ORIGINAL ARTICLE



Reduced-intensity conditioning is effective for allogeneic hematopoietic stem cell transplantation in infants with *MECOM*-associated syndrome

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Abstract

Mutations in the *MECOM* encoding EVI1 are observed in infants who have radioulnar synostosis with amegakaryocytic thrombocytopenia. *MECOM*-associated syndrome was proposed based on clinical heterogeneity. Allogeneic hematopoietic stem cell transplantation (HSCT) is a curative treatment for progressive bone marrow failure. However, data regarding allogeneic HSCT for this rare disease are limited. We retrospectively assessed overall survival, conditioning regimen, regimen-related toxicities and long-term sequelae in six patients treated with allogeneic HSCT. All patients received a reduced-intensity conditioning (RIC) regimen consisting of fludarabine, cyclophosphamide or melphalan, and rabbit anti-thymocyte globulin and/or low-dose total body/thoracic-abdominal/total lymphoid irradiation, followed by allogeneic bone marrow or cord blood transplantation from unrelated donors between 4 and 18 months of age. All patients survived and achieved stable engraftment and complete chimerization with the donor type. Moreover, no patient experienced severe regimen-related toxicities, and only lower grades of acute graft-versus-host disease were observed. Three patients treated with low-dose irradiation had relatively short stature compared to three patients not treated with irradiation. Therefore, allogeneic HSCT with RIC is an effective and feasible treatment for infants with *MECOM*-associated syndrome. Future studies are needed to evaluate the use of low-dose irradiation to avoid risks of other long-term sequelae.

Keywords Inherited bone marrow failure syndrome \cdot Radio-ulnar synostosis with amegakaryocytic thrombocytopenia \cdot *MECOM*-associated syndrome \cdot Reduced-intensity conditioning \cdot Allogeneic hematopoietic stem cell transplantation

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Introduction

Radioulnar synostosis with amegakaryocytic thrombocytopenia (RUSAT) is an inherited bone marrow failure syndrome (IBMFS). This condition is characterized by thrombocytopenia, which progresses to pancytopenia, and congenital proximal fusion of the radius and ulna [1]. In a previous report, two unrelated families presented with RUSAT caused by *HOXA11* mutations [2]. However, not all cases of RUSAT are due to *HOXA11* mutations, and additional genetic loci are also responsible for this condition [3].

We initially reported three patients with RUSAT who presented with heterozygous missense mutations in the *MECOM* encoding the oncoprotein EVI1. These missense mutations were clustered within the 8th zinc finger motif, localized at the C-terminus of the *MECOM*. Moreover, functional assays revealed the critical role of EVI1 in normal hematopoiesis and the development of forelimbs and fingers in humans [4].

The *MECOM*-associated syndrome, a recently discovered disease, was proposed based on clinical findings. That is, patients with *MECOM* mutations have clinical phenotypic heterogeneity for BMF and proximal radioulnar synostosis (RUS). A previous study presented 12 patients, including familial and sporadic patients, with germline mutation in the *MECOM*, and their broad clinical spectrum ranged from isolated RUS with or without mild hematological abnormalities to severe IBMFS without evident skeletal abnormalities [5].

Allogeneic hematopoietic stem cell transplantation (HSCT) is a curative treatment for progressive bone marrow failure in patients with *MECOM*-associated syndrome. However, the appropriate conditioning regimens for this condition is yet to be determined, and the characteristics of early infants treated with HSCT have not been evaluated.

Hence, in this study, we analyzed the clinical outcomes of six patients treated with reduced-intensity conditioning (RIC) regimens and allogeneic HSCT. We provided insights on the effectivity of these regimens as well as their associated risks for infants with *MECOM*-associated syndrome.

Patients and methods

WE retrospectively summarized the clinical and genetic profiles of six patients with *MECOM*-associated syndrome who were treated with allogeneic HSCT and reported in literatures or abstracts in Japan. These included family history, sex, weeks of gestation, initial clinical findings, presence of bone and other abnormalities, hematological data, transfusion dependency, age at progression to pancytopenia, type of *MECOM* mutation, and alterations in the EVI1 protein.

