

GVHD-model mice. Many models have been developed to manipulate, and to clarify, human T-cell-mediated acute GVHD in vivo [28]. The model that we used is optimal for evaluation of the killing ability of ASHmAbs, because it is very simple; it is reproducible; and most of all, the immune response is typically more robust to xenografts than to allografts.

We thought that graft failure requiring second transplantation was a potential adverse effect of kASHmAb administration, because the antibody might target for elimination all donor-type HLA-expressing cells, in particular, HSCs (which are known to express high levels of class I major histocompatibility complex) [21,22]. Unexpectedly, human PBMCs reappeared in mouse peripheral blood 2 to 3 weeks after kASHmAb administration at very high doses. This suggests that the cytotoxic effect of kASHmAb may preferentially injure mature PBMCs, sparing hematopoietic stem progenitor cells. The mechanism of this preference is not clear at present. It could be due to the fact that most HSCs in the bone marrow niche are in a quiescent state and resistant to cell damage such as apoptosis [29,30]. Although the concern persists that patients given kASHmAb are at risk of graft failure, optimization of kASHmAb dosages may solve this issue.

So far, we have generated kASHmAbs against HLA-A2 and HLA-A24 (Supplementary Figure E6, online only, available at www.exphem.org). These two kASHmAbs can cover approximately 23.1% of HLA-mismatched transplants (data from our institute). More kASHmAbs are needed to cover other HLA-mismatched transplants. For example, if we establish 12 more kASHmAbs (HLA-A*02:03, HLA-A*02:06, HLA-A*02:07, HLA-A*24:02, HLA-A*24:20, HLA-A*11:01, HLA-A*26:02, HLA-A*31:01, HLA-A*33:03, HLA-B*35:01, HLA-B*40:02, and HLA-B*51:01), at least 72.3% of GVHD cases ($n = 199$) in our institute can be treated. As an alternative to class I kASHmAbs, we have succeeded in generating several HLA class II ASHmAbs. These mAbs can potentially target activated T cells and antigen presenting cells that play a central role in eliciting GVHD. Administration of them alone or in combination with class I kASHmAbs may further enhance the anti-GVHD effect but minimize side effects.

Furthermore, if the diagnosis of GVHD can be made earlier using recently discovered biomarkers of early GVHD [31], for example, we may be able to treat GVHD safely with low doses of kASHmAb. Should target cells evade kASHmAbs by internalizing HLA molecules [32], kASHmAbs can be labeled with cell-damaging agents such as anti-cancer drugs and radio-isotopes [33,34], thereby inducing cell death. If HLA expression is downregulated [35,36], cell surface expression of HLA can be induced with drugs such as interferon- γ [37]. An additional concern in treatment of GVHD with kASHmAbs is their influence on the graft-versus-tumor (GVT) effect. Although little is known about how GVT and GVHD effects differ,

clearly in most cases there will be no GVT effect without GVHD. To observe the GVT effect even after treatment of GVHD with either class I or class II kASHmAbs will pose intriguing questions.

In conclusion, kASHmAb is an antibody that accurately discriminates between donor and recipient cells and induces target cell death. As use of HLA-mismatched cord blood transplantation and haplo-identical HSCT increases, so may the incidence of GVHD. ASHmAbs may provide an effective treatment for GVHD, favorably influencing the outcome of allo-HSCT. We believe that kASHmAbs, ready-made agents created through well-designed antibody-processing technologies, have great potential in the clinical treatment of GVHD.

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Author contributions

YN and SY designed the study. YN conducted animal experiments, contributed to data analysis and wrote the first draft of the manuscript. JU and YO were responsible for histopathologic analysis of the mice. YN, SY, SCN, NW, ST, and HN contributed to the writing of the article. This work is part of the doctoral thesis of YN.

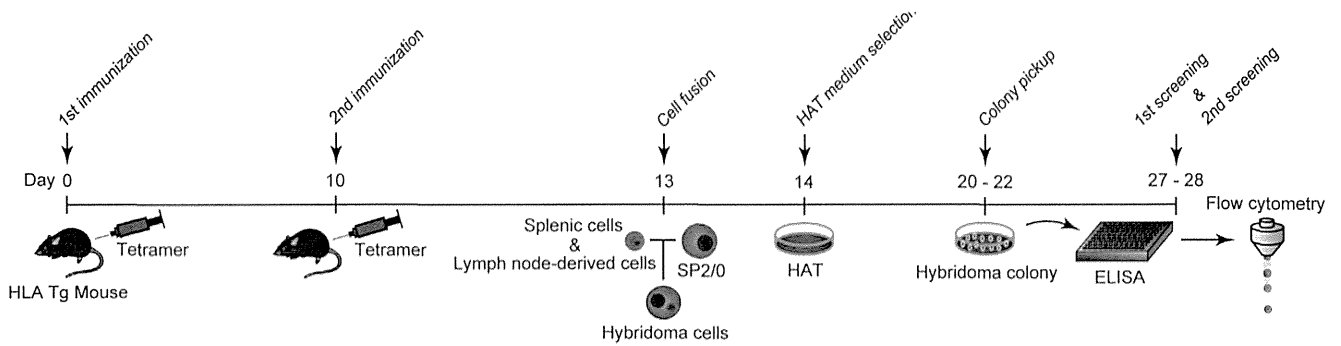
Conflict of interest disclosure

No financial interest/relationships with financial interest relating to the topic of this article have been declared.

