

Characterization and Prognosis of Secondary Acute Myeloid Leukemia in an Asian Population: AML With Antecedent Hematological Disease Confers Worst Outcomes, Irrespective of Cytogenetic Risk

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Abstract. *Background/Aim:* Acute myeloid leukemia (AML) is a biologically heterogeneous disease that can be classified into *de novo* AML and secondary AML. Secondary AML can be further divided into therapy-related AML (t-AML) or AML evolving from antecedent hematological disorder (AHD-AML). This study evaluated the characteristics and prognosis of secondary AML in a homogeneous East Asian population who are often under-represented. *Patients and Methods:* This was a retrospective, longitudinal cohort study of Korean AML patients over 18 years old treated between January 2000 and December 2013. A total of 437 *de novo* AML (80.3%), 41 t-AML (7.5%), and 66 AHD-AML (12.1%) were evaluated. *Results:* First, we found that secondary AML constituted about 19.7% of all AML cases, and t-AML was more prevalent than AHD-AML. Second, we determined AHD-AML as a prognostic factor for inferior survival, independent of other risk factors (HR=2.137, 95%CI=1.534-2.977, $p<0.001$). The induction response rates correlated well with the overall survival. Furthermore, AHD-AML was associated

with worst treatment outcomes and prognosis regardless of cytogenetic risk or age. Interestingly, t-AML was generally associated with better outcomes compared to AHD-AML despite the similarities in treatment schema. *Conclusion:* Secondary AML represents a broad spectrum of diseases and t-AML should be addressed separately from AHD-AML.

Acute myeloid leukemia (AML) is a biologically heterogeneous disease that can be classified into two distinct categories: 1) *de novo* AML that arises in the absence of an identified exposure or prodromal stem cell disorder; and 2) secondary AML. Secondary AML can be further divided into therapy-related AML (t-AML) due to previous exposure to leukemogenic therapies including chemotherapy and radiotherapy, or AML evolving from antecedent hematological disorder (AHD-AML) including myelodysplastic syndrome (MDS), myeloproliferative neoplasm (MPN), or aplastic anemia (AA) (1-3). Secondary AML is generally associated with poor clinical outcomes compared to *de novo* AML (4, 5). However, there is still debate as to whether secondary AML itself is an independent risk factor, as secondary AML patients tend to be older and more fragile (6). Unfortunately, secondary AML patients are commonly excluded from clinical trials, limiting the understanding of the disease. Another unresolved issue is the discrepancies in response and survival rates of secondary AMLs: despite harboring a seemingly targetable mutation, JAK2, post-MPN AML is typically associated with abysmal prognosis, while selected t-AML patients show survivals similar to their *de novo* counterparts (2, 4). Despite continued attempts to genetically characterize secondary AML (7, 8), whether distinct somatic mutations underlie the different treatment responses remain largely unknown. As such, we carried out this study in an effort to refine the characteristics

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Key Words: Acute myeloid leukemia, secondary, therapy related, survival, mutations.

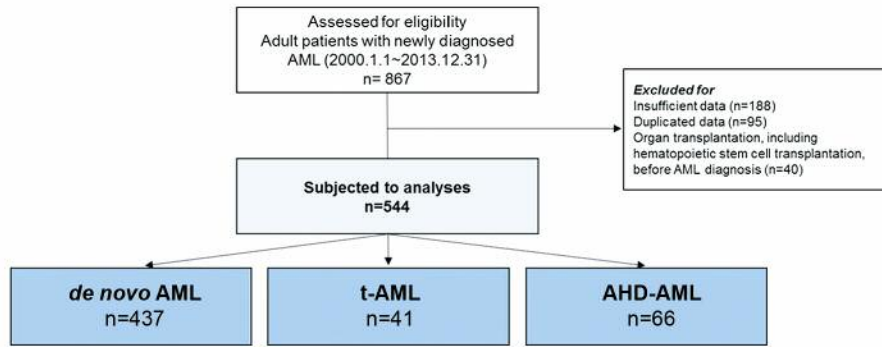


Figure 1. CONSORT diagram.

and prognosis of secondary AML patients, for whom there exist a critical unmet need of new therapeutic options. Specifically, we focused on a homogeneous East Asian population, which is often under-represented.