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Moreover, we summarized the data on overall survival rate, age at transplantation, source of HSC, human leukocyte antigen (HLA) compatibility, type of conditioning regimen, number of total infused nuclear cells, status of bone marrow chimera, administration of graft-versus-host disease (GVHD) prophylaxis, grades of acute GVHD, chronic GVHD, presence of regimen-related toxicities, and longterm sequelae.

Statistical analysis was performed using Student's t test, and a p value < 0.01 was considered statistically significant.

This study was approved by ethics committee of Tohoku University Graduate School of Medicine, and written informed consent was obtained from the patients' parents.

Results

Patients' characteristics

The clinical and genetic profiles of the six patients (Pts) with MECOM-associated syndrome enrolled in this study are shown in Table 1. Pts 5 and 6 had a family history of RUS, and with Pt 6 having maternal history of mild hematological abnormality. Meanwhile, four patients had de novo mutations in MECOM. Initial findings included petechiae, pulmonary bleeding, severe anemia, and fatal distress. Pts 1-3 presented with RUS and bone abnormality. Meanwhile, Pts 4-6 had no RUS and Pt 5 had bone abnormality at birth. Pts 2 and 3 had hearing disability. All patients rapidly progressed to severe pancytopenia or bicytopenia between 0 and 5 months of age, and all of them required repeated transfusion to prevent severe bleeding and anemia. Moreover, they were at a high risk of life-threatening infections due to severe neutropenia. Thus, prophylactic antibiotics and antifungal agents were required.

All heterozygous missense and splice-site mutations were clustered within the 8th zinc finger motif, localized at the C-terminus of the *MECOM*, as described previously (Fig. 1). Among the four patients with de novo mutations in MECOM gene, Pts 1-3 had heterozygous de novo missense mutations (c.2248C>T [p.Arg750Trp], c.2252A>G [p.His751Arg], and c.2266A > G [p.Thr756Ala]) [4], while Pt 4 had a heterozygous de novo mutation (c.2248C>T [p.Arg750Trp]) and somatic mosaicism in the MECOM [6]. Pt 5 and her brother, father, and uncle had heterozygous splice-site mutations (c.2208-4A > G), resulting in p.Cys735_Arg736insSer (CAG insertion) of the EVI1 protein. Further, Pt 6 and her mother had a heterozygous splice-site mutation (c.2285 + 1G > A), resulting in skipping of exon 11 including the 8th zinc finger motif and insertion of intron 11, and somatic loss of heterozygosity (LOH) which reduced the allele fraction of the mutation in blood cells [7].