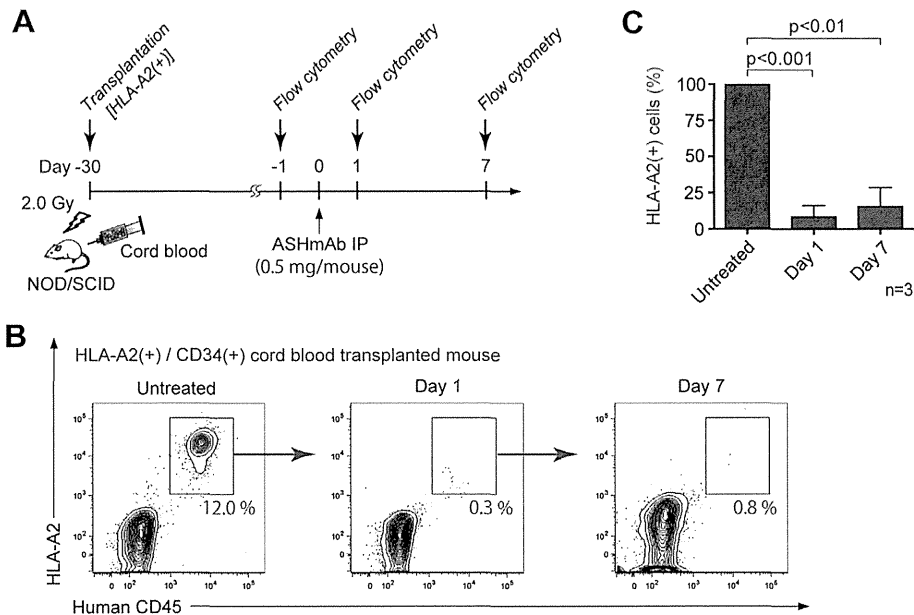
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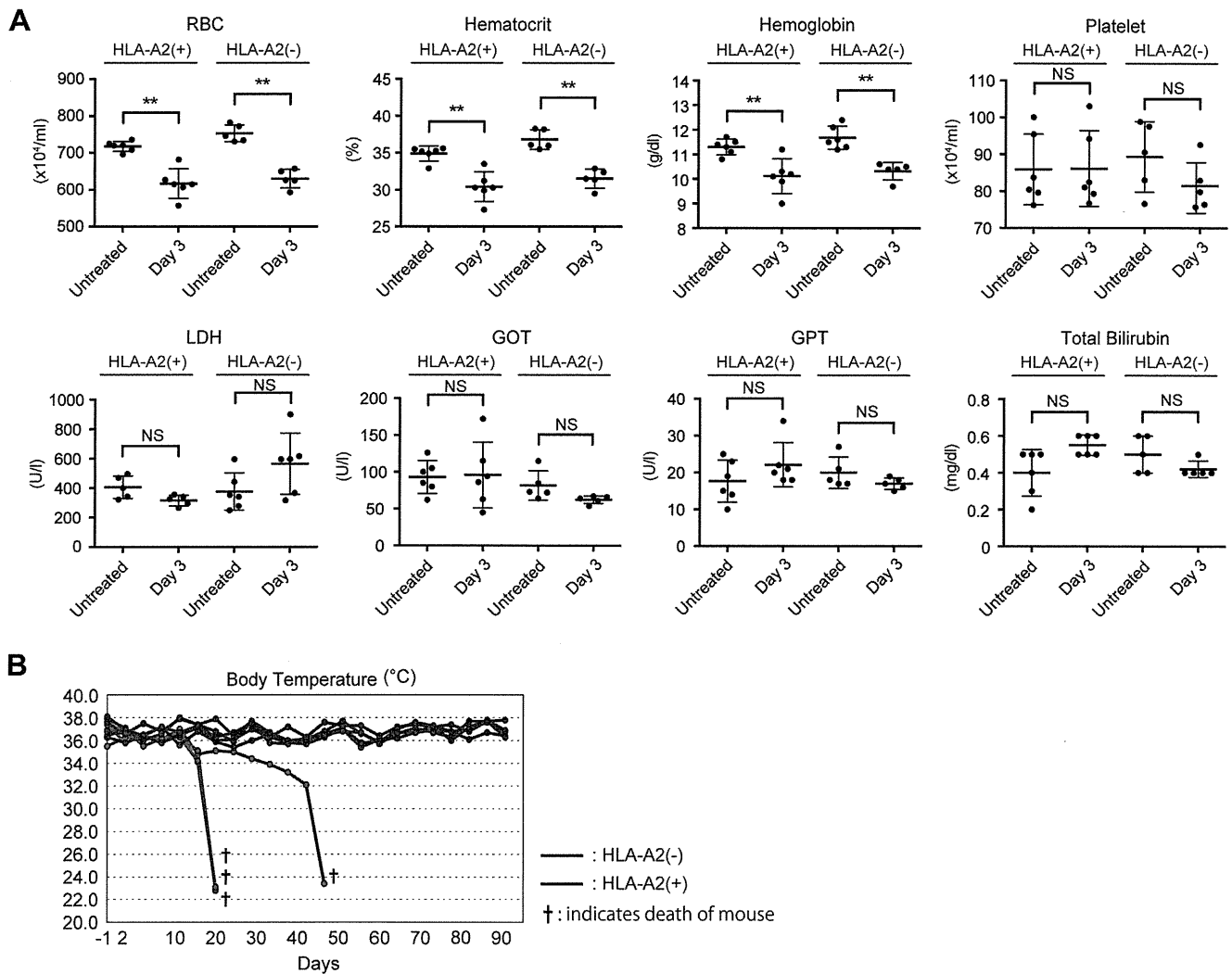
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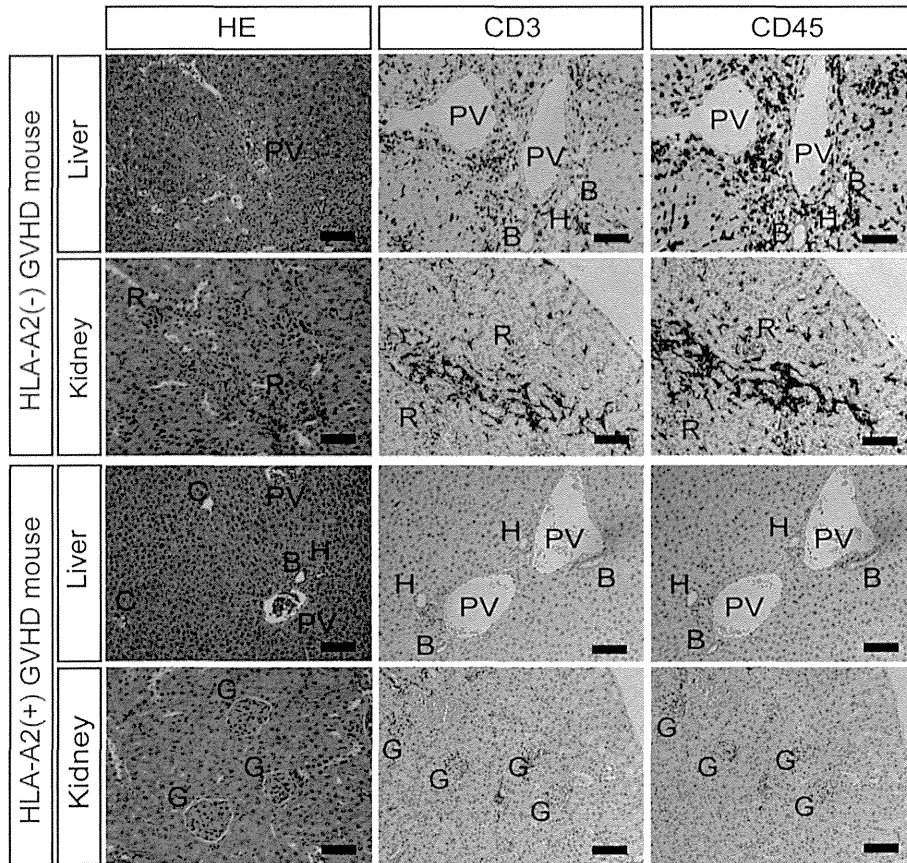
Supplementary Figure E1. Schedule for ASHmAb generation and screening. Day 0: First immunization of HLA-transgenic (Tg) mouse with HLA tetramer. Day 9: Mouse serum assessed for antibodies. Day 10: Second immunization (boost injection) of HLA-Tg mouse with HLA tetramer. Day 13: Fusion of spleen- and lymph node-derived B cells with SP2/0 myeloma cells. Fused cells were cultured in medium without aminopterin for 24 hours. Day 14: Suspension of fused cell-daughter cell candidates in methylcellulose-based HAT medium in 10-cm dishes for culture pending selection. Days 20-22: Selection of hybridoma colonies replating into 96-well plates filled with Dulbecco’s modified Eagle medium. Days 27-28: Screening of supernatants from hybridoma culture; initially by enzyme-linked immunosorbent assay, secondarily by flow cytometry. ELISA = enzyme-linked immunosorbent assay; HAT = hypoxanthine-aminopterin-thymidine; HLA = human leukocyte antigen.



