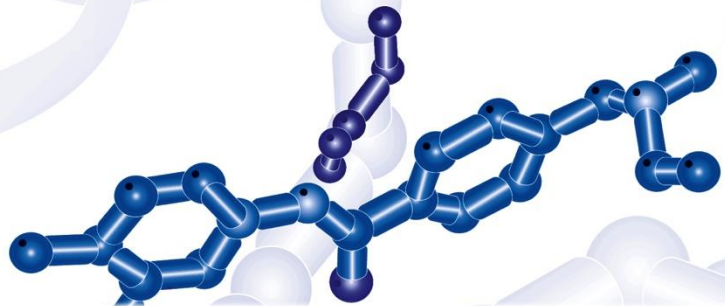




ALMA MATER STUDIORUM  
UNIVERSITÀ DI BOLOGNA  
DIPARTIMENTO DI  
SCIENZE MEDICHE E CHIRURGICHE

POLICLINICO DI  
**SANT'ORSOLA**

SERVIZIO SANITARIO REGIONALE  
EMILIA-ROMAGNA  
Azienda Ospedaliero - Universitaria di Bologna



# New Drugs in Hematology

## Menin inhibitors in AML and ALL

Cristina Papayannidis, MD, PhD  
IRCCS Azienda Ospedaliero Universitaria di Bologna

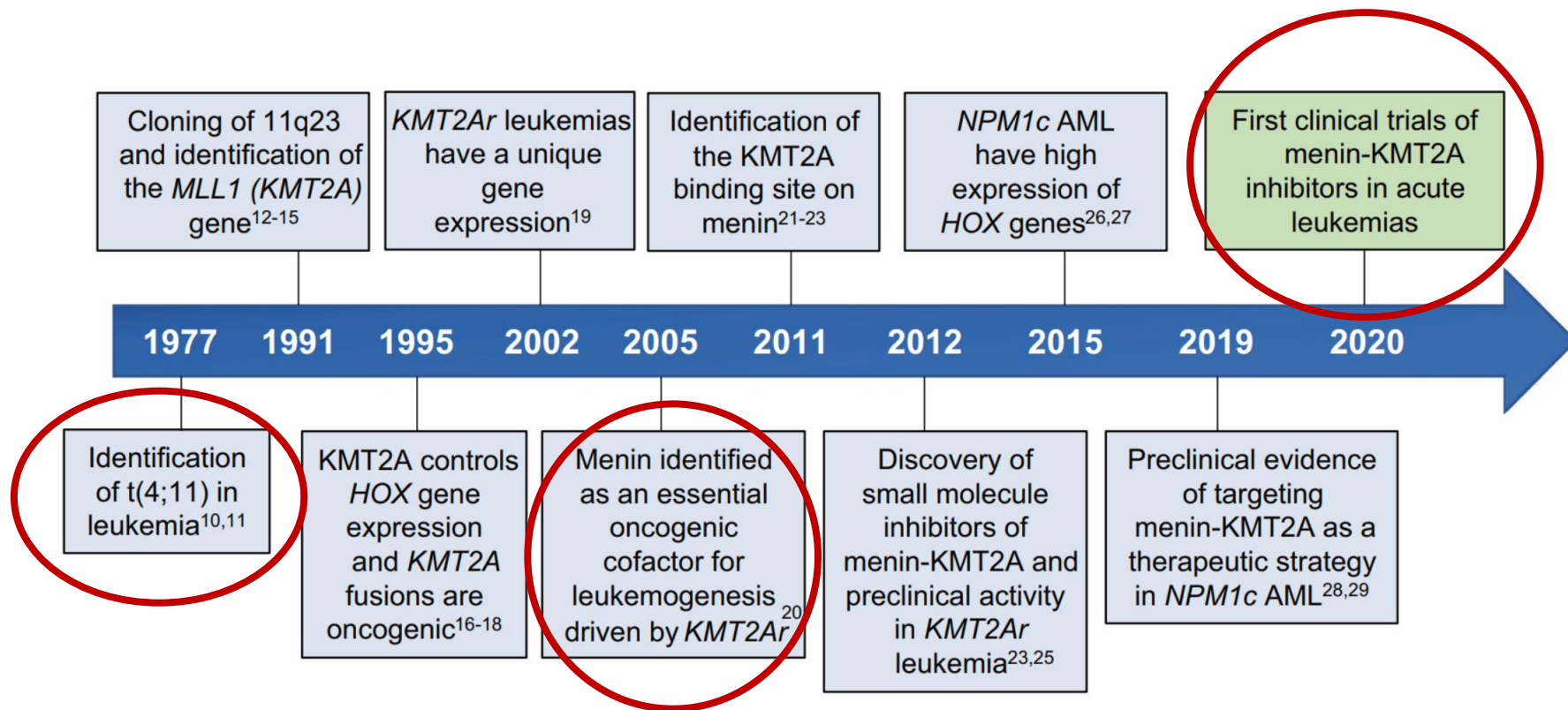
President: Pier Luigi Zinzani

**Bologna,**  
**Royal Hotel Carlton**  
**May 18-19-20, 2026**

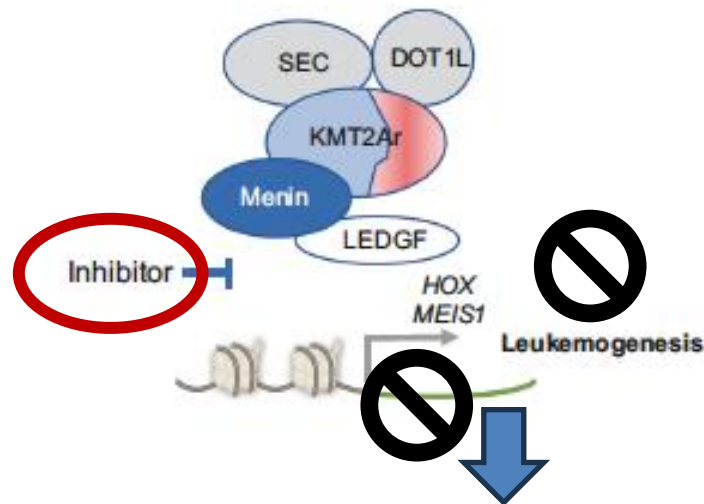
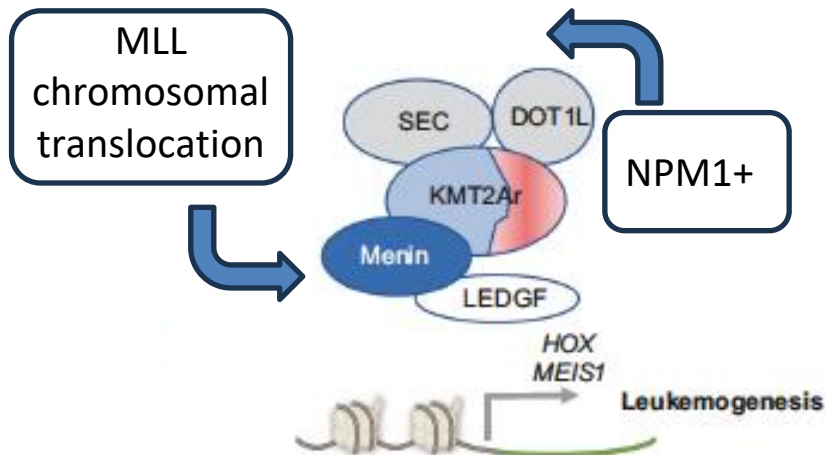
## Disclosures of Cristina Papayannidis

