# MAYO CLINIC









# Acute Myeloid Leukemia in 2024

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- <u>Research Support</u> Astellas, Astex, Celgene, Chordia, Novartis, Actinium, Gilead, H3Biosciences (Roivant/Hemavant), Kura, Sellas, and Takeda
- <u>Advisory Board</u> Intrinsiq (AmeriSouce Bergen), Lava Therapeutics, Targeted Oncology, Treadwell, National Cancer Institute & NHLBI
- <u>Off-Label Use</u> I<sup>131</sup>-Apamistamab & AlloHCT for rel/ref AML; Magrolimab for TP53<sub>mut</sub> AML



# Happy Groundhog Day!



## Alas, 6 more weeks of winter...





# AML Updates in 2024

- Evolving Treatment Landscape
- Advances with Lower Intensity Therapy
- High quality remission & Measurable Residual Disease
- New Targeted Strategies
  - Mutation-based
  - Radioimmunotherapy
  - Targeting TP53

Menin inhibitors for KMT2A & NPM1 I<sup>131</sup>-Apamistamab

Magralimah Allagonaia Tran

Magrolimab, Allogeneic Transplantation

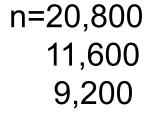


# 2024 AML Estimates in United States

-	Male	1		Fei	male		
Pro	ostate	299,010	29%	Breast	310,720	32%	
Lui	ng & bronchus	116,310	11%	Lung & bronchus	118,270	12%	
Se Co	olon & rectum	81,540	8%	Colon & rectum	71,270	7%	
5 Uri	inary bladder	63,070	6%	Uterine corpus	67,880	7%	Casa
Me Me	elanoma of the skin	59,170	6%	Melanoma of the skin	41,470	4%	Cases
ž Kic	dney & renal pelvis	52,380	5%	Non-Hodgkin lymphom	a 36,030	4%	
-	on-Hodgkin lymphoma	44,590	4%	Pancreas	31,910	3%	Male
E Ora	al cavity & pharynx	41,510	4%	Thyroid	31,520	3%	
E Lei	ukemia	36,450	4%	Kidney & renal pelvis	29,230	3%	• Ferr
Pa	increas	34,530	3%	Leukemia	26,320	3%	
All	sites	1,029,080		All sites	972,060		
	Male			Fei	male		
Lui	ng & bronchus	65,790	20%	Lung & bronchus	59,280	21%	Death
	ostate	35,250	11%	Breast	42,250	15%	
Co	olon & rectum	28,700	9%	Pancreas	24,480	8%	Male
S Pa	increas	27,270	8%	Colon & rectum	24,310	8%	
Liv	ver & intrahepatic bile duct	19,120	6%	Uterine corpus	13,250	5%	Fem
D Lei	ukemia	13,640	4%	Ovary	12,740	4%	
Pai Liv Let Esc Uri No	ophagus	12,880	4%	Liver & intrahepatic bile	duct 10,720	4%	
E Uri	inary bladder	12,290	4%	Leukemia	10,030	3%	
	on-Hodgkin lymphoma	11,780	4%	Non-Hodgkin lymphoma		3%	
Bra	ain & other nervous system	10,690	3%	Brain & other nervous sy	stem 8,070	3%	
All	sites	322,800		All sites	288,920		

Estimates are rounded to the nearest 10, and cases exclude basal cell and squamous cell skin cancers and in situ carcinoma except urinary bladder. Estimates do not include Puerto Rico or other US territories. Ranking is based on modeled projections and may differ from the most recent observed data.

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- ale

n=11,220 6,290 4,930



Siegel et al, CA Cancer J Clin 74:12-49, 2024

# Myeloid Classification: from 2016 to 2023

### Arber DA et al. Blood 2016;127(20): 2391-2405

## The 2016 revision to the World Health Organization classification of myeloid neoplasms and acute leukemia

Daniel A. Arber,<sup>1</sup> Attilio Orazi,<sup>2</sup> Robert Hasserjian,<sup>3</sup> Jürgen Thiele,<sup>4</sup> Michael J. Borowitz,<sup>5</sup> Michelle M. Le Beau,<sup>6</sup> Clara D. Bloomfield,<sup>7</sup> Mario Cazzola,<sup>8</sup> and James W. Vardiman<sup>9</sup>

<sup>1</sup>Department of Pathology, Stanford University, Stanford, CA; <sup>2</sup>Department of Pathology, Weill Cornell Medical College, New York, NY; <sup>3</sup>Department of Pathology, Massachusetts General Hospital, Boston, MA; <sup>4</sup>Institute of Pathology, University of Cologne, Cologne, Germany; <sup>5</sup>Department of Pathology, Johns Hopkins Medical Institutions, Baltimore, MD; <sup>6</sup>Section of Hematology/Oncology, University of Chicago, Chicago, IL; <sup>7</sup>Comprehensive Cancer Center, James Cancer Hospital and Solove Research Institute, The Ohio State University, Columbus, OH; <sup>8</sup>Department of Medicular Medicine, University of Pavia, and Department of Hematology Oncology, Fondazione IRCCS Policlinico San Matteo, Pavia, Italy; and <sup>9</sup>Department of Pathology, University of Chicago, Li

### Arber DA et al. Blood 2022;140(11): 1200-

### International Consensus Classification of Myeloid Neoplasms and Acute Leukemias: integrating morphologic, clinical, and genomic data

Daniel A. Arber,<sup>1</sup> Attilio Orazi,<sup>2</sup> Robert P. Hasserijan,<sup>3</sup> Michael J. Borowitz,<sup>4</sup> Katherine R. Calvo,<sup>5</sup> Hans-Michael Kvasnicka,<sup>6</sup> Sa A. Wang,<sup>7</sup> Adam Bagg,<sup>8</sup> Tiziano Barbui,<sup>9</sup> Susan Branford,<sup>10</sup> Carlos E. Bueso-Ramos,<sup>7</sup> Jorge E. Cortes,<sup>11</sup> Paola Dal Cin,<sup>12</sup> Courtney D. DiNardo,<sup>7</sup> Hervé Dombret,<sup>13</sup> Eric J. Duncavage,<sup>14</sup> Benjamin L. Ebert,<sup>15</sup> Elihu H. Estey,<sup>16</sup> Fabio Facchetti,<sup>17</sup> Kathryn Foucar,<sup>18</sup> Naseema Gangat,<sup>19</sup> Umberto Gianelli,<sup>20</sup> Lucy A. Godley,<sup>1</sup> Nicola Gökbuget,<sup>21</sup> Jason Gotlib,<sup>22</sup> Eva Hellström-Lindberg,<sup>23</sup> Gabriela S. Hobbs,<sup>3</sup> Ronald Hoffman,<sup>24</sup> Elias J. Jabbour,<sup>7</sup> Jean-Jacques Kiladjian,<sup>13</sup> Richard A. Larson,<sup>1</sup> Michelle M. Le Beau,<sup>1</sup> Mignon L.-C. Loh,<sup>25</sup> Bob Löwenberg,<sup>26</sup> Elizabeth Macintyre,<sup>27</sup> Luca Malcovati,<sup>28</sup> Charles G. Mullighan,<sup>29</sup> Charlotte Niemeyer,<sup>30</sup> Olatoyosi M. Odenike,<sup>1</sup> Seishi Ogawa,<sup>31</sup> Alberto Orfao,<sup>32</sup> Elli Papaemmanuil,<sup>33</sup> Francesco Passamonti,<sup>28</sup> Kimmo Porkka,<sup>34</sup> Ching-Hon Pui,<sup>29</sup> Jerald P. Radich,<sup>35</sup> Andreas Reiter,<sup>36</sup> Maria Rozman,<sup>37</sup> Martina Rudelius,<sup>38</sup> Michael R. Savona,<sup>39</sup> Charles A. Schiffer,<sup>40</sup> Annette Schmitt-Graeff,<sup>41</sup> Akiko Shimamura,<sup>15,42</sup> Jorge Sierra,<sup>43</sup> Wendy A. Stock,<sup>1</sup> Richard M. Stone,<sup>15</sup> Martin S. Tallman,<sup>44</sup> Jürgen Thiele,<sup>45</sup> Hwei-Fang Tien,<sup>46</sup> Alexandar Tzankov,<sup>47</sup> Alessandro M. Vannucchi,<sup>48</sup> Paresh Vyas,<sup>49</sup> Andrew H. Wei,<sup>50</sup> Olga K. Weinberg,<sup>51</sup> Agnieszka Wierzbowska,<sup>52</sup> Mario Cazzola,<sup>28</sup> Hartmut Döhner,<sup>53</sup> and Ayalew Tefferi<sup>19</sup>