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| Pt No | 1 | 2 | 3 | 4 | 5 | 6 |
|--|--|---|--|--|---|--|
| Family history | no | no | no | no | RUS in father and uncle Clinodac- tyly in brother, father, and uncle | RUS, congenital left clubfoot, bilateral CDH, chronic thrombocytopenia and transient leu- kopenia in mother |
| Weeks of gestation (weeks) | 35 | 37 | 31 | 40 | 37 | 38 |
| Birth weight (g) | 2160 | 2058 | 2180 | 2936 | 2368 | 3414 |
| Gender | Female | Female | Male | Female | Female | Female |
| Initial findings | Fetal distress Systemic Pete- chiae Severe anemia | Systemic pete- chiae | Fetal hydrops Severe anemia | Pulmonary bleed- ing Severe anemia Thrombocytopenia | Systemic pete- chiae Severe anemia | Systemic petechiae Severe anemia ICH, Convulsion |
| Bone abnormali- ties | RUS Bilateral bony defect of the intermediate phalanges of the fifth digits | RUS Bilateral clinodac- tyly of the fifth digits | RUS Overlapping fingers | No | Bilateral clinodac- tyly of the fifth digits | No |
| Hearing | Normal | Sensorineural hearing impair- ment: Rt 55 dB, Lt 34 dB | Prelingual senso- rineural hearing impairment: Rt 60 dB, Lt 25 dB | Normal | Normal | Normal |
| Leukocyte count at birth (/mm ³) | 6780 | 17,100 | 3220 | 14,220 | 10,700 | 7600 |
| Hemoglobin count at birth (g/dL) | 4.0 | 12.9 | 2.7 | 7.2 | 7.3 | 6.3 |
| Platelet count at birth (/mm ³) | 5000 | 8000 | 89,000 | 9000 | 4000 | 7000 |
| Transfusion dependency | RBC, PC | RBC, PC | RBC, PC | RBC, PC | RBC, PC | RBC, PC \Rightarrow PC only |
| Progression to pancytopenia (months) | 2 | 5 | at birth | 2 | 4 | Bicytopenia only |
| Heterozygous mutations of MECOM gene | c.2266A>G | c.2252A>G | c.2248C>T | c.2248C>T, somatic mosai- cism | c.2208-4A>G | c.2285+1 G>A, LOH |
| Alteration of EVI1 protein | p.Thr756Ala | p.His751Arg | p.Arg750Trp | p.Arg750Trp | p.Cys735-Arg736 ins Ser | exon 11 skipping and ins intron 11 |

Table 1 Clinical and genetic profiles of six patients with MECOM-associated syndrome in this study

Pt patient, RUS radioulnar synostosis, CDH congenital dislocation of the hip joint, ICH intracranial hemorrhage, Rt right, Lt left, RBC red blood cell concentrate, PC platelet concentrate LOH loss of heterozygosity

Donor, stem cell source and GVHD prophylaxis

Data on RIC and allogeneic HSCT are shown in Table 2. The patient's age at HSCT was between 4 and 18 months. The sources of donor cells were bone marrow from an unrelated donor in three patients and unrelated cord blood in the other three patients. The number of total infused nuclear cells was sufficient for engraftment in all patients. HLA compatibility was 8/8 or 7/8 matched in alleles in unrelated bone marrow transplantations and 7/8 or 4/8 matched in alleles in cord blood transplantation (CBT). Regarding GVHD prophylaxis,

five patients received tacrolimus (FK506) and short-term methotrexate (MTX) while one patient received cyclosporin A (CyA) and short-term MTX.

Overall transplant outcomes: engraftment, complications and GVHD

The overall survival rate after receiving HSCT was 100% (Fig. 2).

Neutrophil (Neut) and platelet (Plt) engraftments were successfully achieved in all patients between days + 6

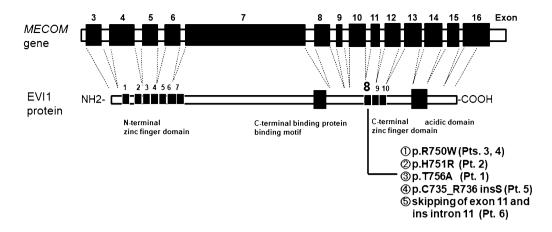


Fig. 1 Summary of MECOM mutations in six patients with MECOM-associated syndrome enrolled in this study [4, 6, 7]

and + 22, and between days + 22 and + 35, respectively. All patients achieved complete chimera of the donor type and independence from transfusion. No severe regimen-related toxicities were observed except grade 1 mucositis and veno-occlusive disease, which were treated with conventional therapies. Two patients presented with grade II acute GVHD of the skin that was easily controlled with 1 mg/kg prednisolone. None of the patients developed chronic GVHD.

Hematological profiles and HSCT regimens of each patient

The hematological profiles and conditioning regimens for each patient are presented in Tables 1 and 2, respectively.