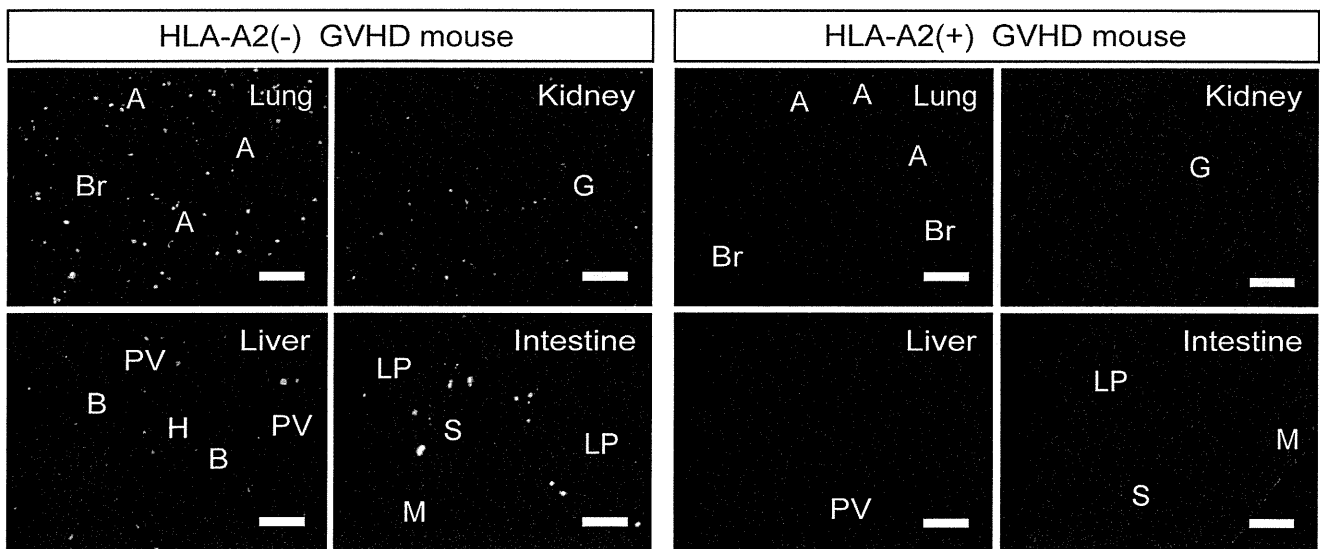
Supplementary Figure E2. Allele-specific cytotoxicity of HLA-A2-killing ASHmAb (kASHmAb) when injected intraperitoneally (IP). (A, B) Schedule and representative examples of in vivo assay of IP-administered HLA-A2 kASHmAb. Cord blood-derived CD34(+) / HLA-A2(+) cells were transplanted into irradiated NOD/SCID mice to create human-mouse bone marrow chimeric mice. One month after transplantation, flow-cytometric peripheral blood analysis was performed just before IP kASHmAb injection (0.5 mg, single dose) and on days 1 and 7 thereafter. Flow-cytometric analysis results show percentages of HLA-A2(+) / human CD45(+) cells in the marked gate. Doublets and dead cells were excluded. (C) Results of statistical analysis of (B). Blue bars represent HLA-A2(+) cells among humanized-mouse peripheral blood mononuclear cells (PBMC). Data shown as mean \pm s.d (n=3, p < 0.05 by Student’s t-test). ASHmAb = allele-specific anti-HLA monoclonal antibody; HLA = human leukocyte antigen.