## WHO Classification of Tumours of Haematopoietic and Lymphoid Tissues

Steven H. Swerdlow, Elias Campo, Nancy Lee Harris, Elaine S. Jaffe, Stefano A. Pileri, Harald Stein, Jürgen Thiele, Daniel A. Arber, Robert P. Hasserjian, Michelle M. Le Beau, Attilio Orazi, Reiner Siebert

### International Agency for Research on Cancer

Lyon, 2017

Khoury JD et al. Leukemia 2022;36(7):1703-1719 The 5th edition of the World Health Organization Classification of Haematolymphoid Tumours: Myeloid and Histiocytic/ Dendritic Neoplasms

Joseph D. Khoury <sup>154</sup>, Eric Solary <sup>258</sup>, Oussama Abla<sup>3</sup>, Yassmine Akkari, <sup>4</sup>, Rita Alaggio<sup>5</sup>, Jane F. Apperley <sup>6</sup>, Rafael Bejar, <sup>7</sup>, Emilio Berti<sup>8</sup>, Lambert Busque <sup>9</sup>, John K. C. Chan<sup>10</sup>, Weina Chen <sup>11</sup>, Xueyan Chen<sup>12</sup>, Wee-Joo Chng<sup>13</sup>, John K. Choi <sup>14</sup>, Isabel Colmenero <sup>15</sup>, Sarah E. Coupland<sup>16</sup>, Nicholas C. P. Cross <sup>17</sup>, Daphne De Jong<sup>18</sup>, M. Tarek Elghetany<sup>19</sup>, Emiko Takahashi <sup>20</sup>, Jean-Francois Emile <sup>21</sup>, Judith Ferry<sup>22</sup>, Linda Fogelstrand<sup>23</sup>, Michaela Fontenay<sup>24</sup>, Ulrich Germing<sup>25</sup>, Sumeet Gujral<sup>26</sup>, Torsten Haferlach <sup>27</sup>, Claire Harrison<sup>78</sup>, Jennelle C. Hodge<sup>29</sup>, Shimin Huo <sup>1</sup>, Joop H. Jansen<sup>30</sup>, Rashmi Kanagal-Shamanna <sup>5</sup>, Hagop M. Kantarjian <sup>31</sup>, Christian P. Kratz <sup>32</sup>, Xiao-Qiu Li<sup>33</sup>, Megan S. Lim<sup>34</sup>, Keith Loeb<sup>35</sup>, Sanam Loghavio <sup>1</sup>, Andrea Marcogliese<sup>19</sup>, Soheil Meshinch<sup>36</sup>, Phillip Michaels<sup>37</sup>, Kikkeri N. Naresh <sup>33</sup>, Yasodha Natkunam <sup>38</sup>, Reza Nejati<sup>39</sup>, German Ott<sup>40</sup>, Eric Padron <sup>41</sup>, Keyur P. Patel<sup>1</sup>, Nikhili Patkar <sup>42</sup>, Jenfifer Picarsic<sup>43</sup>, Uwe Platzbecker <sup>44</sup>, Irene Roberts<sup>45</sup>, Anna Schuh <sup>46</sup>, William Sewell<sup>47</sup>, Cecilia Yeung <sup>35</sup> and Andreas Hochhaus <sup>322</sup>.

## ICC – MDS/AML (10-19% blasts, replaces RAEB-2)



In its final version

To be published as WHO Blue Book 5<sup>th</sup> edition by end of 2023 (current Beta version on IARC website)

# Despite a Wealth of Innovative Options, More Work Needs to Be Done to Expose Patients to Effective Therapy

RW analysis of 629 newly diagnosed AML patients from a comprehensive health system in the Midwest United States, including metropolitan and rural populations (2011-2018)

- 66% of patients aged ≥75 years did not receive any chemotherapy or alternative treatment
- Only 13% of patients had evidence of a genomic report, although it has been used for prognostication for at least the last decade

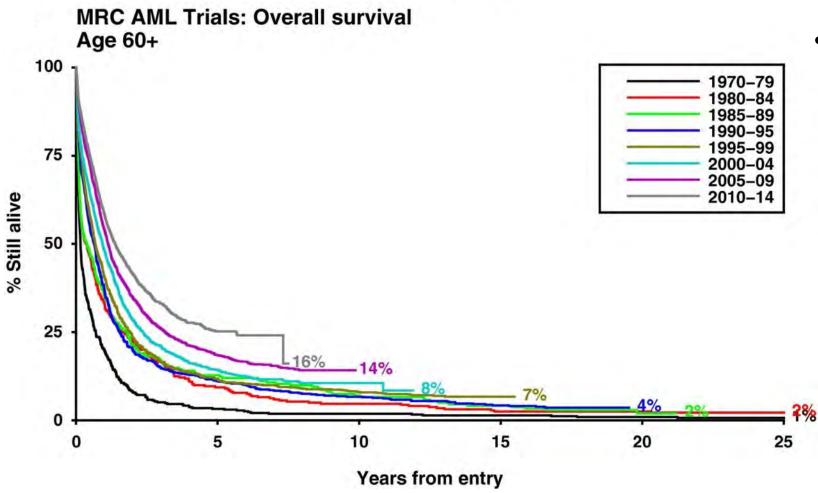
## EMR data from 2,133 AML patients to determine the effect of COVID-19 on AML care

- Compared with the pre-COVID-19 cohort, post-COVID-19 patients were significantly less likely to receive HCT
- Longer HCT waiting times suggest the pandemic affected access to timely transplantation



Byrd JC et al. (ASH 2021). Abstract 3383.

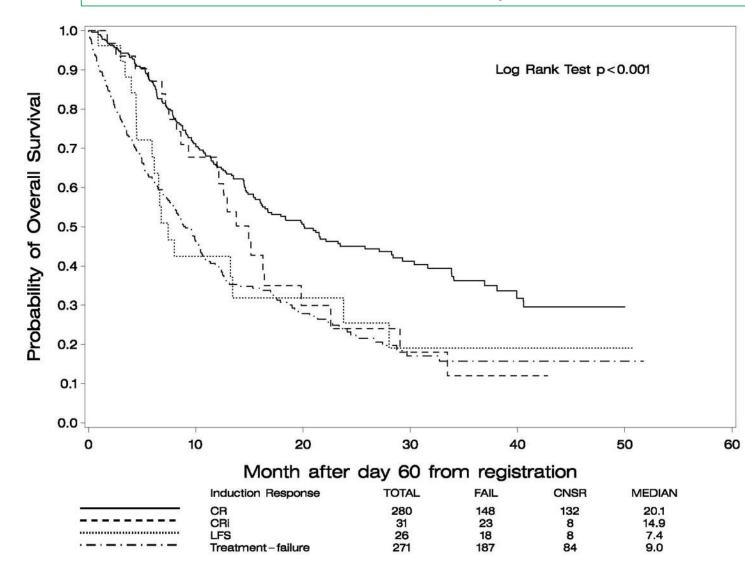
Chen C et al. (ASH 2021). Abstract 4139.



 More intensive chemotherapy combinations have been facilitated by better supportive care, including support from other medical disciplines, and nursing expertise.

Twenty five years of UK trials in acute myeloid leukaemia: what have we learned? Burnett AK, British Journal of Haematology, 188:86-100, 2020

## E2906 Randomized Study in Fit Older Adults with AML



Randomized 1:1 to Intensive therapy ("7&3", HiDAC x2 consolidation), vs. single agent Clofarabine

## \* Superior OS with 7&3

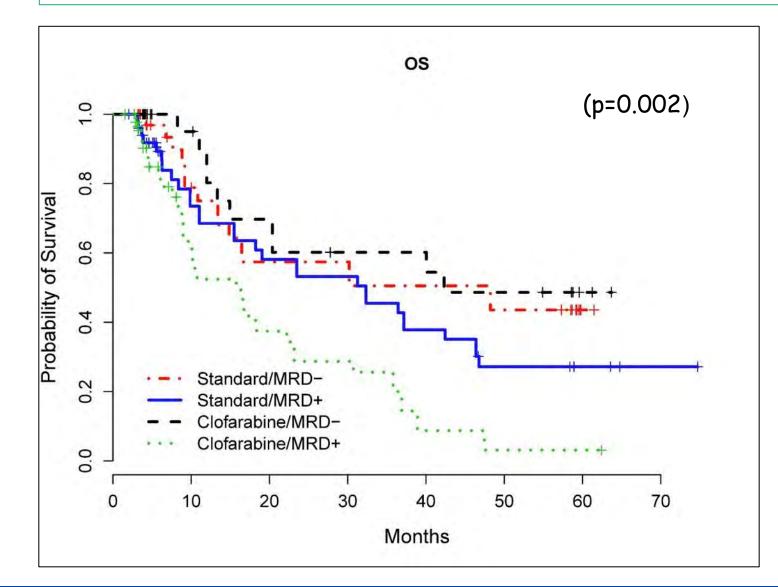
Advantage in achieving CR vs. CR/CRi

- Diminished Benefit CRi after 12 months
- MLFS survival is similar to treatment failure