Pt 1 presented with severe fetal distress at birth (35 weeks and 6 days of gestation), and her birth weight was 2160 g. She appeared extremely pale due to severe anemia, and extensive petechiae were observed on her lower abdomen. Laboratory data revealed a normal white blood cell (WBC) count (6780/mm³) with neutropenia (Neut count 594/mm³), severe anemia (hemoglobin [Hb] level: 4.0 g/dL), and a low Plt count (5000/mm³). The patient required mechanical ventilation, red blood cell (RBC) transfusion, platelet concentrate (PC) transfusion, and treatment for hypovolemic shock. The patient's general condition improved after treatment. Her bone marrow showed low cellularity without excess of blasts, absence of megakaryocytes or dysplasia. Radiographic images showed RUS of the bilateral forearms and bilateral bony defect of the intermediate phalanges of the fifth digits [4]. Neutropenia rapidly progressed, and PC transfusion was required twice a week. Hence, the patient immediately underwent allogeneic HSCT to prevent lifethreatening infections at 4 months of age. Clinical course of allogeneic cord blood transplantation is shown in Fig. 3. Since there were no suitable conditioning regimens for CBT, fludarabine (FLU) (0.83 mg/kg for 5 days), melphalan (L-PAM) (2.3 mg/kg for 2 days) and rabbit anti-thymocyte globulin (rATG) (single dose of 1.25 mg/kg) were administered. HLA 1 allele-mismatched (DR) cord blood was selected, which contained enough total nuclear $(21.2 \times 10^7/$ kg) and CD34 + $(3.6 \times 10^5/$ kg) cells for engraftment. Oral cyclosporine A and short-term intravenous MTX were administered as GVHD prophylaxis due to limited blood access. After completing the conditioning regimen, the patient's WBC count decreased to 0/mm³. She then achieved Neut and Plt engraftments on days + 14 and + 22, respectively. She developed grade 2 acute GVHD of the skin. However, the exanthema disappeared after administering 1 mg/ kg of prednisolone. She achieved complete chimera of the donor type with sufficient recovery of megakaryocytes in the bone marrow on day + 35 and became independent from transfusion.

Pt 2 presented with massive systemic petechiae at birth (37 weeks of gestation), and her birth weight was 2058 g. Laboratory data revealed the following: 17,100/mm³ WBC count; 12.9 g/dL Hb level; and 8000/mm³ Plt count. Low Plt levels (< 10,000/mm³) persisted for 5 months and progressed to pancytopenia, requiring repeated RBC and PC transfusions. The patient's radiographic image showed bilateral RUS and bilateral fifth digit clinodactyly, which caused limitations in forearm supination and pronation. Bone marrow examination revealed low cellularity and absence of megakaryocytes. She received allogeneic bone marrow transplantation (BMT) from an HLA full-matched donor with a sufficient total nuclear cells $(6.9 \times 10^8/\text{kg})$ at the age of 18 months. The conditioning regimen comprised FLU $(25 \text{ mg/m}^2 \text{ for 4 days})$, cyclophosphamide (CY) (50 mg/kg for 4 days), rATG (2.5 mg/kg for 4 days), and total lymphoid irradiation (3 Gy). We administered FK506 and short-term MTX for GVHD prophylaxis. Neut and Plt engraftments were achieved on days + 16 and + 27, respectively. The patient presented with grade 2 acute GVHD of the skin, which was successfully treated with prednisolone [4, 8].