Company name	Research support	Employee	Consultant	Stockholder	Speakers bureau	Advisory board	Other
Pfizer					X	X	
Amgen						X	
Astellas					X		
Abbvie					X		
Menarini Stemline						X	
Servier					X		
Incyte					X		
Janssen						X	
Syndax						X	
Blueprint					X	X	
GSK						X	
Istituto Gentili					X	X	
Jazz Pharmaceuticals					X	X	

## Menin inhibitors: timeline



# Menin is an essential oncogenic cofactor for leukemogenesis driven by NPM1 mut or r-KMT2A



**Growth arrest and differentiation**

## Agenda

- For which patients today?
- How many menin inhibitors do we have today?
- What about safety with menin inhibitors as single agents?
- Which results can we achieve with menin inhibitors?
  - R/R single agents
- What about resistance?
- How can we overcome these mechanisms?
- For which patients, tomorrow?

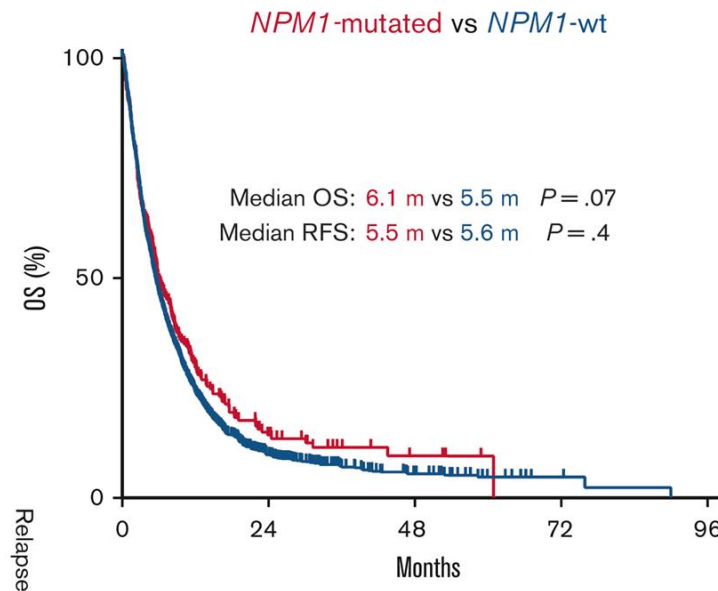


## Background (II)

Risk category†	Genetic abnormality
Favorable	<ul style="list-style-type: none"> <li>t(8;21)(q22;q22.1)/RUNX1::RUNX1T1†,‡</li> <li>inv(16)(p13.1;q22) or t(16;16)(p13.1;q22)/CBFB::MYH11†,‡</li> <li>Mutated NPM1†,§ without FLT3-ITD</li> <li>bZIP in-frame mutated CEBPA  </li> </ul>
Intermediate	<ul style="list-style-type: none"> <li>Mutated NPM1†,§ with FLT3-ITD</li> <li>Wild-type NPM1 with FLT3-ITD (without adverse-risk genetic lesions)</li> <li>t(9;11)(p21.3;q23.3)/MLL2A†,¶</li> <li>Cytogenetic and/or molecular abnormalities not classified as favorable or adverse</li> </ul>
Adverse	<ul style="list-style-type: none"> <li>t(6;9)(p23.3;q34.1)/DEK::NUP214</li> <li>t(v;11q23.3)/KMT2A-rearranged#</li> <li>t(9;22)(q34.1;q11.2)/BCR::ABL1</li> <li>t(8;16)(p11.2;p13.3)/KAT6A::CREBBP</li> <li>inv(3)(q21.3q26.2) or t(3;3)(q21.3;q26.2)/GATA2, MECOM(EV11)</li> <li>t(3q26.2;v)/MECOM(EV11)-rearranged</li> <li>-5 or del(5q); -7; -17/abn(17p)</li> <li>Complex karyotype,** monosomal karyotype††</li> <li>Mutated ASXL1, BCOR, EZH2, RUNX1, SF3B1, SRSF2, STAG2, U2AF1, and/or ZRSR2‡‡</li> <li>Mutated TP53³</li> </ul>