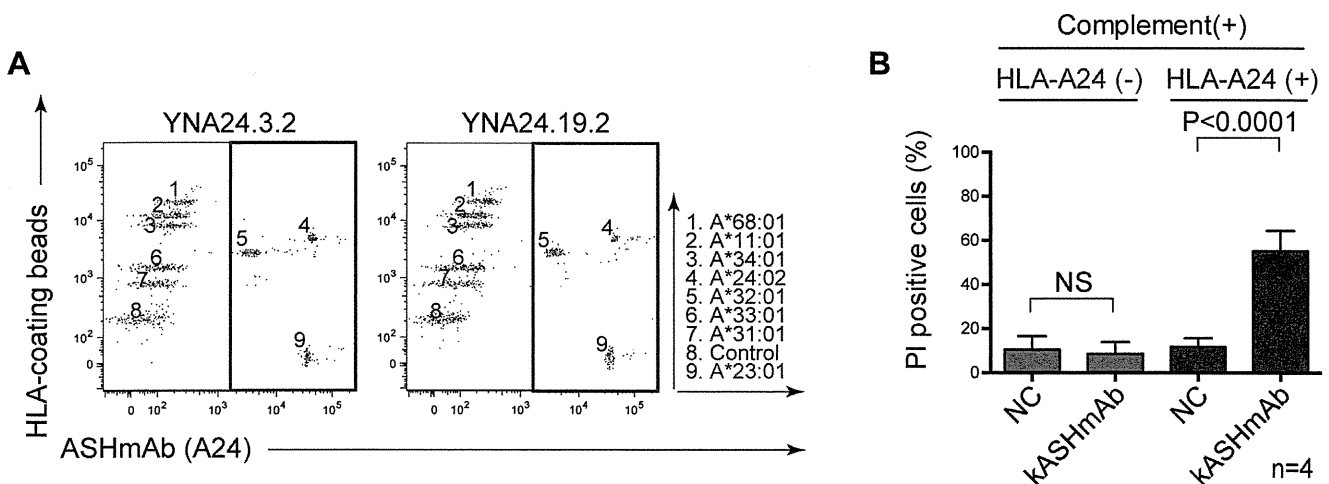
Supplementary Figure E3. Results of hematologic and biomarker studies of GVHD-model mice before and after kASHmAb treatment. (A) Blood was sampled from the retroorbital venous plexus and examined on days -1 (just before PBMC transplantation) and 3 (just before kASHmAb injection). We determined complete blood counts (CBC; hematocrit with red blood cell, hemoglobin, and platelet concentrations), serum total bilirubin concentration, LDH, GOT, and GPT activities in 5 HLA-A2(-) PBMC-transplanted mice and 6 HLA-A2(+) PBMC-transplanted mice. All mice were 7-8 weeks old. Paired t-test, $*P < 0.05$; $**P < 0.01$; NS: not significant. (B) Body temperature of GVHD-model mice before and after kASHmAb treatment [Red line; HLA-A2(-) PBMC-transplanted mice = GVHD(+), Blue line; HLA-A2(+) PBMC-transplanted mice = GVHD(-)]. Body temperature was determined twice weekly between 1500h and 1900h at room temperature 22°C and ambient humidity 55%. Percentage change from initial weight is shown. HLA = human leukocyte antigen; RBC = red blood cells; LDH = lactate dehydrogenase; GOT = glutamic oxaloacetic transaminase; GPT = glutamic pyruvic transaminase; GVHD = graft-versus-host disease; PBMC = peripheral blood mononuclear cell.



Supplementary Figure E4. Tissues from GVHD-model mice immunostained for human antigens (kidney and liver). Three weeks after transplantation, liver and kidneys were obtained from NOD/Shi-scid/IL-2R γ null mice intravenously transplanted with human PBMCs [HLA-A2(-) or HLA-A2(+)]. Sections of routinely processed formalin-fixed, paraffin-embedded material were stained with anti-human CD3 and CD45 antibodies with 3,3'-diamino-benzidine (DAB) substrate as chromogen and hematoxylin as nuclear counterstain. While HLA-A2(+) GVHD mouse shows no human derived-cells, HLA-A2(-) GVHD mouse organs show human CD3 or CD45 cells; magnification, original image, x200, liver; x400, kidney). Scale bar 400 μ m (kidney: 200 μ m). B = Bile duct; C = Central vein; G = Glomerulus; GVHD = graft-versus-host disease; H = Hepatic artery; HE = hematoxylin and eosin; HLA = human leukocyte antigen; PBMC = peripheral blood mononuclear cell ; PV = Portal vein; R = Renal corpuscle.



Supplementary Figure E5. Apoptosis in sections of organs of HLA-A2(-) and HLA-A2(+) GVHD-model mice after TdT-mediated dUTP nick end-labeling (TUNEL). Using routinely processed formalin-fixed, paraffin-embedded material, we evaluated apoptosis in sections of liver, lungs, intestine, and kidneys of HLA-A2(-) and HLA-A2(+) GVHD-model mice, obtained 3 weeks after transplantation, by light microscopy after TdT-mediated dUTP nick end-labeling. Green dots indicate apoptotic cells. Magnification, original images, x200. Scale bars 400 μ m. A = Alveoli; B = Bile duct; Br = Bronchiole; G = Glomerulus; GVHD = graft-versus-host disease; H = Hepatic artery; HLA = human leukocyte antigen; LP = Lamina propria; M = Muscularis; PV = Portal vein; S = submucosa.



Supplementary Figure E6. Allele-specific cytotoxicity of HLA-A24-killing ASHmAb (in vitro). (A) To establish ASHmAb-producing clones, on day 28 hybridomas (Supplementary Fig. 1) selected as producing anti-HLA antibody at initial screening were secondarily screened with FlowPRA, using flow cytometry (YNA24.3.2 and YNA24.19.2). Hybridoma-culture supernatants were incubated with HLA-coated beads. After incubation, the beads were washed and stained with secondary antibodies. Bead fluorescence intensities were measured using a flow cytometer and specificities of anti-HLA monoclonal antibodies were determined. Representative flow cytometry data are shown for a combination of FlowPRA beads coated with HLA-A*68:01, A*11:01, A*34:01, A*24:02, A*32:01, A*33:01, A*31:01, and control antigen or A*23:01. (B) Statistical analysis of in vitro killing assay using A24 ASHmAb with baby-rabbit complement. HLA-A24(-) cells (red bar) and HLA-A24(+) cells (blue bar) from healthy donors were each cultured with isotype control or A24 ASHmAb hybridoma supernatant. Percentages of dead cells were determined by propidium iodide staining and flow cytometry. Doublets and dead cells were excluded from flow-cytometry data. NC; Negative control. NS; Not significant. Data shown as mean \pm s.d (n=4, p < 0.05 by Student's t-test). ASHmAb = allele-specific anti-HLA monoclonal antibody; HLA = human leukocyte antigen; kASHmAb = killing ASHmAb; NC = negative control; NS = nonsignificant; PI = propidium iodide.