Patients and Methods

Study design and subjects. This was a retrospective, longitudinal cohort study of Korean AML patients over 18 years old treated between January 2000 and December 2013 at the Seoul National University Hospital. Initially, a total of 867 patients were identified (Figure 1). After excluding those with insufficient data, those who underwent organ transplantation including hematopoietic stem cell transplantation (HSCT) and duplicated data, 544 patients with complete set of bone marrow examination data and cytogenetic information were enrolled. In the end, there were 437 patients with *de novo* AML (80.3%), 41 with t-AML (7.5%), and 66 with AHD-AML (12.1%). Their medical records were reviewed and analyzed for demographics and baseline disease characteristics including cytogenetics, chemotherapy dosing and schedule, factors related to HSCT, response to chemotherapy and HSCT, adverse events, and survival outcomes.

This study was conducted according to the Declaration of Helsinki and was approved by the institutional review board of Seoul National University Hospital (IRB No. H-1607-198-782). All authors had access to the study data and reviewed and approved this study.

AML diagnosis and risk stratification. The diagnosis of AML was made according to the WHO Classification of Hematopoietic Neoplasms, which requires identification of 20% or more leukemic blasts in the bone marrow (9). Cytogenetic studies were performed at each center, whose satisfactory performance was monitored by a national external quality assurance scheme. Bone marrow cells were cultured for 24 h and then the karyotype was analyzed using the standard G-banding technique. The karyotypes were constructed and chromosomal abnormalities were reported in accordance with the International System for Human Cytogenetic Nomenclature (10). Prognostic grouping of cytogenetics was performed according to Southwest Oncology Group (SWOG) criteria (11). Fms-related tyrosine kinase 3 (FLT3) internal tandem duplication (ITD) and nucleophosmin-1 (NPM1) mutations were analyzed using DNA samples obtained at initial diagnosis and multiplex polymerase chain

reaction (PCR). Complex karyotype was defined as any karyotype with at least 3 chromosome aberrations, regardless of their type and the individual chromosomes involved. Monosomal karyotype was defined as 2 or more distinct autosomal chromosome monosomies or 1 single autosomal monosomy in addition to structural abnormalities.

Statistical analysis. The overall survival (OS) curves were estimated using the Kaplan-Meier method. OS was defined as the time from the date of AML diagnosis to death from any cause. If patients survived without death, the survival was censored on the last date of follow-up. Cox proportional hazards model and logistic regression were used to identify significant prognostic indicators for survival. Differences between groups were assessed using Student’s t-test, one-way analysis of variance for continuous variables, or Pearson chi-square test for categorical variables, as appropriate. All data were analyzed using the Statistical Package for the Social Sciences software (IBM® SPSS® statistics, version 22.0, Armonk, NY, USA). p-Values of <0.05 was considered to be statistically significant.

Results

Patient characteristics. Table I represents the baseline characteristics of all patients. While sex distribution was uniform across different AML subtypes, patients with AHD-AML tended to be older (median: 61.5, range=22-77 years) compared to patients with *de novo* AML (median: 51, range=18-85 years; $p=0.001$) and t-AML (median: 53, range=19-74 years; $p=0.439$). Poor cytogenetics were most frequently observed in patients with AHD-AML (*de novo* AML vs. t-AML, $p=0.179$; *de novo* AML vs. AHD-AML, $p<0.001$; t-AML vs. AHD-AML, $p=0.321$).

Among t-AML patients, the most common diagnosis was lymphoma (n=12, 30%) and breast cancer (n=7, 17%), together constituting approximately half of the group (Figure 2A). The median latency between diagnosis of the primary disease and the diagnosis of AML was 47 months (range=31-235 months).

As shown in Figure 2B, among AHD-AML patients, the most common diagnosis was MDS (n=41, 62%) followed by

Table I. Patient characteristics.

