



## The 65th ASH Annual Meeting Abstracts

## POSTER ABSTRACTS

## 613.ACUTE MYELOID LEUKEMIAS: CLINICAL AND EPIDEMIOLOGICAL

**Clinical Characteristics and Outcomes of Myeloid Neoplasms with *Mecom* Rearrangement: Results from a Nationwide Multicenter Study**

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*MECOM* rearrangements are detected in myeloid neoplasms [myelodysplastic neoplasm (MDS) and acute myeloid leukemia (AML)] which are associated with dismal prognosis. Classic *MECOM* rearrangements include *inv(3)(q21q26.2)* and *t(3;3)(q21;q26)* while non-classic subtypes are *3q26.2/MECOM* rearrangement with other partners. Both classic and non-classic rearrangements result in *MECOM* overexpression which involves in the process of leukemogenesis. Recently, the World Health Organization classification 2022 categorized myeloid neoplasms with these genetic abnormalities as "AML with *MECOM* rearrangement" regardless of blast count. We aim to explore frequency, clinical characteristics and outcomes including treatment response in this AML subtype among Thai myeloid neoplasms.

AML data was collected from a national registry which were conducted by Thai acute leukemia working group. Other than AML with *MECOM* rearrangements, AML was categorized as favorable, intermediate, and adverse risk groups according to European Leukemia Network 2022. MDS data was collected from a multicenter study group involving 4 medical centers. Other than MDS with *MECOM* rearrangements, MDS was categorized into 5 risk groups according to R-IPSS score system (very low, low, intermediate, high and very high-risk groups). Myeloid neoplasms with *MECOM* rearrangements were analyzed among their diseases and grouped together and compared to MDS and AML cohorts.

A total of 9 cases of myeloid neoplasms with *MECOM* rearrangement were detected. Among non-M3 AML cases, there were 6 cases with *MECOM* rearrangement from 746 non-M3 AML (0.8%) while 3 cases were detected in 163 MDS (1.8%). Seven of 9 cases (78%) were female gender. Five of 9 cases were classic *MECOM* rearrangement [1 case with *inv(3)(q21q26.2)* and 4

cases with t(3;3)(q21;q26)] while the other 4 cases were non-classic rearrangements [3 cases with t(3;21)(q26.2;q22) and 1 case with t(3;7)(q26;q21)].

In the AML cohort, AML with *MECOM* rearrangement showed lower white blood cell, but higher platelet counts compared to other groups (favorable, intermediate, and adverse risk groups) ( **Table 1**). Among AML cases receiving intensive chemotherapy, *MECOM* rearrangement subgroup showed lower complete response (CR) rate compared to others favorable, intermediate, and adverse risk groups. (0% vs. 77.3% vs. 37.6% vs. 23.8%;  $p < 0.001$ ). Of note, among 6 AML with *MECOM* rearrangement, there were 4 patients who received intensive chemotherapy but none of them responded to the treatment.

In the MDS cohort, MDS with *MECOM* rearrangement showed lower hemoglobin and platelet counts compared to other groups ( **Table 2**). Among 3 MDS with *MECOM* rearrangement, one patient received azacitidine with investigational drug (sabatolimab/placebo) and achieved complete hematologic response. He eventually progressed after 12 cycles of the treatments and subsequently died.

When combining 3 MDS and 6 AML with *MECOM* rearrangement as one group and compared survival rate with others: survival rate of 9 myeloid neoplasms with *MECOM* rearrangement is worse than the adverse group of AML and the very high risk group MDS with a 1-year survival rate of 22% ( **Figure 1 and 2**).

In conclusion, myeloid neoplasms with *MECOM* rearrangements are rare with the frequency of 0.8% in non-M3 AML and 1.8% in MDS. This subtype is more common in female gender. The prognosis of myeloid neoplasms with *MECOM* rearrangement is dismal with a 1-year survival rate of 16.7% in AML and 6-month survival rate of 33% in MDS. Chemotherapy should be avoided in this subtype due to non-responsiveness, hypomethylating agent showed benefit and can be considered as a bridging treatment before stem cell transplantation. Novel therapy targeting *MECOM* gene should be further explored to improve outcomes in this AML subtype.

