

MENIN INHIBITORS vs FLT3 INHIBITOR-BASED THERAPY

IN NPM1-MUTATED RELAPSED/REFRACTORY AML: COMPARATIVE EFFICACY, MRD DEPTH, AND SAFETY PROFILES

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BACKGROUND

NPM1-mutated relapsed/refractory AML is driven by HOXA/MEIS transcriptional addiction and frequent FLT3 co-mutation. Two therapeutic paradigms have emerged: FLT3 inhibitor (FLT3i)-based kinase blockade and menin inhibition (MENi) targeting transcriptional dependency.



OBJECTIVE

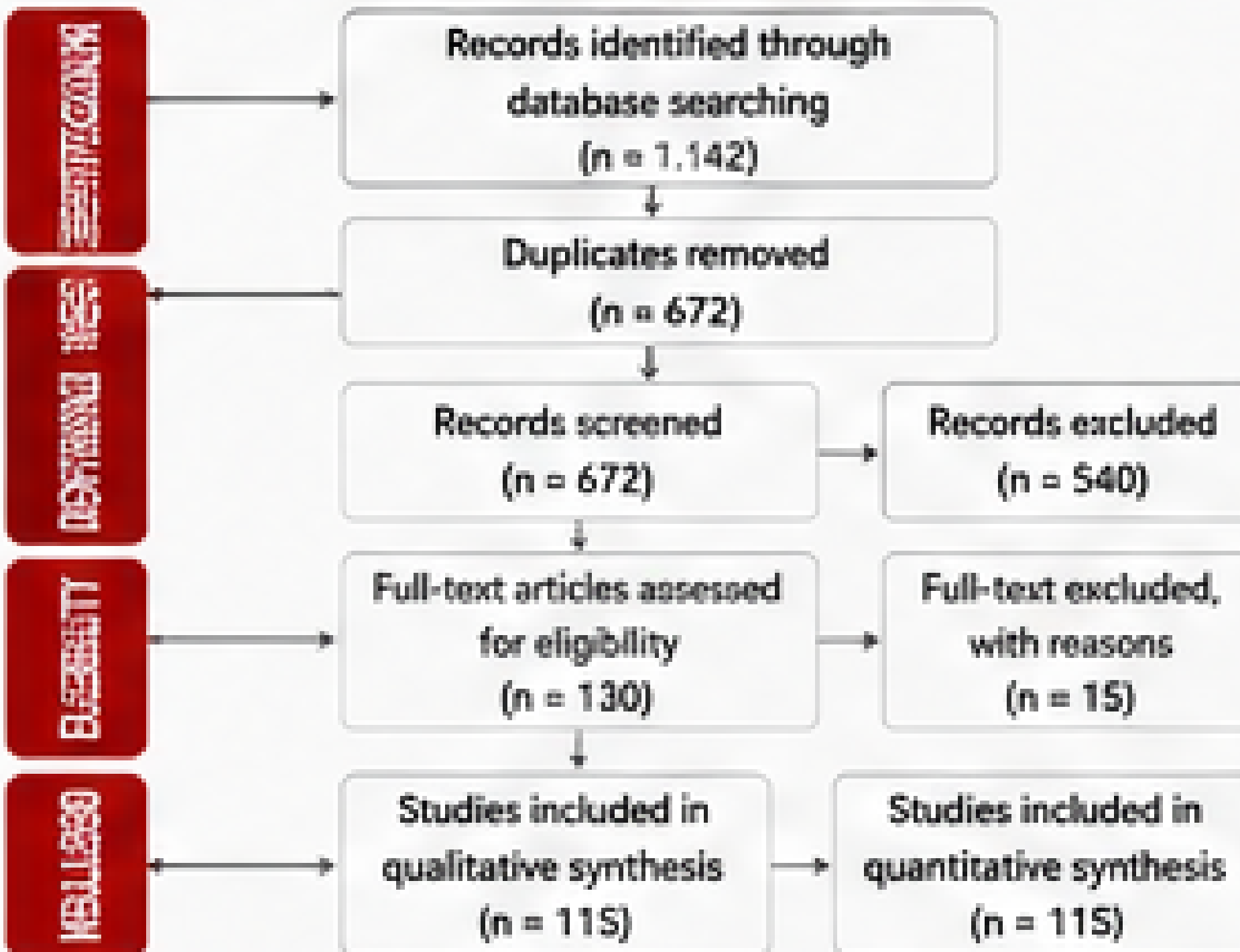
To compare efficacy and safety of MENi vs FLT3i-based regimens in NPM1-mutated relapsed/refractory AML using a systematic review of early clinical data.