%	<i>de novo</i> (N=437)	t-AML (N=41)	AHD-AML (N=66)
Year of diagnosis			
2000-2004	82 (18.8)	9 (22.0)	16 (24.2)
2005-2009	173 (39.6)	18 (43.9)	21 (31.8)
2010-2013	182 (41.6)	14 (34.1)	29 (43.9)
Age at dx (years)			
Median [range]	51 (18-85)	53 (19-74)	61.5 (22-77)
<65 years old	345 (78.9)	27 (65.9)	42 (63.6)
≥65 years old	92 (21.1)	14 (34.1)	24 (36.4)
Gender (male)			
	261 (59.7)	21 (51.2)	38 (57.6)
Cytogenetic risk group			
Favorable	107 (24.5)	7 (17.1)	0
Intermediate	234 (53.5)	19 (46.3)	40 (60.6)
Poor	96 (22.0)	15 (36.6)	26 (39.4)
Extramedullary involve			
	37 (8.5)	6 (14.6)	0
Lab findings at dx ¹			
BM blast (%)	57.1 (27.5)	56.1 (27.4)	37.4 (20.4)
WBC count (10 ³ /l)	31.4 (83.7)	27.4 (50.0)	13.8 (20.9)
Platelet count (10 ⁹ /l)	74.0 (79.3)	50.7 (63.4)	149.5 (98.0)
Hb (g/dl)	8.3 (2.0)	8.1 (1.7)	8.1 (1.6)
Treatment			
Intensive chemotherapy	424 (97.0)	36 (87.8)	56 (84.8)
Low intensity chemotherapy ²	5 (1.1)	1 (2.4)	4 (6.1)
No treatment	8 (1.4)	4 (9.8)	6 (9.1)

t-AML: Therapy related AML; AHD: antecedent hematologic disease; dx: diagnosis; NA: not applicable; BM: bone marrow; WBC: white blood cell; Hb: hemoglobin. ¹Values are presented as mean (±standard deviation); ²Low intensity chemotherapy includes hypomethylating agents and low-dose cytarabine treatment.

MPN (n=18, 27%). When MDS-AML and MPN-AML were compared, there were no differences in the number of patients with poor cytogenetics (MDS-AML 39% vs. MPN-AML 41.2%, $p=0.879$). Patients with AHD-AML were associated with significantly lower white blood cell (WBC) count at diagnosis compared with patients with *de novo* AML ($p=0.001$).

Treatment and response according to AML subtypes. In total, 516 out of 544 patients (94.9%) underwent intensive chemotherapy (Table I). Per subtype, 97% (424/437) of *de novo* AML patients were induced, while 87.8% (36/41) of t-AML patients and 84.8% (56/66) of AHD-AML patients were induced. Patients with *de novo* AML were more likely to achieve complete remission (CR) than patients with other subtypes of AML, while patients with AHD-AML were least likely to achieve CR ($p<0.001$, Table II). Such lower CR rate for AHD-AML was observed across the cytogenetic risk groups. Only 19.6% of the AHD-AML patients were subsequently subjected to consolidation chemotherapy, while 68.2% of the *de novo* AML patients and 47.2% for the t-AML patients received consolidation. With reference to age,