| Pt. No (age at transplant) | Donor source | Conditioning regimen | Infused total cell counts | HLA com- patibility | GVHD prophy- laxis | Neutro- phils > 500/mm ³ | Plate- lets $> 50,000/$ | Chimerism | Regimen- related | aGVHD | cGVHD o | Age at last | Body height at last follow up | Body weight at last follow up [kg | Other |
|---|--------------------------|--|-------------------------------------|---|--|---|---|--|---|----------------------------|--------------------------------------|-------------------------------|----------------------------------|--|---------------------------------------|
| | | | (/kg) | | | | mm³ | | toxicity | | | follow up (y) | [cm (SD)] | (SD)] | |
| 1 (4 months) | Unrelated CB | FLU 0.83 mg/kg×5 L-PAM 2.3 mg/kg×2 rATG 1.25 mg/kg×1 | 21.2×10 ⁷ | Allele 7/8 match DR 1 locus mismatch | Oral CyA short- term MTX | Day+14 | Day + 22 | BM complete chimera at day + 35 | Mucositis grade 1 | Skin stage 3 (grade 2) | No | 8.0 | 116.0 (- 1.61) | 17.0 (- 2.85) | |
| 2 (18 months) | Unrelated BM | FLU 25 mg/m2×4 CY 50 mg/kg×4 rATG 2.5 mg/kg×4 TLI 3 Gy | 6.9×10 ⁸ | Allele 8/8 match | FK506 Short-term MTX | Day + 16 | Day + 27 | BM complete chimera at day + 60 | Generalized convul- sion at day 1 MRSA sepsis at day 60 | Skin stage 3 (grade 2) | No | 14.1 | 139.5 (- 3.01) | 32.3 (- 3.15) | |
| 3 (8 months) | Unrelated BM | FLU 0.83 mg/kg×5 CY 50 mg/kg×4 TAI 2 Gy | 2.7×10^{8} | Allele 7/8 match DR 1 locus mismatch | FK506 short-term MTX | Day + 6 | Day + 23 | BM complete chimera at day + 23 | ou | Skin stage 1 (grade 1) | No | 11.7 | 134.8 (~ 1.61) | 28.0 (- 1.85) | GH replace- ment therapy (+) |
| 4 (5 months) | Unrelated CB | FLU 1 mg/kg×5 L-PAM 2.3 mg/kg×2 TBI 3 Gy | 12.7×10^{7} | Allele 7/8 match DR 1 locus mismatch | FK506 short-term MTX | Day + 18 | Day + 35 | BM complete chimera at day + 36 | VOD grade 1 | Skin stage 2 (grade 1) | No | 4.2 | 87.8 (- 3.08) | 11.6 (– 1.82) | |
| 5 (8 months) | Unrelated CB | FLU 0.83 mg/kg×5 L-PAM 2.3 mg/kg×2 rATG 1.25 mg/kg×1 | 17.2×10 ⁷ | Allele 4/8 match DR 2 loci, C 2 loci mismatch | FK506 short-term MTX | Day + 22 | Day + 34 | BM complete chimera at day + 30 | оц | Skin stage 1 (grade 1) | No | 3.7 | 97.9 (+ 0.21) | 13.0 (- 0.26) | |
| 6 (14 months) | Unrelated BM | FLU 25 mg/m2×5 L-PAM 90 mg/m2×2 rATG 1.25 mg/kg×2 | 3.6×10 ⁸ | Allele 7/8 match C 1 locus mismatch | FK506 short-term MTX | Day + 18 | Day + 30 | BM complete chimera at day + 33 | ы | oN | No | 4.2 | 105.0 (+1.14) 17.2(+0.85) | 17.2(+0.85) | |
| <i>RIC</i> reduce GVHD, <i>cG</i> mocytes alo | d-intensity VHD chron | conditioning iic GVHD, 2 total lympho | g, <i>HSCT</i> hen SD standard (| natopoietic deviation, (| RIC reduced-intensity conditioning, HSCT hematopoietic stem cell transplantation, Pt patient, No number, HLA human leukocyte antigen, GVHD graft-versus-host disease, aGVHD acute GVHD, cGVHD chronic GVHD, SD standard deviation, CB cord blood, BM bone marrow, FLU fludarabine, L-PAM 1-phenylalanine mustard, CY cyclophosphamide, rATG rabbit anti-thy- mocytes olohulin, TII total lymphoid irradiation, TAI thoracic-abdominal irradiation, TBI total hody irradiation, CvA evclosnorin A. MTX methorrexate. FK506 factoritims. MRSA methicillin- | splantation, <i>P</i> , <i>BM</i> bone ma | <i>t</i> patient, <i>No</i> urrow, <i>FLU</i> fl <i>RI</i> total body | number, <i>HL</i> udarabine, <i>L</i> - | A human I -PAM 1-pho 2va evelos | leukocyte a enylalanine | Intigen, G mustard, TTX methoo | VHD gra CY cycl trevate | aft-versus-ho ophosphami | RIC reduced-intensity conditioning, HSCT hematopoietic stem cell transplantation, Pt patient, No number, HLA human leukocyte antigen, GVHD graft-versus-host disease, aGVHD acute GVHD, cGVHD chronic GVHD, SD standard deviation, CB cord blood, BM bone marrow, FLU fludarabine, L-PAM 1-phenylalanine mustard, CY cyclophosphamide, rATG rabbit anti-thy- mocytes of bullin, TH total humboid irrediation, TM thoracic abdominal irrediation, TM total body irrediation, CA evoloscovin, A MTY methorescue, EK506 recolutions, MPSA methodilin, | VHL vit au |