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**Table 1 Characteristics of AML patients categorized by cytogenetic risk groups**  
Data present n (%), median (IQR), a using Chi-square test and b using Kruskal-Wallis test, rearr=rearrangement

|   | MECOM rearr     | Favorable        | Intermediate     | Adverse         | Total            | P-value             |
|---|-----------------|------------------|------------------|-----------------|------------------|---------------------|
| <b>N (%)</b>                              | 6 (0.8%)        | 75 (10.1%)       | 493 (66.1%)      | 172 (23.1%)     | 746 (100)        |                     |
| <b>Sex</b>                                |                 |                  |                  |                 |                  | 0.084 <sup>a</sup>  |
| Female                                    | 6 (100.0%)      | 45 (60.0%)       | 263 (53.3%)      | 89 (51.7%)      | 403 (54.0%)      |                     |
| Male                                      | 0 (0.0%)        | 30 (40.0%)       | 230 (46.7%)      | 83 (48.3%)      | 343 (46.0%)      |                     |
| <b>Age at diagnosis</b>                   | 53 (41- 62)     | 43 (27- 55)      | 54 (41- 63)      | 52.5 (40- 63)   | 53 (40- 62)      | <0.001 <sup>b</sup> |
| <b>WBC (x10<sup>3</sup>cells/uL)</b>      | 6.9 (3.0- 28.6) | 27.7 (8.3- 57.3) | 27.2 (6.4- 78.0) | 7.9 (2.8- 36.1) | 22.1 (5.2- 64.9) | <0.001 <sup>b</sup> |
| <b>Hb (g/dL)</b>                          | 8.4 (7.4- 8.8)  | 8.0 (6.3- 9.3)   | 7.9 (6.7- 9.4)   | 7.9 (6.8- 9.1)  | 7.9 (6.7- 9.3)   | 0.846 <sup>b</sup>  |
| <b>Platelet (x10<sup>3</sup>cells/uL)</b> | 135 (77-184)    | 41 (19- 77)      | 49.7 (22- 90)    | 49 (25- 87)     | 48 (23- 88)      | 0.025 <sup>b</sup>  |
| <b>Response</b>                           |                 |                  |                  |                 |                  | <0.001 <sup>a</sup> |
| Not CR                                    | 4 (66.7%)       | 8 (10.7%)        | 104 (21.1%)      | 53 (30.8%)      | 169 (22.7%)      |                     |
| CR  | 0 (0.0%)        | 58 (77.3%)       | 185 (37.6%)      | 41 (23.8%)      | 284 (38.1%)      |                     |
| Not evaluate                              | 2 (33.3%)       | 9 (12.0%)        | 203 (41.3%)      | 78 (45.3%)      | 292 (39.2%)      |                     |

**Table 2 Characteristics of MDS patients categorized by R-IPSS groups**  
Data present n (%), median (IQR), a using Chi-square test and b using Kruskal-Wallis test, rearr=rearrangement