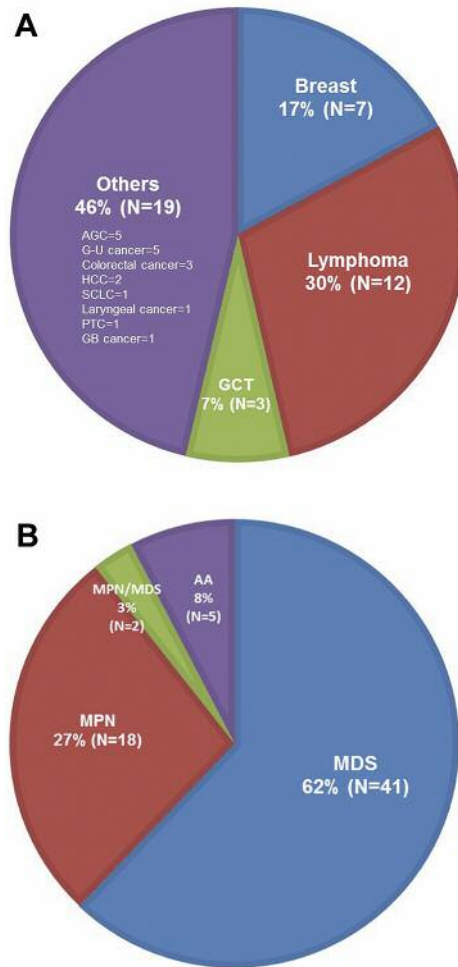


Figure 2. Subclassification of secondary AML. AML: Acute myeloid leukemia; MDS: myelodysplastic syndrome; MPN: myeloproliferative neoplasm; AA: aplastic anemia. (A) Therapy-related AML. (B) AML with antecedent hematological disease.

older patients showed trends towards lower CR rate in all 3 AML subtypes, however the difference was only significant in *de novo* AML (86.3% CR rate in patients <65 years old vs. 72.0% in patients ≥65 years old, $p=0.002$).

Death occurred most frequently in AHD-AML (Table II). Specifically, the majority of AHD-AML patients expired before 1st relapse, suggesting high rates of treatment related mortality.

Overall survival. The median OS of all 544 patients was 20 months (Figure 3A), with improvement in survival evident per year of diagnosis. Overall, *de novo* AML patients were associated with best OS (median: 29 months) followed by t-AML patients (median: 14 months) and AHD-AML patients

Table II. Treatment schema and outcomes of patients undergoing intensive chemotherapy.

%	de novo (N=424)	t-AML (N=36)	AHD-AML (N=56)
Induction CR	354 (89.4)	24 (66.7)	18 (32.1)
Cytogenetic risk group			
Favorable	102/107 (95.3)	6/7 (85.7)	NA
Intermediate	189/226 (83.6)	13/18 (72.2)	12/36 (33.3)
Poor	64/91 (70.3)	5/11 (45.5)	7/20 (35.0)
Age			
<65 years old	295/342 (86.3)	19/26 (73.1)	15/37 (40.5)
≥65 years old	59/82 (72.0)	5/10 (50.0)	3/19 (15.8)
Consolidation chemotherapy	289 (68.2)	17 (47.2)	11 (19.6)
Number (median, range)	2 (1-3)	2 (1-3)	2 (1-3)
Upfront HSCT	65 (15.3)	13 (14.6)	12 (21.4)
HSCT donor			
Related/unrelated	43 (66.2)/ 22 (33.8)	10 (76.9)/ 3 (23.1)	5 (41.7)/ 7 (58.3)
Match/mismatch	50 (76.9)/ 15 (23.1)	8 (61.5)/ 5 (38.5)	9 (75.0)/ 3 (25.0)
HSCT conditioning			
Reduced intensity	45 (69.2)	10 (76.9)	8 (66.7)
Myeloablative	20 (30.8)	3 (23.1)	4 (33.3)
Death			
Any cause of death	232 (54.7)	23 (63.9)	44 (78.6)
Death before 1 st relapse ¹	68 (29.3)	8 (34.8)	24 (54.5)
Related to induction	40	6	16
Related to consolidation	11	0	1
Related to upfront HSCT	17	2	7

t-AML: Therapy related AML; AHD: antecedent hematologic disease; CR: complete remission; NA: not applicable; consol: consolidation; HSCT: hematopoietic stem cell transplantation. ¹Percentage out of total death.

(median: 7 months, Figure 3B). The difference in survival between *de novo* AML and t-AML did not reach statistical significance, but there was significant difference between *de novo* AML vs. AHD-AML ($p<0.001$) and t-AML vs. AHD-AML ($p=0.018$). When MDS-AML (median OS: 7 months) and MPN-AML (median OS: 6 months) were compared, there was no difference in OS ($p=0.349$, data not shown). Figure 3C shows the OS of patients per cytogenetic risk group. AHD-AML patients were associated with worst OS outcomes, irrespective of cytogenetic risk group.