Dohner H et al., Blood 2022

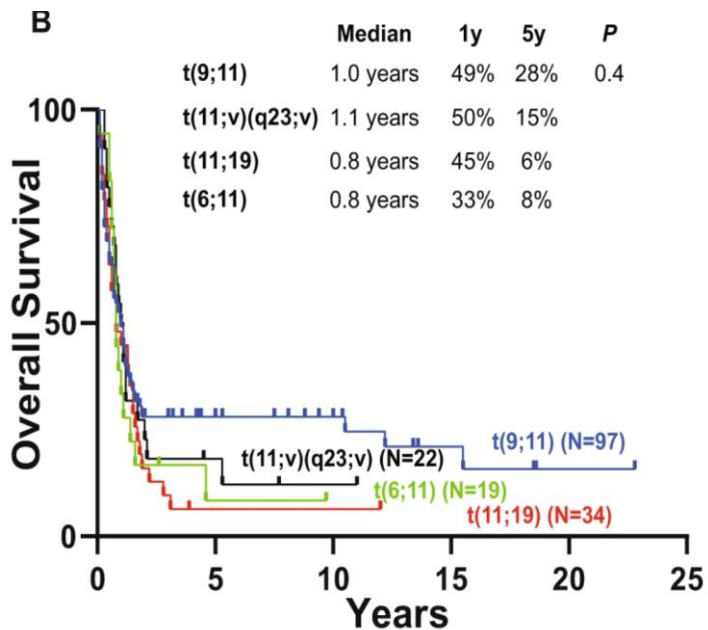
## R/R NPM1-mut OVERALL SURVIVAL



Issa GC et al., Blood Adv 2023

## r-KMT2A AML outcome

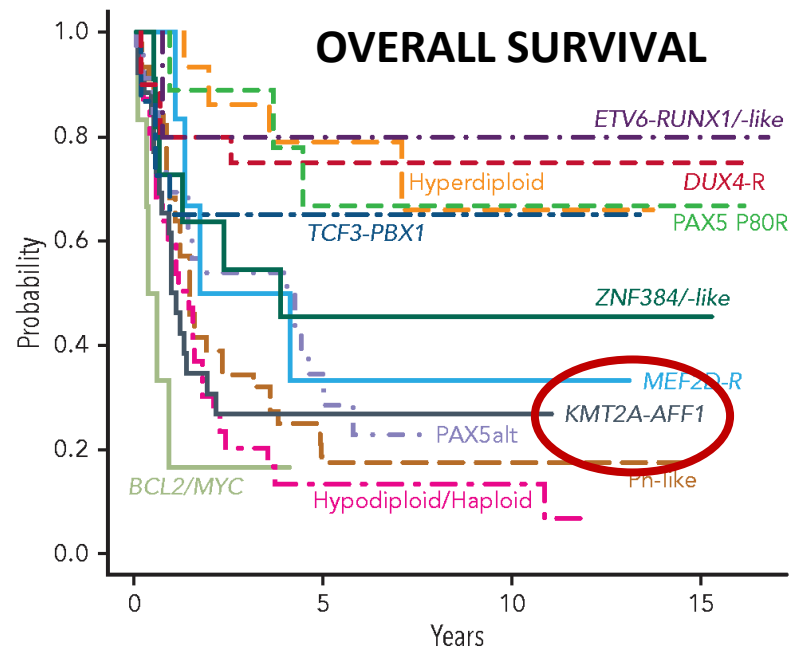
### OVERALL SURVIVAL



Issa G C et al, Blood Cancer Jour 2021

## r-KMT2A ALL outcome

### OVERALL SURVIVAL





Paietta E et al, Blood 2021

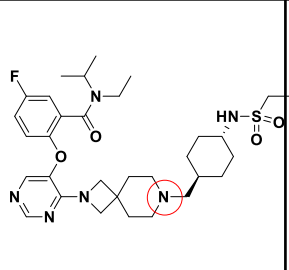
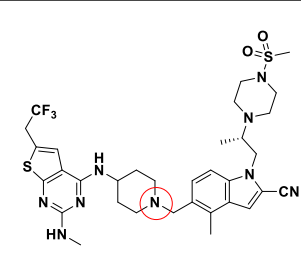
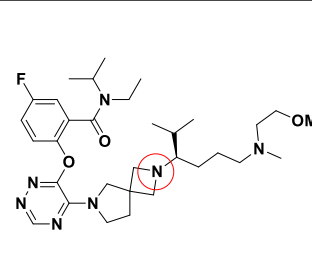
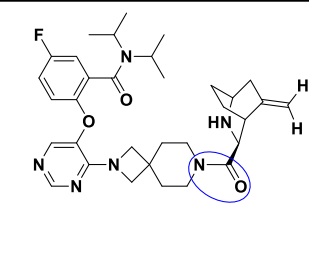
## Agenda

- For which patients today?
- How many menin inhibitors do we have today?
- What about safety with menin inhibitors as single agents?
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  - R/R single agents
- What about resistance?
- How can we overcome these mechanisms?
- For which patients, tomorrow?

## Menin inhibitors in clinical development

Trial name (NCT)	Agent (route)	Phase 1/ 2 expansion cohorts For relapsed/refractory disease	Phase /# pts	Current status
 AUGMENT-101 (NCT04065399)	<b>Revumenib</b> (SNDX-5613) PO BID	(a) ALL or MPAL with <i>KMT2Ar</i> (b) AML with <i>KMT2Ar</i> ; (c) <i>NPM1</i>	Phase 1 /2 (n=186)	<b>NPM1<sup>mut</sup> r-KMT2A</b> FDA approval Nov 15, 2024
 KOMET-001 (NCT04067336)	<b>Ziftomenib</b> (KO-539) PO QD	(a) AML with <i>KMT2Ar</i> (b) AML with <i>NPM1c</i>	Phase 1 /2 (n=199)	<b>NPM1 mut FDA approved in 2025</b>
NCT04811560	<b>Bleximenib</b> (JNJ-75276617) PO QD	(a) AML/ALL with <i>KMT2Ar</i> (b) AML with <i>NPM1c</i>	Phase 1 (n=110)	<b>Phase 1 (EHA 2024)</b> Recruiting in combination with chemo
NCT04988555	<b>Enzomenib</b> (DSP-5336) PO QD	RR-AML/RR-ALL Ph2: <i>NPM1/KMT2Ar</i>	Phase 1/2 (n=70)	<b>Phase 1 (EHA 2024)</b> Recruiting
COVALENT-101 (NCT05153330)	BMF-219 PO	(a) AML/ ALL ( <i>KMT2Ar</i> , <i>NPM1</i> ) (b)DLBCL; (c) MM; (d) CLL/SLL	Phase 1 (n=177)	Multiple cohorts Actively enrolling

## Menin inhibitors are not the same with differences in chemical structure and physiochemical properties

	Revumenib	Ziftomenib	Bleximenib	Enzomenib
Structure*				
	Tertiary amine bond	Tertiary amine bond	Tertiary amine bond	<b>Amide bond</b>

\*National Center for Biotechnology Information (2024). PubChem Compound Summary for CID 132212657, 138497449, 156498110, 146430058

- Each menin inhibitor has a different chemical structure and different physiochemical properties such as polar surface area, lipophilicity, and basicity that may impact safety and efficacy<sup>1,2</sup>

Chemical Property*	Polar Surface Area <sup>1,2,3</sup>	Lipophilicity <sup>1,2</sup>	Basicity <sup>1,2</sup>
Target Value	PSA > 75	ClogP < 3.0	Pka < 7.4
<b>Enzomenib</b>	<b>89.8</b>	<b>2.9</b>	<b>6.68</b>
Revumenib	106.9	3.13	9.27
Bleximenib	85.6	3.63	9.48
Ziftomenib	119.7	5.88	8.85

## Agenda

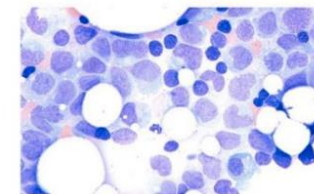
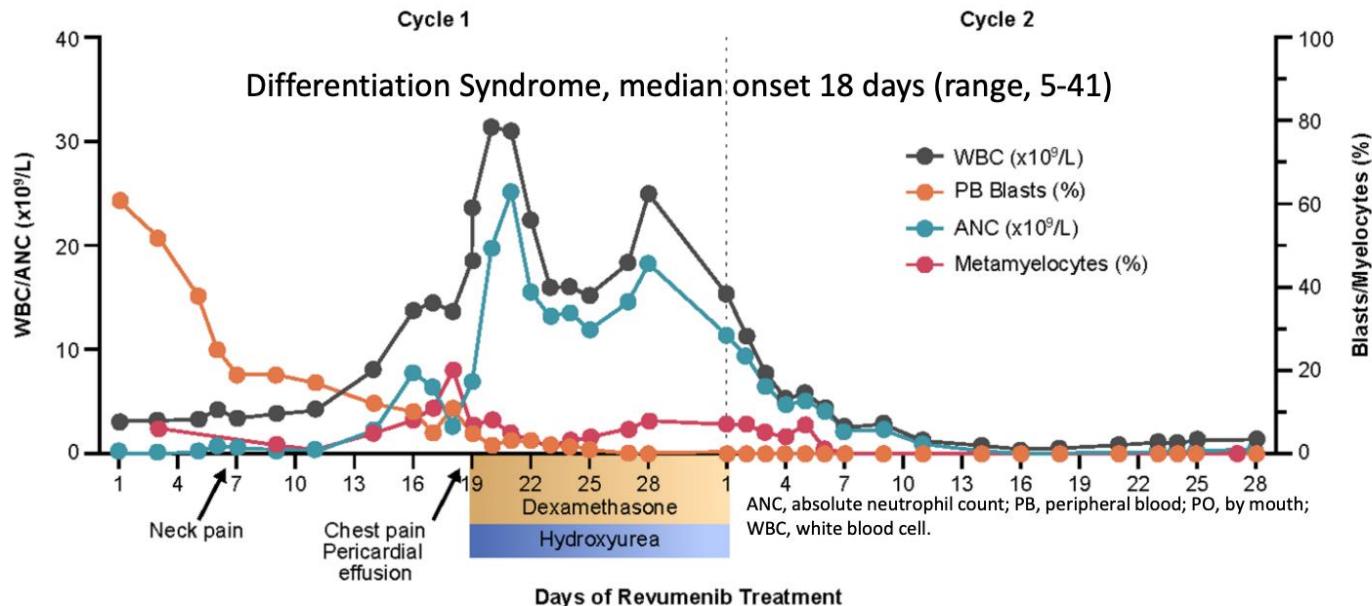
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- How can we overcome these mechanisms?
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## Safety profile