Foran et al, ASH 2016

## Prospective MRD Evaluation in E2906 using Multiparameter Flow Cytometry



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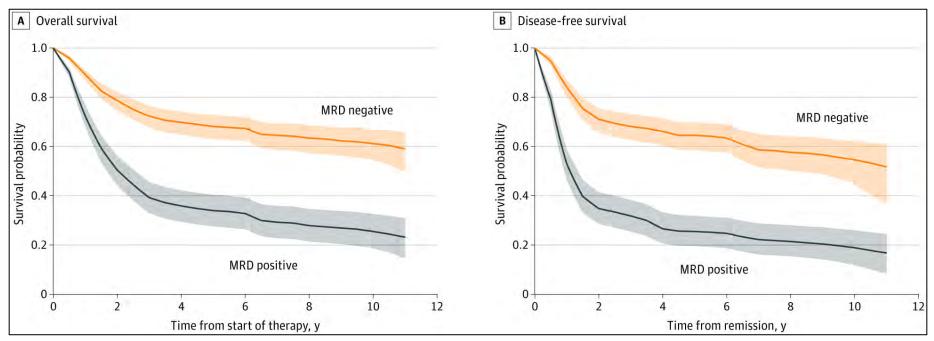
- CR/CRi 50%
- MRD-negative remission in 41% (n=161 evaluable)





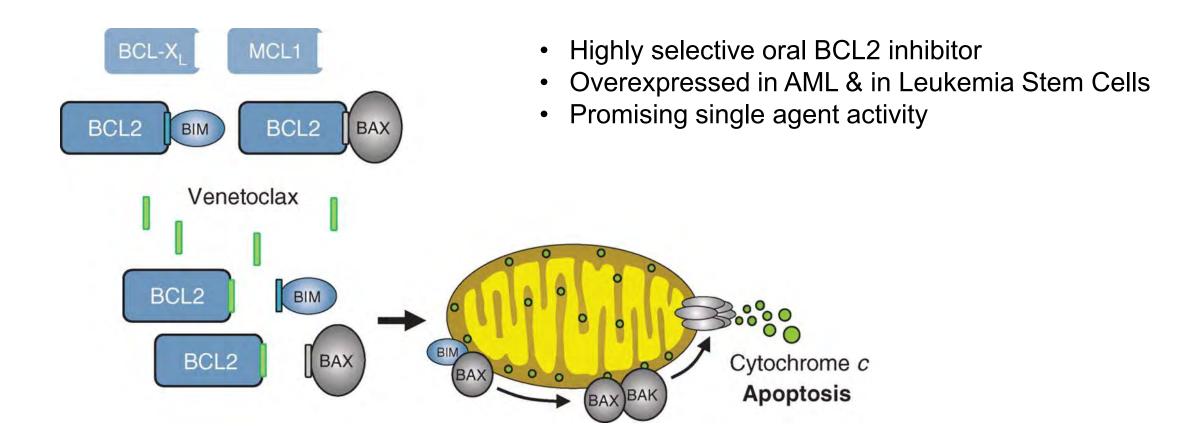
# From: Association of Measurable Residual Disease With Survival Outcomes in Patients With Acute Myeloid Leukemia: A Systematic Review and Meta-analysis

Short et al, JAMA Oncol. 2020;6(12):1890-1899.



- N=11,151 from 81 publications
- MRD by PCR, MFC, NGS, cytogenetics and/or FISH, or others
- Average Hazard Ratio for achieving MRD negativity:
  - OS **HR 0.36** (95% Bayesian credible interval [CrI], 0.33-0.39)
  - DFS **HR 0.37** (95% Crl, 0.34-0.40)

# **Mechanism of action of Venetoclax**







AACR Anterior Association for Cancer Research

CANCER DISCOVERY

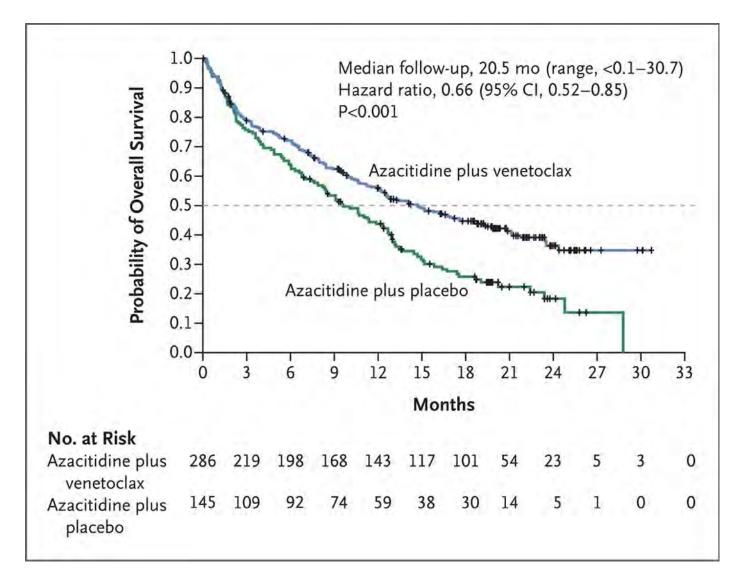
### ORIGINAL ARTICLE

# Azacitidine and Venetoclax in Previously Untreated Acute Myeloid Leukemia

Courtney D. DiNardo, M.D., Brian A. Jonas, M.D., Ph.D., Vinod Pullarkat, M.D., Michael J. Thirman, M.D., Jacqueline S. Garcia, M.D., Andrew H. Wei, M.B., B.S., Ph.D., Marina Konopleva, M.D., Ph.D., Hartmut Döhner, M.D., Anthony Letai, M.D., Ph.D., Pierre Fenaux, M.D., Ph.D., Elizabeth Koller, M.D., Violaine Havelange, M.D., Ph.D., Brian Leber, M.D., Jordi Esteve, M.D., Ph.D., Jianxiang Wang, M.D., Vlatko Pejsa, M.D., Ph.D., Roman Hájek, M.D., Ph.D., Kimmo Porkka, M.D., Ph.D., Arpád Illés, M.D., D.Sci., David Lavie, M.D., Roberto M. Lemoli, M.D., Kazuhito Yamamoto, M.D., Ph.D., Sung-Soo Yoon, M.D., Ph.D., Jun-Ho Jang, M.D., Su-Peng Yeh, M.D., Mehmet Turgut, M.D., Wan-Jen Hong, M.D., Ying Zhou, Ph.D., Jalaja Potluri, M.D., and Keith W. Pratz, M.D.

- Patients age >75 years, or *ineligible* for standard induction therapy
  - CHF or chronic stable angina
  - Pulmonary Disease (DLCO or FEV1 <65%)
  - ECOG performance-status score >1
- n=431, 2:1 randomization





## Viale A: Phase 3 randomized study AZA-Venetoclax vs. AZA-Placebo

- Improved OS overall
- Higher CR rates across all subgroups
- Some able to proceed to 'curative' AlloHCT

\* <u>Venetoclax Dosing recommendations</u>: DiNardo & Wei, Blood 135:85, 2020 *"How I treat AML in the era of new drugs"* 