When age was taken into consideration (Figure 4), patients ≥65 years old (median OS: 12 months) were generally associated with poorer outcomes compared to patients <65 years old (median OS: 31 months, $p<0.001$). AHD-AML was associated with worst outcomes, irrespective of age (Figure 4A). In patients <65 years old, AHD-AML showed inferior survival compared to *de novo* AML in each of the 3 cytogenetic risk groups (Figure 4B), indicating that poor

Table III. Multivariable analysis for survival in patients undergoing intensive chemotherapy.

	Overall survival	
	HR (95%CI)	p-Value
Age		
≥65 vs. <65	1.766 (1.360-2.295)	<0.001
Gender		
Male vs. Female	1.350 (1.065-1.712)	0.013
Type of AML		
<i>de novo</i>	ref	ref
t-AML	1.347 (0.871-2.083)	0.180
AHD-AML	2.137 (1.534-2.977)	<0.001
Year of diagnosis		
2000-2004	ref	ref
2005-2009	0.839 (0.627-1.122)	0.236
2010-2013	0.665 (0.486-0.884)	0.006
Cytogenetic risk group		
Favorable	Ref	ref
Intermediate	1.743 (1.248-2.434)	0.001
Poor	2.610 (1.796-3.792)	<0.001

outcomes in AHD-AML depend on karyotype. In patients ≥65 years old, while AHD-AML showed inferior survival to *de novo* AML ($p=0.034$) in the intermediate risk group, there were no differences in survival between the 3 AML subtypes in the high-risk group (Figure 4C). There were only 8 patients with favorable cytogenetic risk (7 *de novo* AML, 1 t-AML), thus survival analysis was not performed.

In multivariate analyses (Table III), AHD-AML was recognized as independent prognostic factor for survival (HR=2.137, 95%CI=1.534-2.977, $p<0.001$), while t-AML was not ($p=0.180$). Other prognostic factors included older age, male sex and intermediate/poor cytogenetic risk.

Discussion

Secondary AML is generating substantial economic and social burdens as the number of cancer survivors is rapidly increasing (12). Recognizing the relative paucity of data from Asian populations, we carried out this study to investigate the characteristics and prognosis of secondary AML hoping of establishing ground works for more innovative therapeutic approaches for this often-neglected population of patients. First of all, through our study, we found that secondary AML constituted about 19.7% of all AML cases. Second, we determined AHD-AML as a prognostic factor for inferior survival, independent of other risk factors. AHD-AML was associated with worst treatment outcomes and prognosis regardless of cytogenetic risk or age. Interestingly, t-AML was generally associated with better outcomes compared to AHD-AML despite the similarities in treatment schema.