Single-Unit Cord Blood Transplantation after Granulocyte Colony-Stimulating Factor–Combined Myeloablative Conditioning for Myeloid Malignancies Not in Remission



Takaaki Konuma^{1,*}, Seiko Kato¹, Jun Ooi²,
Maki Oiwa-Monna¹, Yasuhiro Ebihara¹, Shinji Mochizuki¹,
Koichiro Yuji¹, Nobuhiro Ohno¹, Toyotaka Kawamata¹,
Norihide Jo¹, Kazuaki Yokoyama¹, Kaoru Uchimaru¹,
Shigetaka Asano³, Arinobu Tojo¹, Satoshi Takahashi¹

¹ Department of Hematology/Oncology, Institute of Medical Science, University of Tokyo, Tokyo, Japan

² Department of Hematology/Oncology, Teikyo University School of Medicine, Tokyo, Japan

³ System Medical Biology Laboratory, School of Advanced Science and Engineering, Waseda University, Tokyo, Japan

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ABSTRACT

High disease burden in myeloablative allogeneic hematopoietic stem cell transplantation is associated with adverse outcomes in patients with acute myelogenous leukemia (AML) and myelodysplastic syndrome (MDS). Quiescent leukemia stem cells could be induced to enter cell cycle by granulocyte colony-stimulating factor (G-CSF) administration and become more susceptible to chemotherapy. We report on the outcome of unrelated cord blood transplantation (CBT) using a conditioning regimen of 12 Gy total body irradiation, G-CSF–combined high-dose cytarabine, and cyclophosphamide in 61 adult patients with AML or advanced MDS not in remission. With a median follow-up of 97 months, the probability of overall survival and cumulative incidence of relapse at 7 years were 61.4% and 30.5%, respectively. In multivariate analysis, poor-risk cytogenetics and high lactate dehydrogenase values at CBT were independently associated with inferior survival. These data demonstrate that CBT after G-CSF–combined myeloablative conditioning is a promising curative option for patients with myeloid malignancies not in remission.

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INTRODUCTION

The prognoses of patients with acute myelogenous leukemia (AML) and advanced myelodysplastic syndrome (MDS) who have not achieved remission after chemotherapy have been poor. Although allogeneic hematopoietic stem cell transplantation (allo-HSCT) is the only potentially curative therapy for such patients, high disease burden has been reported to be associated with increased relapse or poor survival rate after allo-HSCT [1–9]. Recently, cord blood (CB) has been considered an acceptable alternative as a source of hematopoietic stem cells in unrelated allo-HSCT for adult patients without HLA-identical related or unrelated donors [9–16]. In comparison with other sources of allo-HSCT, one of the main advantages of using CB for patients with a high disease burden who require urgent transplantation is its rapid and convenient availability. Because it was shown that administration of granulocyte colony-stimulating factor (G-CSF) increased the susceptibility of cell-cycle-specific agent cytarabine in leukemia cells in vitro [17], we administered G-CSF–combined high-dose cytarabine in myeloablative conditioning for allo-HSCT [18,19] and reported that a G-CSF–combined conditioning regimen provided better engraftment and survival results in cord blood

transplantation (CBT) for myeloid malignancies [13–16]. The objective of this retrospective study was to confirm the effects of CBT after G-CSF–combined myeloablative conditioning in adult patients with myeloid malignancies not in remission and to identify variables influencing long-term outcomes.

PATIENTS AND METHODS

Patients and Transplantation Procedures

This retrospective study included 61 consecutive adult patients who underwent unrelated transplantation using single-unit CB for AML or advanced MDS not in remission at our institute between 1998 and 2013. Thirty-two patients were included in our previous study [15,16] and extended the follow-up. The diagnoses of AML and MDS were made according to the World Health Organization classification. Advanced MDS was defined as having refractory anemia with excess blasts type 1 or refractory anemia with excess blasts type 2 by World Health Organization classification. Myeloid malignancies not in remission were defined as more than 5% blasts in the bone marrow (BM), or circulating blasts in peripheral blood (PB) or central nervous system. The cytogenetic subgroups were defined according to the Southwest Oncology Group/Eastern Cooperative Oncology Group criteria for AML [20] and International Prognostic Scoring System criteria for MDS [21]. All patients received 12 Gy total body irradiation (TBI) in 4 divided fractions on days –8 and –7, cytarabine on days –5 and –4 (total dose 12 g/m², and 3 g/m² every 12 hours for 2 days) with 5 µg/kg G-CSF (lenograstim) from 12 hours before the first dose of cytarabine to the end of cytarabine dosing, and cyclophosphamide (total dose 120 mg/kg) on days –3 and –2 [15,16]. Fifty-eight patients received cyclosporine (CSP) (3 mg/kg/day) with a short course of methotrexate (15 mg/m² on day +1 and 10 mg/m² on days +3 and +6), and 3 patients received CSP only as graft-versus-host disease (GVHD) prophylaxis. CB units were obtained from the Japanese Cord Blood Bank Network. Donor-recipient HLA-matching status was based on antigen level HLA-A and -B and on allele level HLA-DRB1 typing. All patients received similar supportive care and CB units were

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* Correspondence and reprint requests: Takaaki Konuma, Department of Hematology/Oncology, Institute of Medical Science, University of Tokyo, 4-6-1, Shirokanedai, Minato-ku, Tokyo 108-8639, Japan.

E-mail address: tkonuma@ims.u-tokyo.ac.jp (T. Konuma).

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Table 1
Characteristics of Patients, Cord Blood Units, and Transplantation

Characteristic	Value
No. of patients	61
Sex	
Male	36 (59)
Female	25 (41)
Age, median (range), yr	41 (18–55)
CMV serostatus	
Positive	54 (86)
Negative	7 (11)
Disease type	
De novo AML	24 (39)
AML secondary to MDS	24 (39)
Advanced MDS [*]	13 (21)
Cytogenetics [†]	
Good	1 (2)
Intermediate	27 (44)
Poor	30 (49)
Unknown	3 (5)
Bone marrow blasts at CBT, median (range), %	17.7 (1.4–86.0) [¶]
< 25%	39
≥ 25%	22
Peripheral blood blasts at CBT, median (range), %	6.5 (0–68.5)
Absent	12
Present	49
LDH at CBT	
≤ ULN	41 (67)
> ULN	20 (33)
Disease status at CBT [‡]	
Untreated	31 (51)
Primary refractory	14 (23)
Refractory relapse	16 (26)
Time from diagnosis to CBT, median (range), mo	7 (1–219)
Conditioning regimen	
TBI12Gy+Ara-C/G-CSF+CY	61
GVHD prophylaxis	
CyclosporineA+methotrexate	58 (95)
CyclosporineA	3 (5)
Number of nucleated cells, median (range), ×10 ⁷ /kg	2.43 (1.32–5.50)
Number of CD34 ⁺ cells, median (range), ×10 ⁵ /kg	1.03 (.21–2.27)
HLA disparities [§]	
1	13 (21)
2	32 (52)
3	14 (22)
4	2 (3)

CMV indicates cytomegalovirus; AML, acute myelogenous leukemia; MDS, myelodysplastic syndrome; CBT, cord blood transplantation; LDH, lactate dehydrogenase; ULN, upper limit of normal; TBI, total body irradiation; Ara-C, cytosine arabinoside; G-CSF, granulocyte colony-stimulating factor; CY, cyclophosphamide; GVHD, graft-versus-host disease; HLA, human leukocyte antigen.

