

# Real-World Treatment Patterns and Outcomes Among Patients With Newly Diagnosed *NPM1*-Mutated Acute Myeloid Leukemia in the United States

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- Although there are large datasets focused on real-world outcomes in newly diagnosed *NPM1m* AML, real-world evidence that captures academic and community sites and longitudinally reports on relapse is limited<sup>1</sup>
- *NPM1* mutations occur in ~30% of adult AML cases<sup>2</sup>
  - Patients with *NPM1m* AML without *FLT3*-ITD mutations are assigned to the favorable-risk cohort,<sup>3</sup> but outcomes in R/R *NPM1m* AML are poor<sup>4</sup>
  - Less is known regarding long-term treatment outcomes among patients with newly diagnosed *NPM1m* AML treated in community and academic centers

The objective of this study is to describe treatment patterns and outcomes among patients with newly diagnosed *NPM1m* AML in the United States

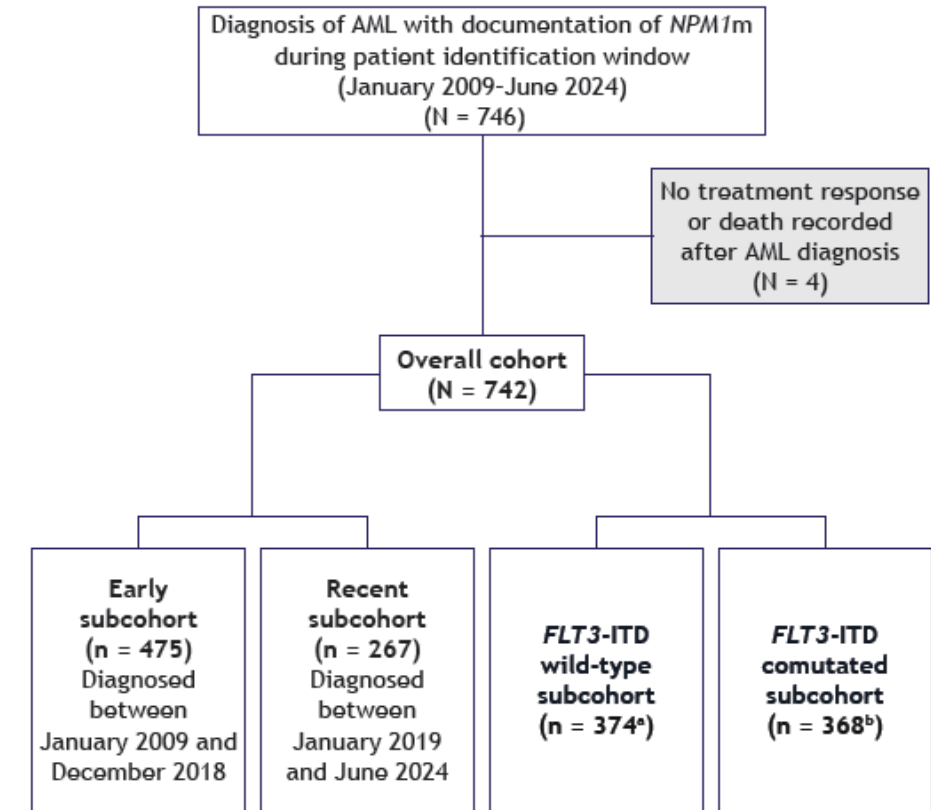
*FLT3*-ITD, fms-like receptor tyrosine kinase 3 internal tandem duplication.

1. Othman J, et al. *Blood*. 2024;144(7):714-728. 2. Papaemmanuil E, et al. *N Engl J Med*. 2016;374(23):2209-2221. 3. Döhner H, et al. *Blood*. 2022;140(12):1345-1377. 4. Issa GC, et al. *Blood Adv*. 2023;7(6):933-942.

# Study design

- This retrospective study used real-world data from COTA Healthcare collected from adults with an initial AML diagnosis between January 2009 and June 2024 and had documented *NPM1m* status
- Patients were grouped into subcohorts by diagnosis year to evaluate the impact of venetoclax approval in November 2018
  - Early: diagnosed between January 2009 and December 2018
  - Recent: diagnosed between January 2019 and June 2024
- Patients were also grouped into subcohorts by *FLT3*-ITD comutation status

Figure 1. Patient selection



<sup>a</sup>Without *FLT3*-ITD comutation at any time during follow-up or unknown. <sup>b</sup>With *FLT3*-ITD comutation at any time during follow-up. ITD, internal tandem duplication.