METHODS

- Systematic review of PubMed and Embase (2019–2026) following PRISMA guidelines
- Screening performed using Rayyan.ai
- Phase 1/2 trials and retrospective cohorts reporting NPM1-mutated outcomes included
- Median follow-up: 4–18 months
- Data synthesized using descriptive study-level aggregation
- Settings: Multicenter academic and tertiary referral centers across North America, Europe, and Asia

PRISMA FLOW DIAGRAM



CONCLUSIONS

MENi and FLT3i-based regimens demonstrate comparable remission rates in NPM1-mutated relapsed/refractory AML but differ in response depth. MENi shows higher MRD negativity. FLT3i provides broader disease control with higher infectious burden, while MENi is associated with differentiation syndrome. These data support optimized sequencing strategies in NPM1-mutated AML.

STUDY AT A GLANCE



115

TOTAL PATIENTS



68

MENi

(Revumenib, Ziftomenib)

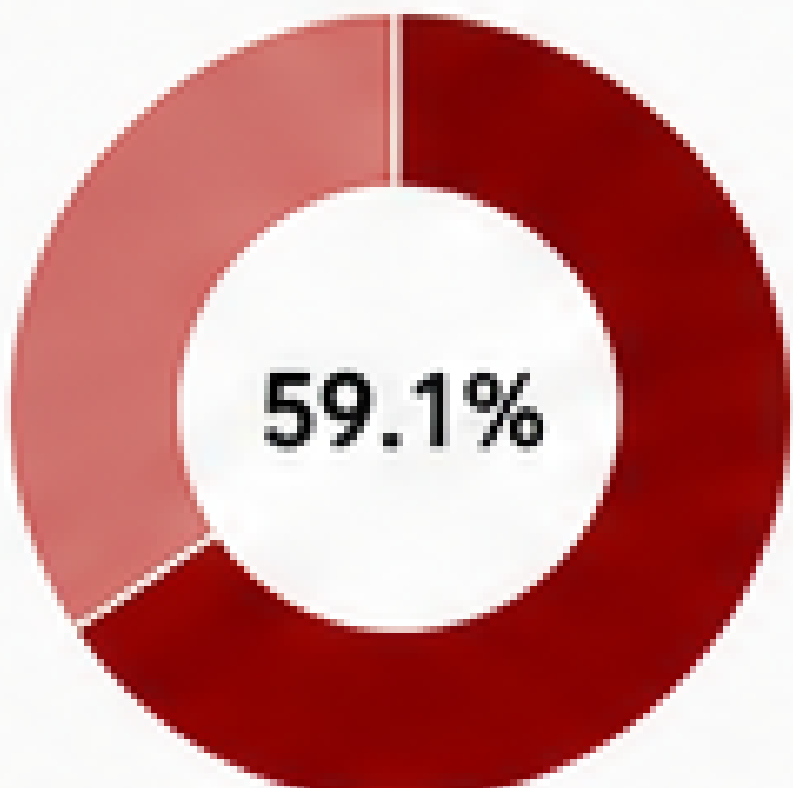


47

FLT3i-BASED

(Gilteritinib ± combo)

PATIENTS & DISEASE CHARACTERISTICS



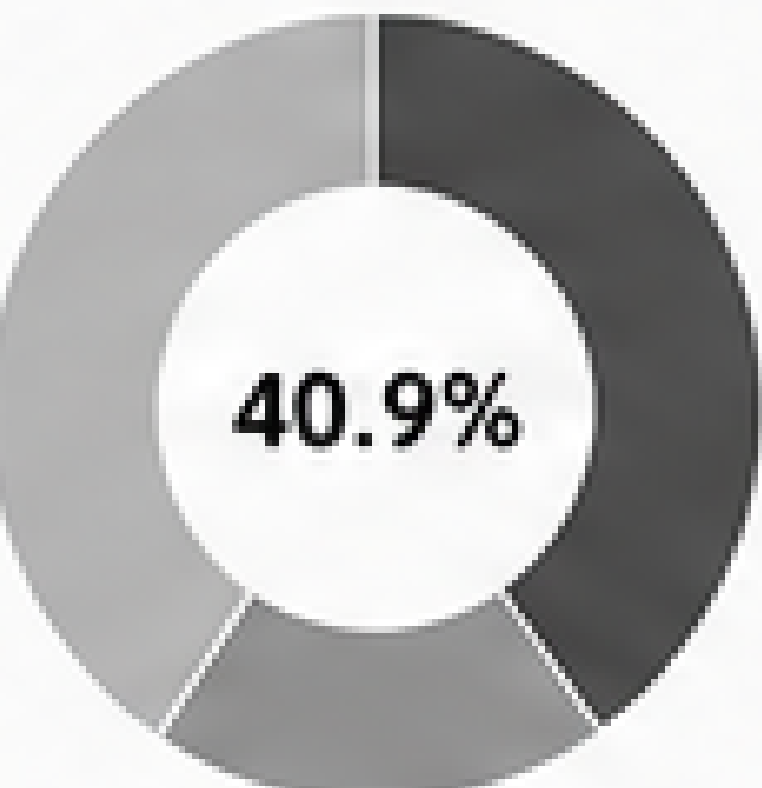
MENi (n = 68)