Subgroup	Azacitidine plus Venetoclax	Azacitidine plus Placebo	Haza	rd Ratio for Death (95% CI)	
	no. of events/	total no. (%)			
All patients	161/286 (56.3)	109/145 (75.2)			0.64 (0.50-0.82)
Sex					
Female	61/114 (53.5)	41/58 (70.7)			0.68 (0.46-1.02)
Male	100/172 (58.1)	68/87 (78.2)			0.62 (0.46-0.85)
Age					
<75 yr	66/112 (58.9)	36/58 (62.1)	·		0.89 (0.59-1.33)
≥75 yr	95/174 (54.6)	73/87 (83.9)			0.54 (0.39-0.73)
Geographic region					
United States	27/50 (54.0)	21/24 (87.5)			0.47 (0.26-0.83)
Europe	70/116 (60.3)	46/59 (78.0)	)		0.67 (0.46-0.97)
China	9/24 (37.5)	5/13 (38.5)			1.05 (0.35-3.13)
Japan	10/24 (41.7)	9/13 (69.2)	+ <b>-</b>	-1	0.52 (0.20-1.33)
Rest of world	45/72 (62.5)	28/36 (77.8)		4	0.73 (0.45-1.17)
Baseline ECOG score					and a start a
Grade <2	89/157 (56.7)	65/81 (80.2)	Hard S		0.61 (0.44-0.84)
Grade ≥2	72/129 (55.8)	44/64 (68.8)			0.70 (0.48-1.03)
Type of AML					
De novo	120/214 (56.1)	80/110 (72.7)			0.67 (0.51-0.90)
Secondary	41/72 (56.9)	29/35 (82.9)			0.56 (0.35-0.91)
Cytogenetic risk					
Intermediate	84/182 (46.2)	62/89 (69.7)			0.57 (0.41-0.79)
Poor	77/104 (74.0)	47/56 (83.9)		+	0.78 (0.54-1.12)
Molecular marker					
FLT3	19/29 (65.5)	19/22 (86.4)		-	0.66 (0.35-1.26)
IDH1	15/23 (65.2)	11/11 (100.0)			0.28 (0.12-0.65)
IDH2	15/40 (37.5)	14/18 (77.8)		8	0.34 (0.16-0.71)
IDH1 or IDH2	29/61 (47.5)	24/28 (85.7)			0.34 (0.20-0.60)
TP53	34/38 (89.5)	13/14 (92.9)			0.76 (0.40-1.45)
NPM1	16/27 (59.3)	14/17 (82.4)		_	0.73 (0.36-1.51)
AML with myelodysplasia-related	changes				
Yes	56/92 (60.9)	38/49 (77.6)		à.	0.73 (0.48-1.11)
No	105/194 (54.1)	71/96 (74.0)			0.62 (0.46-0.83)
Bone marrow blast count					
<30%	46/85 (54.1)	28/41 (68.3)		-	0.72 (0.45-1.15)
30 to <50%	36/61 (59.0)	26/33 (78.8)			0.57 (0.34-0.95)
≥50%	79/140 (56.4)	55/71 (77.5)			0.63 (0.45-0.89)
			1 1	0 10.0	
			Angeltiding plus	Anacitidina plus	
			Azacitidine plus Venetoclax Better	Azacitidine plus Placebo Better	

# Viale-A <u>SubGroup</u> Analysis

- Significantly higher CR rates in all subgroups
- Some subgroups <u>not</u> significant for OS (post hoc)
  - Age <75 years
  - Poor Risk cytogenetics
  - Some mutation groups (e.g. TP53, NPM1, FLT3)

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## DiNardo. N Engl J Med 383:617, 2020

## **Original Study**

Check for updates

## Induction Therapy and Survival for Acute Myeloid Leukemia in Hispanic Adults from Puerto Rico

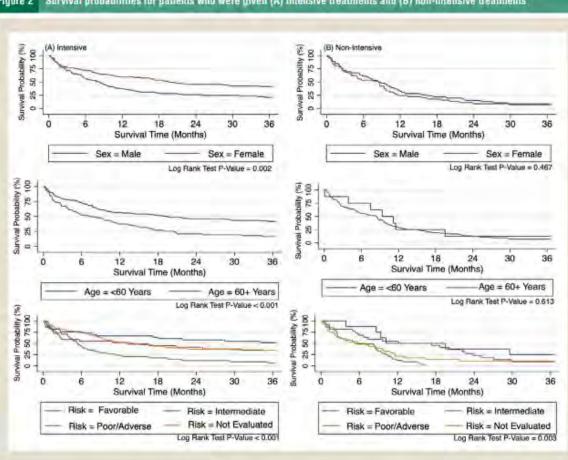
Maira A. Castaneda-Avila,<sup>1,#</sup> Tonatiuh Suárez Ramos,<sup>2,#</sup> Carlos R. Torres-Cintrón,<sup>2,#</sup> Luis A. Cotto-Santana,<sup>3</sup> Guillermo Tortolero-Luna,<sup>2,4,#</sup> Karen J. Ortiz-Ortiz<sup>2,4,5,#</sup>

### Abstract

We described the first-line therapy and survival of Hispanics from Puerto Rico with acute myeloid leukemia. Age, risk, and comorbidities were associated with induction therapy type. Among patients who received intensive therapy, those of  $\geq$ 60 years had a higher risk of death. The low survival and the disparities observed highlight the need to examine further new treatment options for older and comorbid patients.

**Background:** Acute myeloid leukemia (AML) is the most common type of leukemia in adults. There are no previous studies evaluating AML treatment patterns in Puerto Rico. We describe the first-line therapy patterns and survival of patients diagnosed with AML in Puerto Rico using the Puerto Rico Central Cancer Registry Health Insurance Linkage Database (2011-2015). **Methods:** We describe patient characteristics according to intensive, non-intensive, and non-treatment status. We used Cox proportional hazard models to evaluate the factors associated with the risk of death stratified by intensive and non-intensive therapy. For this study, 385 patients with AML were included. **Results:** The mean age was 67 years old and 50.1% were female. Nearly half of AML patients (46.8%) received intensive treatment, 23.6% received non-intensive treatment, and 26.2% did not receive treatment. The overall 3-year survival rate was 17.9%. Among those who received intensive therapy, the risk of death among females was lower than males (hazard ratio [HR]: 0.64, 95% confidence interval [CI]: 0.44-0.93). Patients 60 years or older who received intensive treatment had a higher risk of death than younger patients (HR: 1.67, 95% CI: 1.09-2.55). Patients with poor/adverse risk receiving intensive (HR: 3.43, 95% CI: 1.76-6.69) or non-intensive (HR: 4.32, 95% CI: 1.66-11.28) treatment had a higher risk of death than patients with a favorable risk category. **Conclusion:** Our findings are the first step to monitor the quality of care of patients with AML in Puerto Rico, particularly related to the administration of appropriate induction therapies, which is one of the most important predictors of AML survival.

Clinical Lymphoma, Myeloma and Leukemia, Vol. 22, No. 10, e922–e930 © 2022 Published by Elsevier Inc. Keywords: AML, Hazard ratio, Prognostic markers, Leukemia treatment, Risk of death



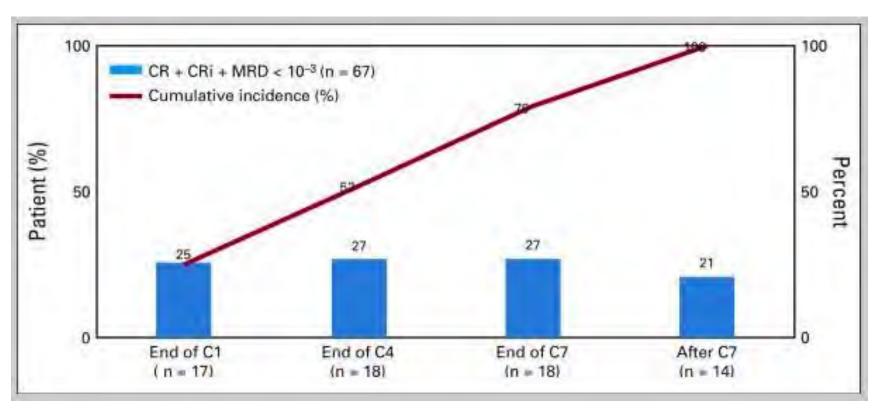


Castaneda-Avila et al, Clinical Lymphoma, Myeloma & Leukemia. 22:e922-930, 2022

### Maira A. Castaneda-Avila et al

Figure 2 Survival probabilities for patients who were given (A) intensive treatments and (B) non-intensive treatments

