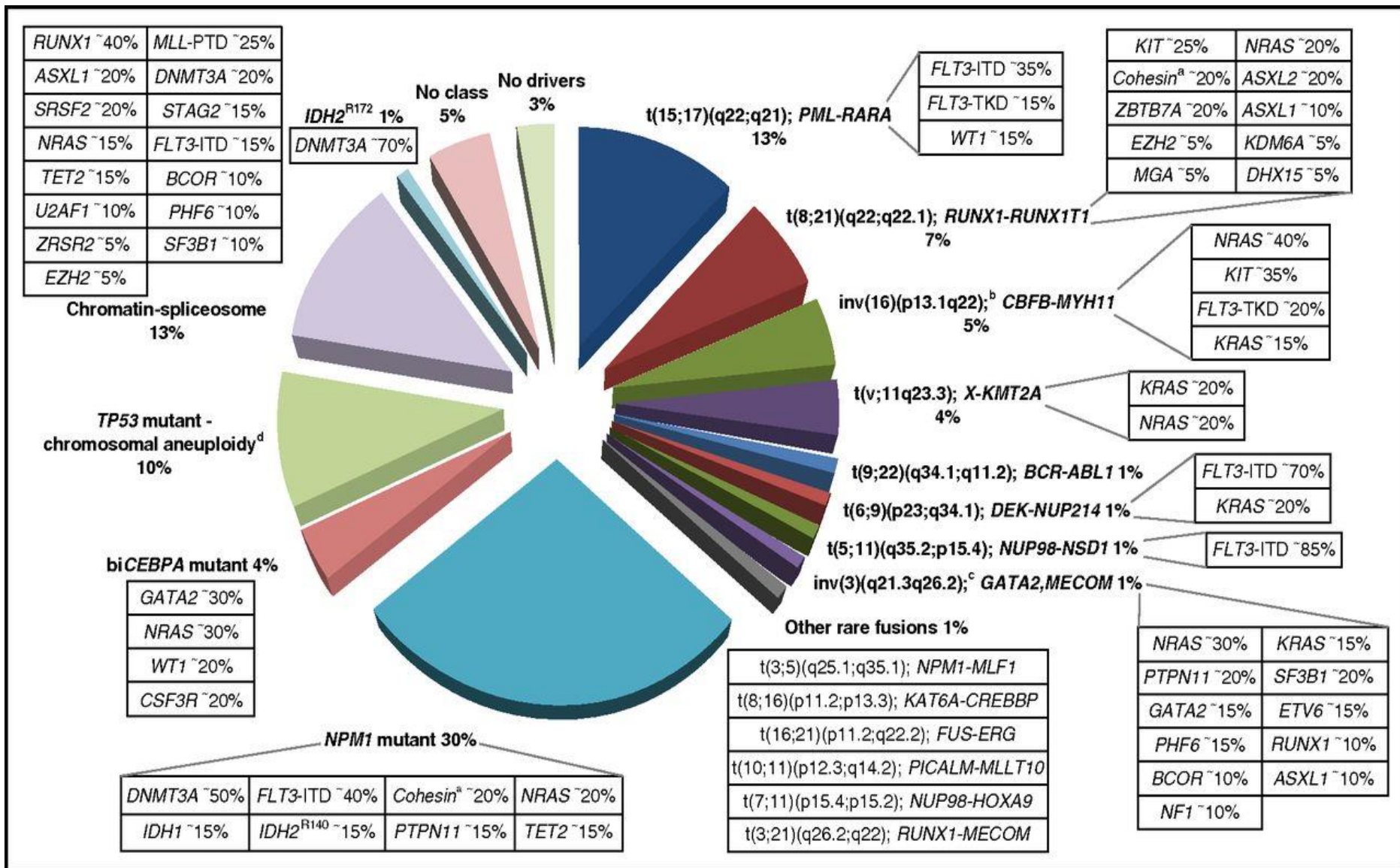
*Thomas et al. NEJM, 1979; Mayer et al. NEJM, 1994; Fernandez et al. NEJM, 2009; Stone et al. NEJM, 2017*

# Acute Myeloid Leukemia

## State-of-the-Art 2018

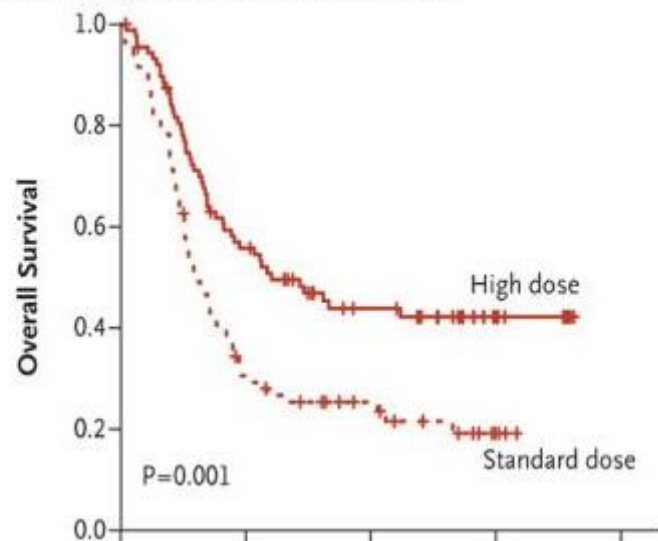
- **Defined by cytogenetic and molecular interactions**
  - Intensified induction/less intensive consolidation
  - Increased importance of minimal residual disease
  - Expanded availability of allogeneic transplantation
  - Paradigm shift in older patients
  - Incorporation of novel agents
-

# Molecular Classes of AML and Recurrent Gene Mutations

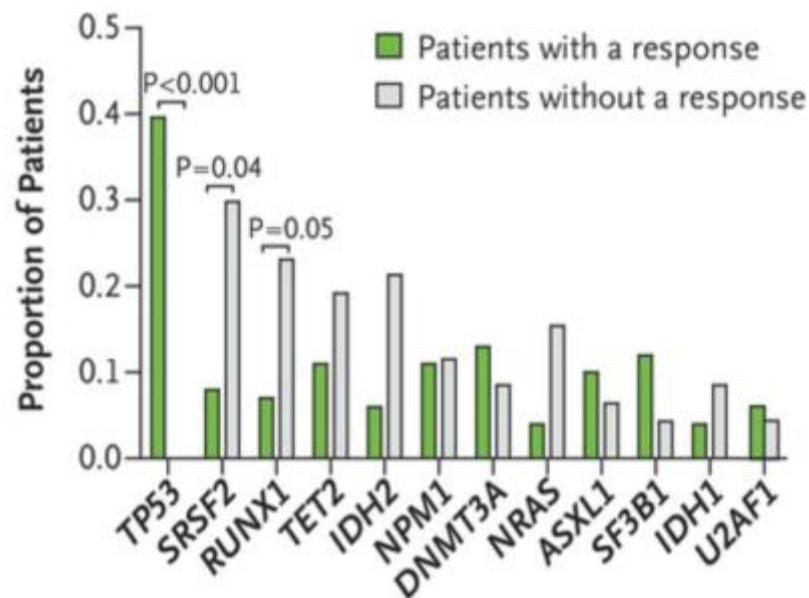


# Risk-Stratification and Prognostication of AML Informed by Mutational Profile

Mutant *DNMT3A* or *NPM1*, or *MLL* Translocation



*Patel et al. NEJM, 2012*



*Welch et al. NEJM, 2016*

# Mutation Patterns in Older Adults Predict Response to Chemotherapy

## Good Risk

- CR 81%: *NPM1* plus
  - Chromatin mutations
  - Cohesin mutations
  - *FLT3-TKD*
  - Spliceosome mutations
  - *RAS* pathway mutations
  - *FLT3-ITD*<sup>wt</sup>
- DFS 46%: *NPM1* plus
  - *ASXL1*
  - *SF 1*
  - *SMC1A*
  - *SRSF2*
- OS 45%: *NPM1* plus
  - Chromatin mutations
  - *IDH2* mutation
  - *SF 1*
  - *SRSF2*

## Poor Risk

- CR 32%
  - U2AF1*
  - WT1*
  - Complex karyotype
- DFS 2%
  - FLT3*
  - RUNX1*
  - TP53, U2AF1*
- OS 4%
  - BCOR*
  - FLT3-ITD*
  - U2AF1, WT1*
  - t(9;11), complex karyotype*



# Gene Mutations Important in Everyday Practice

## “Clinically Actionable”

<u>Gene</u>	<u>Incidence</u>	<u>Associations</u>	<u>Impact</u>
<i>FLT3-ITD/(TKD)</i>	25%	<i>NPM1</i>	Unfavorable
<i>NPM1</i>	33%	<i>FLT3</i>	Favorable
<i>dCEBP<math>\alpha</math></i>	8%	<i>FLT3</i>	Favorable
<i>C-KIT</i>	15%	<i>CBF</i>	Unfavorable [in t(8;21), but less clear in inv(16)]; <sup>1</sup> D816 worse than others
<i>IDH1 and 2</i>	22%	<i>NPM1</i>	Favorable
<i>P53</i>	7%	t-AML, Complex karyotype (60%)	Unfavorable

<sup>1</sup>Yui et al. ASH abstr 2785, 2016

# RATIFY (C10603) Trial Schema

P  
R  
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N  
A  
R  
Y  
R  
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G  
I  
S  
T  
E  
R

Stratify\*  
*FLT3*  
ITD  
or  
TKD

R  
A  
N  
D  
O  
M  
I  
Z  
E

DNR  
ARA-C  
Midostaurin

CR

HiDAC  
Midostaurin

X 4

Midostaurin  
MAINTENANCE  
12 months

DNR  
ARA-C  
Placebo

CR

HiDAC  
Placebo

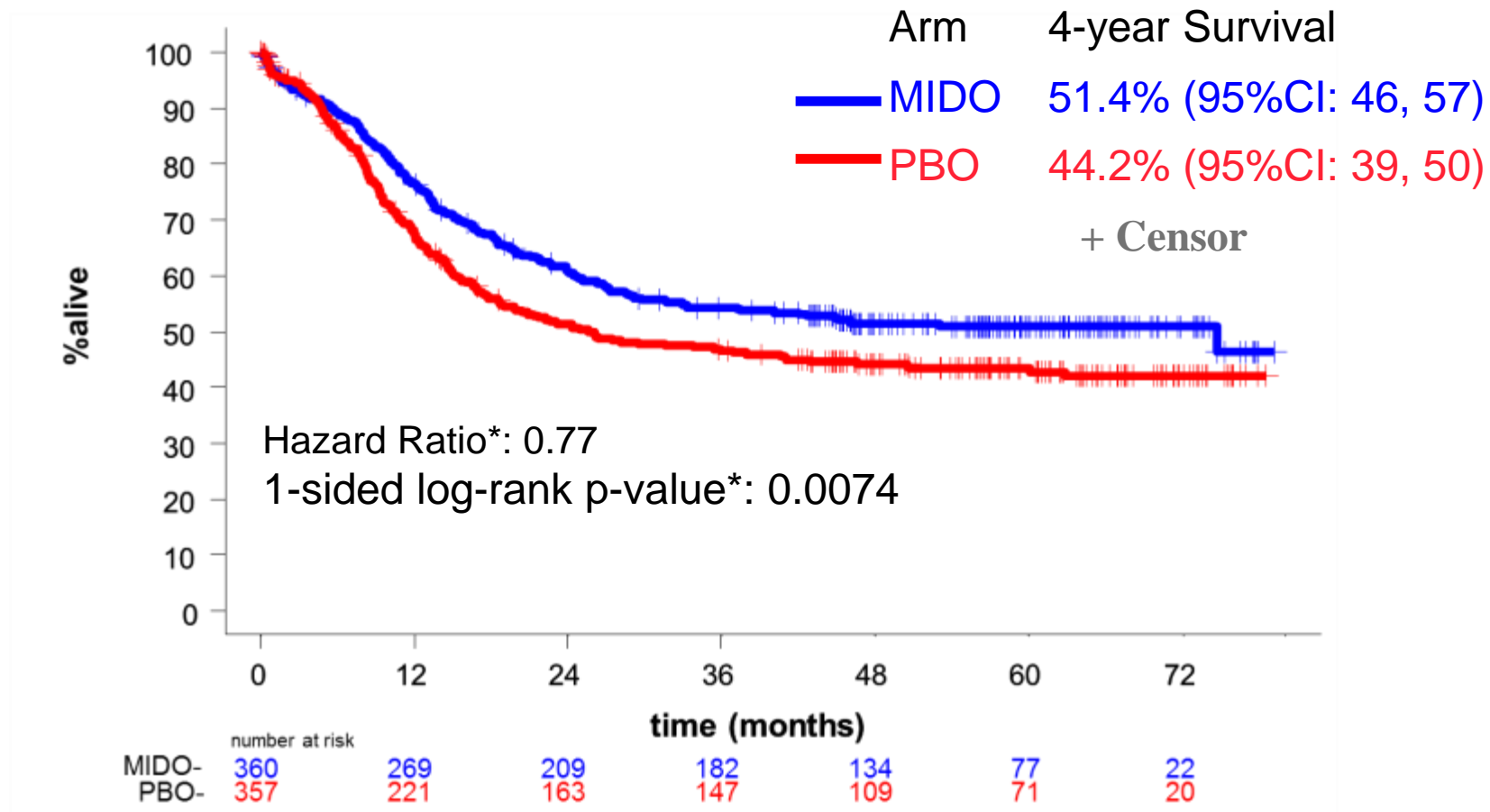
X 4

Placebo  
MAINTENANCE  
12 months

Stratification: TKD; ITD with allelic ratio  $<0.7$  'vs'  $\geq 0.7$

# Overall Survival

23% reduced risk of death in the Mido arm



# Midostaurin in AML

- First agent with (sustained) regulatory approval in 40 years
- BUT, will it be practice changing? Will it have a true (clinically meaningful) impact?
  - OS increase only 7%
  - Benefit more in *FLT3*-TKD than ITD
  - Which phase of treatment important?
  - Among least potent *FLT3* inhibitors
  - Role in maintenance unclear<sup>1</sup>
  - Beneficial effect of Midostaurin most pronounced in *NPM1*<sup>wt</sup>/*FLT3*<sup>high</sup> group<sup>2</sup>