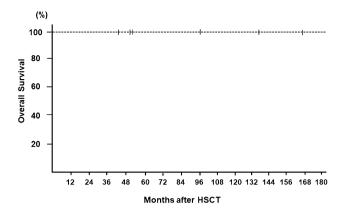


Fig. 2 Overall survival rate after allogeneic hematopoietic stem cell transplantation in six patients with *MECOM*-associated syndrome

Pt 3 was born at 31 weeks of gestation, weighing 2180 g. The patient presented with severe pancytopenia at birth. Laboratory results showed 3220/mm³ WBC count, 48/mm³ Neut count, 2.7 g/dL Hb level, and 89,000/mm³ Plt count. Severe neutropenia (<100/mm³) persisted and the patient's platelet count decreased to < 20,000/mm³, requiring repeated RBC and PC transfusions. Moreover, antibiotics, antifungal agents, and immunoglobulin via intravenous infusion were administered to treat prolonged and repeated infections. The patient's radiographic image showed bilateral RUS and overlapping fingers, which caused limitations in the forearm supination and pronation. Bone marrow examination revealed an absence of megakaryocytes. The patient received allogeneic BMT from an HLA 1 allele-mismatched (DR) donor with sufficient CD34 + cells $(10.4 \times 10^6/\text{kg})$ at the age of 8 months. The conditioning regimen comprised

Fig. 3 Clinical course of alloge-

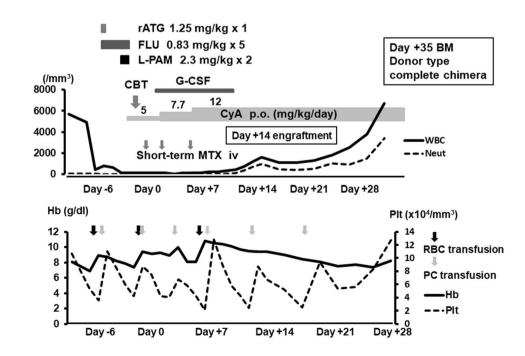
neic cord blood transplantation

in Pt 1

FLU (0.83 mg/kg for 5 days), CY (50 mg/kg for 4 days), and thoracic-abdominal irradiation (2 Gy). FK506 and shortterm MTX were administered for GVHD prophylaxis. Neut and Plt engraftments were achieved on days + 6 and + 23, respectively. The patient only presented with grade 1 acute GVHD of the skin and was treated with a steroid ointment [4, 9].