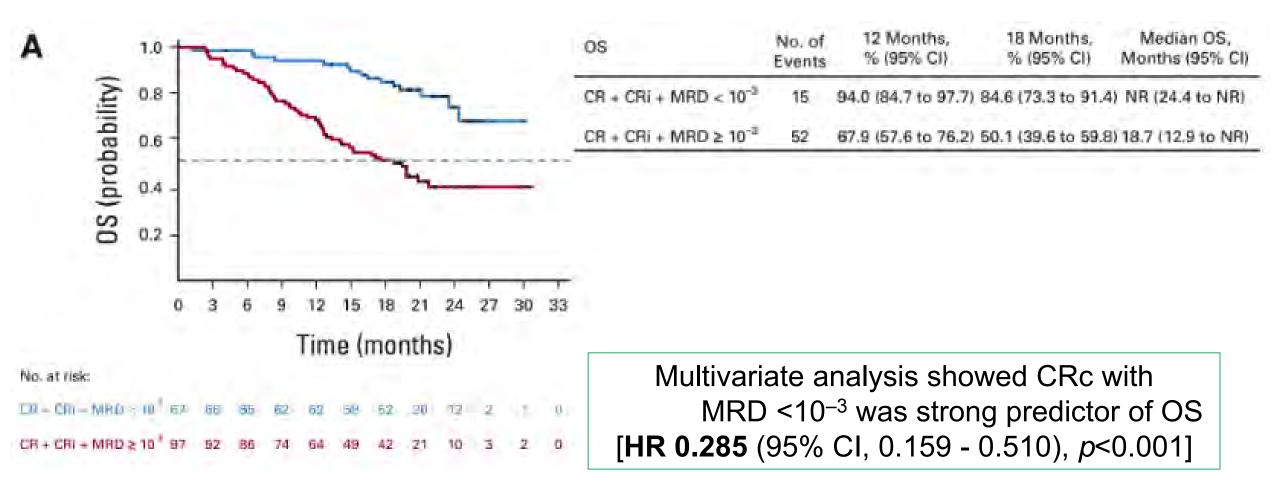
# MRD with Azacitidine & Venetoclax in Viale-A Trial



- Prospective central (LabCorp) MRD testing by MFC
  - 86% of CRc patients evaluable
- Threshold 10<sup>-3</sup> (i.e. <0.1%) residual leukemia blasts
- 41% MRD-neg CR/CRi



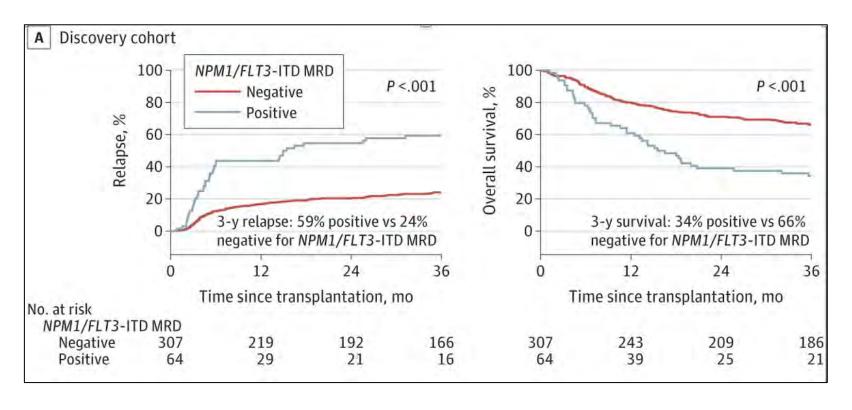
# MRD with Azacitidine & Venetoclax in Viale-A Trial





Pratz et al, JCO 40:855, 2022

## *Pre-MEASURE* Observational Study of FLT3<sub>m</sub> and NPM1<sub>m</sub> in Remission Prior to Allogeneic Transplantation



- 17.3% with residual NPM1\_and/or FLT3-ITD variants
- <u>Higher rates of relapse</u> at 3 years
   68% vs 21%
- HR 4.32 [95% Cl, 2.98-6.26]; P < .001)</li>
- Decreased survival at 3 years 39% vs 63%
- HR, 2.43 [95% CI, 1.71 3.45]; P < .001).</li>

- Variants in FLT3 & NPM1 from Blood from AML patients
- 111 CIBMTR sites from 2013 through 2019. Clinical data were collected through May 2022
- Targeted error-corrected DNA sequencing at a variant allele fraction (VAF) of 0.01%



# **Evaluating MRD in AML**

<u>Timing</u>

CR/CRi; Later if low intensity...

**Prior to Transplant** 

Surveillance

## **Platform**

.. MFC (LAIP, DFN, both)

PCR (NPM1, FLT3, CBF), or NGS PCR (NPM1, CBF), or NGS

# Goals & Challenges:

- Increase MRD-negative CR rate
  - Addition targeted inhibitors, immunotherapy (clinical trials)
- Improve outcomes in MRD-positive remission
  - e.g. intensify HiDAC consolidation (older adults); maintenance with oral azacitidine (e.g. NPM1<sub>m</sub>); addition of targeted agents (Gilteritinib in MRD+ AML with FLT3-ITD after AlloHCT), etc.



# Advances Since 2017, and the Rapidly Developing Future of AML Care



## **Emerging advances center on**

- Integrating novel targeted doublets and triplets
- Development novel targeted agents, and strategies for challenging subgroups (*e.g. TP53*-mutated AML)
- Determining role for immunotherapy in AML and increasing benefit of allogeneic transplantation

- CPX-351 (sAML, AML-MRC)
- Midostaurin,
   Gilteritinib &
   Quizartinib
- Ivosidenib,
   Olutasidenib,
   and Enasidenib
- Ivosidenib + azacitidine combo

- Re-emergence of gemtuzumabozogamicin
- ✓ Venetoclax
- Oral azacitidine as maintenance
- Glasdegib



## Sept 10, 2009

TH NEW ENGLAND JOUGNAL of MEDICINE

### ORIGINAL ARTICLE

### Recurring Mutations Found by Sequencing an Acute Myeloid Leukemia Genome

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### ABSTRACT

### BACKGROUND

The full complement of UNA mutations that are responsible for the pathogenesis of acute myeloid leukemia (AML) is not yet known.

#### METHODS

We used massively parallel DNA sequencing to obtain a very high level of coverage (approximately 98%) of a primary, cytogenetically normal, de novo genome for AML with minimal maturation (AML-MI) and a matched normal skin genome.

### PESSIATS

We identified 12 acquired (somatic) mutations within the coding sequences of genes and 52 somatic point mutations in conserved or regulatory portions of the genome. All mutations appeared to be heterozygous and present in nearly all cells in the tumor sample. Four of the 64 mutations occurred in at least 1 additional AML sample in 188 samples that were tested. Mutations in NRAS and NPMI had been identified previously in patients with AML, but two other mutations had not been identified. One of these mutations, in the IDH1 gene, was present in 15 of 187 additional AML genomes tested and was strongly associated with normal cytogenetic status; it was present in 13 of 80 cytogenetically normal samples (16%). The other was a nongenic mutation in a genomic region with regulatory potential and conservation in higher mammals; we detected it in one additional AML rumor. The AML genome that we sequenced contains approximately 750 point mutations, of which only a small fraction are likely to be relevant to pathogenesis.

#### CONCLUSIONS

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By comparing the sequences of tumor and skin genomes of a patient with AML-M1, we have identified recurring mutations that may be relevant for pathogenesis.

NENGLI MED 36171 NEIM.ORG SEPTEMBER 10, 2009

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Cancer	Cell
8 - 1	5
ALL	101

### The Common Feature of Leukemia-Associated IDH1 and IDH2 Mutations Is a Neomorphic Enzyme Activity Converting a-Ketoglutarate to 2-Hydroxyglutarate

10

Patrick S, Ward,<sup>1</sup> Jav Patel,<sup>3</sup> David R, Wise,<sup>1</sup> Omar Abdel-Wahab,<sup>3</sup> Bryson D, Bennett,<sup>6</sup> Hilary A, Coller,<sup>6</sup> Justin R, Cross,<sup>1</sup> Valeria R. Fantin," Cyrus V. Hedvat," Alexander E. Perl, Joshua D. Rabinowitz,<sup>6</sup> Martin Carroll, Shinsan M. Su," Kim A. Sharp,<sup>2</sup> Ross L. Levine,<sup>3</sup> and Craig B. Thompson<sup>1</sup> Abramson Cancer Center, Division of Hematology and Oncology, Department of Medicine Department of Biochemistry and Biophysica University of Pennsylvania, Philadelphia, PA 19104, USA Human Oncology and Pathogenesis Program \*Department of Pathology Memorial Sloan-Kettering Cancer Center, New York, NY 10065, USA <sup>2</sup>Department of Chemistry and Integrative Genomics Department of Molecular Biology Princeton University, Princeton, NJ 08544, USA Agios Pharmaceuticals, Cambridge, MA 02139, USA Correspondence: craig@mail.med.upenn.edu DOI 10.1016/j.ecr.2010.01.020

#### SUMMARY

The somatic mutations in cytosolic isocitrate dehydrogenase 1 (IDH1) observed in gliomas can lead to the production of 2-hydroxyglutarate (2HG). Here, we report that tumor 2HG is elevated in a high percentage of patients with cytogenetically normal acute myeloid leukemia (AML). Surprisingly, less than half of cases with elevated 2HG possessed IDH1 mutations. The remaining cases with elevated 2HG had mutations in IDH2, the mitochondrial homolog of IDH1. These data demonstrate that a shared feature of all cancer-associated IDH mutations is production of the oncometabolite 2HG, Furthermore, AML patients with IDH mutations display a significantly reduced number of other well characterized AML-associated mutations and/or associated chromosomal abnormalities, potentially implicating IDH mutation in a distinct mechanism of AML pathogenesis.