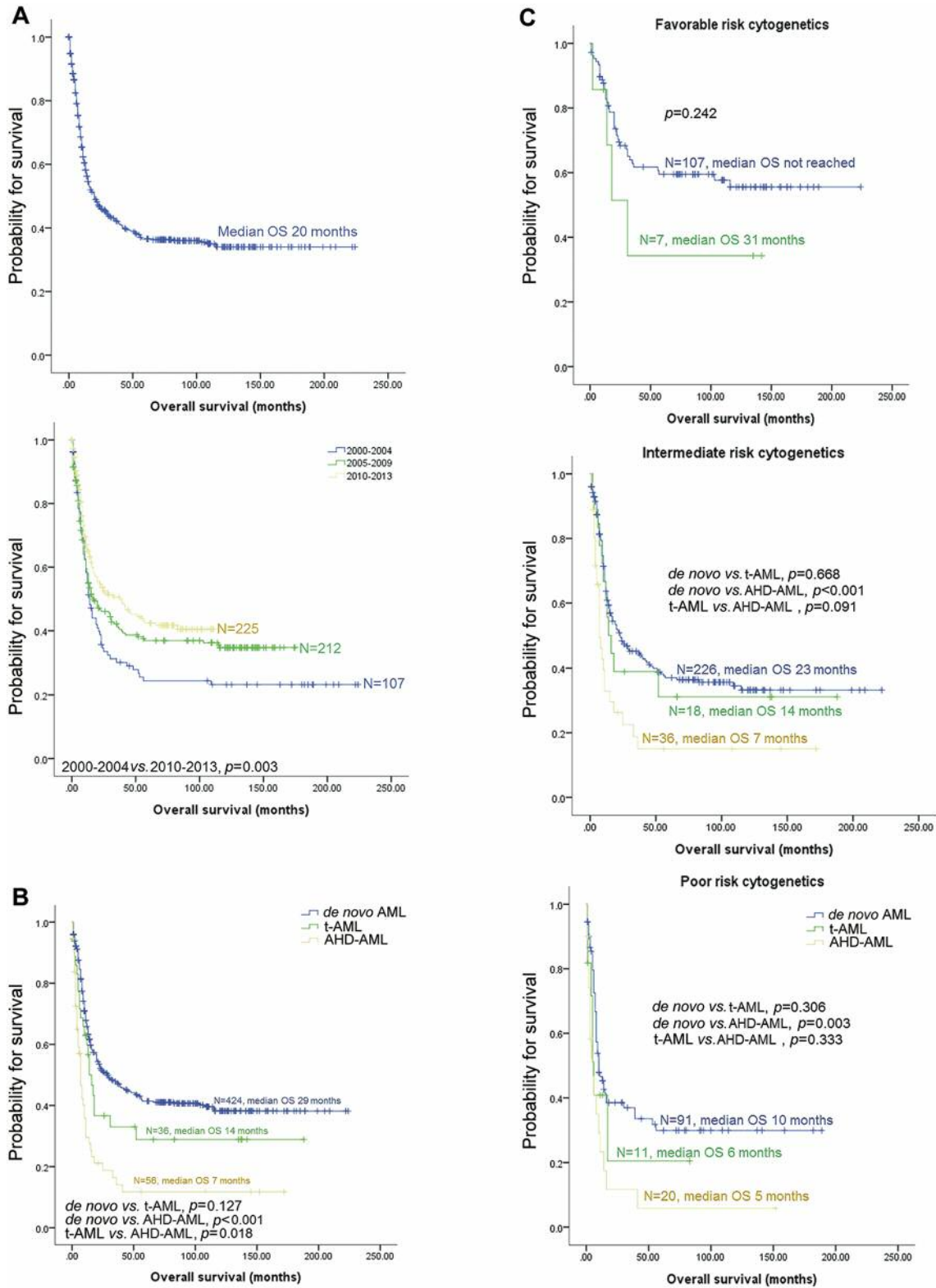


Figure 3. Overall survival according to year of diagnosis, AML subtypes and cytogenetic risk group. AML: Acute myeloid leukemia; t-AML: therapy related AML; AHD-AML: AML with antecedent hematological disease. (A) Left, all patients; Right, according to year of diagnosis. (B) According to AML subtypes, for patients undergoing intensive chemotherapy. (C) According to cytogenetic risks groups, for patients undergoing intensive chemotherapy. Left, favorable risk group; Middle, intermediate risk group; Right, poor risk group.