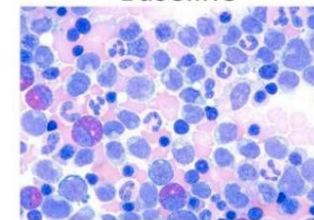
Agent (# pts)	Revumenib (n=199)	Ziftomenib (n=112)	Bleximenib (n=146)	Enzomenib (n=84)
Trial	Phase 2	Phase 1b/2	Phase 1	Phase 1
DLT (Y/N)	Yes	Yes	Yes	No
DLT	QTC prolongation	DS	DS	Not reported
DS (all)	KMT2Ar:27.7%; NPM1:19%	24%	14.4% all doses	10.7%
DS (≥Gr3)	13-16%	13%	6.8%	None reported
Febr Np (≥Gr3)	34.5-38.3%	22%	19.2%	23.8%
Neutrop (≥Gr3)	29.8%	Not reported	28%	17.9%
Thromb (≥Gr3)	16.7-23.4%	20%	36%	23.8%
QTc PR (any)	25.5%-42.9%	3%	0%	1%
QTc PR (≥Gr3)	13.8-22.6%	2%	0%	1%

*Issa GC et al JCO 2025; Wang ES et al JCO 2025; Searle E et al Blood 2024; Zeidner J et al Blood 2024*

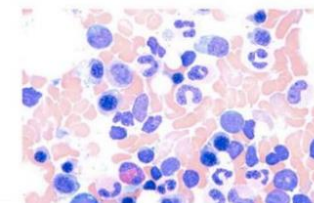
## Differentiation Syndrome



Baseline



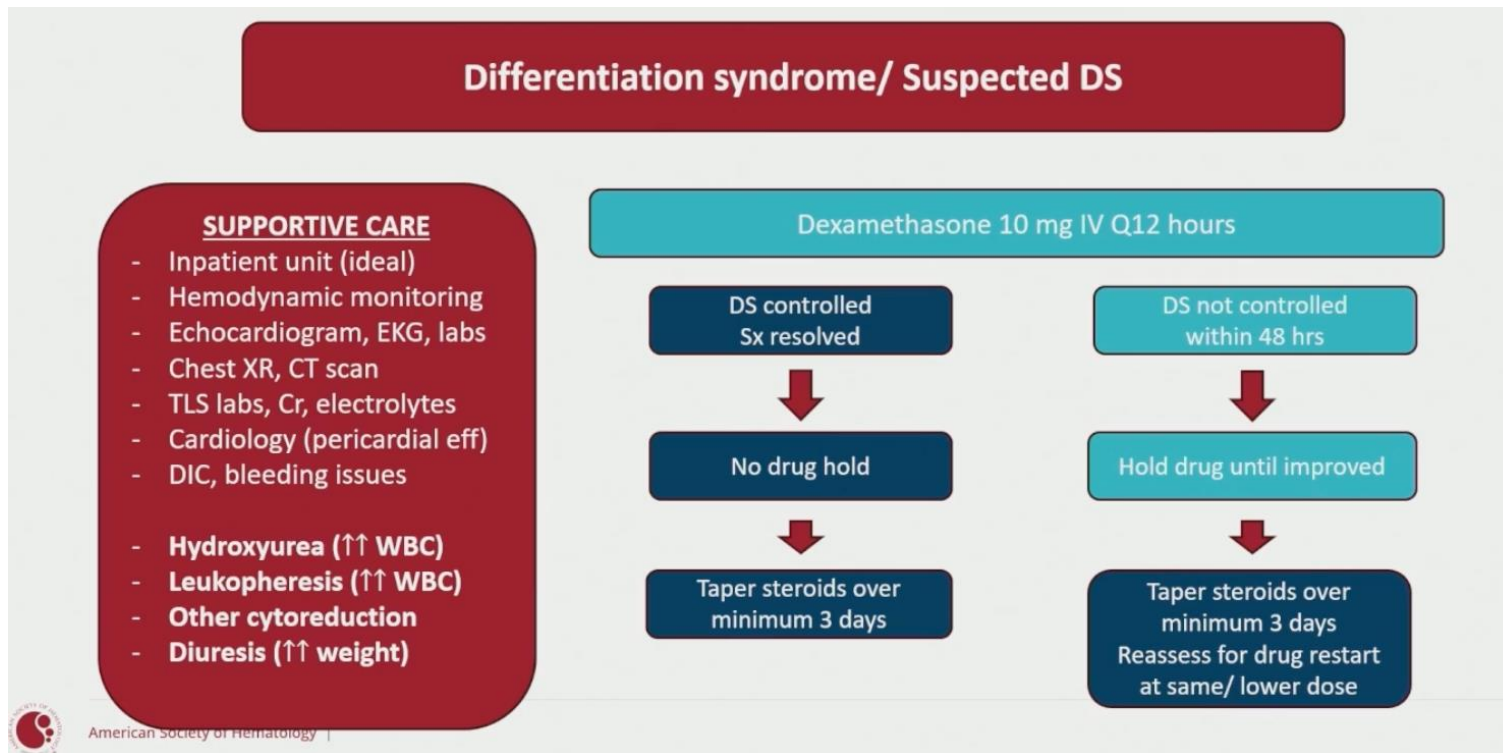
C2 D1



C3 D1

- 71-year-old with *KMT2Ar* AML relapsed after an allogeneic stem cell transplant
- Received revumenib at 339 mg PO q12h (Arm A), and achieved CRh, MRD negative

# How to manage Differentiation Syndrome



## Safety profile

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*Issa GC et al JCO 2025; Wang ES et al JCO 2025; Searle E et al Blood 2024; Zeidner J et al Blood 2024*

## Agenda

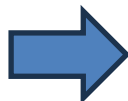
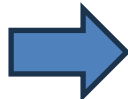
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## Menin inhibitors in R/R setting (single agents)