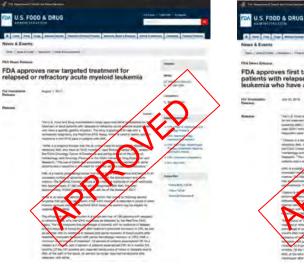


## First patient treated **Sept 2013**



News & Event

## July 20, 2018





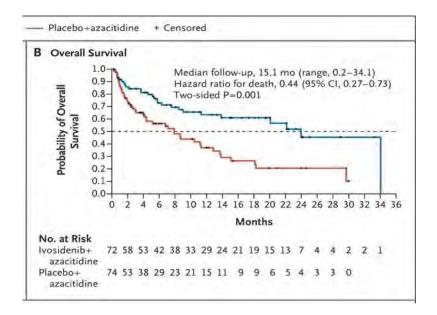
\*2022 Olutasidenib

Enasidenib

### ORIGINAL ARTICLE

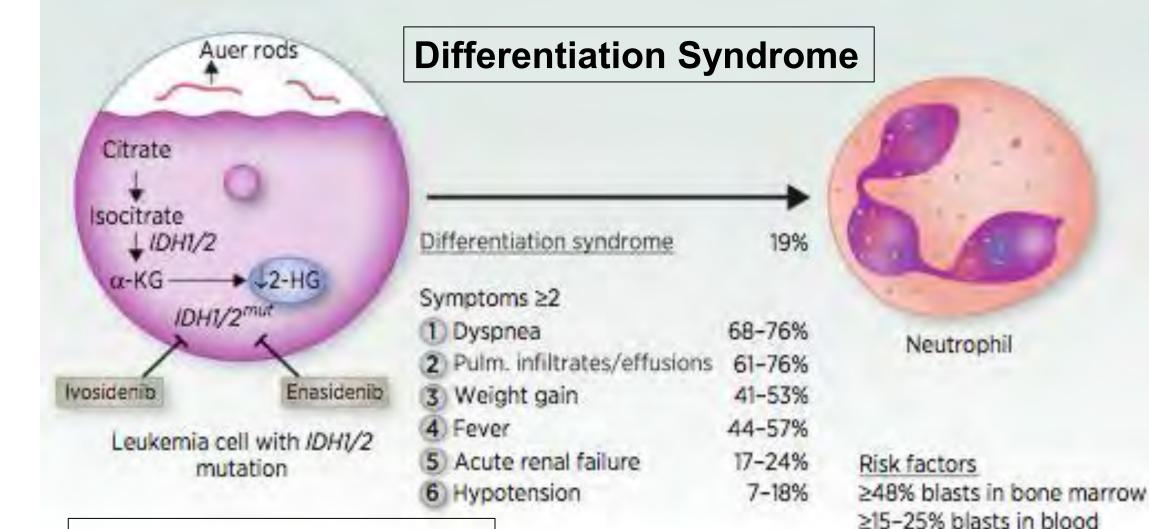
## Ivosidenib and Azacitidine in IDH1-Mutated Acute Myeloid Leukemia

Pau Montesinos, M.D., Ph.D., Christian Recher, M.D., Ph.D., Susana Vives, M.D., Ewa Zarzycka, M.D., Jianxiang Wang, M.D., Giambattista Bertani, M.D., Michael Heuser, M.D., Rodrigo T. Calado, M.D., Ph.D., Andre C. Schuh, M.D., Su-Peng Yeh, M.D., Scott R. Daigle, M.S., Jianan Hui, Ph.D., et al.



- Median survival 24.0 vs. 7.9 months
  - HR 0.44 (95% CI, 0.27-o 0.73; p=0.001)
- CR/CRh 53% vs.18% (p<0.001)
- DS 14 %
- FDA Approval 1<sup>st</sup> line Aza combination 12/22





Dexamethasone 10mg IV BID

Norsworthy, Clin Cancer Res 26:4280, 2020

Hydroxyurea

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Continue Agent if possible

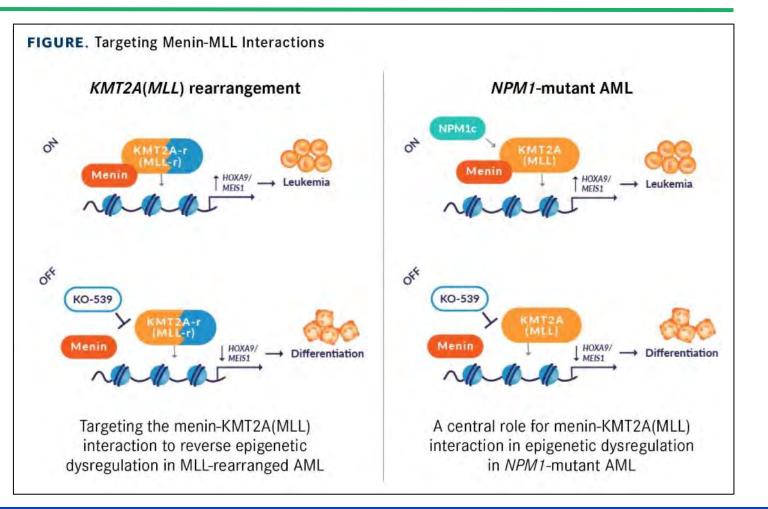
Zeidner, Clin Cancer Res 26:4174, 2020

Concomitant TET2, SRSF2

# New Oral agents in AML Menin Inhibitors - NPM1<sub>mut</sub> & MLL rearrangement

- Target HOXA9/Meis1

   overexpression in AML, which
   leads to AML
   proliferation/renewal
- AML with  $NPM1_{mut}$  or MLLr
- Promising agents in clinical development (KO-539, SNDX-5613, BMF-219, others...)
- DS as a clinical toxicity/concern



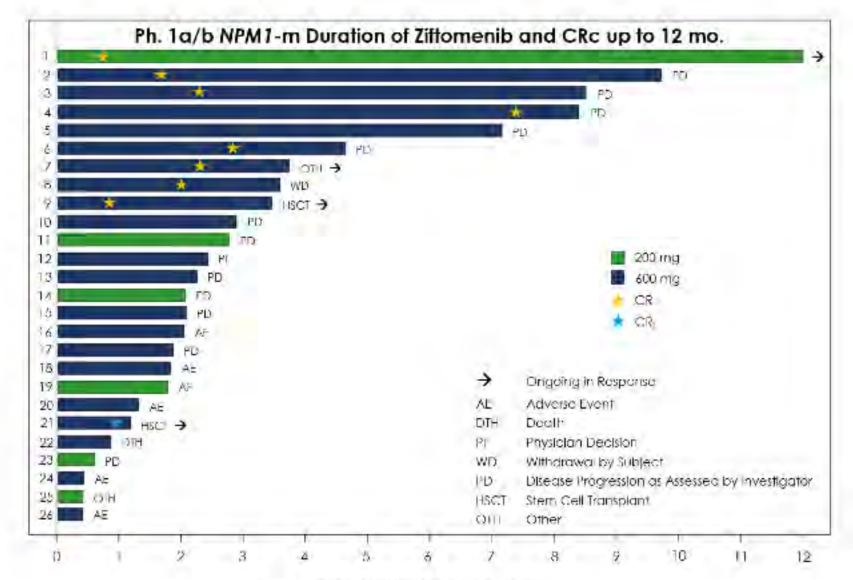


## 5%-10% KMT2A Translocation

## 25-30% NPM1 mutation

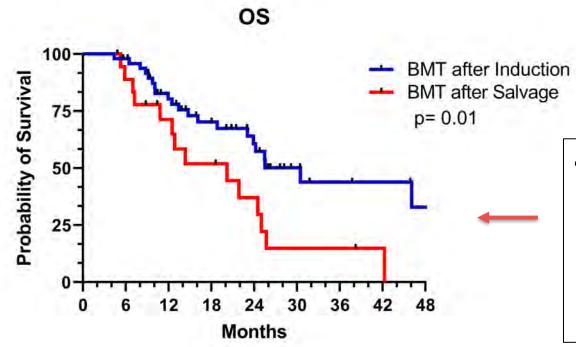
## Ziftomenib Monotherapy Drives Durable Responses