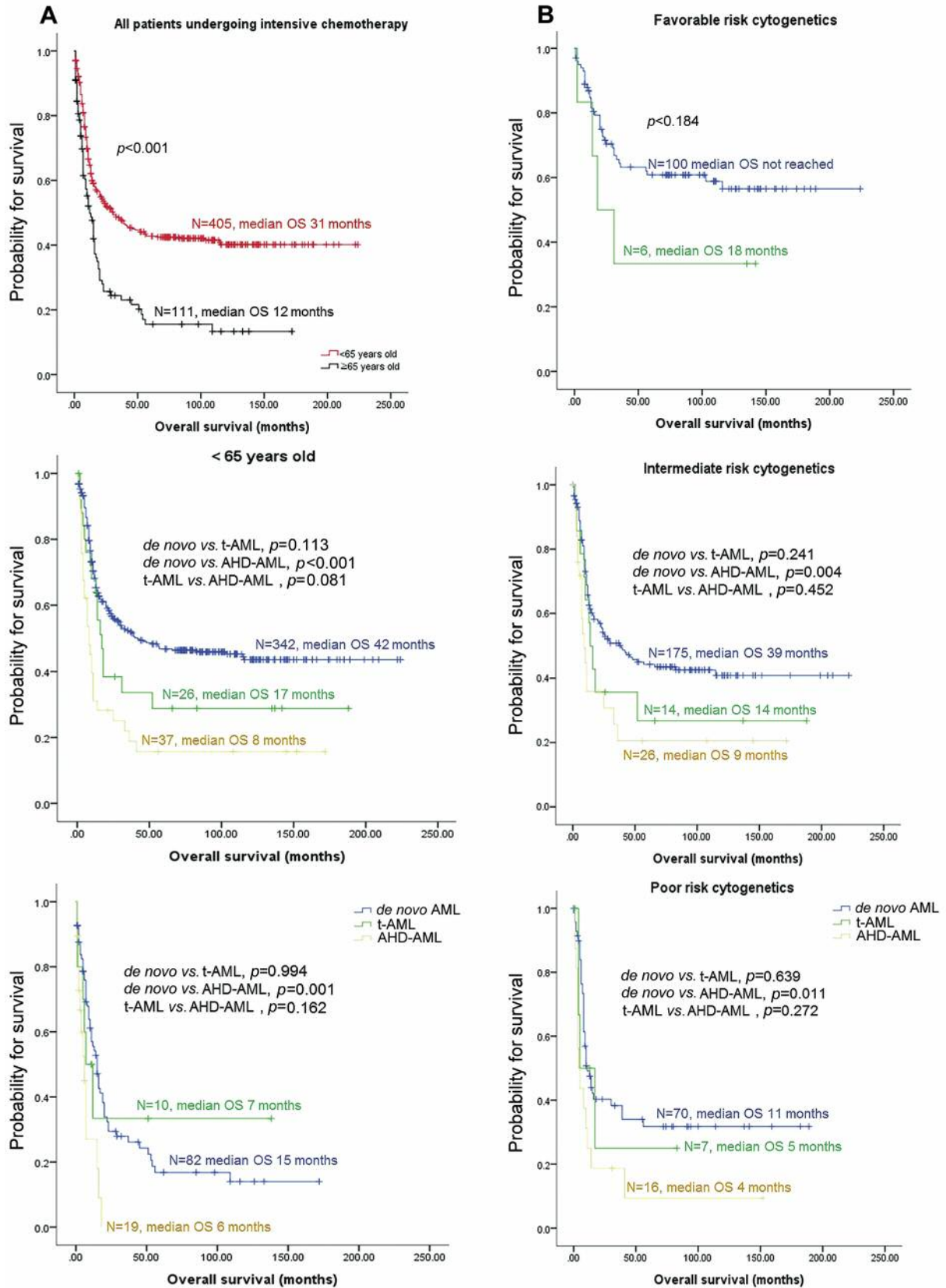


Figure 4. Continued

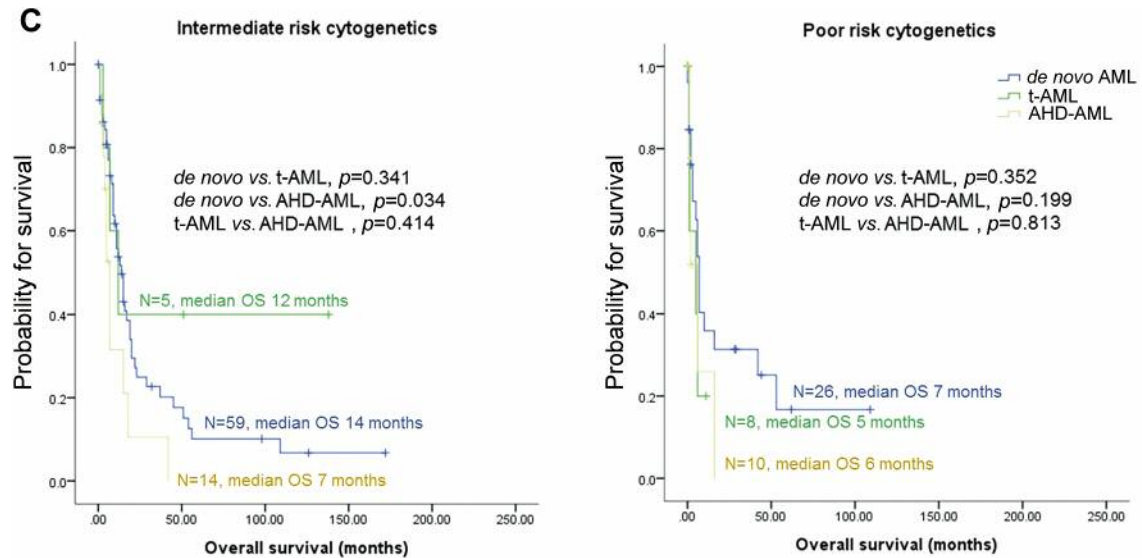


Figure 4. Overall survival according age. AML, acute myeloid leukemia; t-AML, therapy related AML; AHD-AML, AML with antecedent hematological disease. (A) According to age group, for patients undergoing intensive chemotherapy. Left, all patients; Middle, patients <65 years old; Right, patients ≥ 65 years old. (B) Patients <65 years old undergoing intensive chemotherapy. Left, favorable risk group; Middle, intermediate risk group; Right, poor risk group. (C) Patients ≥ 65 years old undergoing intensive chemotherapy. Left, intermediate risk group; Right, poor risk group.

As in previous reports (2, 13), lymphoma was the most common primary malignancy followed by breast cancer in t-AML group. The survival correlated with response rates. Higher induction CR rate was observed in *de novo* AML across all cytogenetic risk groups, followed by t-AML and AHD-AML. The difference in induction CR rates translated into difference in the number of patients undergoing consolidation treatment. Notably, the number of consolidation cycles and percentage of patients receiving upfront HSCT were similar between the 3 groups. Based on our data and Nilsson et al.'s report (14), we suggest that upfront HSCT is the only realistic curative option for patients with AHD-AML. However, since treatment related mortality was highest in AHD-AML, careful selection of patients and adaptive modifications during HSCT process is crucial. Also, this observation suggests more innovative therapeutic approaches are needed for AHD-AML patients.