115

TOTAL PATIENTS

NPM1-MUTATED RELAPSED/REFRACTORY AML



FLT3i-Based (n = 47)

PATIENT CHARACTERISTICS (OVERALL)



Median Age
61
years
(range 22–78)



61.2%

Median Prior
Lines of Therapy

2

(range 1–5)

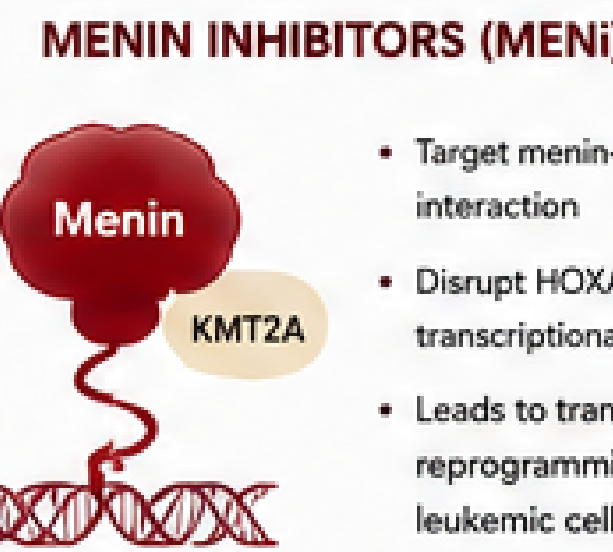
Median Follow-up

10.2

months
(range 4–18)

MECHANISM OF ACTION

MENIN INHIBITORS (MENi)



• Target menin-KMT2A/HOXA interaction
• Disrupt HOXA/MEIS-driven transcriptional dependency
• Leads to transcriptional reprogramming and leukemic cell death

FLT3 INHIBITOR-BASED (FLT3i)