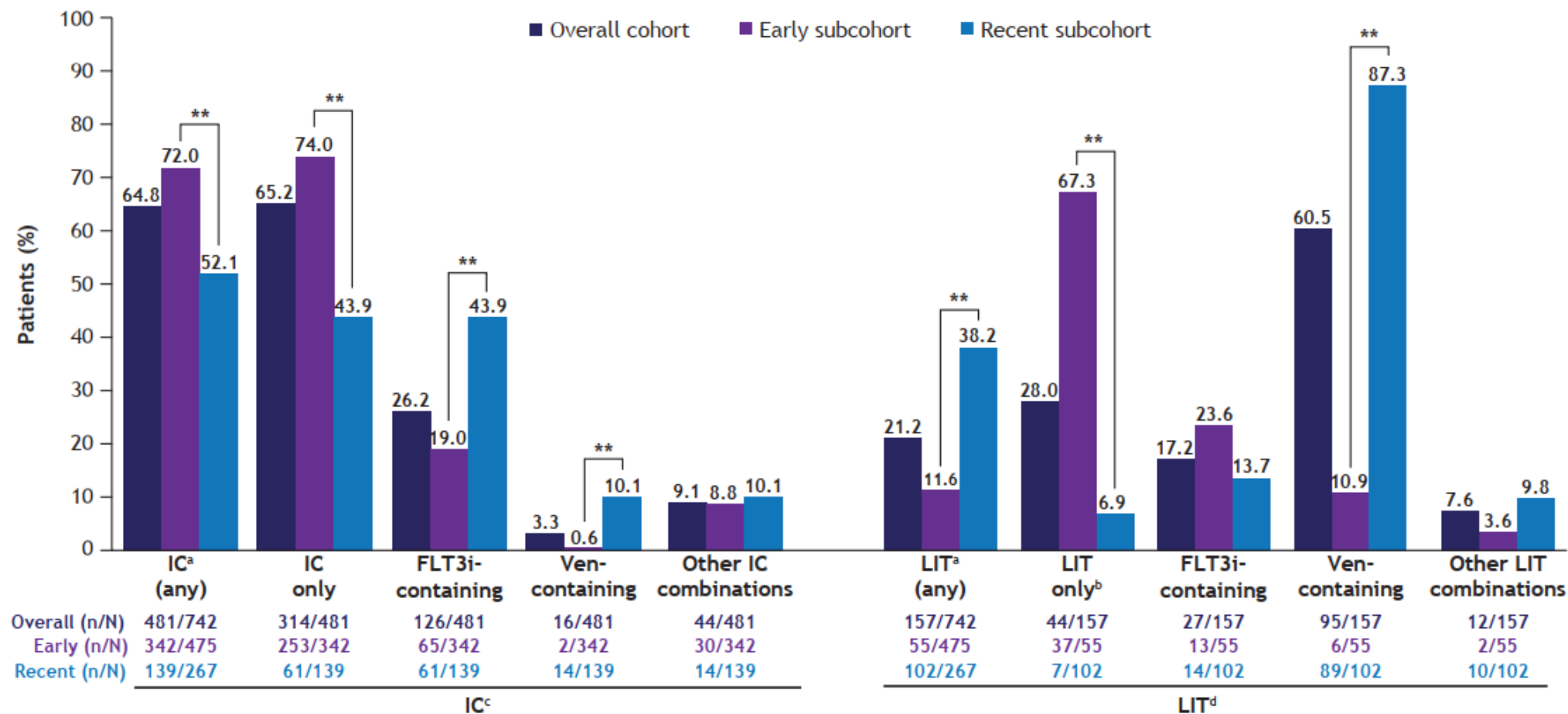
Table 1. Demographic and baseline characteristics<sup>a</sup>