One of the most obvious limitations of this study is the lack of detailed genetic sequencing data. Since the study period was set between year 2000 and 2013, genetic sequencing data was not available in most patients. Whole genome sequencing data was available in 140 patients (112 *de novo* AML, 9 t-AML, 19 AHD-AML) but sequencing was performed at different time points using various platforms, and since the quality of sequencing could not be guaranteed, comparative analyses were not performed. As specific mutations can distinguish sAML like disease within t-AML and elderly *de novo* AML (7, 15, 16), a follow up study with detailed genetic data is needed for better classification of Asian sAML. Another pitfall is the small

number of patients. Although the number of patients enrolled in this study is not small by any means for a single center, it is not big enough to be readily extrapolated as national data. However, being a single center data, there is the advantage of uniformity (*i.e.* patients received same sequence of diagnostics and treatment) which assures the quality of our data. Nevertheless, a larger epidemiologic study, preferably of East Asian descents, should ensue for corroboration of our findings.

In conclusion, we provide a better understanding of AML landscape in the Korean population. Since secondary AML is a rather heterogenous disease, t-AML should be addressed separately from AHD-AML. Continuous efforts to better characterize and treat secondary AML are crucial.

Conflicts of Interest

No conflicts of interest to disclose regarding this study.

Authors' Contributions

Designed the study: Ja Min Byun, Junshik Hong. Patient enrollment and data collection: all the authors. Analyzed the data: Ja Min Byun, Sheehyun Kim. Wrote the paper: Ja Min Byun, Sheehyun Kim. Revised the paper: all the authors.

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