Pt 4 was born at 40 weeks of gestation and had no remarkable family history of any illness. Her birth weight was 2936 g. She presented with massive pulmonary bleeding and developed respiratory insufficiency, requiring intubation and mechanical ventilation. Laboratory data revealed severe anemia (Hb level 7.2 g/dL) and thrombocytopenia (Plt count 9000/mm³) at birth. The patient presented with severe neutropenia that progressed to pancytopenia at 2 months of age. Bone marrow examination showed hypocellular marrow without megakaryocytes or dysplasia. Bone abnormalities were not observed. At the age of 5 months, we performed allogeneic CBT from HLA 1 allele-mismatched (DR1) cord blood due to recurrent life-threatening bacterial infection and transfusion dependency. The conditioning regimen consisted of FLU (1 mg/kg for 5 days), L-PAM (2.3 mg/kg for 2 days), and total body irradiation (3 Gy). FK506 and shortterm MTX were administered as GVHD prophylaxis. Neut and Plt engraftments were achieved on days + 18 and + 35. respectively [6].

Pt 5 presented with systemic petechiae at birth (37 weeks of gestation), and her birth weight was 2368 g. She had severe bicytopenia (WBC count 10,700/mm³; Hb level 7.3 g/dL; and Plt count 4000/mm³). Hence, the patient required weekly PC transfusion since birth. Her father and uncle had RUS, and her brother, father, and



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uncle presented with clinodactyly of the fingers, but had no hematological abnormalities. Bone marrow examination revealed the absence of megakaryocytes, no excess of blasts and extremely low cellularity (3000/mm³). The patient presented with bilateral clinodactyly of the fifth digit. However, RUS was not observed. Emergent CBT was required due to transfusion dependency at the age of 8 months. The HLA compatibility of CB was DR 2 antigen mismatch (A/B/DR 4/6 match) and 4 allele mismatches (A/B/DR/C 4/8 match). The total and CD34 + cell countsat 17.21×10^7 /kg and 6.01×10^5 /kg, respectively, were sufficient for engraftment. The conditioning regimen in this patient was same as that of Pt 1. FK506 and short-term MTX were administered as GVHD prophylaxis. Neut and Plt engraftments were achieved on days + 22 and + 34, respectively. The patient only presented with grade 1 acute GVHD of the skin [7].

Pt 6 was born at 38 weeks and 6 days of gestation, and her birth weight was 3414 g. She had a maternal history of chronic thrombocytopenia (Plt count 41,000/mm³), transient leukopenia, bilateral RUS, congenital left clubfoot, and bilateral congenital disposition of the hip. The patient developed petechiae at birth, and laboratory data revealed severe anemia (Hb level 6.3 g/dL) and thrombocytopenia (Plt count 7000/mm³). No bone abnormalities were observed in the patient, and RBC and PC transfusions once per week were initially required. The transfusion dependency improved gradually, probably depending on somatic LOH in blood cells. Since Pt 6 remained PC transfusiondependent, she was treated with allogeneic BMT from an HLA 1 allele-mismatched (C) unrelated donor at the age of 14 months. The conditioning regimen consisted of FLU $(25 \text{ mg/m}^2 \text{ for 5 days}), \text{ L-PAM } (90 \text{ mg/m}^2 \text{ for 2 days}),$ and rATG (1.25 mg/kg for 2 days). FK506 and short-term MTX were administered as GVHD prophylaxis. Neut and

Fig. 4 Risk of short stature in patients who received reduced-intensity conditioning regimens with low-dose irradiation. Standard deviations (SDs) of the mean body height before and 3 years after HSCT among patients who received low-dose irradiation (radiation group, n=3) and those who did not (non-radiation group, n=3) compared to age-matched healthy infants

Plt engraftments were achieved on days + 18 and + 30, respectively. The patient did not present with any symptoms of acute GVHD, and all lineages of hematopoietic cells recovered well [7].