Menin inhibitor	Revumenib	Ziftomenib	Bleximenib	Enzomenib
Phase Trial	Ph 2	Ph 1b/2	Ph 1	Ph 1
Dose/frequency	163 mg bid	600 mg qd	90-100 mg bid	200-300 mg bid
KMT2Ar (n)	97	None	15	23
NPM1m (n)	83	93	14	17
CR	KMT: 15.5%; NPM1: 18.8%	NPM1: 18%	N/A	N/A
CR/CRh	KMT: 22.7%; NPM1m: 23.4%	NPM1: 35%	38%	37.5%
ORR (CRc+PR + MLFS)	KMT: 63.9%; NPM 46.9%	NPM1: 35%	47%	62.5%
MRD-neg CR/CRh	KMT: 61%; NPM: 64%	NPM1: 65%	N/A	N/A
Median DOR CR/CRh	KMT: 6.4; NPM1: 4.7 mo	NPM1: 3.7 mo	6 mo	KMT: NR; NPM1 7.0mo
Median OS (mo)	KMT: 8.0; NPM1: 4.0 mo	NPM1: 6.1 mo	N/A	N/A

*Aldoss I et al EHA abstract 2025; Arellano et al EHA abstract 2025; Wang E et al JCO Nov 2025; Zeidner J et al ASH abstract 2025*



## Enzomenib response rates

### KMT2Ar AL (n = 39)

- Optimal RP2D as monotherapy is 300 mg po bid (n=15)
- At RP2D
  - Overall Response rate (CR/CRh/CRi/MLFS) 73.3%
  - Composite CR rate (CR/CRh/CRi) 60%
  - CR + CRh rate 40%

### NPM1m AML (n = 25)

- Dose optimization is ongoing at 200, 300 and 400 mg po bid and initial activity is similar across dose levels

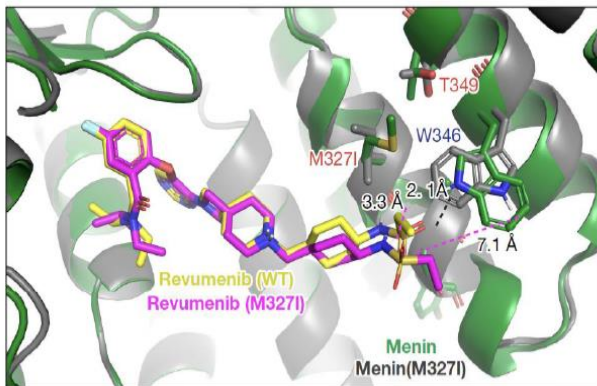
Response Category*	NPM1m		
	200 mg BID (n = 10)	300 mg BID (n = 7)	400 mg BID (n = 8)
Overall Response rate (CR/CRh/CRi/MLFS)	60% (6/10)	57.1% (4/7)	37.5% (3/8)
Composite CR rate (CR/CRh/CRi)	50% (5/10)	42.9% (3/7)	37.5% (3/8)
CR/CRh rate	50% (5/10)	42.9% (3/7)	37.5% (3/8)

Daver N et al, ASH 2025

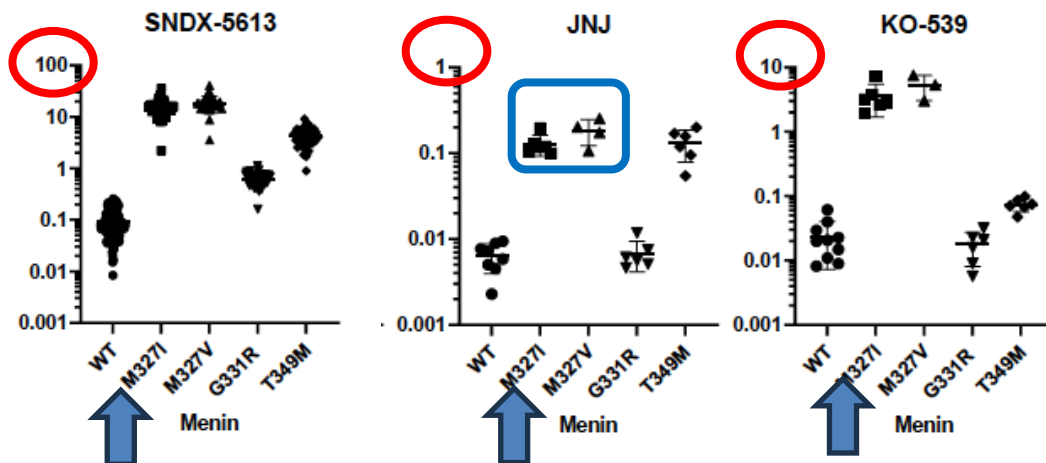
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- How can we overcome these mechanisms?
- For which patients, tomorrow?

# MEN1 mutations generate structural perturbations that impact small-molecule binding



Loss of hydrogen bond interactions between revumenib in the **M327I mutant** and the menin protein



- Binding affinities of all compounds reduced by I/V mutations at M327
- T349M reduces binding for most chemotypes
- G331R change has variable effects across chemotypes

## Other resistance mechanisms

Resistance mechanism	Genetic/epigenetic alterations	Consequences	Frequency/detection method	Therapeutic strategy
Mutations in menin-binding domain	<b>MEN1 mutations (e.g. M3271, T349M, G331R, E368K)</b>	Reduced menin inhibitor binding	39-40% after revumenib by RNA sequencing	<b>Alternative menin inhibitors; combination therapy</b>
Alternative proliferation pathways	<b>RAS pathway mutations (NRAS, KRAS, PTPN11)</b>	Leukemic proliferation indept of menin inhibition	Common acquired pathways by DNA/RNA NGS	<b>Combination therapy with MAPK, FLT3, or other inhibitors</b>
Apoptotic and differentiation resistance	<b>TP53 mutations, MCL-1 upregulation, BCL-xL</b>	Reduced apoptosis and differentiation response	Frequently co-occurring in high-risk AML	<b>Combine with MCL-1 inhibitors or or BCL-2 inhibitors</b>
Epigenetic reprogramming and MYC activation	<b>Loss of PRC1.1, MYC upregulation</b>	Blocked differentiation, altered epigenetics	Identified by chromatin & transcriptome assays	<b>Combine with epigenetic modifiers</b>

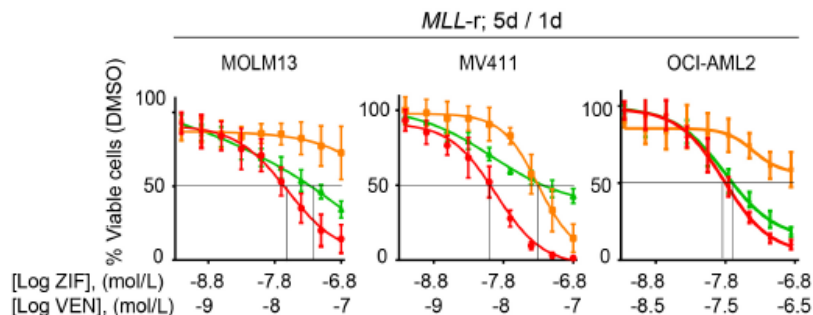
*Thakur R and Wang ES. ASH Education Bk 2025*

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## Preclinical data support menin inhibitors combinations in r-*KMT2A* leukemia