|                                                                                            | Overall cohort<br>(N = 742) | Early subcohort<br>(N = 475) | Recent subcohort<br>(N = 267) | <i>FLT3</i> -ITD wild-type subcohort <sup>b</sup><br>(N = 374) | <i>FLT3</i> -ITD commutated subcohort<br>(N = 368) |
|--------------------------------------------------------------------------------------------|-----------------------------|------------------------------|-------------------------------|----------------------------------------------------------------|----------------------------------------------------|
| Age at initial AML diagnosis, median (IQR), y                                              | 62.0 (51.0-71.0)            | 60.0 (50.0-69.5)             | 66.0** (56.0-73.0)            | 63.0 (52.0-72.0)                                               | 61.0 (50.0-70.0)                                   |
| Age group , n (%)**                                                                        |                             |                              |                               |                                                                |                                                    |
| 19-59 y                                                                                    | 317 (42.7)                  | 227 (47.8)                   | 90 (33.7)                     | 149 (39.8)                                                     | 168 (45.7)                                         |
| ≥60 y                                                                                      | 425 (57.3)                  | 248 (52.2)                   | 177 (66.3)                    | 225 (60.2)                                                     | 200 (54.3)                                         |
| Male, n (%)                                                                                | 322 (43.4)                  | 205 (43.2)                   | 117 (43.8)                    | 176 (47.1)                                                     | 146 (39.7) <sup>†</sup>                            |
| Race, n (%)                                                                                |                             |                              |                               |                                                                |                                                    |
| White                                                                                      | 570 (76.8)                  | 376 (79.2)                   | 194 (72.7)                    | 281 (75.1)                                                     | 289 (78.5)                                         |
| Black or African American                                                                  | 34 (4.6)                    | 24 (5.1)                     | 10 (3.7)                      | 15 (4.0)                                                       | 19 (5.2)                                           |
| Asian                                                                                      | 30 (4.0)                    | 17 (3.6)                     | 13 (4.9)                      | 17 (4.5)                                                       | 13 (3.5)                                           |
| Other/multiple/missing                                                                     | 108 (14.6)                  | 58 (12.2)                    | 50 (18.7)                     | 61 (16.3)                                                      | 47 (12.8)                                          |
| Clinician practice setting at start of AML treatment, n (%)                                |                             |                              |                               |                                                                |                                                    |
| Academic                                                                                   | 452 (60.9)                  | 287 (60.4)                   | 165 (61.8)                    | 227 (60.7)                                                     | 225 (61.1)                                         |
| Community                                                                                  | 290 (39.1)                  | 188 (39.6)                   | 102 (38.2)                    | 147 (39.3)                                                     | 143 (38.9)                                         |
| Presence of positive comutations, n (%)                                                    |                             |                              |                               |                                                                |                                                    |
| <i>FLT3</i> -ITD                                                                           | 368 (49.6)                  | 243 (51.2)                   | 125 (46.8)                    | 0                                                              | 368 (100) <sup>‡</sup>                             |
| <i>FLT3</i> -TKD                                                                           | 110 (14.8)                  | 70 (14.7)                    | 40 (15)                       | 57 (15.2)                                                      | 53 (14.4)                                          |
| <i>DNMT3A</i>                                                                              | 214 (28.8)                  | 113 (23.8)                   | 101 (37.8)**                  | 97 (25.9)                                                      | 117 (31.8)                                         |
| <i>IDH2</i>                                                                                | 118 (15.9)                  | 61 (12.8)                    | 57 (21.3)**                   | 69 (18.4)                                                      | 49 (13.3)                                          |
| <i>IDH1</i>                                                                                | 103 (13.9)                  | 56 (11.8)                    | 47 (17.6)*                    | 69 (18.4)                                                      | 34 (9.2) <sup>‡</sup>                              |
| <i>NRAS</i>                                                                                | 75 (10.1)                   | 38 (8)                       | 37 (13.9)*                    | 46 (12.3)                                                      | 29 (7.9) <sup>†</sup>                              |
| Duration of follow-up from AML diagnosis to death or end of study period, median (IQR), mo | 23.8 (9.5-60.2)             | 29.0 (11.6-78.8)             | 17.4 (7.9-39.2)               | 32.7 (12.2-65.1)                                               | 19.7 (8.3-55.4)                                    |

\**P* <0.05 between Early and Recent subcohorts. \*\**P* <0.01 between Early and Recent subcohorts. <sup>†</sup>*P* <0.05 between *FLT3*-ITD wild-type and comutated subcohorts. <sup>‡</sup>*P* <0.01 between *FLT3*-ITD wild-type and comutated subcohorts.