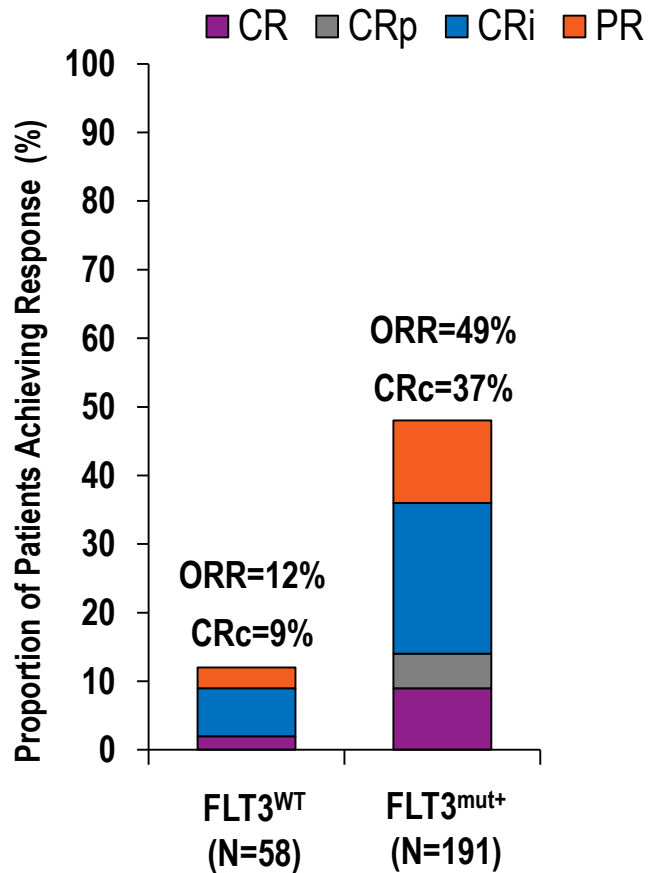
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<sup>1</sup>Larson et al. ASH abstr 145, 2017;

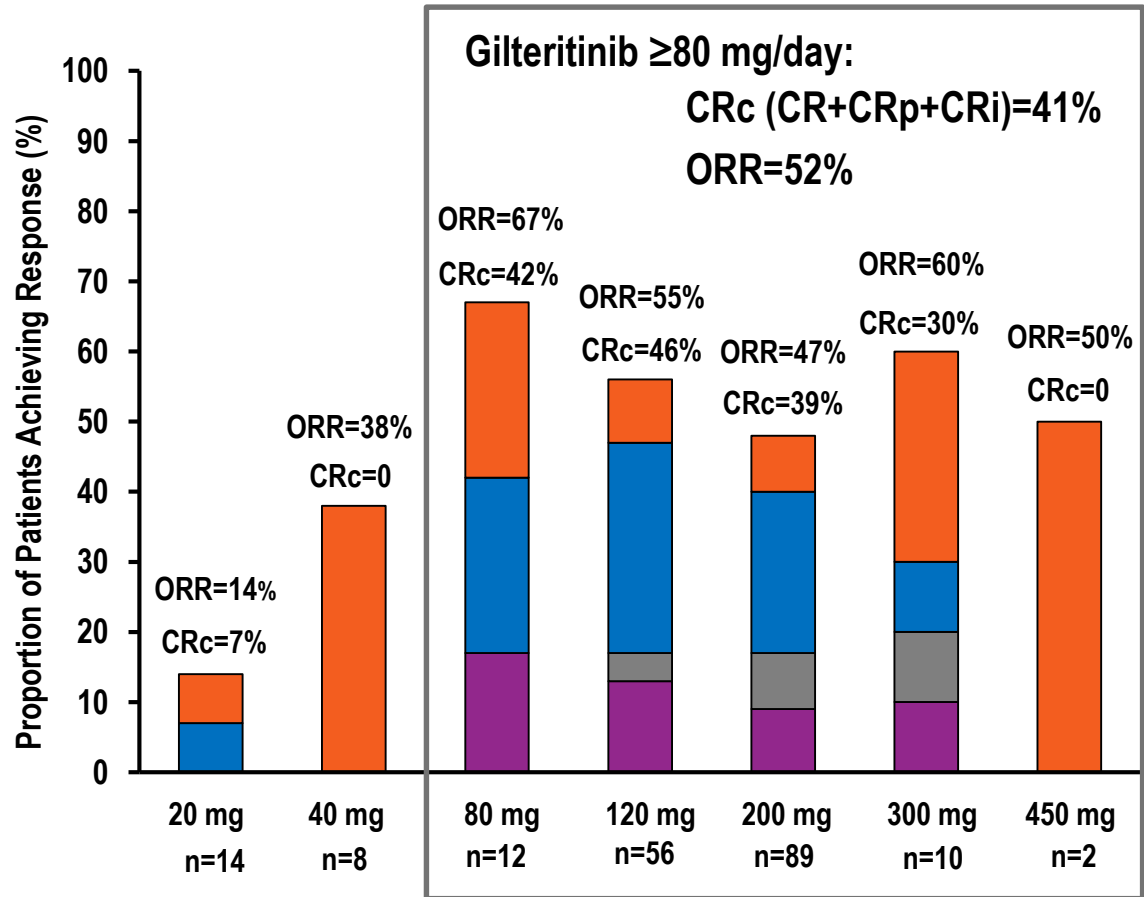
<sup>2</sup>Dohner et al. ASH abstr 467, 2017

# Antileukemic Activity of Gilteritinib

Response in FLT3<sup>mut+</sup> and FLT3<sup>WT</sup> Patients (N=249)



Response in FLT3<sup>mut+</sup> Patients by Gilteritinib Dose (N=191)



# Second Generation *FLT3* Inhibitors

- **Gilteritinib: inhibits *FLT3-ITD* and *D835***
  - rando trial vs Midostaurin + induction chemo
  - vs placebo as maint posttransplant (MORPHO)
  - vs chemo in rel/refr (registration)
  - with 7+3 and HiDAC, CRc 90% in *FLT3* pos<sup>1</sup>
- **Quizartinib: most potent *FLT3* inhibitor**
  - rando trial vs placebo + induction chemo (QuANTUM-First)
  - vs salvage chemo in R/R (QuANTUM-R)
  - with AZA or LoDAC in R/R, high ORR<sup>2</sup>
- **Crenolanib: inhibits *FLT3-ITD*, *D835*, *PDGFa* and *b***
  - with induction chemo CR 83%, 72% with 1 cycle<sup>3</sup>

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<sup>1</sup>Pratz et al. ASH, 2017 (abstr 722); <sup>2</sup>Swaminathan et al. ASH, 2017 (abstr 723); <sup>3</sup>Wang et al. ASH, 2017 (abstr 566)

# *FLT3* Mutations in AML

- Frequent in normal cytogenetic AML
  - Associated with high WBC, packed marrow
  - ITD associated with high relapse rate, poor OS; TKD less so
  - Most common in APL, but appears not prognostic
  - Resistance mechanisms include point mutations, high levels of *FLT3* ligand
-

# Minimal Residual Disease

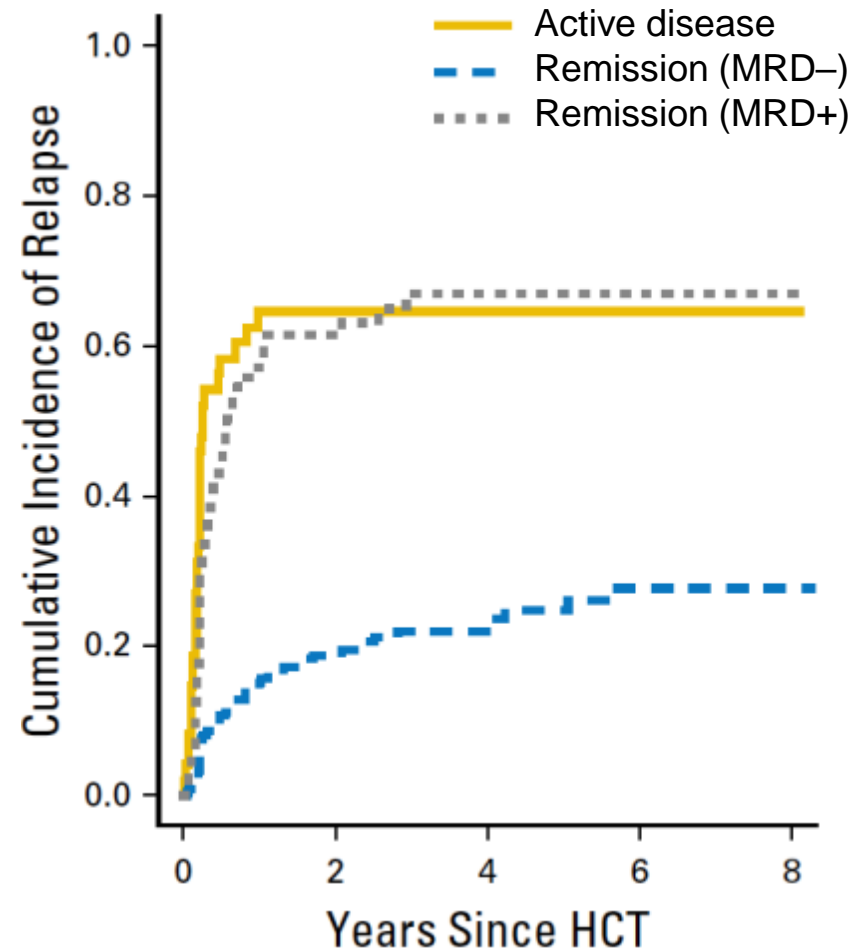
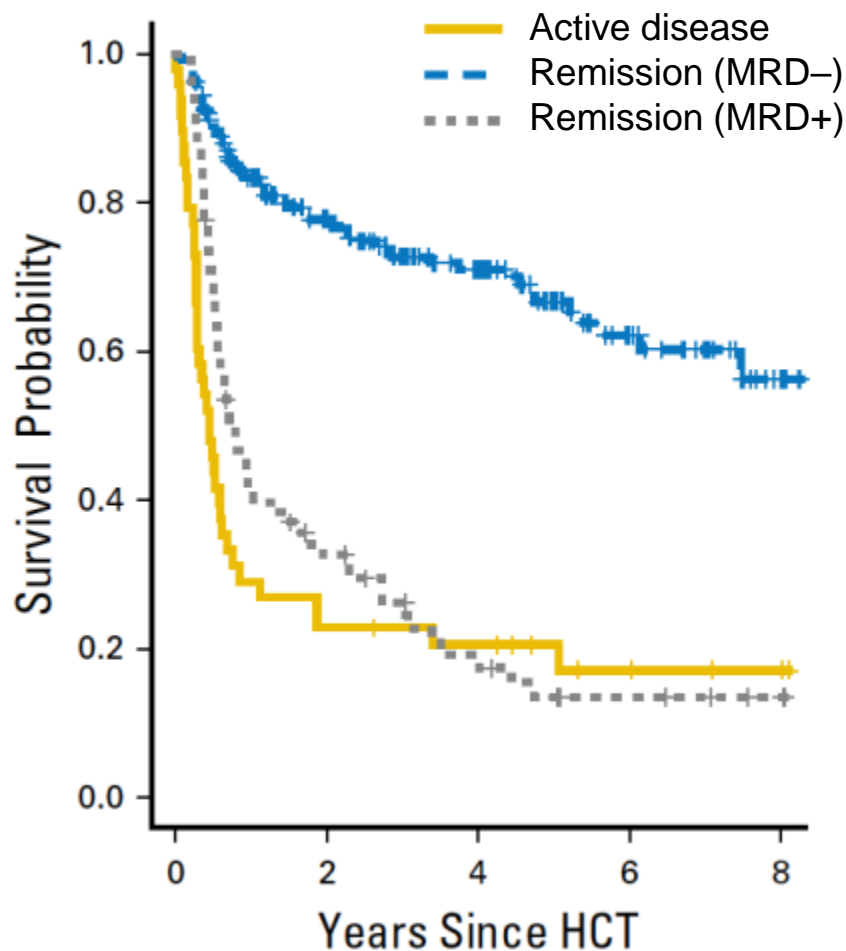
- Detected by increasingly sensitive techniques (immunophenotyping, PCR, sequencing)
- Most studied in patients with *NPM1* mutation and CBF AMLs<sup>1,2</sup>
- Persistence of somatic mutations with VAF>1% in CR assoc. with increased risk of death and relapse<sup>3</sup>
- Complex due to genetic heterogeneity and multiple subclones
- Has prognostic implications following chemotherapy and before allogeneic transplantation
- Will rapidly become incorporated in routine clinical practice