|   | MECOM rearr       | Very low risk      | Low risk            | Intermediate       | High risk          | Very high risk     | Total               | P-value |
|---|-------------------|--------------------|---------------------|--------------------|--------------------|--------------------|---------------------|---------|
| <b>N (%)</b>                              | 3 (1.8%)          | 25 (15.3%)         | 63 (38.7%)          | 30 (18.4%)         | 23 (14.1%)         | 19 (11.7%)         | 163 (100%)          |         |
| <b>Sex</b>                                |                   |                    |                     |                    |                    |                    |                     | 0.631   |
| Female                                    | 1 (33.3%)         | 13 (52.0%)         | 34 (54.0%)          | 15 (50.0%)         | 13 (56.5%)         | 7 (36.8%)          | 83 (50.9%)          |         |
| Male                                      | 2 (66.7%)         | 12 (48.0%)         | 29 (46.0%)          | 15 (50.0%)         | 10 (43.5%)         | 12 (63.2%)         | 80 (49.1%)          |         |
| <b>Age at diagnosis</b>                   | 76.0 (66.0- 81.0) | 73.0 (66.0- 79.0)  | 72.0 (64.0- 81.0)   | 68.0 (54.0- 73.0)  | 69.0 (59.0- 78.0)  | 71.0 (60.0- 77.0)  | 71.0 (63.0- 79.0)   | 0.167   |
| <b>WBC (x10<sup>3</sup>cells/uL)</b>      | 1.9 (1.8- 7.0)    | 3.9 (3.0- 5.2)     | 4.4 (3.2- 6.8)      | 3.3 (2.2- 5.4)     | 3.2 (2.0- 6.1)     | 2.1 (1.6- 4.8)     | 3.7 (2.4- 6.0)      | 0.663   |
| <b>Hb (g/dL)</b>                          | 5.3 (4.6- 6.7)    | 10.6 (10.0- 11.9)  | 8.3 (7.2- 9.3)      | 8.1 (7.1- 10.0)    | 8.1 (6.9- 8.9)     | 7.7 (6.6- 8.3)     | 8.3 (7.1- 10.0)     | <0.001  |
| <b>Platelet (x10<sup>3</sup>cells/uL)</b> | 18.0 (7.0- 415.0) | 93.0 (71.0- 208.0) | 156.0 (69.0- 292.0) | 84.0 (46.0- 127.0) | 84.0 (43.0- 128.0) | 94.0 (39.0- 117.0) | 100.0 (59.0- 189.0) | 0.002   |

**Figure 1 Survival rate of AML with MECOM rearrangement among other AML subgroups by cytogenetic risk**

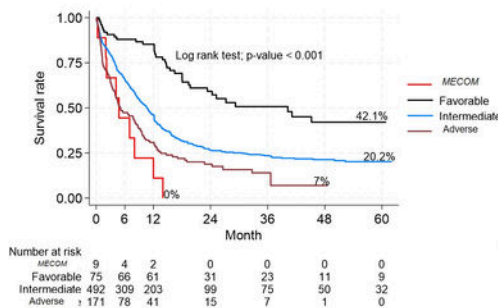


Table Hazard ratio for Cytogenetic risk in for AML patients.  
HR: Hazard ratio was estimated by Cox regression

| Cytogenetic risk | HR (95%CI)        | P-value |
|------------------|-------------------|---------|
| Favorable        | 1                 | Ref     |
| Intermediate     | 2.35 (1.66-3.33)  | <0.001  |
| Adverse          | 3.58 (2.46-5.21)  | <0.001  |
| MECOM            | 5.55 (2.65-11.60) | <0.001  |

**Figure 2 Survival rate of MDS with MECOM rearrangement among other MDS subgroups by R-IPSS**

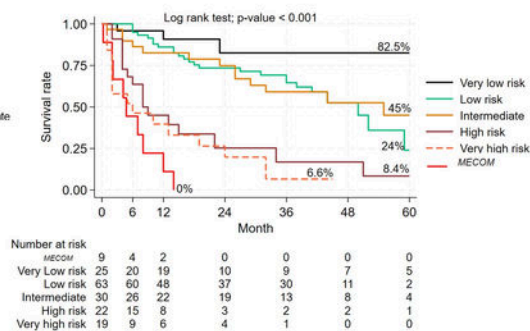


Table Hazard ratio for IPSS risk in for MDS patients  
HR: Hazard ratio was estimated by Cox regression

| R-IPSS risk    | HR (95%CI)          | P-value |
|----------------|---------------------|---------|
| Very low risk  | 1                   | Ref     |
| Low risk       | 3.49 (1.05-11.59)   | 0.041   |
| Intermediate   | 3.36 (0.96-11.79)   | 0.059   |
| High risk      | 12.35 (3.59-42.44)  | <0.001  |
| Very high risk | 18.04 (3.59-42.44)  | <0.001  |
| 3Q             | 34.42 (9.01-131.48) | <0.001  |

**Figure 1**

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