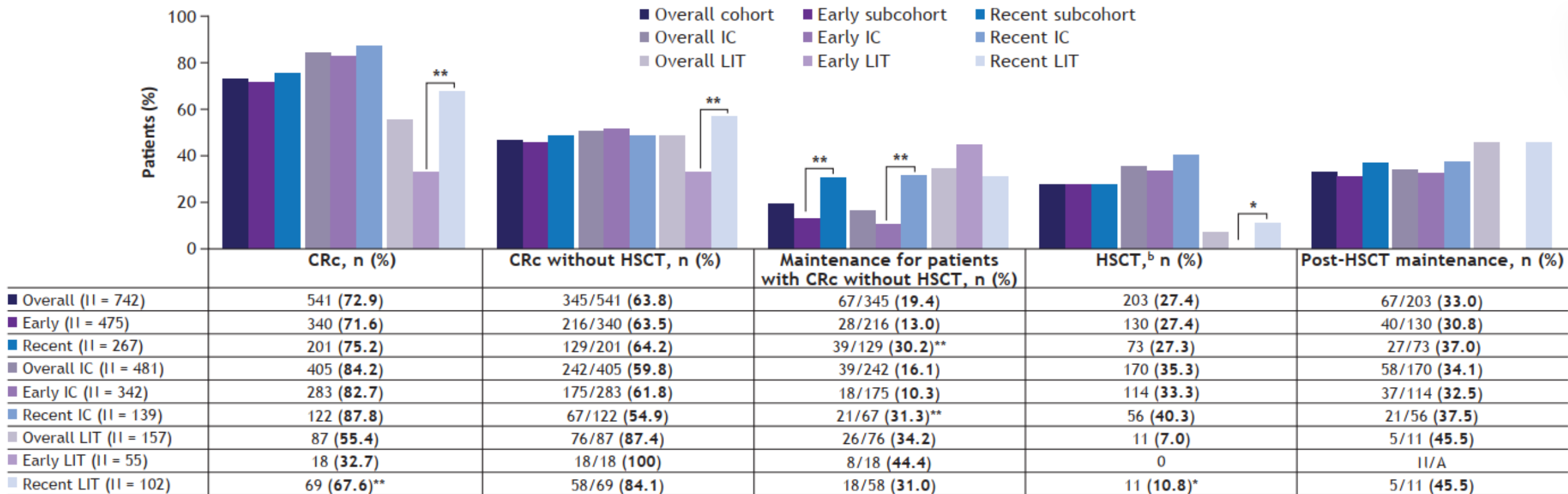
<sup>a</sup>At initial AML diagnosis. <sup>b</sup>Documented negative or unknown, or undocumented testing.

IQR, interquartile range; TKD, tyrosine kinase domain.