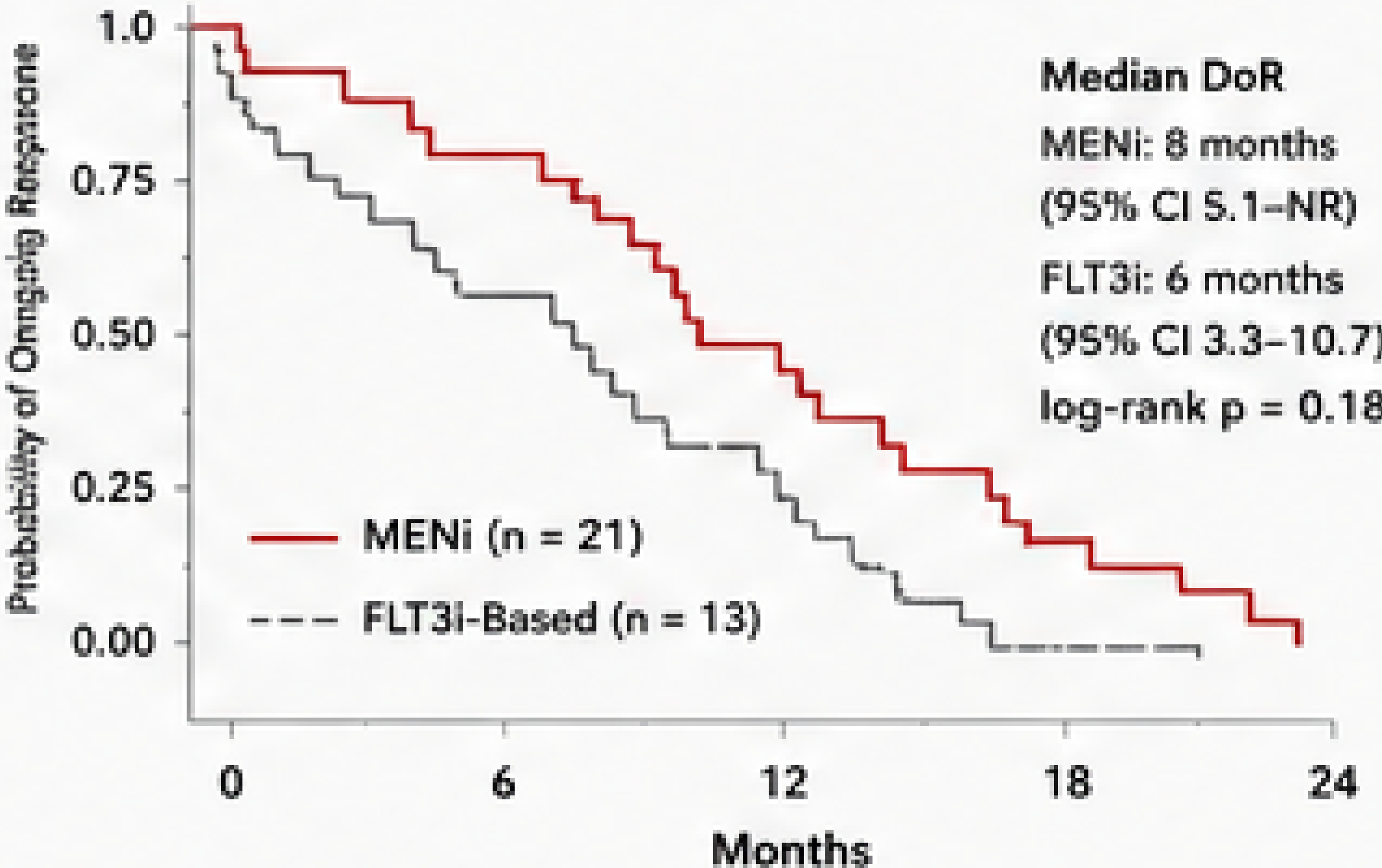


• Inhibit FLT3 signaling pathway
• Block proliferation and survival signals
• Result in disease control

EFFICACY OUTCOMES

MENi (n = 68)	OUTCOME	FLT3i-BASED (n = 47)
32/68 (47.1%)	Overall Response Rate (ORR)	20/47 (42.6%)
21/68 (30.9%)	Complete Response/CRi	13/47 (27.7%)
12/21 (57.1%)	MRD Negativity (among evaluable responders)	6/20 (30.0%)
8 months	Duration of Response (DoR, median)	6 months

DURATION OF RESPONSE (DoR)



No. at risk	0	6	12	18	24
MENi	21	16	11	6	3
FLT3i	13	9	6	3	1

SAFETY & TOXICITY

INCIDENCE OF GRADE ≥3 ADVERSE EVENTS

■ MENi (n = 68) ■ FLT3i-Based (n = 47) p-value



Any Grade ≥3
Adverse Event

48.5%
53.2%

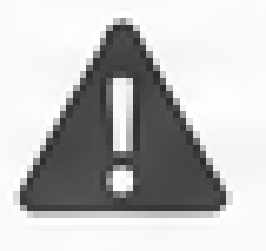
p = 0.60



Infectious Events
(Grade ≥3)

29.4%
40.4%

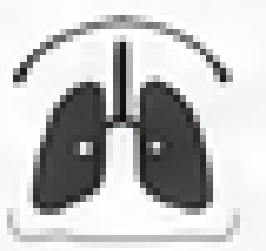
p = 0.21



Differentiation
Syndrome

13.2%
0%

p = 0.004



Hematologic Toxicity
(Grade ≥3)

26.5%
27.7%

p = 0.88

OTHER SAFETY OUTCOMES

TREATMENT DISCONTINUATION RATE



7.4% (5/68)
8.5% (4/47)

ICU ADMISSIONS (GRADE ≥3 EVENTS)



5.9% (4/68)
8.5% (4/47)

p = 0.72

KEY TAKEAWAY



MENi and FLT3i-based regimens show comparable remission rates in NPM1-mutated relapsed/refractory AML but differ in response depth. MENi achieves deeper MRD negativity, while FLT3i is associated with higher infectious burden and broader disease control. MENi is associated with differentiation syndrome. These data define biologically distinct response trajectories and support optimized sequencing strategies in NPM1-mutated AML.

DISCLAIMER: AI was used to assist with the creation of diagrams.

ABBREVIATIONS

AML: acute myeloid leukemia; CR: complete remission; CRi: complete remission incomplete; DoR: duration of response; FLT3: fms-related tyrosine kinase 3; HOXA: homeobox A; MEIS: MEIS homeobox; MRD: measurable residual disease; NPM1: nucleophosmin 1; ORR: overall response rate; PRISMA: Preferred Reporting Items for Systematic Reviews and Meta-Analyses