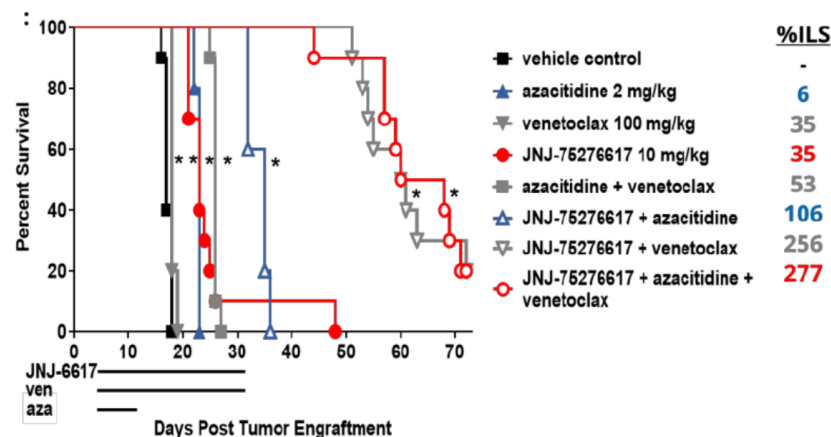
### Ziftomenib



- Ziftomenib
- Venetoclax
- Combination

Rausch J et al, Haematologica 2023

### JNJ-75276617



Chul Kwon M et al, ASH 2023

## Menin inhibitors partners

### Conventional or approved agents

- Standard high or low intensity chemotherapy regimens
- Hypomethylating agents<sup>1</sup>
- BCL2 inhibitors<sup>1,2</sup>
- FLT3 inhibitors<sup>3,4,5</sup>
- IDH inhibitors<sup>6</sup>
- Blinatumomab (CD3;CD19)
- Inotuzumab (ADC: CD22)
- Gemtuzumab (ADC: CD33)
  
- Revlimid<sup>7</sup>
- [...]

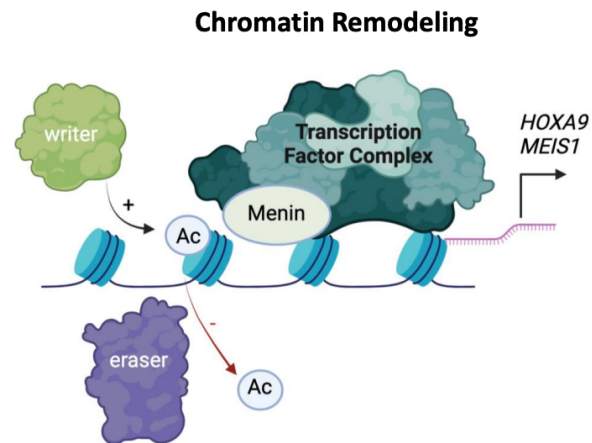
Writers

Readers

Erasers

### Investigational agents/targets

- DOT1L<sup>8,9</sup>
- CBP/p300<sup>10</sup>
- KAT6
- HBO1
- BRD4<sup>10</sup>
- ENL YEATS
- LEDGF
- SGF29
- HDAC<sup>11</sup>
- LSD1
- CDK4/6<sup>12</sup>
- CDK9
- SWI/SNF
- XPO1<sup>13,14,15</sup>
- [...]



### References indicate preclinical evidence of combination with menin inhibition

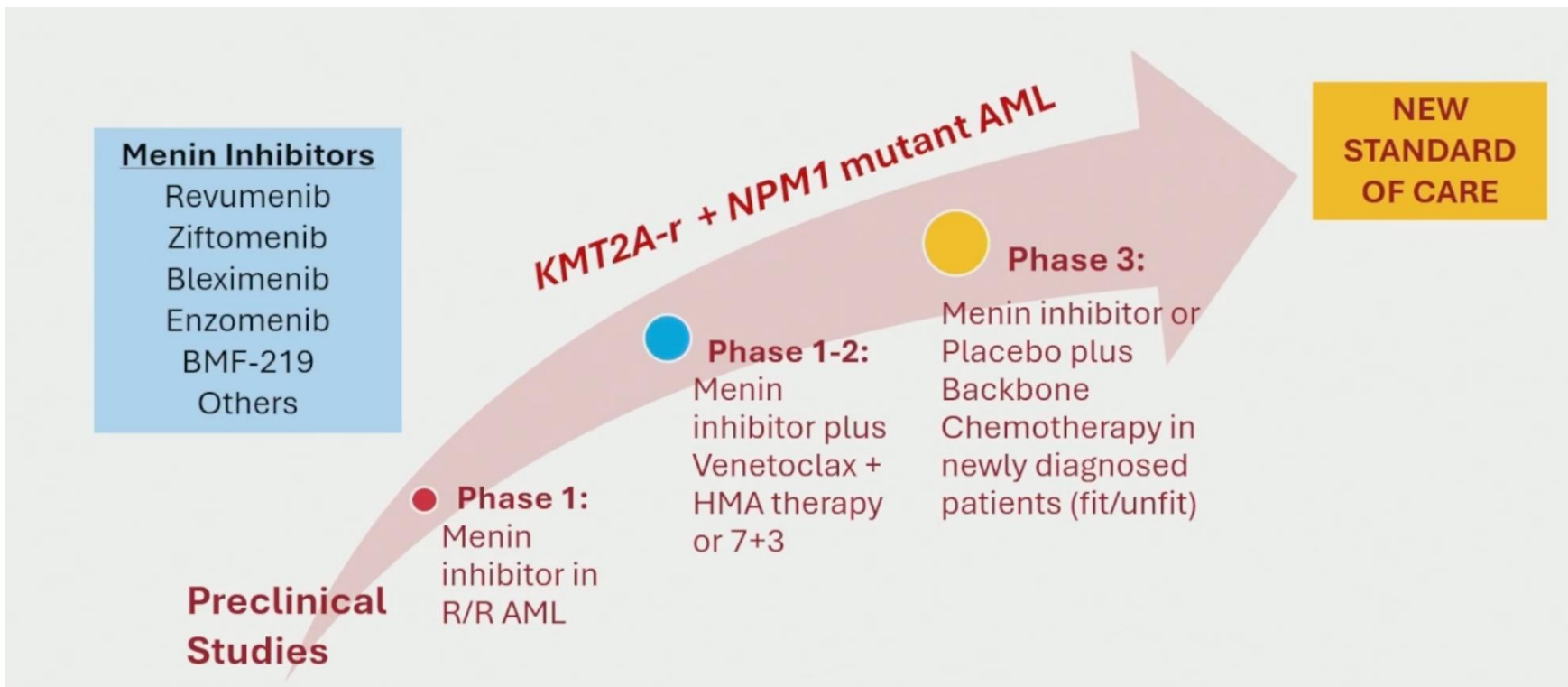
<sup>1</sup>Carter B. et al. Blood (2021) 138 (17): 1637–1641;2; <sup>2</sup>Fiskus W et al. Blood Cancer J. 2022 Jan 11;12(1):5; <sup>3</sup>Dzama MM et al. Blood. 2020;136(21):2442-56. <sup>4</sup>Miao H et al. Blood. 2020;136(25):2958-63. <sup>5</sup>Carter B et al. Haematologica. 2023 Feb 2. <sup>6</sup>Yoon JJ et al. Blood (2022) 140 (Supplement 1): 5946. <sup>7</sup>Aubrey BJ et al. Nat Cancer. 2022 May;3(5):595-613. <sup>8</sup>Kühn MW et al. Cancer Discov. 2016 Oct;6(10):1166-1181. <sup>9</sup>Dafflon C et al. Leukemia. 2017;31:1269–77. <sup>10</sup>Fiskus, W. Et al. Blood Cancer J. 13, 53 (2023). <sup>11</sup>Mill CP et al. Leukemia. 2023 Mar 28. <sup>12</sup>Soto-Feliciano et al Cancer Discov 1 January 2023; 13 (1): 146–169 <sup>13</sup>Brunetti L et al. Cancer Cell. 2018 Sep 10;34(3):499-512.e9. <sup>14</sup>Uckelmann et al. Cancer Discov 1 March 2023; 13 (3): 724–745. <sup>15</sup>Wang XQD et al. Cancer Discov 1 March 2023; 13 (3): 746–765.