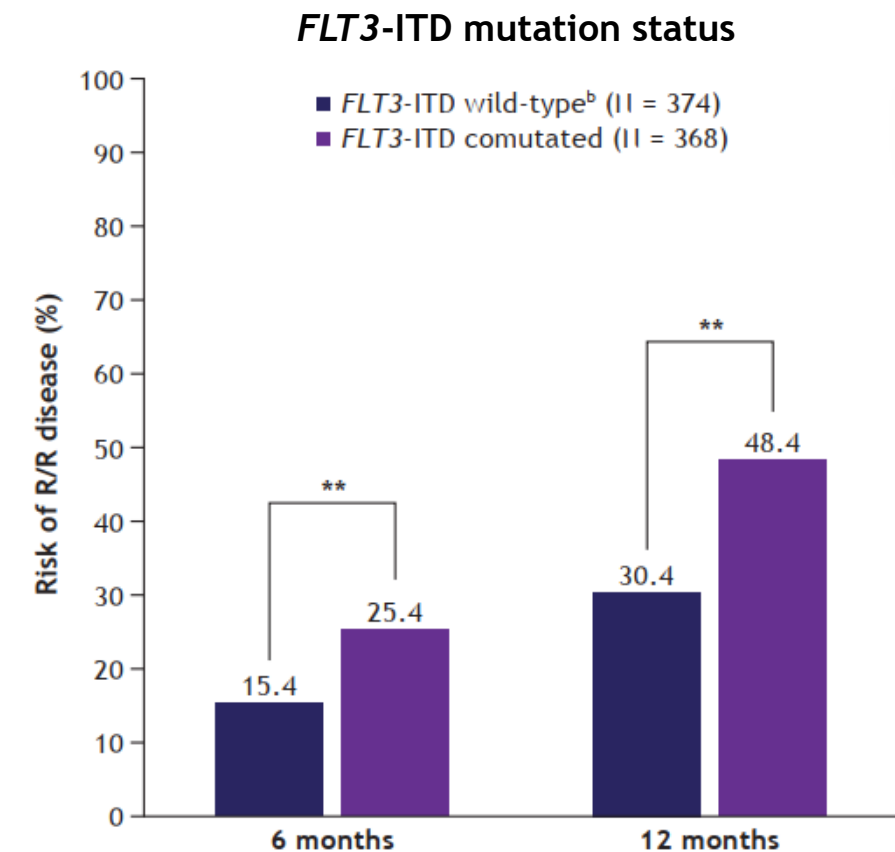
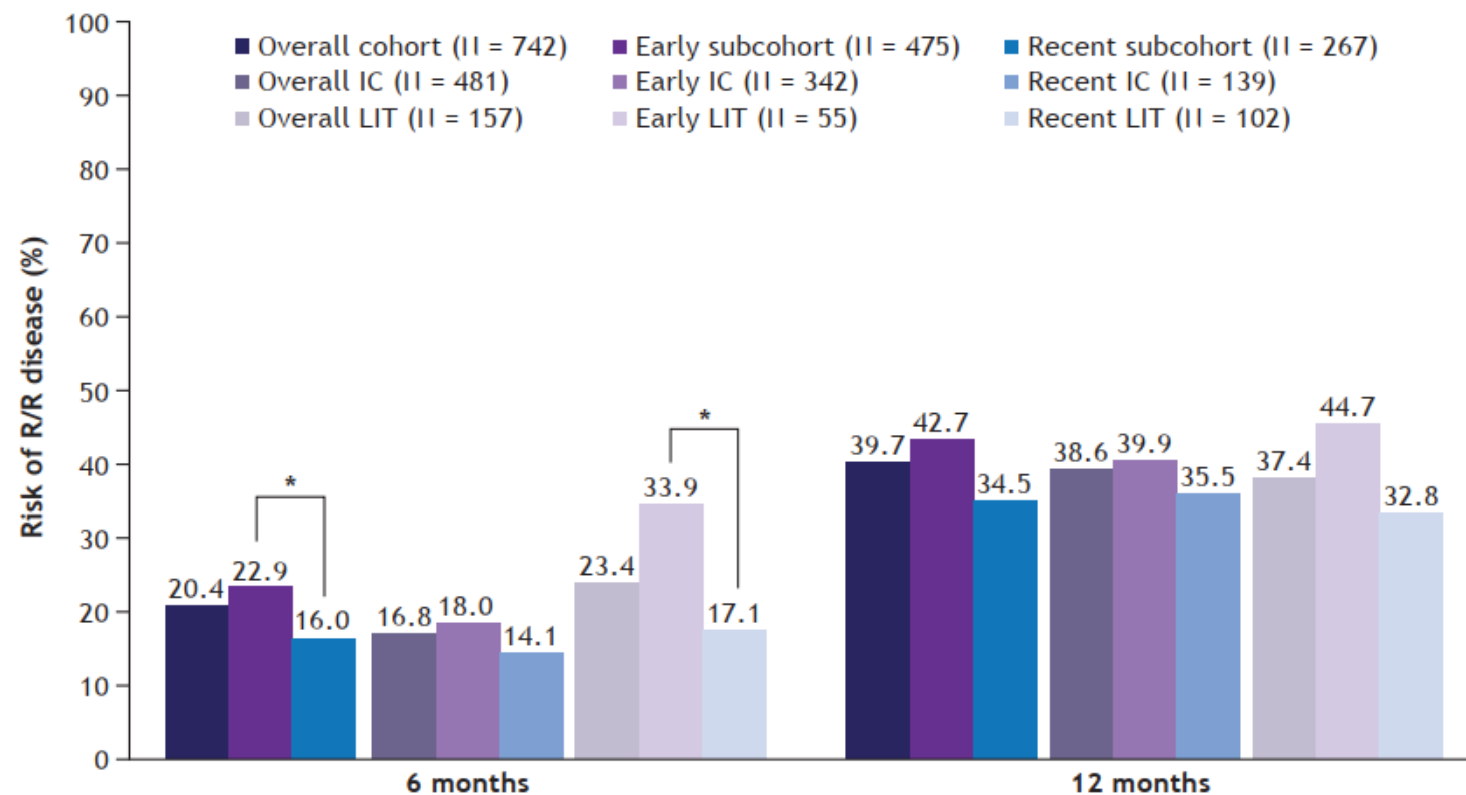
\*\*P < 0.01 between Early and Recent subcohorts.

<sup>a</sup>Subcategories are not mutually exclusive (eg, patients could have received both an FLT3i and venetoclax). <sup>b</sup>LIT only contains hypomethylating agent, gemtuzumab, or LDAC. <sup>c</sup>IC included HiDAC, fludarabine + cytarabine + granulocyte colony-stimulating factor ± idarubicin, and mitoxantrone + etoposide + cytarabine. <sup>d</sup>LIT included venetoclax-based regimens, targeted and hypomethylating agents, and LDAC. HiDAC, high-dose cytarabine; LDAC, low-dose cytarabine; ven, venetoclax.