Data presented are n (%) unless otherwise indicated.

* Advanced MDS are defined as having refractory anemia with excess blasts-1 (RAEB-1) or RAEB-2 by WHO criteria.

† The cytogenetic subgroups according to the Southwest Oncology Group/Eastern Cooperative Oncology Group criteria for AML and International Prognostic Scoring System criteria for MDS.

‡ Untreated was defined as no treatment before conditioning regimen, indicating that the majority of patients with AML secondary to MDS or advanced MDS received CBT as an up-front treatment. Primary refractory was defined as failure to achieve complete remission with induction chemotherapy. Refractory relapse was defined as failure to achieve complete remission with salvage chemotherapy after first or subsequent relapse.

§ The number of HLA disparities, defined as the low resolution for HLA-A and -B and the high resolution for HLA-DRB1.

¶ The 5 patients with less than 5% blasts in the bone marrow included circulating blasts in peripheral blood (n = 3) or central nervous system (n = 2).

selected, as previously reported [15,16]. The institutional review board of the Institute of Medical Science, University of Tokyo approved this study. This study was conducted in accordance with the Declaration of Helsinki.

End Points and Statistical Analysis

The primary study end point was overall survival (OS), defined as time from the date of transplantation to the date of death or last contact. Secondary end points were relapse, including disease progression before engraftment; transplantation-related mortality (TRM); neutrophil and platelet engraftment; acute graft-versus-host disease (aGVHD); and chronic GVHD (cGVHD). Relapse was defined as morphologic evidence of disease in PB, BM, or extramedullary sites. TRM was defined as death during remission. Neutrophil engraftment was defined as the first of 3 consecutive days during which the absolute neutrophil count was at least $.5 \times 10^9/L$. Platelet engraftment was achieved on the first of 3 days when the platelet count was higher than $50 \times 10^9/L$ without transfusion support. Both aGVHD and cGVHD were graded according to the previously published criteria [22,23].

The incidence of aGVHD was evaluated in all engrafted patients, whereas the incidence of cGVHD was evaluated in engrafted patients surviving more than 100 days.

The probability of OS was estimated according to the Kaplan-Meier method, and the groups were compared using the log-rank test. The probabilities of relapse, TRM, neutrophil and platelet engraftment, and acute and chronic GVHD were estimated based on a cumulative incidence method to accommodate competing risks [24]. Multivariate analysis was performed with a Cox proportional hazard model adjusted for OS and Fine and Gray proportional hazards model for relapse [25]. The following variables were considered: age (< 45 versus ≥ 45 years), disease type (de novo AML versus AML secondary to MDS versus advanced MDS), cytogenetic risk (other than poor versus poor), proportion of blasts in BM (< 25 versus ≥ 25%), the presence of blasts in PB (absent versus present), lactate dehydrogenase (LDH) at CBT (≤ upper limit of normal versus > upper limit of normal), disease status at CBT (untreated versus primary refractory versus refractory relapse), cord blood nucleated cell count (< 2.5 versus ≥ 2.5 × 10⁷/kg), and HLA disparities based on antigen level HLA-A and -B and allele level

HLA-DRB1 (≤ 2 versus ≥ 3). All statistical analyses were performed with EZR (Saitama Medical Center, Jichi Medical University, Saitama, Japan), a graphical user interface for R 2.13.0 (R Foundation for Statistical Computing, Vienna, Austria) [26]. $P < .05$ was considered significant. Analysis of data was performed in August 2013.

RESULTS

Patient and CB unit characteristics are shown in Table 1. The median age was 41 years (range, 18 to 55 years), the median number of nucleated cells was $2.43 \times 10^7/\text{kg}$ (range, 1.32 to $5.50 \times 10^7/\text{kg}$), and the median number of CD34+ cells was $1.03 \times 10^5/\text{kg}$ (range, $.21$ to $2.27 \times 10^5/\text{kg}$). Disease types were de novo AML in 24 patients, AML secondary to MDS in 24, and advanced MDS in 13. The majority of patients with de novo AML with multilineage dysplasia ($n = 2$), AML secondary to MDS ($n = 19$), or advanced MDS ($n = 10$) received CBT as an up-front treatment, which was classified as untreated group ($n = 31$). Among patients with primary refractory status ($n = 14$), 3 patients received CBT after the first cycle of induction chemotherapy. The median number of prior chemotherapy treatments before CBT for primary refractory status was 3 (range, 1 to 5). The median time from diagnosis to CBT was 7 months (range, 1 to 219 months), and the median period of follow-up for survivors after CBT was 97 months (range, 5 to 181 months).

The cumulative incidence of neutrophil recovery was 93.4% (95% confidence interval [CI], 81.0% to 97.8%) at 60 days after CBT with a median time to achieve greater than $.5 \times 10^9/\text{L}$ neutrophils of 22 days (range, 18 to 41 days). Disease progression before engraftment occurred in 2 patients. The cumulative incidence of platelet recovery was 78.7% (95% CI, 65.7% to 87.2%) at 100 days after CBT with a median time to an untransfused platelet count greater than $50 \times 10^9/\text{L}$ of 50 days (range, 30 to 179 days). The cumulative incidences of grade II to IV acute GVHD and extensive chronic GVHD were 62.3% (95% CI, 48.7% to 73.2%) at 100 days and 32.9% (95% CI, 21.4% to 44.9%) at 3 years after CBT, respectively. The probability of OS at 7 years was 61.4% (95% CI, 47.1% to 72.9%). The cumulative incidence of relapse at 7 years was 30.5% (95% CI, 19.2% to 42.6%). The cumulative incidence of TRM at 100 days and at 1 year was 6.6% (95% CI, 2.1% to 14.7%) and 8.2% (95% CI, 3.0% to 16.9%), respectively (Figure 1).

In multivariate analysis, poor-risk cytogenetics (hazard ratio [HR], 7.14; 95% CI, 2.33 to 21.80; $P < .001$) and high LDH value (HR, 4.00; 95% CI, 1.33 to 12.07; $P = .013$) were associated with inferior survival (Figure 2, Table 2). De novo AML (HR, 9.66; 95% CI, 1.06 to 87.75; $P = .044$), primary refractory status at CBT (HR, 6.47; 95% CI, 1.86 to 22.51; $P = .003$), and high LDH value (HR, 3.75; 95% CI, 1.11 to 12.57; $P = .032$) were associated with an increased relapse incidence (Table 3, Supplemental Figure 1). In contrast, the proportion of blasts

in BM and the presence of blasts in PB did not show any impact on survival and relapse incidence.