## Menin inhibitors in triplets for R/R AML patients

Agent (# pts)	Bleximenib + Ven/Aza	Revumenib + Ven/Aza (SAVE)	Ziftomenib + Ven/Aza (KOMET-007)	Enzomenib + Ven/Aza
Trial	Phase 1	Phase 1	Phase 1	Phase 1
Number Pts	N=125	N =33	N=83	N=22
Start Menin Inhib	C1 D4	C1 D1	C1 D8	C1 D1
DS	6% (5 Gr3)	9% ((1 Gr3)	1%	10% (no Gr3)
QTc prolongation	4%	64%; 9% ≥Gr3	0%	10% (no Gr 3)
CR	Not reported	39%	27%/6%	Not reported
CR/CRh	40.9%	48%	NPM 40%/KMT 22%	Not reported
CRc (CR/CRh/CRi)	59.1%	60%	NPM 48%/KMT 28%	50%
ORR	81.8%	82%	NPM 65%/KMT 41%	77%

*Wei et al EHA 2025; Issa et al ASH 2024; Issa G et al ASH 2025; Watts J et al ASH 2025*

## Let's move Menin inhibitors to the first line

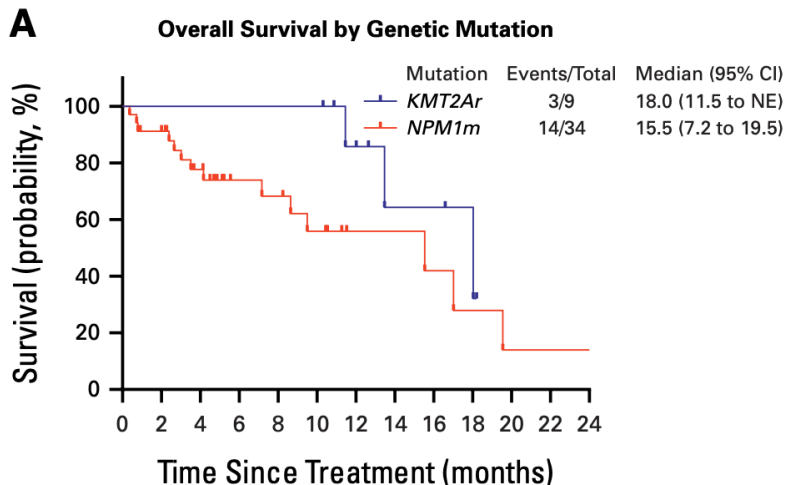


## Menin inhibitors in triplets for ND unfit AML patients

Agent (# pts)	Bleximenib + Ven/Aza	Revumenib + Ven/Aza (SAVE)	Ziftomenib + Ven/Aza (KOMET-007)
Trial	Phase 1	Phase 1	Phase 1
Number Pts	N=125	N =21	N=40
Start Menin Inhib	C1 D4	C1 D1	C1 D8
DS	6% (5 Gr3)	19% (10% Gr 3)	3% (no Gr3)
QTc prolongation	4%	43% (No $\geq$ Gr3)	3% (1 Gr 3)
CR	Not reported	76%	73%
CR/CRh	65%	81%	78%
CRc (CR/CRh/CRi)	75%	86%	86%
ORR	90%	86%	89%

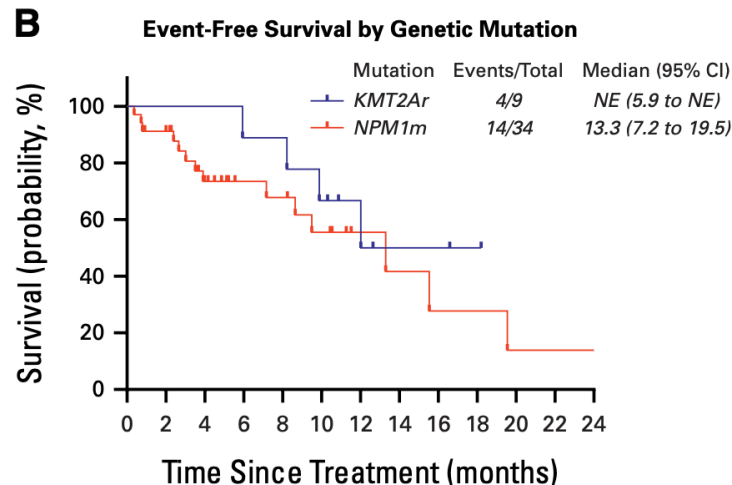
Wei A et al EHA 2025; Jen W et al ASH 2025; Roboz G et al ASH 2025

## Acacitidine+Venetoclax+Revumenib: Survival



Number at risk:

	0	2	4	6	8	10	12	14	16	18	20	22	24
<i>KMT2Ar</i>	9	9	9	9	9	9	6	3	3	2	0	0	0
<i>NPM1m</i>	34	30	19	13	12	9	4	3	2	2	1	1	1



Number at risk:

	0	2	4	6	8	10	12	14	16	18	20	22	24
<i>KMT2Ar</i>	9	9	9	8	8	6	4	2	2	1	0	0	0
<i>NPM1m</i>	34	30	19	13	12	9	4	3	2	2	1	1	1