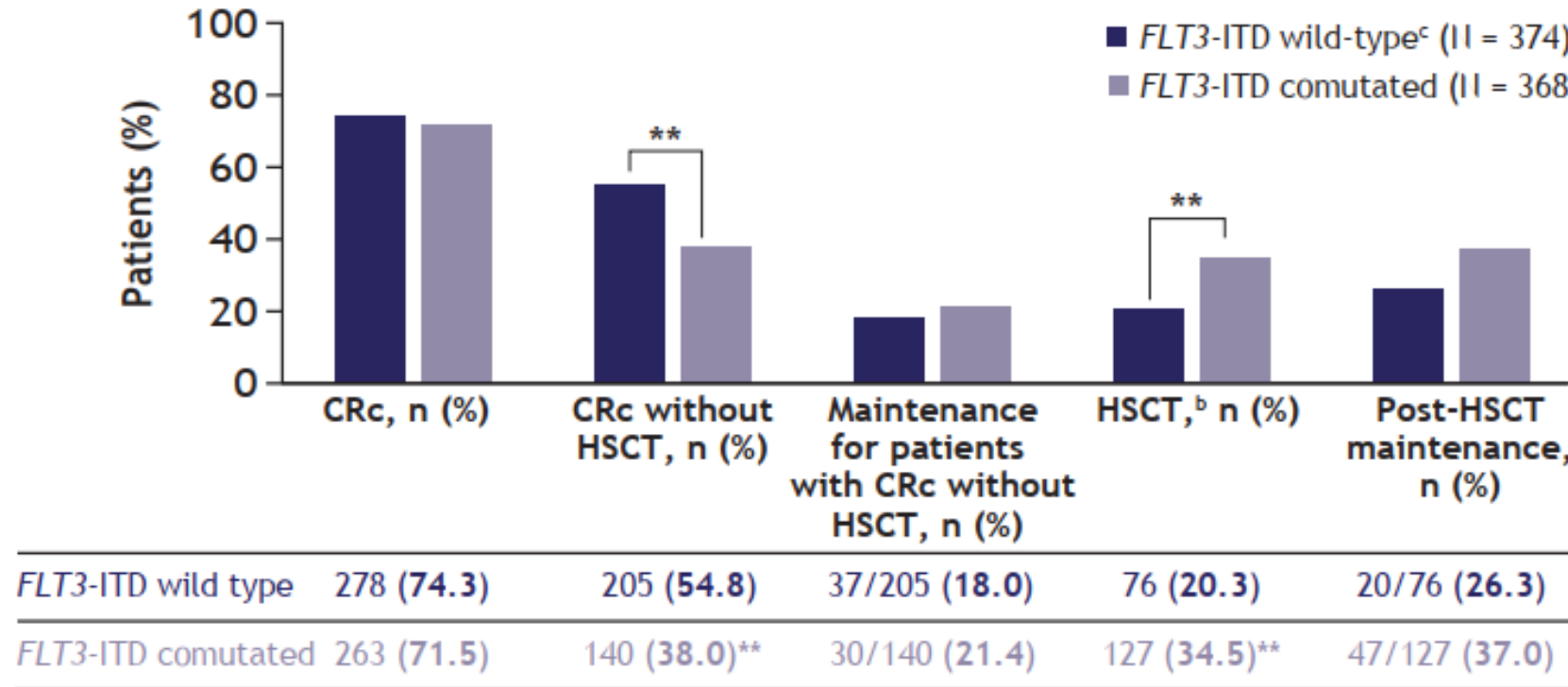
\* $P < 0.05$  between Early and Recent subcohorts. \*\* $P < 0.01$  between Early and Recent subcohorts.

<sup>a</sup>Percentages calculated among the total patient numbers for each subgroup shown, unless otherwise specified. <sup>b</sup>Seven patients (6 in Early subcohort and 1 in Recent subcohort) had HSCT without CRc. CRc, composite complete remission.



\* $P < 0.05$  between Early and Recent subcohorts. \*\* $P < 0.01$  between Early and Recent subcohorts.