DISCUSSION

Previous reports have suggested that the only potentially curative therapy for patients with myeloid malignancies not in remission is allo-HSCT. However, the incidence of relapse has been reported to be high, and several reports showed long-term survival rates of only 10% to 30% [1–6]. Several factors, including blasts in BM or PB, cytogenetics, and donor availability, have been associated with outcome. In this study, poor-risk cytogenetics and high LDH value were significantly associated with inferior OS. De novo AML, primary refractory status, and high LDH value were associated with increased relapse. However, we found no impact of disease burden on survival and relapse. In fact, several retrospective studies did not show any advantage of induction chemotherapy before allo-HSCT to reduce the disease burden for patients with advanced MDS or AML secondary to MDS [27–29]. Therefore, the majority of patients with advanced MDS or AML secondary to MDS received G-CSF–combined myeloablative conditioning followed by CBT without prior induction chemotherapy in our institute.

After physicians have decided that allo-HSCT is appropriate for patients with myeloid malignancy not in remission, the elective timing of the transplantation is the main advantage of CBT. In fact, CBT timing is decided depending on the patient's conditions, such as control of infection and disease burden. Such elective timing of CBT might have contributed to disease burden not being shown to influence outcome in our study. On the other hand, the use of CB as a source of hematopoietic stem cells could offer the opportunity for patients to receive allo-HSCT without related or unrelated donors. Moreover, the lower incidence of severe GVHD without compromising graft-versus-leukemia effects in CBT may also have contributed to long-term survival in our study.

Relapse is the most important cause of treatment failure after allo-HSCT, particularly in patients with myeloid malignancies not in remission. This is mainly due to the residual leukemic cells that have escaped the cytotoxic effect of conditioning before transplantation. To reduce disease relapse, the role of a more intense conditioning regimen has been analyzed extensively [30]. Since chemosensitization of leukemia cells with G-CSF enhances the cytotoxicity of the cell-cycle–specific agent cytarabine [17], we administered G-CSF–combined high-dose cytarabine in the standard conditioning regimen of TBI/cyclophosphamide. The clinical efficacy of concomitant use of G-CSF with chemotherapy has remained controversial in newly diagnosed or relapsed refractory AML and MDS [31,32]. Recently, Pabst et al. reported

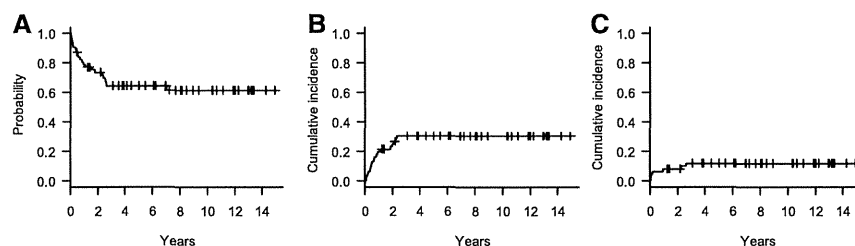


Figure 1. Probability of overall survival and cumulative incidences of relapse and transplant-related mortality after G-CSF–combined myeloablative CBT. Overall survival (A), relapse (B), and transplantation-related mortality (C) in 61 patients with AML or advanced MDS not in remission after G-CSF–combined myeloablative CBT.

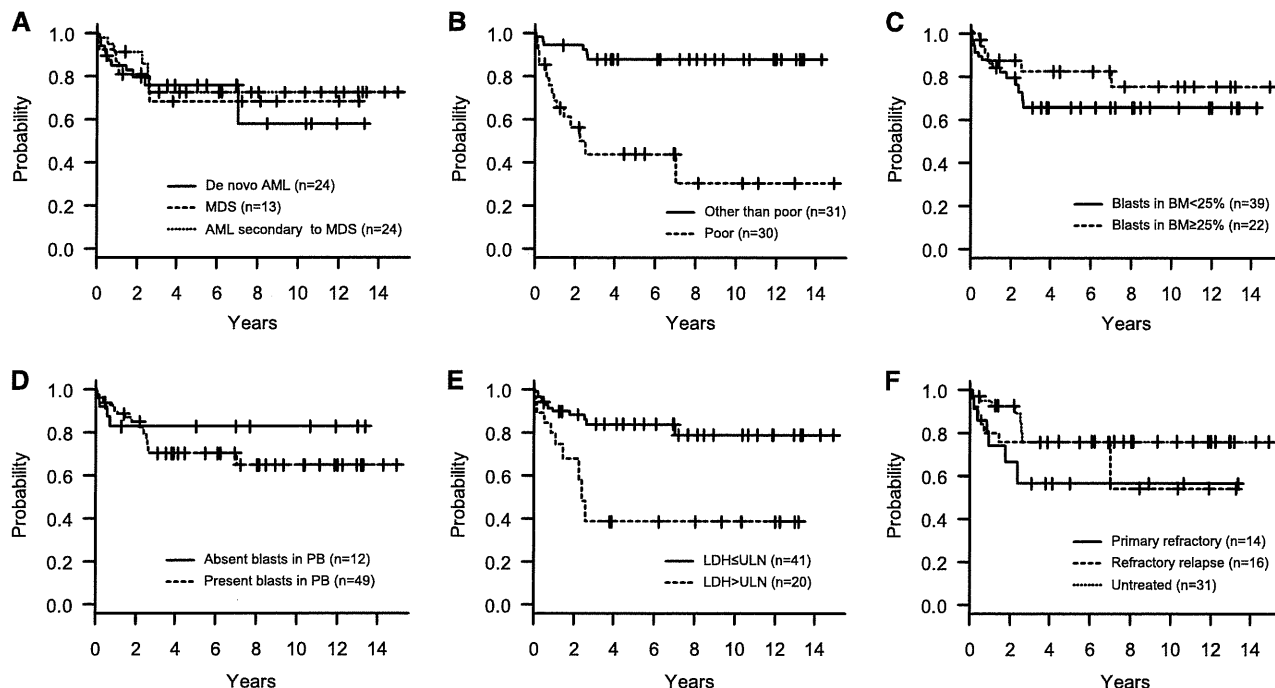


Figure 2. Adjusted probabilities of overall survival in 61 patients with AML and advanced MDS not in remission after G-CSF–combined myeloablative CBT. The adjusted probabilities of overall survival grouped according to the disease type (A), cytogenetic risk (B), the proportion of blasts in bone marrow (BM) (C), the presence of blasts in peripheral blood (PB) (D), the lactate dehydrogenase (LDH) value at cord blood transplantation (CBT) (E), and disease status at CBT (F). Multivariate analysis for overall survival is shown in Table 2.