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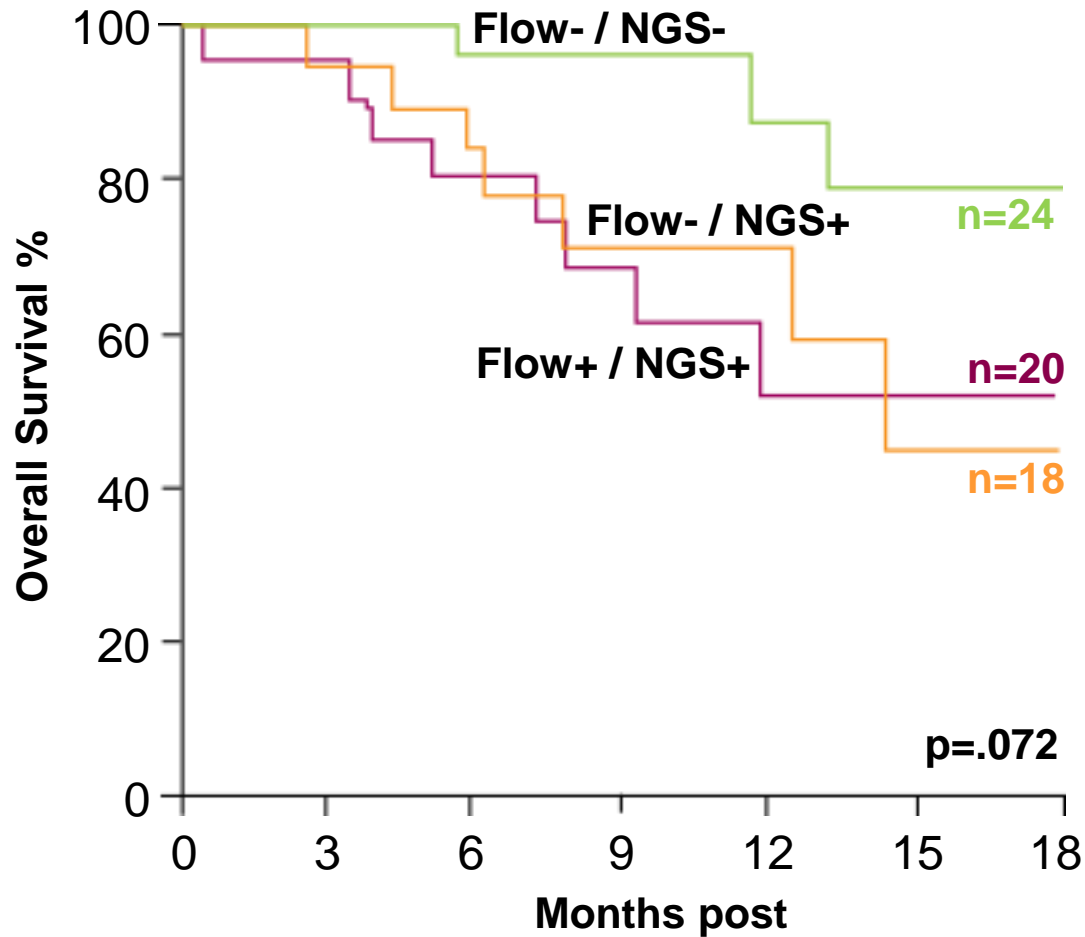
<sup>1</sup>Ivey et al. *NEJM*, 2016; <sup>2</sup>Kapp-Schworer et al. *ASH abstr* 183, 2017; <sup>3</sup>Morita et al. *ASH abstr* 2667, 2017



# Equivalent Post-Transplant Outcomes for Pre-transplant AML MRD (by FC) and Active AML



# Flow Cytometry and NGS in AML Assessment Pre-allogeneic Transplant



# Limitations of MRD Detection in AML

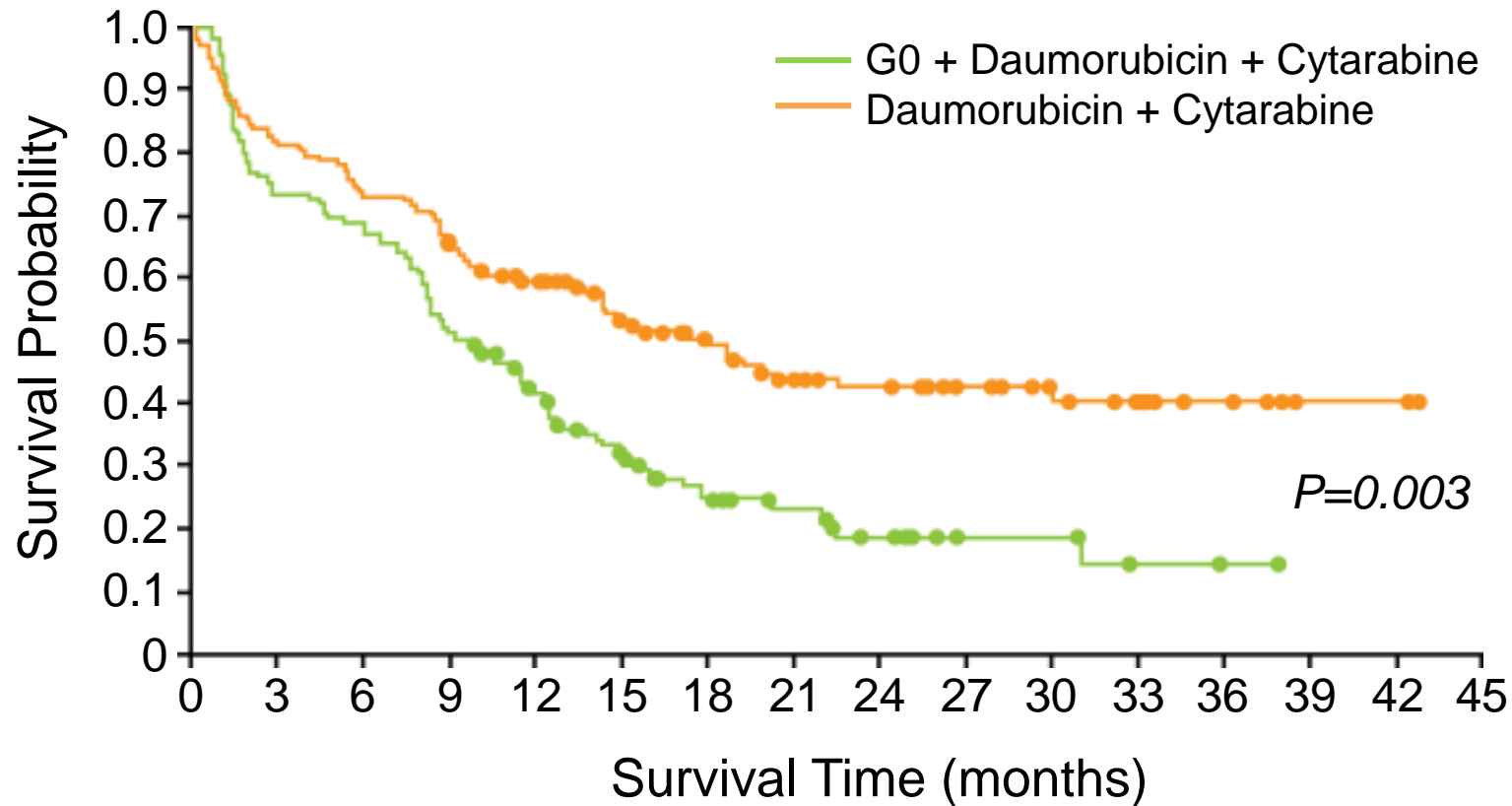
- Methodologies not standardized
  - Thresholds for defining MRD vary
  - Heterogeneity of the disease
  - Clonal hematopoiesis
  - Variable distribution of leukemia cells after treatment
  - Lack of effective agents to target MRD
  - Randomized trials needed to show benefit of intervention
-

# Agents With Regulatory Approval (or Breakthrough Designation)

Agent	Target	Population	Setting
Midostaurin	<i>FLT3</i>	<i>FLT3-ITD</i> or <i>TKD</i>	Treatment naïve w chemo in induc and consol
Gemtuzumab ozogamicin	<i>CD33</i>	CBF and possibly intermed-risk	Treatment naïve CD33+ adults w chemo or single agent or Rel/refr adults and peds
CPX-351	Cytotoxic	t-AML or AML with MRC	Treatment naïve with t-AML or AML with MRC
Enasidenib	<i>IDH2</i>	IDH mutated	Rel/refr AML w m <i>IDH2</i>
Venetoxlax	<i>BCL-2</i>	Elderly adults	Treatment naïve (with LoDAC)

# Gemtuzumab Ozogamicin (Fractionated) in Newly Diagnosed AML Ages 50-70

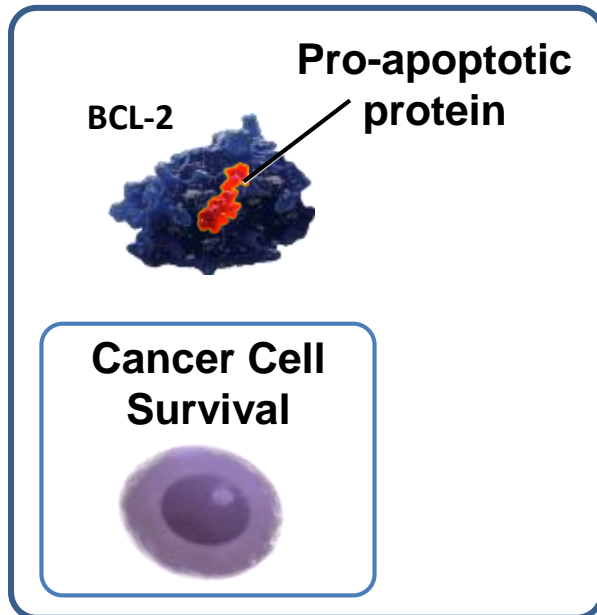
Kaplan-Meier Plot of Event-Free Survival (mITT Population) ALFA-0701 Trial