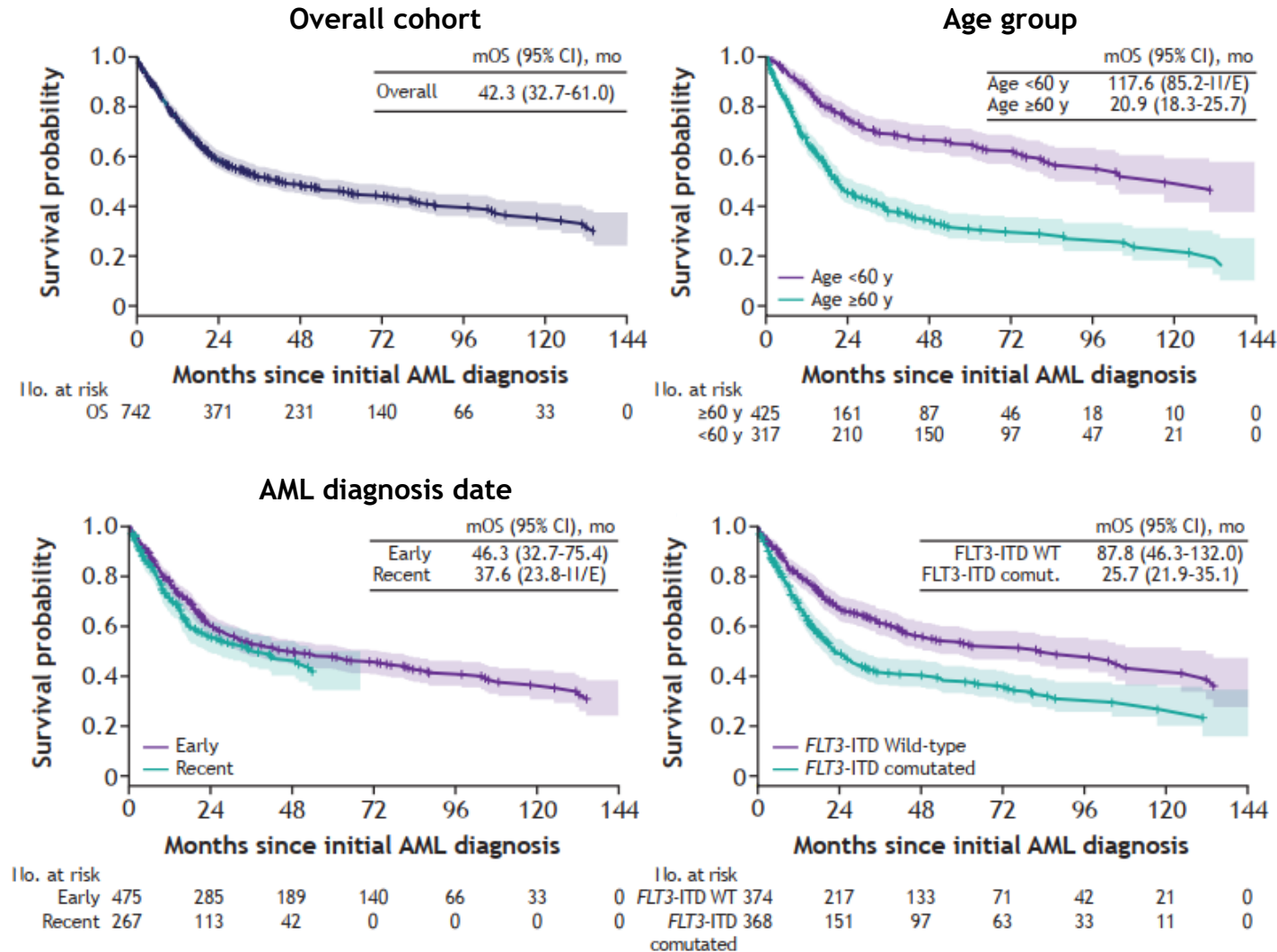
<sup>a</sup>From date of AML diagnosis and includes time to frontline treatment and first assessment. <sup>b</sup>Includes patients with documented negative testing and undocumented testing.