Table 2
Univariate and Multivariate Analysis of Prognostic Factors for Survival

Variable	Univariate Analysis			Multivariate Analysis		
	Number	7-year OS (95% CI)	P	Hazard Ratio*	95% CI	P
Age						
< 45 years	36	63.5 (44.1–77.7)		1		
≥ 45	25	58.7 (36.7–75.4)	.555	.69	.25–1.86	.464
Disease type						
Advanced MDS	13	59.3 (27.5–81.0)		1		
AML secondary to MDS	24	74.4 (51.6–87.6)		.58	.13–2.54	.471
De novo AML	24	47.4 (23.0–68.4)	.234	.97	.18–5.16	.978
Cytogenetics [†]						
Other than poor	31	80.3 (61.3–90.6)		1		
Poor	30	38.9 (18.8–58.6)	.002	7.14	2.33–21.80	<.001
Bone marrow blasts at CBT, %						
< 25	39	58.0 (40.8–71.8)		1		
≥ 25	22	68.2 (41.2–84.7)	.297	.59	.16–2.09	.418
Peripheral blood blasts at CBT						
Absent	12	66.7 (33.7–86.0)		1		
Present	49	60.2 (44.0–73.1)	.983	1.18	.34–4.10	.787
LDH value at CBT						
≤ ULN	41	67.4 (48.9–80.4)		1		
> ULN	20	50.0 (27.1–69.2)	.147	4.00	1.33–12.07	.013
Disease status at CBT						
Untreated	31	71.1 (50.1–84.5)		1		
Primary refractory	14	50.0 (22.9–72.2)		2.76	.78–9.77	.114
Refractory relapse	16	50.0 (20.2–74.1)	.234	1.75	.30–10.22	.530
Number of nucleated cells, ×10 ⁷ /kg						
≥ 2.5	29	59.2 (37.9–75.3)		1		
< 2.5	32	64.1 (44.3–78.4)	.989	.99	.38–2.58	.989
HLA disparities [‡]						
≤ 2	45	60.3 (43.7–73.4)		1		
≥ 3	16	65.0 (35.1–83.7)	.597	.98	.30–3.18	.975

MDS indicates myelodysplastic syndrome; AML, acute myelogenous leukemia; CBT, cord blood transplantation; LDH, lactate dehydrogenase; ULN, upper limit of normal; HLA, human leukocyte antigen; OS, overall survival; CI, confidence interval.

* Hazards ratio for overall mortality.

[†] The cytogenetic subgroups according to the Southwest Oncology Group/Eastern Cooperative Oncology Group criteria for AML and International Prognostic Scoring System criteria for MDS.

[‡] The number of HLA disparities defined as the low resolution for HLA-A and -B and the high resolution for HLA-DRB1.

Table 3
Univariate and Multivariate Analysis of Prognostic Factors for Relapse

Variable	Univariate Analysis			Multivariate Analysis		
	Number	7-year Relapse (95% CI)	P	Hazard Ratio	95% CI	P
Age						
< 45	36	29.3 (15.0–45.2)		1		
≥ 45	25	32.0 (14.9–50.6)	.567	1.62	.50–5.17	.420
Disease type						
Advanced MDS	13	7.7 (.4–30.5)		1		
AML secondary to MDS	24	29.8 (12.9–49.0)		4.37	.38–49.80	.230
De novo AML	24	43.4 (22.4–62.7)	.096	9.66	1.06–87.75	.044
Cytogenetics ^c						
Other than poor	31	23.0 (9.9–39.2)		1		
Poor	30	38.2 (20.5–55.7)	.163	2.33	.90–5.97	.078
Bone marrow blasts at CBT, %						
< 25	39	26.0 (13.3–40.6)		1		
≥ 25	22	39.2 (18.0–59.9)	.397	1.72	.57–5.16	.330
Peripheral blood blasts at CBT						
Absent	12	16.7 (2.3–42.8)		1		
Present	49	33.8 (20.6–47.4)	.309	3.08	.40–23.70	.280
LDH value at CBT						
≤ ULN	41	25.6 (13.1–40.1)		1		
> ULN	20	40.0 (18.5–60.8)	.240	3.75	1.11–12.57	.032
Disease status at CBT						
Untreated	31	17.8 (6.3–34.1)		1		
Primary refractory	14	50.0 (21.4–73.3)		6.47	1.86–22.51	.003
Refractory relapse	16	37.5 (14.5–60.7)	.043	1.36	.26–7.05	.71
Number of nucleated cells, ×10 ⁷ /kg						
≥ 2.5	29	35.5 (18.3–53.1)		1		
< 2.5	32	25.3 (11.7–41.5)	.525	.54	.14–2.12	.380
HLA disparities [†]						
≤ 2	45	34.0 (20.4–48.1)		1		
≥ 3	16	20.3 (4.5–43.9)	.306	.53	.11–2.49	.420

MDS indicates myelodysplastic syndrome; AML, acute myelogenous leukemia; CBT, cord blood transplantation; LDH, lactate dehydrogenase; ULN, upper limit of normal; HLA, human leukocyte antigen; CI, confidence interval.

^c The cytogenetic subgroups according to the Southwest Oncology Group/Eastern Cooperative Oncology Group criteria for AML and International Prognostic Scoring System criteria for MDS.

[†] The number of HLA disparities defined as the low resolution for HLA-A and -B and the high resolution for HLA-DRB1.

significantly improved survival with concomitant use of G-CSF with escalated-dose, but not with conventional-dose cytarabine [31]. In the setting of allo-HSCT, the conditioning regimen consisting of G-CSF–combined high-dose cytarabine and TBI 12 Gy was feasible and might reduce post-transplantation relapse in patients with AML [18,19]. The presence of quiescent leukemia stem cells (LSCs), which are thought to be resistant to chemotherapy, might contribute to relapse after treatment. Recently, a xenograft model demonstrated that cytarabine with G-CSF recruited quiescent LSCs into a phase of the cell cycle, leading to enhanced elimination of LSCs within the niche [33]. This effect might have contributed to reduced relapse in our study. Although these findings should be confirmed in prospective studies, the combination of G-CSF–combined myeloablative conditioning with CBT offered a promising curative option for patients with myeloid malignancies not in remission.

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SUPPLEMENTARY DATA

Supplementary data related to this article can be found at <http://dx.doi.org/10.1016/j.bbmt.2013.12.555>.

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