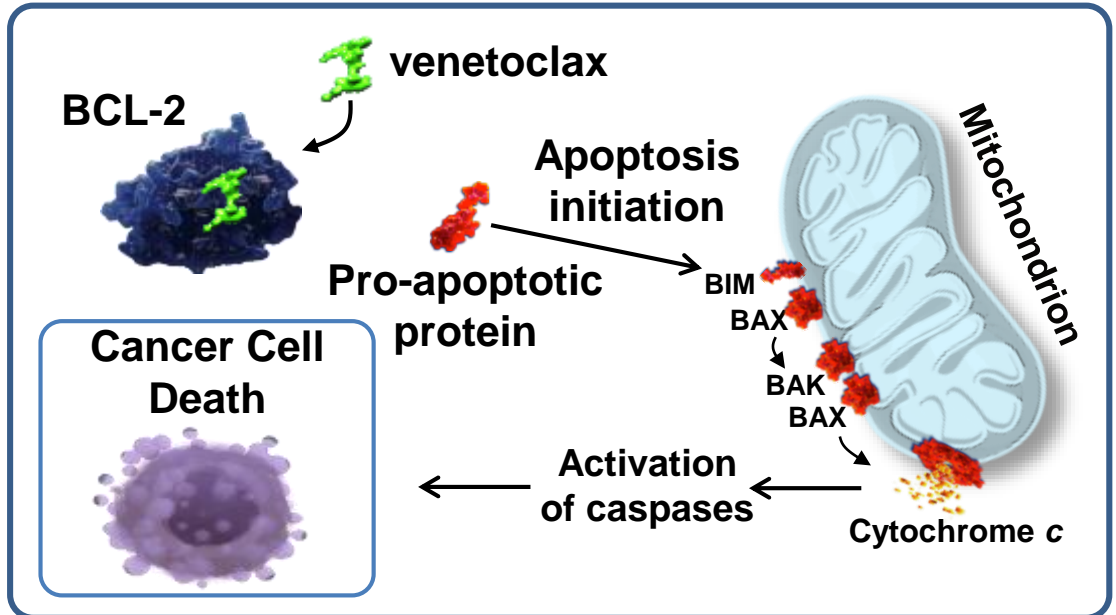
# Gemtuzumab Ozogamicin: Reapproved

- First ab-drug conjugate approved for human use-2000
- Withdrawn, lack of OS benefit and toxicity-2010
- Reapproved for adults with new AML and pts > age 2 with R/R disease-2017
- CD33 single nucleotide polymorphism rs121459419 C→T may be biomarker for response
- OS benefit in fav-risk and trend in intermed-risk
- Risk of SOS/VOD 8% in 146 pts (69 with prophylaxis: heparin or ursodiol or defibrotide) after allograft
- Expression of CD33 blast expression impacts outcome

# Venetoclax: Promotes Apoptosis Through Selective Inhibition of BCL-2



**BCL-2 overexpression allows cancer cells to evade apoptosis by sequestering pro-apoptotic proteins.<sup>1-3</sup>**



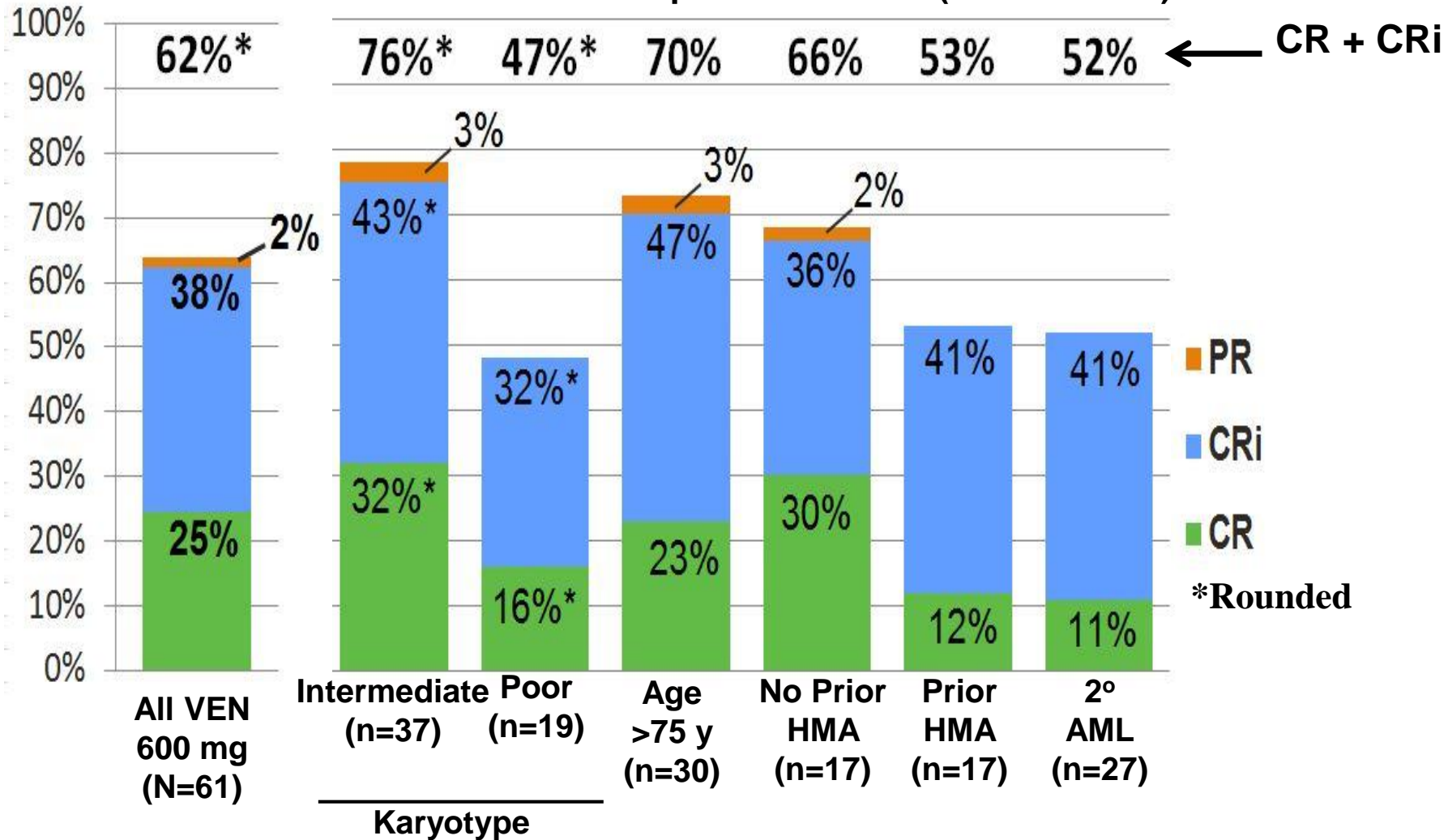
**Venetoclax binds selectively to BCL-2, freeing pro-apoptotic proteins that initiate programmed cell death (apoptosis).<sup>4-6</sup>**

1. Levenson et al. *Sci Transl Med* 2015; 2. Czabotar, et al. *Nature Reviews* 2014; 3. Plati et al. *Integr Biol (Camb)* 2011; 4. Certo et al. *Cancer Cell*. 2006; 5. Souers et al. *Nat Med*. 2013; 6. Del Gaizo Moore V et al. *J Clin Invest*. 2007

# CR/CrI Rates

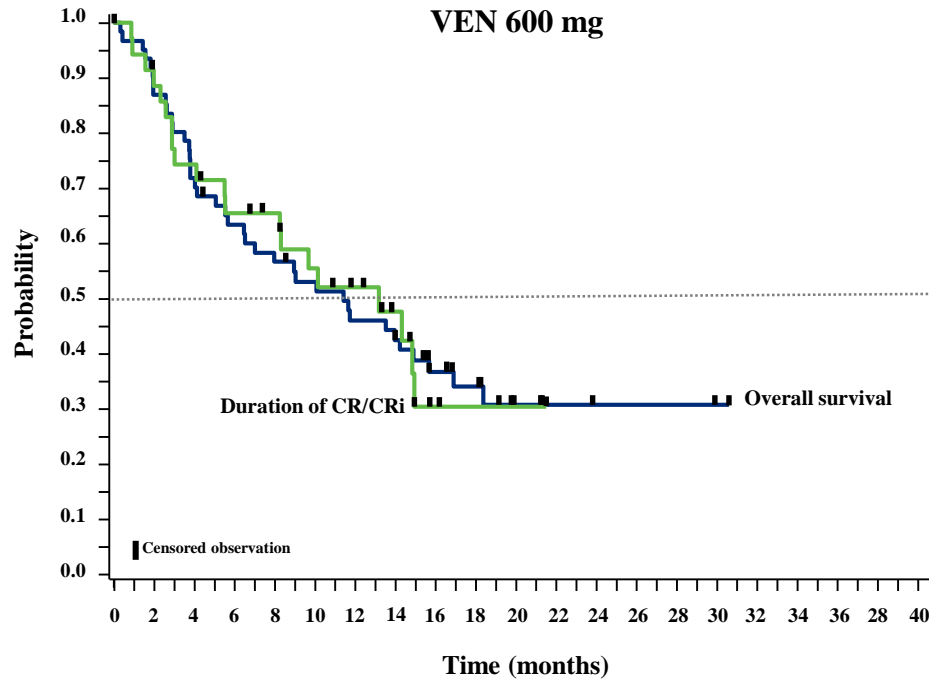
## LoDAC + Venetoclax

Median time to response: 1 month (<1–9 months)



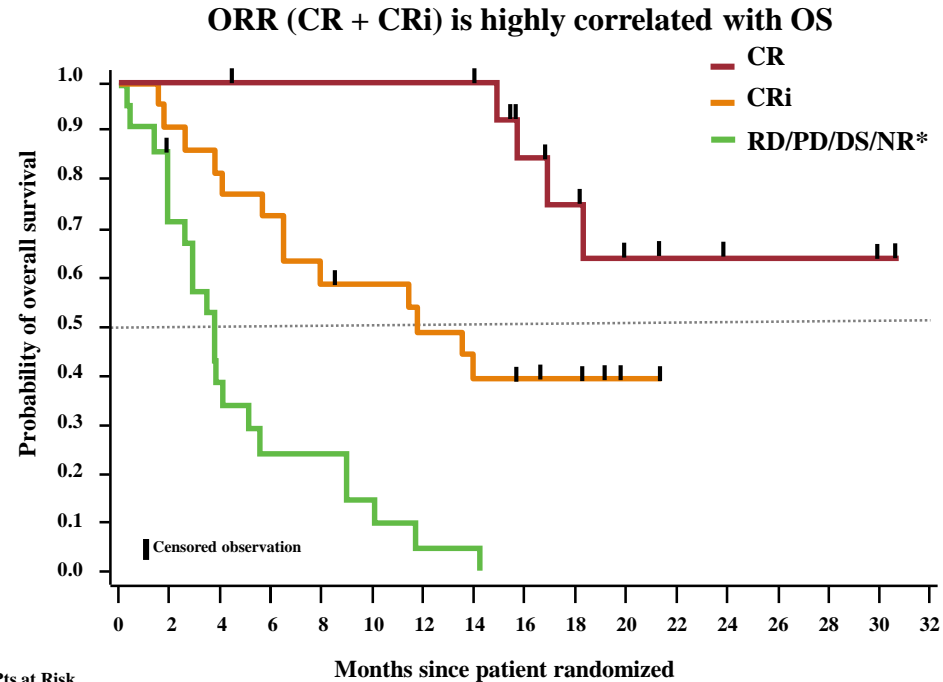


# DOR, Survival, and Survival by Response



Pts at Risk

	38	33	32	27	26	24	22	21	20	17	16	14	13	12	9	4	3	2	2	2	1	1
	61	52	43	37	33	30	26	24	16	13	6	3	2	2	2	2	1					