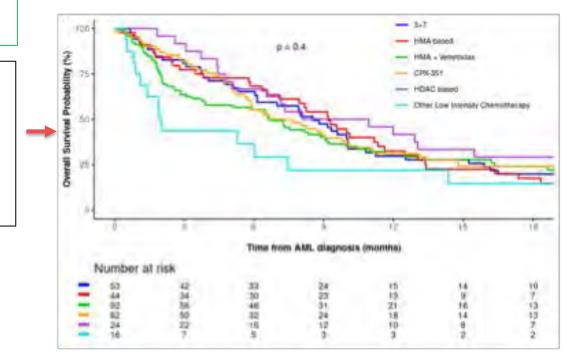
- Median DoR 8.2 months (med. follow up 8.8 months)
- 35% CR rate (7/20) at 600mg dose (RP2D)
  - 33% (2/6) with FLT3 comutations, and 50% (4/8) with IDH co-mutations achieved CR
- DS: 15% (grade 1/2) and 5% (grade 3)
- Ziftomenib less likely to induce Menin resistance mutations



# AML with TP53 Mutation

 Despite surge in use of venetoclax-based therapy, no improvement in duration response or overall survival in "Real World" dataset (Badar, ASH 2023)

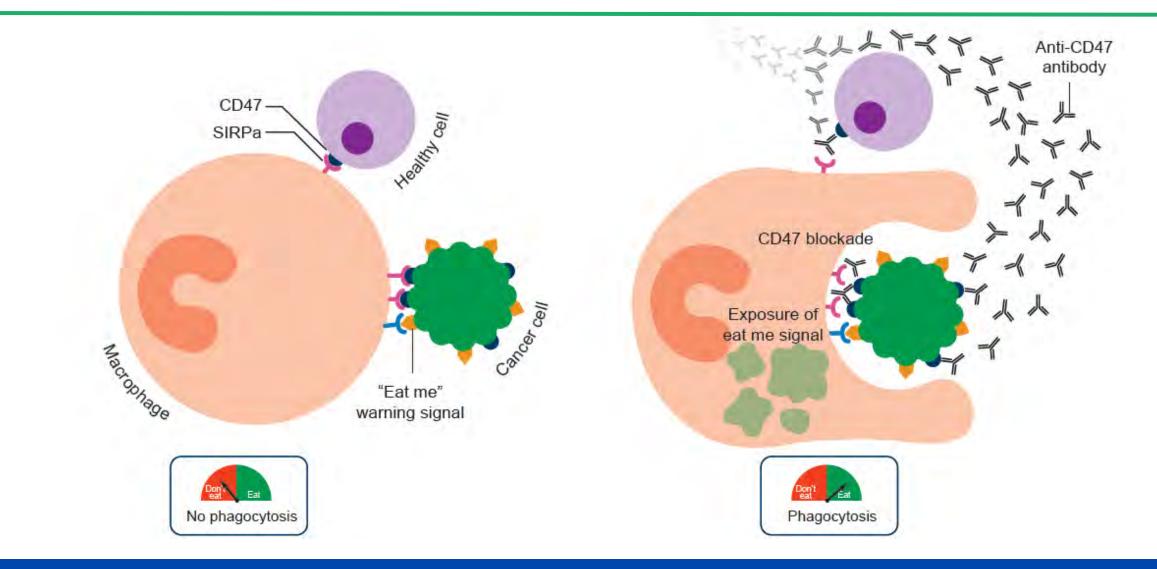




 Impact of Allogeneic Transplantation, esp. in 1<sup>st</sup> line (Med. OS 30.5 mos. vs. 20.2 mos.), again demonstrated in RWD analysis (Badar, ASH 2023)
 HR: 0.15 (95% CI 0.04-0.40), p=0.002



## Mechanism of Action of CD47-Blocking Antibodies as Novel Checkpoint Inhibitors





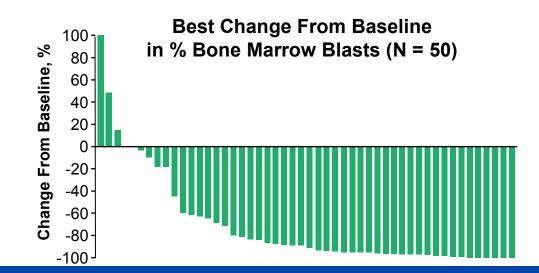
### Chao MP et al. *Front Oncol* 9:1380, 2019

# Magrolimab + Azacitidine Appeared "Efficacious" Against TP53 AML in Early Phase Trial

- No significant cytopenias, infections, or immune-related AEs observed
- On-target hemolytic anemia
- Median OS 10.8 months

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Outcome	Patients With TP53 AML (N = 72)
ORR, % (95% CI)	48.6 (36.7, 60.7)
<b>CR, % (95% CI)</b> MRD- CR <sup>a</sup> , % (95% CI)	<b>33.3 (22.7, 45.4)</b> (n = 24/72) 50.0 (29.1, 70.9) (n = 12/24)
CRi/CRh, n (%)	6 (8.3)
PR, n (%)	4 (5.6)
MLFS, n (%)	1 (1.4)
DOR, median (95% Cl), mo	8.7 (6.5, 10.4)



Frontline *TP53*-mutated AML in the Phase 3 ENHANCE-2 trial (currently recruiting; NCT04778397)

Daver N. EHA2022 Hybrid Congress. Abstract S132.

# Gilead Statement on the Discontinuation of Magrolimab Study in AML with TP53 Mutations

**Foster City, Calif., September 26, 2023** – Gilead Sciences has stopped its ENHANCE-2 study in acute myeloid leukemia (AML) with TP53 mutations. Based on an ad hoc analysis, and following review by an independent data monitoring committee, Gilead concluded that magrolimab is unlikely to demonstrate a survival benefit in AML with TP53 mutations compared to standard of care. There were no new safety signals identified and the safety profile was comparable between treatment arms. ENHANCE-2 is a randomized, open-label, Phase 3 trial to determine if magrolimab plus azacitidine improves overall survival, compared to physician's choice of venetoclax plus azacitidine or intensive chemotherapy in previously untreated AML with TP53 mutations. Gilead is working with study investigators on appropriate next steps for patients enrolled in this study. This decision follows the previously announced partial clinical hold placed on the ENHANCE-2 study.



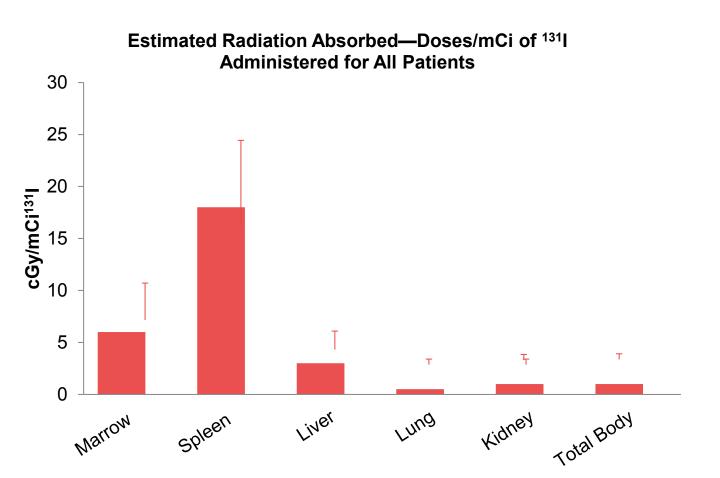
AML with TP53 mutation remains an area of pressing need for novel therapies that improve outcome

# Iomab-B (I<sup>131</sup>-Apamistamab) & CD45: Mechanisms and Biodistribution

- CD45 antigen expressed on virtually all lymphocytes, and 85%-90% of acute leukemias
- Iomab-B (<sup>131</sup>I apamistamab): anti-CD45 mAb targeting lymphohematopoietic cells with β-particle– emitting radionuclide<sup>131</sup>I
- Offers target-specific ablation as HCT conditioning regimen