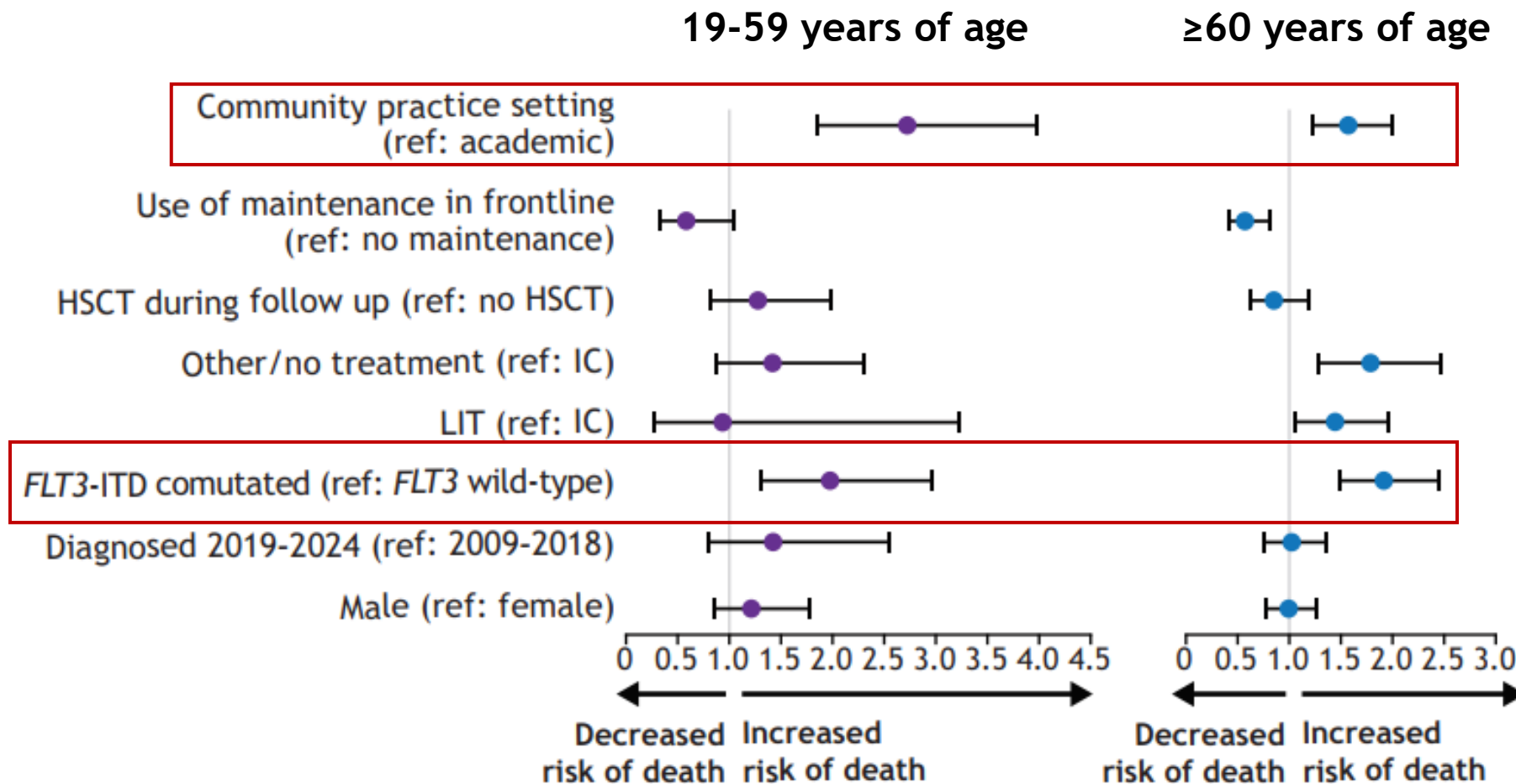
\*\* $P < 0.01$  between *FLT3*-ITD wild-type and comutated subcohorts.

<sup>a</sup>Percentages calculated among the total patient numbers for each subgroup shown, unless otherwise specified. <sup>b</sup>Seven patients (3 in *FLT3*-ITD wild-type subcohort and 4 in *FLT3*-ITD comutated subcohort) had HSCT without CRc.

<sup>c</sup>Includes patients with documented negative testing and undocumented testing.



Comut, comutated; ITD, internal tandem duplication mo, months; mOS, median overall survival; N/E, not evaluable; WT, wild-type.



ITD, internal tandem duplication; ref, reference population.

# Limitations

- Electronic health record data may be incomplete or inaccurate due to missing fields, inconsistent documentation, and variation in coding and recording practices across providers
- Analyses were limited to variables available in the COTA database, which may not include all relevant patient characteristics and outcomes of interest
- Identification of comutations may be impacted by testing patterns over time, with more testing in the Recent subcohort resulting in higher percentages of patients with comutations
- Risk of R/R may be underestimated, as it includes time between diagnosis and treatment and time between treatment and first response
- The study population may not be representative of all adults with *NPM1*m AML in the United States

# Conclusions

- The study showed less use of IC and more use of LIT, primarily with venetoclax, in patients with newly diagnosed *NPM1*m AML in 2019 to 2024 compared with 2009 to 2018
- Among those treated with LIT, risk of R/R at 6 months decreased between the 2 time periods; however, risk of R/R is still unacceptably high, with ~1/3 of patients at 12 months reaching the same rate of R/R as those treated with IC
- No statistically significant differences were detected in OS between the 2 time periods
- Outcomes with frontline therapies remain suboptimal despite recent treatment advances, such as venetoclax, highlighting the need for novel therapeutic approaches, including targeted therapies

# Acknowledgments

- We thank Adan Sosa, PharmD, an employee of Syndax Pharmaceuticals Inc., for review of the data
- Editorial assistance and graphic support was provided by Lumanity Communications Inc., and funded by Syndax Pharmaceuticals Inc.