Pts at Risk

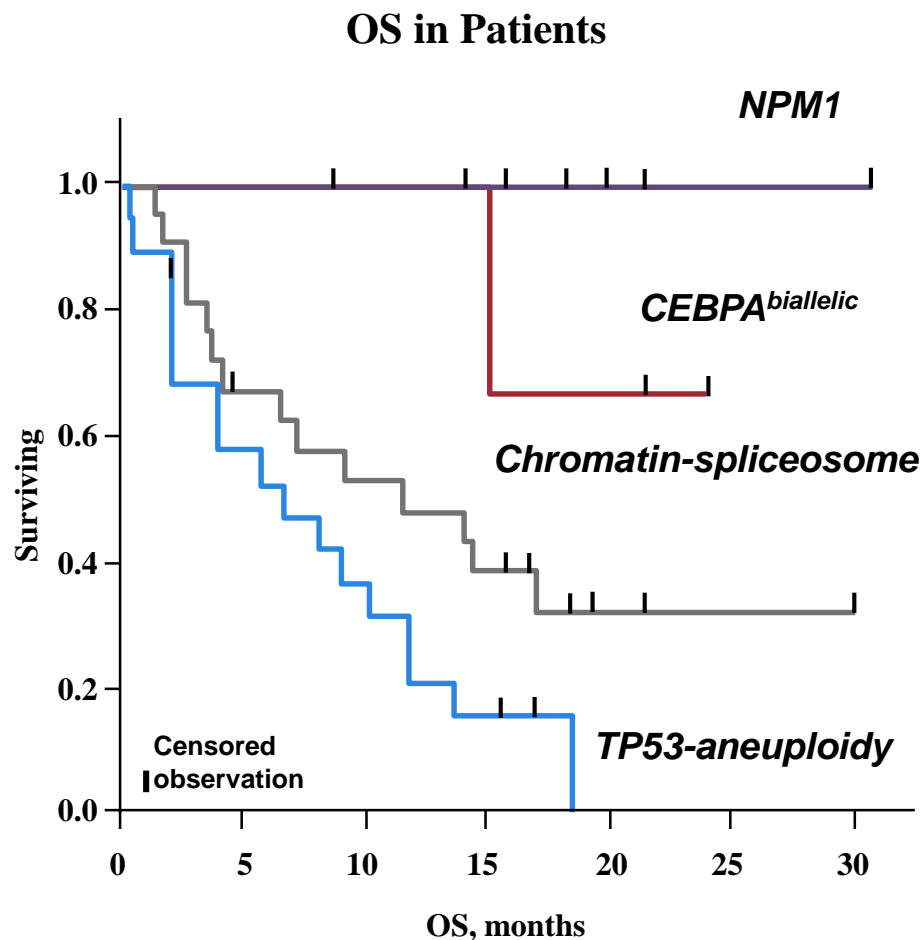
		16	16	16	15	15	15	15	15	10	8	5	3	2	2	2	1	0
CR	16	16	16	15	15	15	15	15	15	10	8	5	3	2	2	1	0	
CRi	22	20	18	16	13	12	10	8	6	5	1	0						
RD/PD/DS/NR	22	15	8	5	5	3	1	1	0									

# Outcomes According to Molecular Drivers of AML

Cytogenetics	ORR (CR + CRi)	Median OS, mo
Intermediate risk n = 37	28 (76%)	15.7
Adverse risk n = 19	9 (47%)	5.7

## Molecular Subgroups

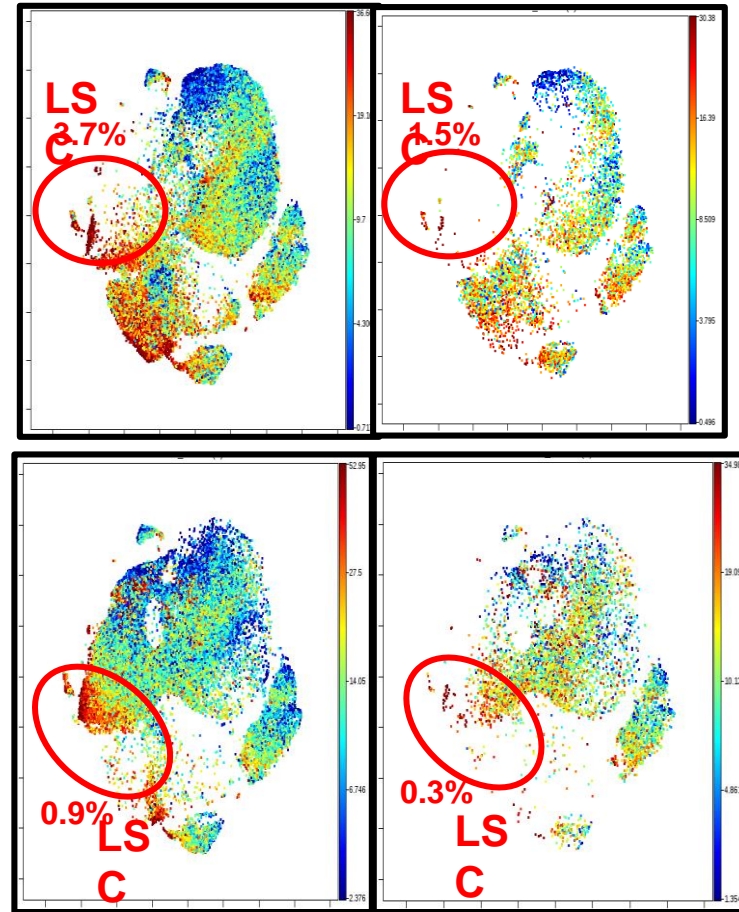
<b>NPM1</b> n = 7*	7 (100%)	NR
<b>CEBPA<sup>biallelic</sup></b> n = 3	3 (100%)	NR
<b>Chromatin-spliceosome</b> n = 22	15 (68%)	11.4
<b>TP53-aneuploidy</b> n = 20	10 (50%)	6.5



# Venetoclax and Azacitidine Results in Rapid Eradication of Blasts and LSCs

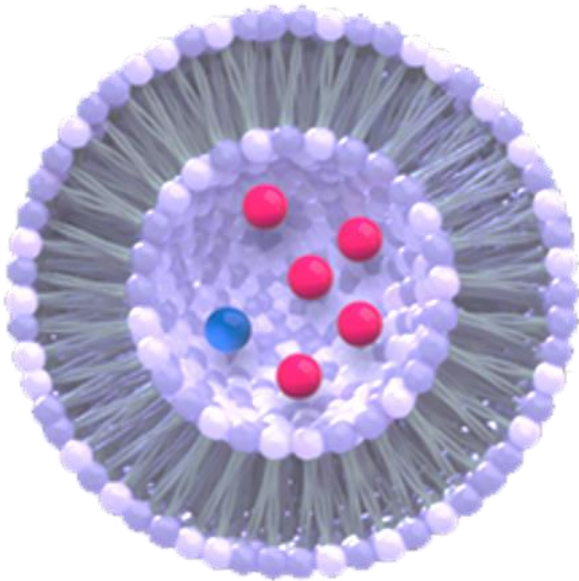
## Peripheral Blood Blasts (%)

	Pre- Treatment	24 Hours Post- Treatment	72 Hours Post- Treatment
Pt 1	71%	50%	16%
Pt 2	81%	72%	34%



LSCs defined as Lin-/CD34+/CD123+/HLA-DR+/CD117+/CD33

# CPX-351 Uses a Nano-Scale Delivery Complex

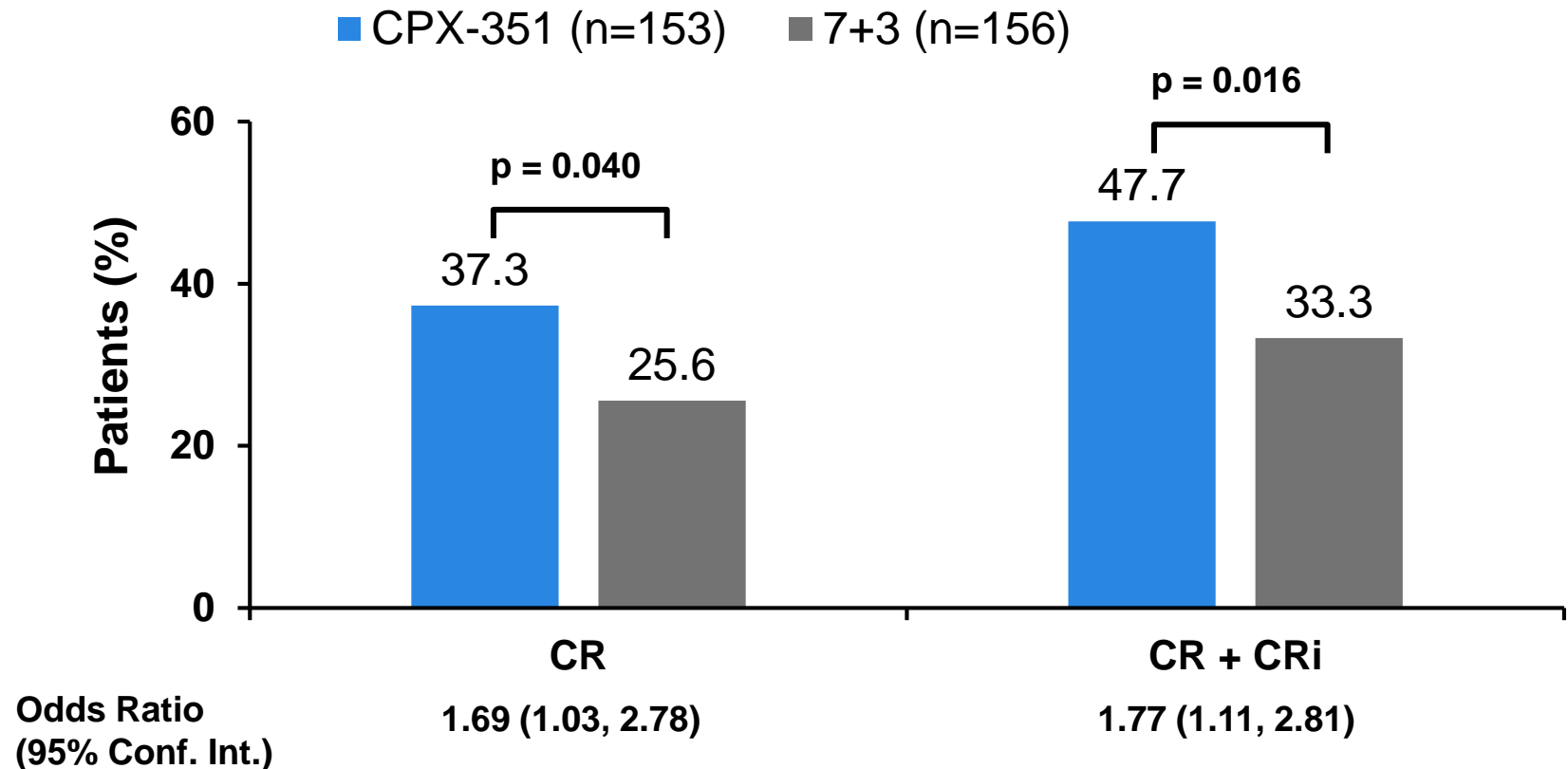


- 100 nm bilamellar liposomes
- 5:1 molar ratio of cytarabine to daunorubicin
- 1 unit = 1.0 mg cytarabine plus 0.44 mg daunorubicin

US FDA Approved August 2017 for t-AML and AML with MRC

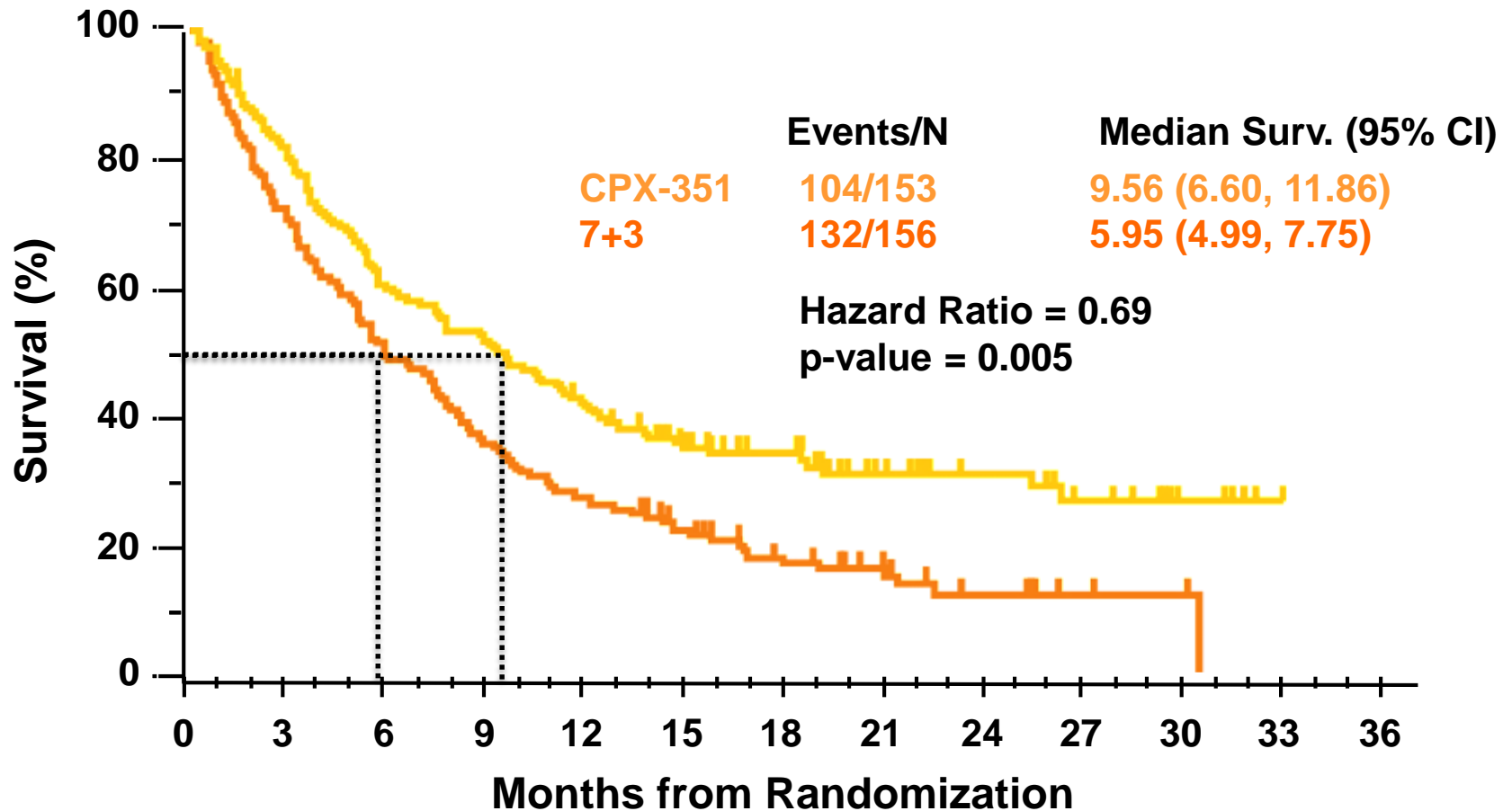
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# Patients Treated With CPX-351 Exhibited Statistically Significant Improvements in Response Rate