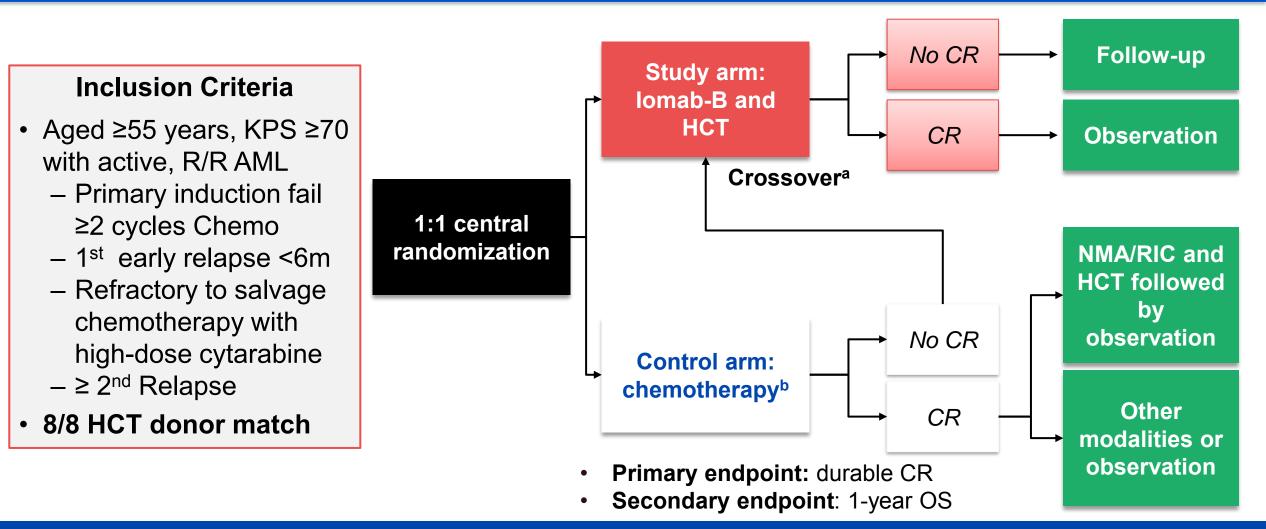
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 Does not bind other normal tissues; directs radiation to leukemic and immune cells





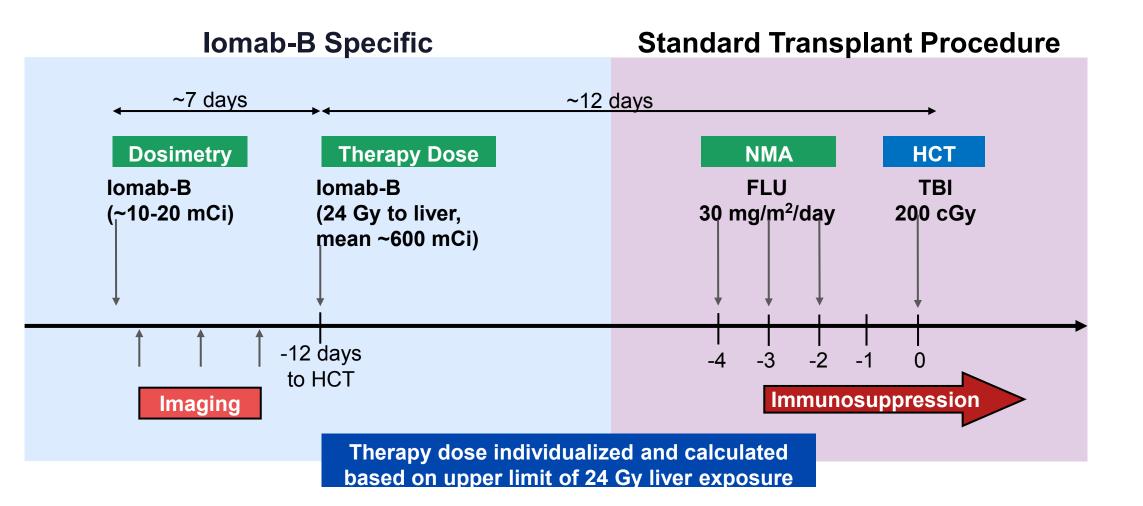
# Phase 3 SIERRA Trial: Iomab-B Prior to HCT vs Chemotherapy in R/R AML





<sup>a</sup> Control arm patients with no CR offered crossover for ethical reasons.
 <sup>b</sup> Physician choice of best salvage chemotherapy using approved products.

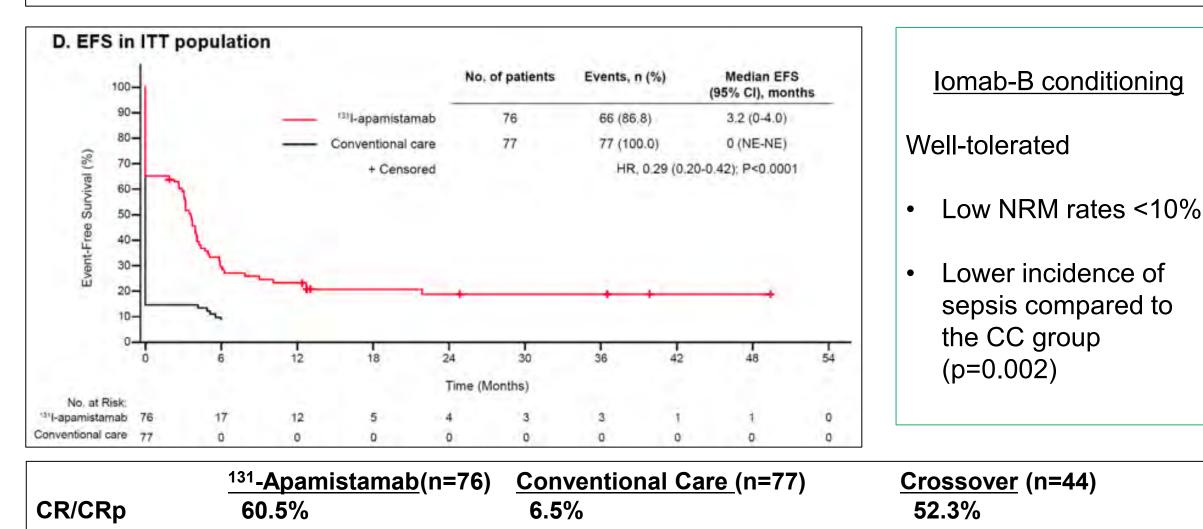
# **SIERRA: Iomab-B Treatment Schedule**





Gyurkocza B et al. ASTCT/CIBMTR (Tandem) 2023 and EHA 2023

## <sup>131</sup>-Apamistamab vs. Conventional Care



0%

(\*p<0.0001)

\*Durable CR

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17.1%

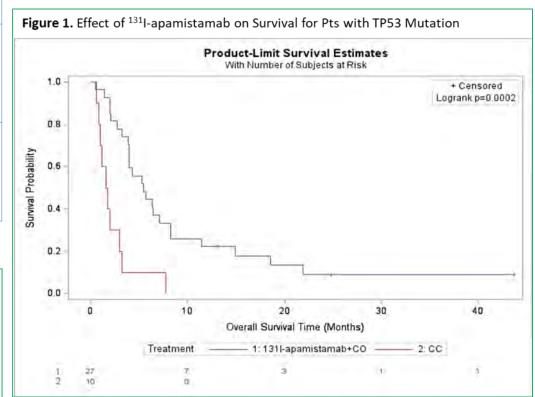
Gyurkocza B et al. TCT (Tandem), LBA 3, 2023

# <sup>131</sup>-Apamistamab vs. Conventional Care in TP53<sub>mut</sub>

Table 2: CR and dCR rates by TP53 mutation status and treatment received

	lor	nab-B + Cros	sover	Conventional Care		
Response Type	N	%	95% CI	N	%	95% CI
TP53 Positive	N = 27			N = 10		
CR	15	55.56	(35.33, 74.52)	0	0	_
Durable CR	4	14.81	(4.19, 33.73)	0	0	_
Wildtype	N = 93			N = 23		
CR	54	58.06	(47.38, 68.22)	4	17.39	(4.95, 38.78)
Durable CR	15	16.13	(9.32, 25.20)	0	0	-

- 24.2% with TP53 mutation in trial
  - Similar CR and dCR with TP53<sub>mut</sub> vs. TP53<sub>wt</sub>
- Excluding CO, Median OS 5.49 mos. vs. 1.66 mos.in pts who did not receive Iomab-B (Figure)
  - HR=0.23; 95% CI [0.10, 0.52]; (p=0.0002)



Foran et al. TCT (Tandem), San Antonio, TX, Feb. 2024

# Conclusions

- •Goal Achievement high quality (MRD-negative) remission
- •Novel targeted agents, novel toxicities (DS, AIHA)
- Targeting TP53 remains a pressing clinical need
  Allogeneic transplantation improves OS for eligible patients, although long-term DFS rates remain poor
- Advances in relapsed & refractory AML, including matched donor AlloHCT for high-risk populations
- •Importance of supporting clinical trials in diverse clinical populations







"The best interest of the patient is the only interest to be considered, and in order that the sick may have the benefit of advancing knowledge, a union of forces is necessary."

1910: Dr. William J. Mayo

Rush Medical College, commencement address