SCT 23%

Zeidner J et al, JCO 2025

## Menin inhibitors in clinical development in 1L AML fit patients

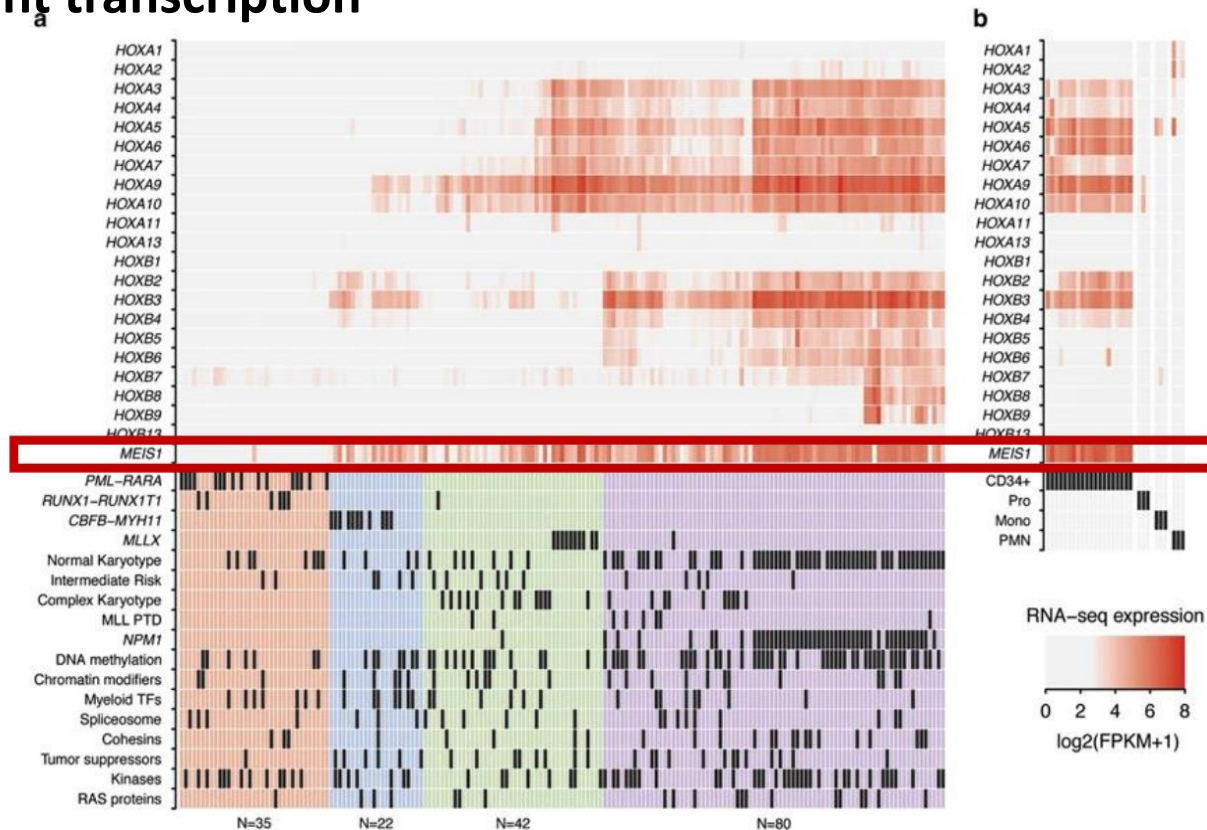
	Revumenib		Ziftomenib	Bleximenib
Trial	Ph 3 REVEAL-ND NPM1 NCT07211958	<i>Ph 2 RAVEN ISS (planned in H2'26)</i>	Ph 3 KOMET-017 NCT07007312	Ph 3 HOVON-181 NCT07223814
Mutation	mNPM1	<i>KMT2Ar</i>	mNPM1, KMT2Ar (one arm)	mNPM1, KMT2Ar (one arm)
Regimen	REV + '7+3'	<i>REV + VEN/AZA</i>	ZIF + '7+3'	BLE + '7+3'
Enrolment	468	<i>n/a</i>	1300	875

## Agenda

- For which patients today?
- How many menin inhibitors do we have today?
- What about safety with menin inhibitors as single agents?
- Which results can we achieve with menin inhibitors?
  - R/R single agents
- What about resistance?
- How can we overcome these mechanisms?
- For which patients, tomorrow?




# Targeting menin-dependent transcription


~ 60% of AML with upregulation of *HOX/MEIS* (associated with other genotypes in addition to *KMT2Ar* and *mNPM1*)





## Targeting menin-dependent transcription

**Table 2** Genetic alterations with overexpression of *HOXA* genes predicted to potentially respond to menin inhibitors.<sup>1</sup>

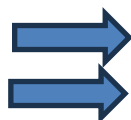
Alteration/mutation	Cytogenetics	Phenotype				References
<i>KMT2Ar</i>	11q23 rearrangements	AML, ALL, MPAL	✓	✓	✓	[26, 132, 133]
<i>KMT2A</i> -PTD	Normal karyotype	AML	✓	✓		[26, 134]
<i>NPM1c</i>	Normal karyotype	AML	✓	✓	✓	[26, 135]
<i>NPM1-MLF1</i>	t(3;5)(q25;q34)	MDS, AML	✓			[136, 137]
<i>NUP98r</i>	11p15 rearrangements	AML, T-ALL, MDS	✓	✓	✓	[122–124]
<i>SET-NUP214</i>	t(9;9)(q34;q34)	AML, T-ALL, AUL	✓		✓	[138]
<i>RUNX1-EVII</i>	t(3;21)(q26;q22)	AML	✓		✓	[139]
<i>MYST3-CREBBP</i>	t(8;16)(p11;p13)	AML	✓			[140]
<i>CDX2-ETV6</i>	t(12;13)(p13;q12)	AML		✓		[141]
<i>CALM-AF10</i>	t(10;11)(p13;q14-21)	T-ALL, AML, MPAL	✓	✓	✓	[142–144]
<i>MN1-ETV6</i>	t(12;22)(p13;q12)	AML, MDS		✓	✓	[145]
<i>EZH2</i>	–	MDS, AML	✓			[146]
<i>IDH1/IDH2</i>	–	MNs			✓	[147, 148]
<i>ASXL1</i>	–	MNs		✓		[149]
<i>CEBPA</i>	–	AML			✓	[150]
	Trisomy 8	MNs	✓			[151]

 Denotes direct examination of patient samples with the corresponding genotype showing upregulation of *HOXA* genes.

 Denotes mouse models of the corresponding genotype leading to upregulation of *Hox* genes.

 Denotes examination of cells lines or other in vitro investigations demonstration a role of *HOX* genes or menin inhibition in the corresponding genotype.

One patient with AML (*SETD2*, *RUNX1*) attained CR, MRD+ with ziftomenib 100 mg.<sup>2</sup>



## Menin inhibitors: take home messages (I)

- **At least 4 agents in development, 2 FDA approved (Revumenib, Ziftomenib)**
- Clinical activity as single agent in R/R AML with NPM1mut and r-KMT2A:
  - ✓ 30-60% ORR
  - ✓ Duration of response: 5-7 months
  - ✓ Overall survival: 6-8 months
- On target drug effect: **Differentiation syndrome**
- Are there **differences between agents**? Yes, due to:
  - ✓ Drug: Half-lives, tissue penetration,azole interactions, QTc prolongation

## Menin inhibitors: take home messages (II)

- **Triplets** are promising
- Reduced DS compared to monotherapy
- R/R: **60-80% ORR**
- ND: **80-90% ORR**
  
- Phase 3 randomized trials ongoing (Ava-Ven/3+7)
- Combination with other targeted agents?
- Efficacy in **other leukemia subtypes?**
- Sequential menin inhibitors treatment? Role of MRD?



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## Thank you!

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Emanuela Ottaviani

Carolina Terragna

Cecilia Monaldi

Sara de Santis

Valentina Robustelli

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