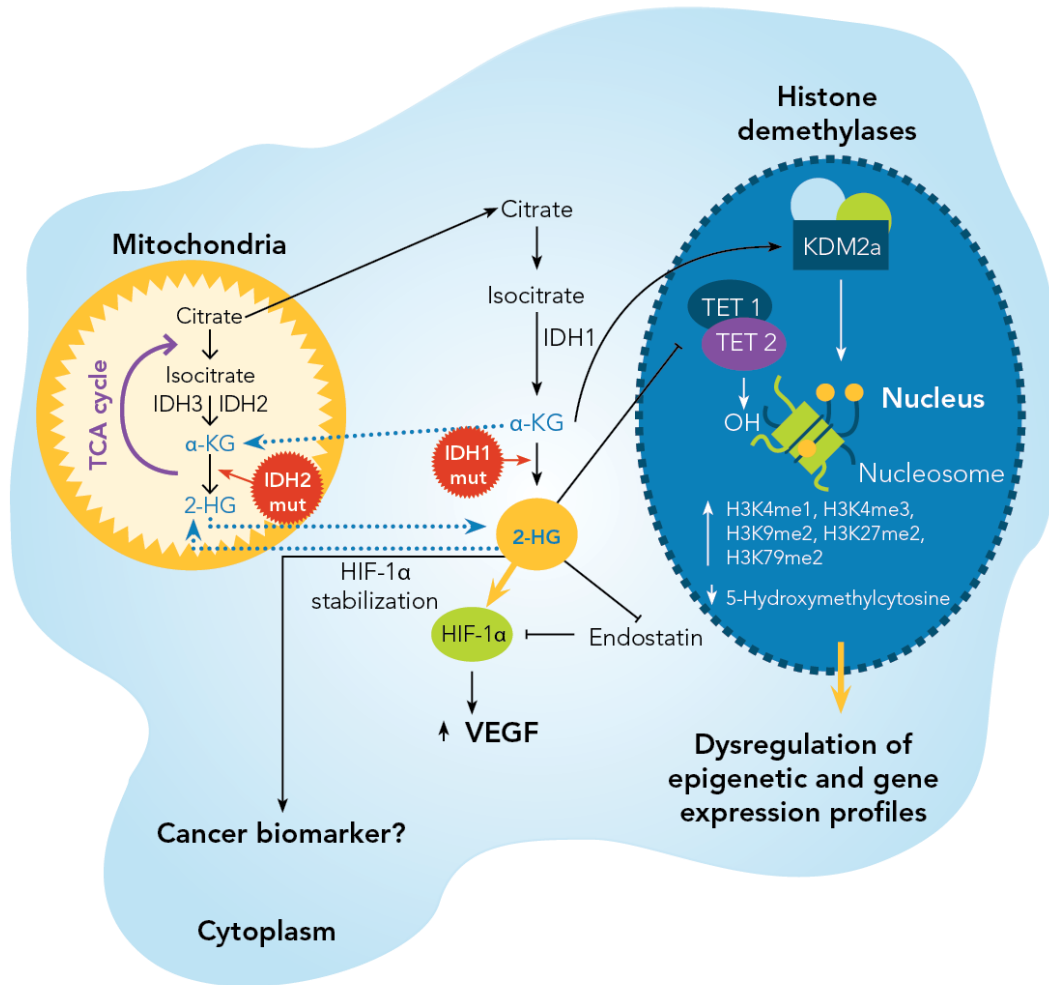


# Overall Survival Was Greater in the CPX-351 Arm Compared to the 7+3 Arm

Kaplan-Meier Curve for Overall Survival ITT Analysis Population

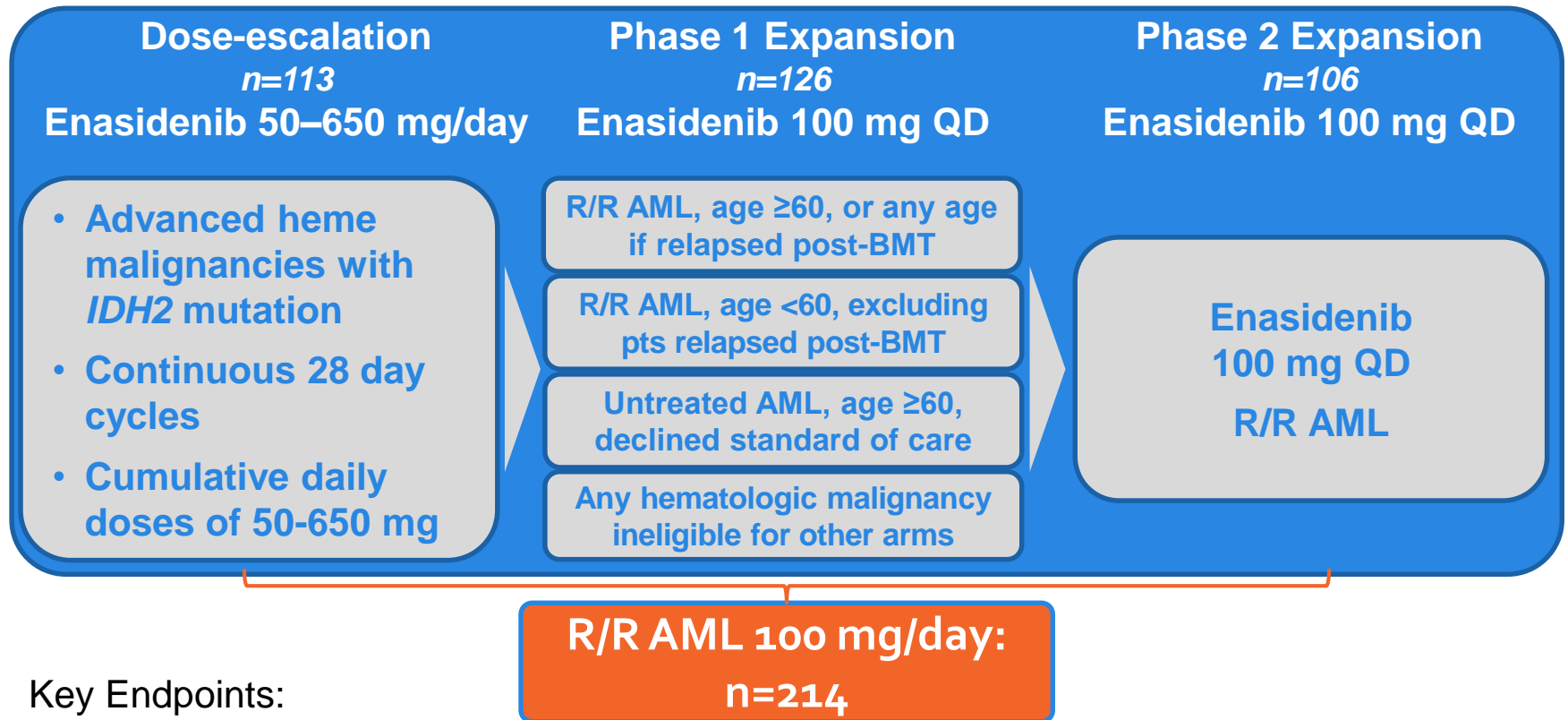


# Role of *IDH* in Malignancy



- IDH is critical metabolic enzyme in the citric acid cycle
- IDH1 in cytoplasm and IDH2 in mitochondria
- Cancer-associated IDHm produces 2-hydroxyglutarate (2-HG) and blocks normal cellular differentiation

# Phase 1/2 Study Design



## Key Endpoints:

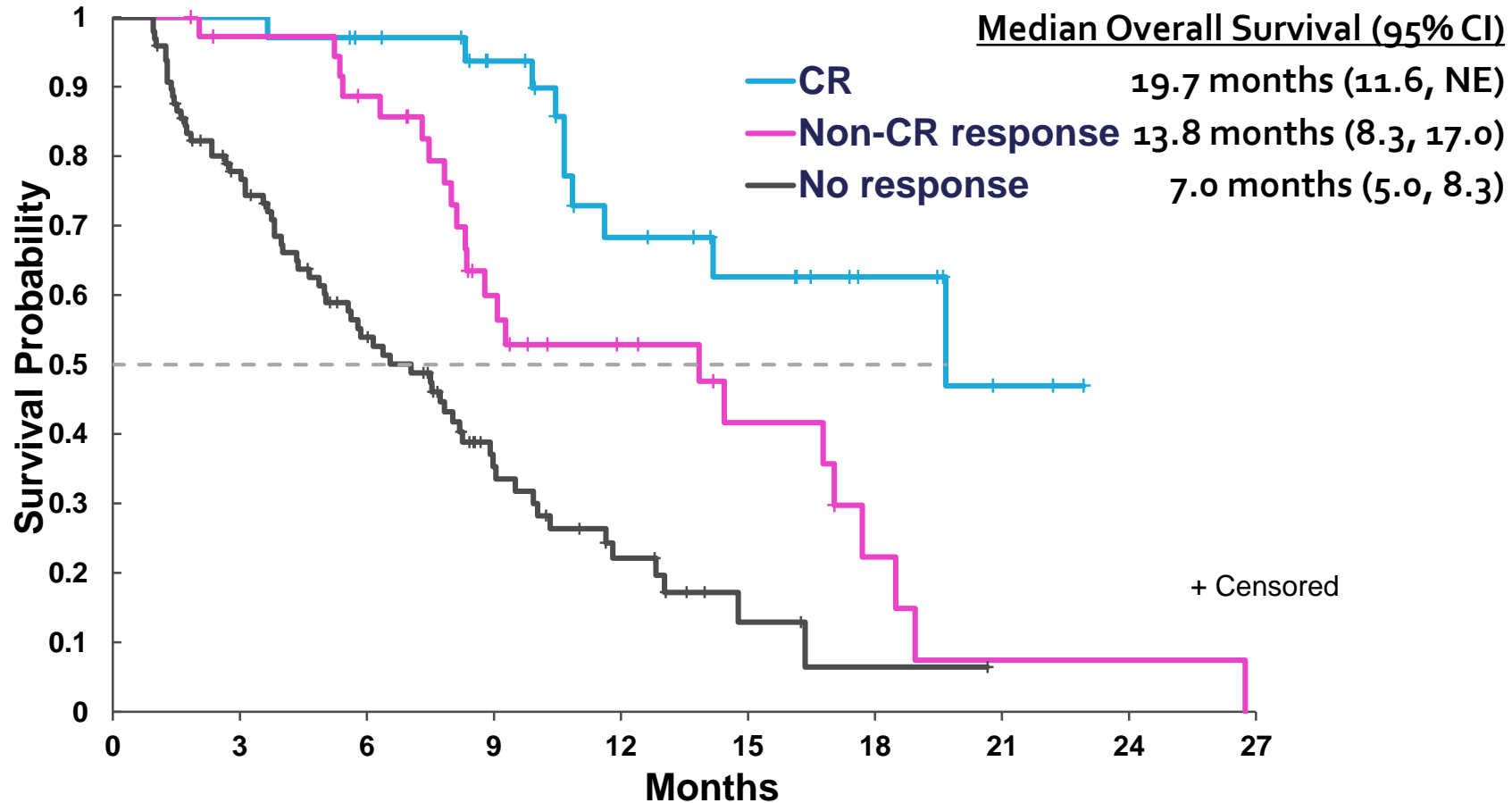
- Safety, tolerability, MTD, DLTs
  - MTD not reached at doses up to 650 mg/day
- Responses assessed by local investigator per IWG criteria<sup>1</sup>
- Assessment of clinical activity, with focus on 100-mg daily dose in patients with R/R AML



# Response in R/R AML

	Relapsed/Refractory AML	
	Enasidenib 100 mg/day (n=214)	All doses (N=281)
<b>Overall response rate, % [n/N]</b> [95% CI]	<b>37% (79/214)</b> [30.4, 43.8]	<b>38% (108/281)</b> [32.7, 44.4]
<b>Best response</b>		
CR, n (%) [95% CI]	<b>43 (20.1)</b> [14.9, 26.1]	<b>55 (19.6)</b> [15.1, 24.7]
CRi or CRp, n (%)	17 (7.9)	22 (7.8)
PR, n (%)	8 (3.7)	16 (5.7)
MLFS, n (%)	11 (5.1)	15 (5.3)
SD, n (%)	110 (51.4)	137 (48.8)
PD, n (%)	11 (5.1)	15 (5.3)
NE, n (%)	2 (0.9)	3 (1.1)
<b>Time to first response (mos), median (range)</b>	1.9 (0.5–11.1)	1.9 (0.5-11.1)
<b>Duration of response (mos), median [95%CI]</b>	5.6 [4.6, 7.4]	5.6 [4.6, 6.5]
<b>Time to CR (mos), median (range)</b>	3.7 (0.7–11.2)	3.8 (0.5-11.2)
<b>Duration of response in pts with CR (mos), median [95%CI]</b>	8.8 [5.6, NR]	7.4 [6.4, 14.7]

# Overall Survival by Best Response



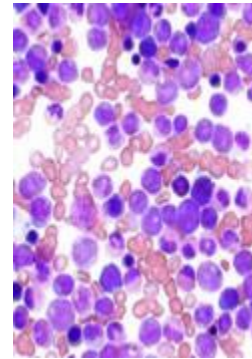
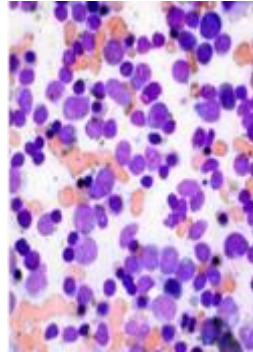
## Morphological evidence of myeloid differentiation

### Patient 1

Screening  
37% blasts

Cycle 1 Day 15  
Evidence of  
cellular  
differentiation

Cycle 3 Day 1  
4% blasts



## FISH evidence of myeloid differentiation

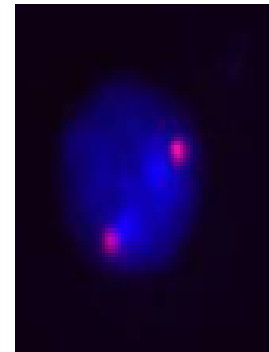
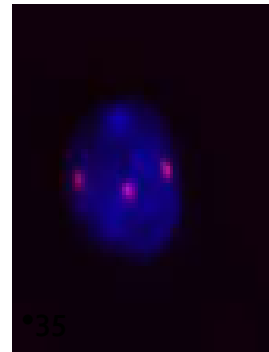
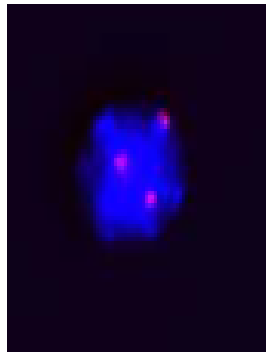
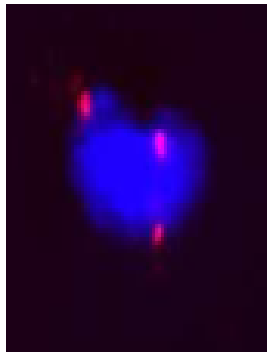
### Patient 2 C2D1, trisomy 8

Blasts

Promyelocytes

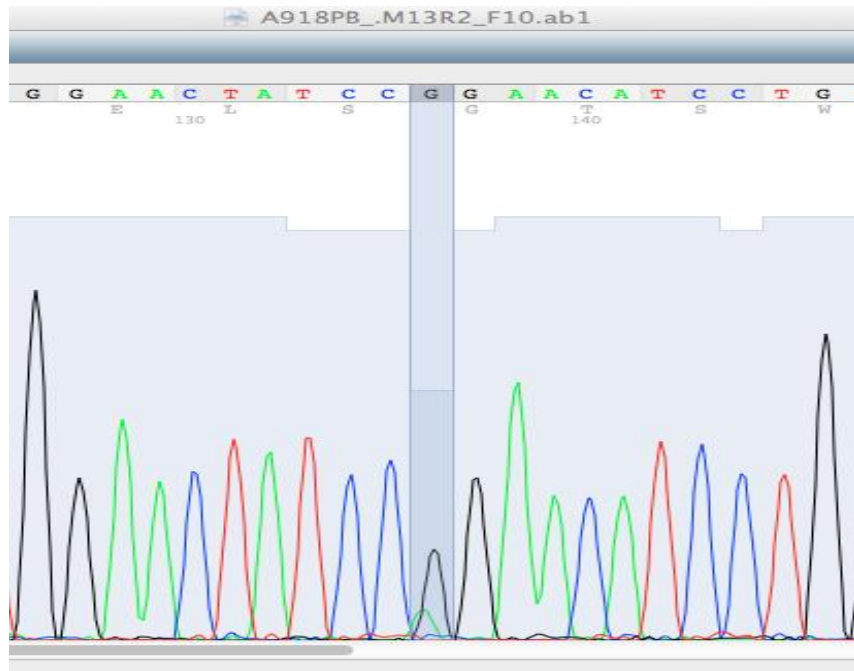
Mature  
Granulocyte

Lymphocytes

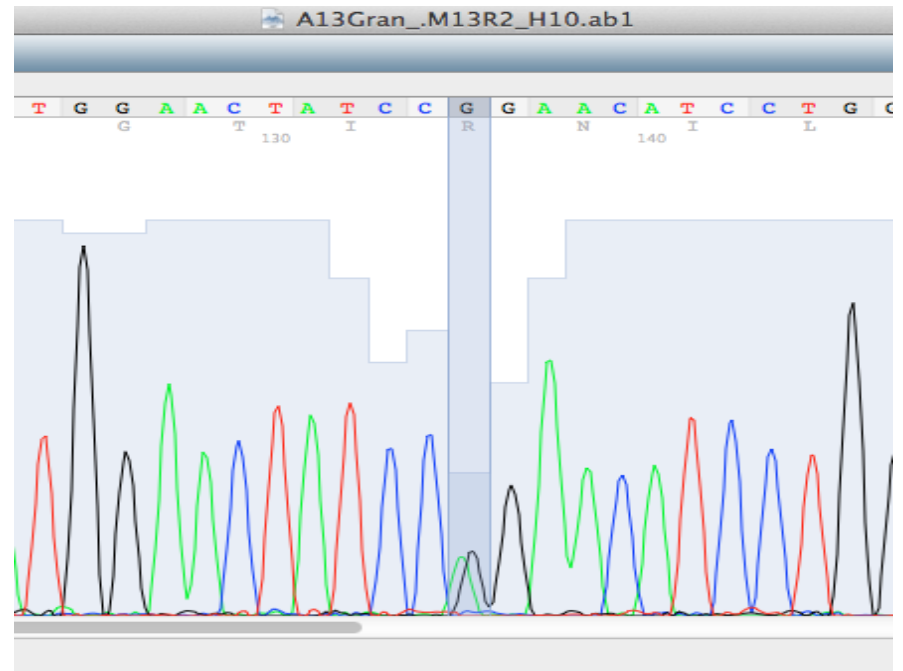


# Molecular Evidence of Differentiation

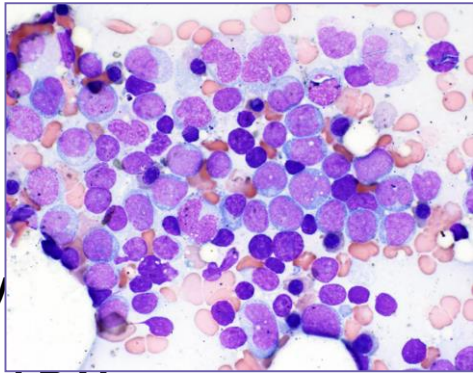
Screening – PBMC



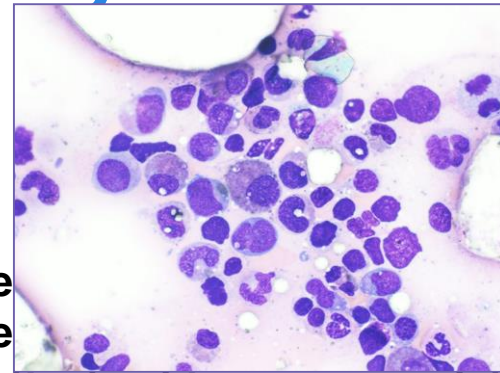
Cycle 3 day 1 – Remission - Granulocytes



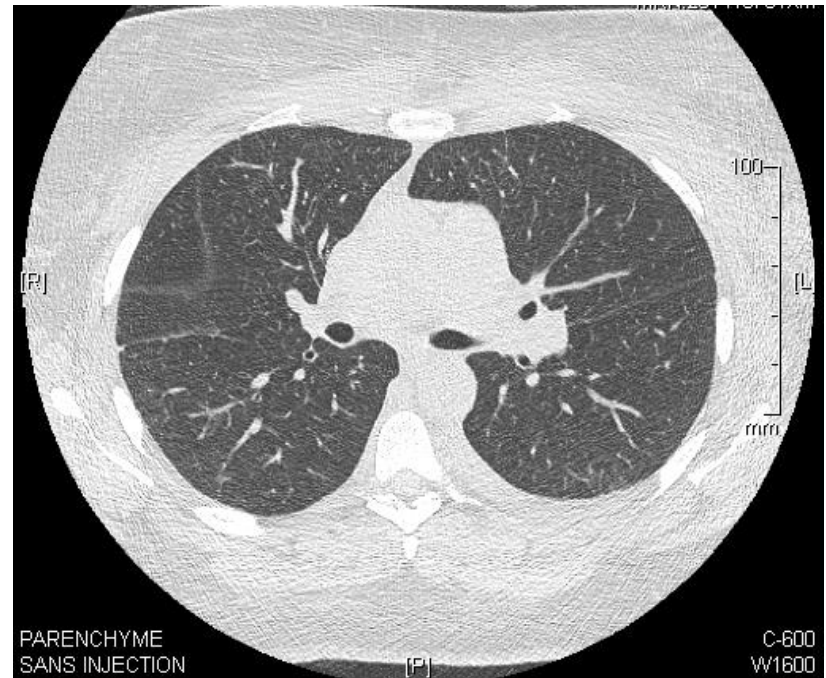
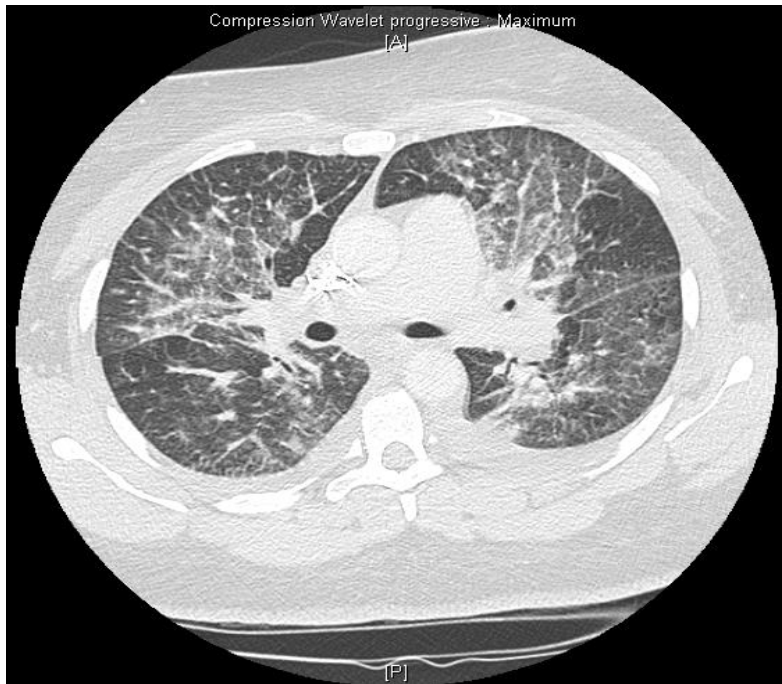
# Differentiation Syndrome



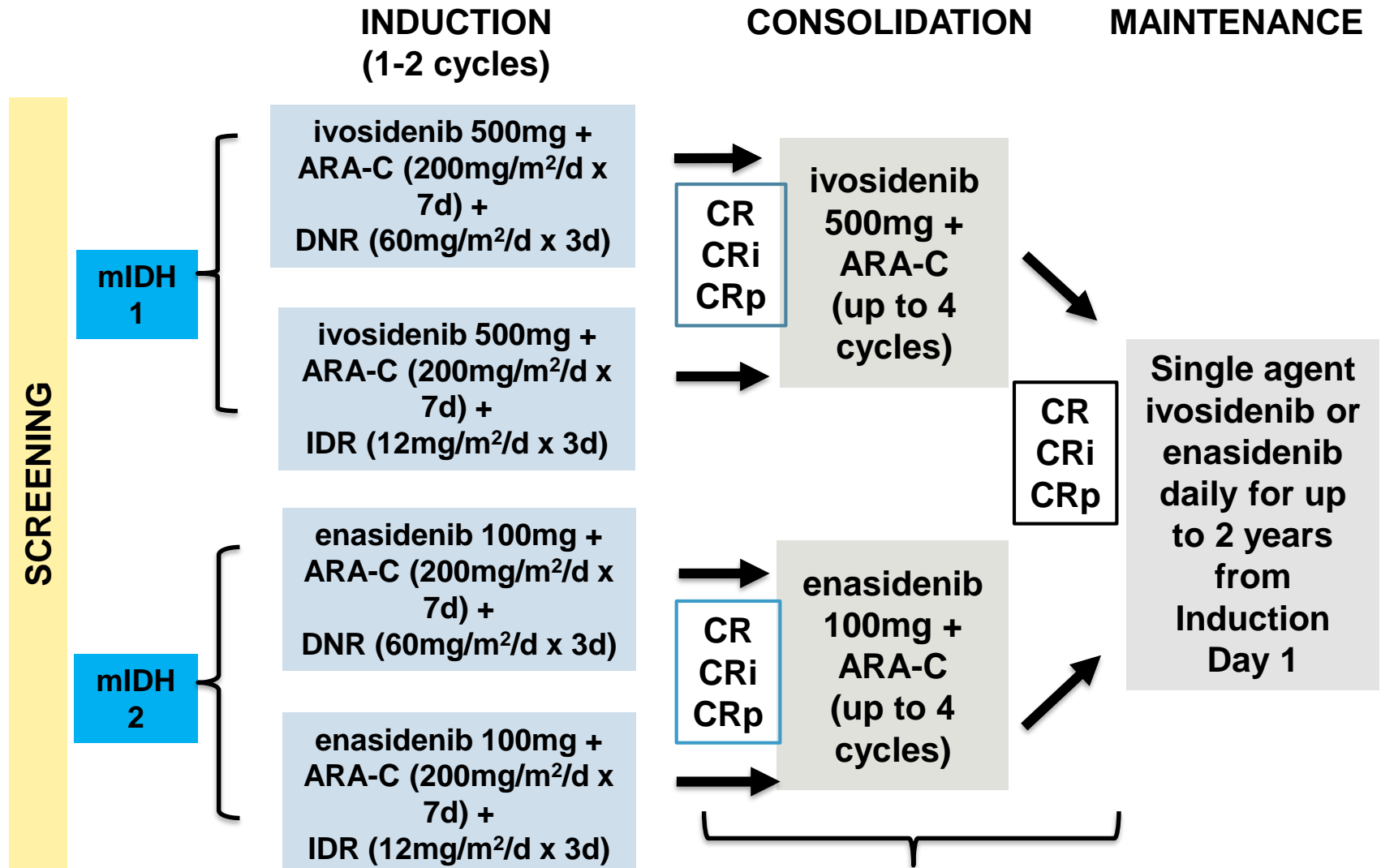
- 21 day
  - Fever,
  - Normal BAL
- daily



- De
  - Re
  - Patient achieves a complete remission
- for 15 days  
toms



# Ivosidenib or Enasidenib Plus Chemotherapy Phase I Trial



# Best Overall Response Summary

Response, n (%)	Ivosidenib (AG-120) + CT			Enasidenib (AG-221) + CT		
	All (n=30)	De novo (n=21)	sAML (n=9)	All (n=50)	De novo (n=27)	sAML (n=23)
CR+CRi/CRp	77	91	44	62	67	57
CR	63	71	44	50	59	39
CRi/CRp	13	19	-	12	7	17
MLFS	3	-	11	20	15	26
PR	7	5	11	-	-	-
Persistent disease	7	5	11	10	7	13
NE	7	-	22	8	11	4



# Novel Agents in AML

Agent	Target
Selinexor	XPO1
Tamibarotene	<i>RAR-alpha</i>
Entospletinib	<i>SYK</i>
Palbociclib	CDK6
Cobimetinib	<i>MAPK</i>
Pevonedistat	NEDD8-activating enzyme
H3B-8800/E7070, E7820 (Spliceosome inhibitors)	SF3B1/RBM39

*Daver et al. ASH abstr 1344, 2017; Drusbosky et al. ASH abstr 3909, 2017; Daver et al. ASH abstr 813, 2017; Guo et al. ASH abstr 3820, 2017; Yoshimi et al. ASH abstr 473, 2017*



# Questions Generated From New Drug Approvals

- Should Gemtuzumab be given to all CBF AMLs and older adults with fav- and intermed-risk?
  - How should transplant strategies be affected by Gemtuzumab in induction?
  - Must Gemtuzumab be given as in ALFA trial with specific induction and chemotherapy regimens (dauno in consol)? For Midostaurin?
  - When a pt has AML with MRC and an *IDH2* mutation, should pt be treated with CPX-351 or on trial with chemotherapy and Enasidenib? If AML-MRC and FLT3 pos: CPX-351 or Mido?
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# AML Treatment Strategies in 2018

AML subgroup	Candidate for intensive chemo	Not a candidate for intensive chemo
All patients	Clinical trial preferred	Clinical trial preferred
CBF	GO + chemo	HMA/LoDAC + Venetoclax*
CD33 pos	GO + chemo, ? If pretransplant	GO or HMA/LoDAC + Venetoclax
t-AML or AML w/MRC (incl complex cyto)	CPX-351 ind/consol, transplant	HMA/LoDAC + Venetoclax*
<i>TP53</i> mutant	Chemo vs decitabine x 10d	Decitabine x5d or x10d
<i>FLT3</i> +	Mido + chemo ind/consol/maint, transplant	?AZA + sorafenib or HMA alone
<i>IDH1/2</i> +	Chemo	HMA/LoDAC + Venetoclax*
Marker -	Chemo	HMA/LoDAC + Venetoclax*

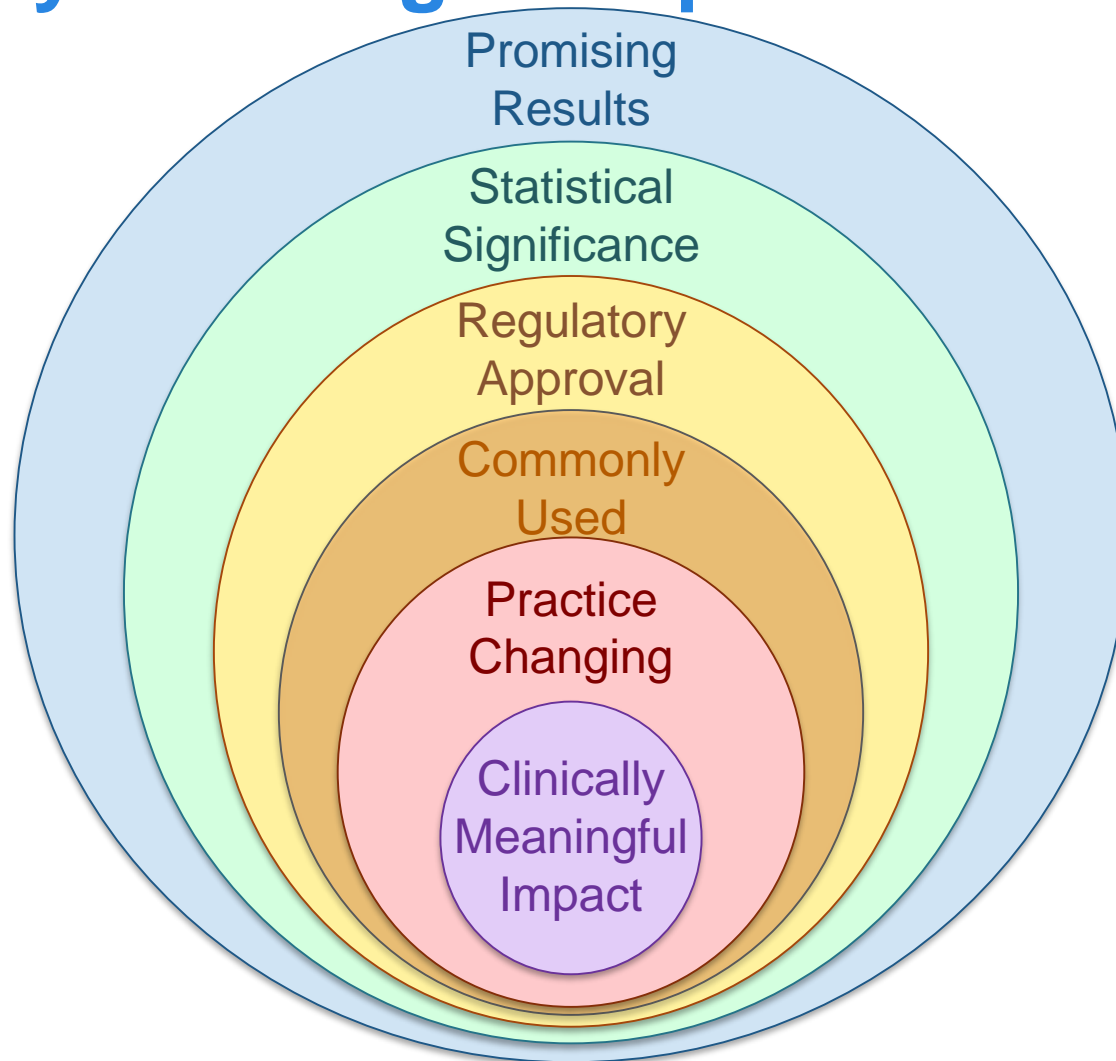
\*HMA/LoDAC + Venetoclax awaiting phase III data

# AML Treatment Strategies in 2018: R/R

AML subgroup	Candidate for intensive chemo	Not a candidate for intensive chemo
All patients	Clinical trial preferred	Clinical trial preferred
R/R <i>IDH2+</i>	Enasidenib	Enasidenib
R/R <i>IDH1+</i>	Clinical trial with ivosidenib preferred	Clinical trial with ivosidenib preferred
R/R <i>FLT3+</i>	Strongly favor clinical trial	Strongly favor clinical trial
R/R <i>TP53</i> mutant	Chemo vs decitabine x 10d	Decitabine x5d or x10d
R/R CD33+	Chemo or GO	HMA/LoDAC + Venetoclax* or GO
R/R post-allo transplant w extramedullary AML	Chemo vs HMA vs ipilimumab	HMA vs ipilimumab
R/R marker -	Chemo vs HMA vs HMA/LoDAC + Venetoclax*	HMA vs HMA/LoDAC + Venetoclax*

\*Lower RR for HMA/LoDAC + Venetoclax in R/R setting  
 (Dinardo et al. Am J Hematol 2018; Goldberg et al. ASH 2017, abstr 1353)

# The Circuitous Road To A Clinically Meaningful Impact Of A New Drug



# Summary and Conclusions

- AML is a heterogeneous disease of diverse somatic genetic mutations
  - Molecular genetics inform classification, prognosis, therapy and depth of remission
  - Era of precision medicine is here
  - Many novel agents with unique mechanisms of action available
  - MRD has emerged an important prognostic factor
  - Therapeutic paradigms are shifting
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