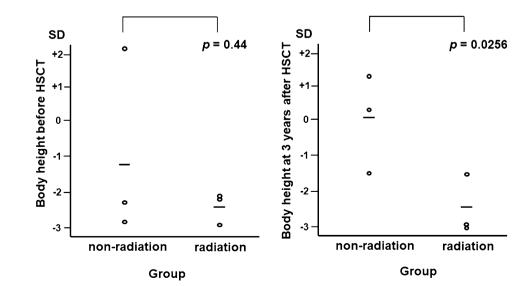
Long-term sequelae after allogeneic HSCT

All patients had good quality of life after allogeneic HSCT. However, there was no improvement in RUS nor in hearing disorders among the affected patients. In terms of long-term sequelae, we evaluated and compared the means and standard deviations (SDs) of body height between three patients who received irradiation (radiation group, n=3) and three patients who did not receive irradiation (non-radiation group, n=3) (Fig. 4). The body height in non-radiation group improved to normal levels of age-matched healthy infants after HSCT. However, the risk of short stature worsened at 3 years after HSCT in the radiation group. Pt 3 in the radiation group received growth hormone replacement therapy after HSCT. Nonetheless, this difference was not statistically significant due to limited number of patients. None of the patients have presented with secondary malignancies 3 years after RIC and allogenic HSCT.

Discussion

RUSAT is a rare disease associated with IBMFS. The EVI1 protein plays an important role in maintaining normal hematopoiesis and hematopoietic stem cell functions. Hence, allogeneic HSCT is considered a reasonable curative treatment for *MECOM*-associated syndrome.

The heterogeneity of *MECOM*-associated syndrome has been a topic of interest because of the recent increase in the number of patients with *MECOM* mutations. In this



case series, we assessed the broad clinical spectrum of *MECOM*-associated syndrome in six patients, including two patients with somatic mosaicism or LOH [6, 7]. In other current study, 6 of 179 children and young patients with undiagnosed IBMFS had *MECOM* mutations. None of the 6 patients had a remarkable family history, and four had no skeletal abnormalities. Moreover, only one had RUS [10].

All patients required allogeneic HSCT to overcome transfusion dependency and to prevent life-threatening infections in early infancy. However, there are two major problems with BMF treatment; the donor source for HSCT and the appropriate conditioning regimen. Regarding the donor source, a related or unrelated bone marrow donor can be chosen if HLA-matched donors are available. Appropriate cord blood is also applicable, as it can be urgently used compared to bone marrow from an unrelated donor in cases of emergent HSCT. Moreover, a sufficient number of infused cells is commonly available for infantile patients. In terms of conditioning regimens, they must be selected based on two conflicting issues, which are as follows: myelosuppressive effects for engraftment as well as lower incidence of regimen-related toxicities and long-term adverse effects, including short stature, endocrinopathy, infertility, and risk of secondary malignancy. Therefore, RIC regimens comprising FLU, alkylating agents, immunosuppressants, such as rATG and campath-1H, and/or low-dose irradiation have been used for nonmalignant diseases. rATG is an extremely strong immunosuppressant that eliminates T lymphocytes. Thus, it is not recommended as a conditioning regimen for CBT due to high mortality caused by delayed immune reconstitution, viral reactivation, and relapse of malignant diseases [11, 12]. The immunosuppressive effects of rATG are believed to be dose-dependent [13, 14], and thus, lowdose rATG was added to prevent long-term adverse effects caused by irradiation in Pts 1, 5, and 6. Three patients who received low-dose rATG for CBT did not show other adverse events such as delayed engraftment and viral reactivation. However, if the patient is at high risk of rejection owing to recipient T cell activation caused by viral infections or hemophagocytic syndrome, use of low-dose TBI and/or urgent second HSCT should be considered.

Long-term sequelae are critical in the management of infants who receive allogeneic HSCT. Irradiation at HSCT was found to be major factor for long-term height loss and relative risk for relevant growth deficiency increased in young patients [15]. Consistent with the previous report, patients treated with low-dose irradiation were at risk of short stature compared with patients without irradiation 3 years after HSCT in this study. However, the statistical significance and the difference among total body, thoracicabdominal or total lymphoid irradiation remained undetermined due to limited number of patients in this case series. Of note, fatal cardiac complications during severe infections were reported in 2 of 6 patients after HSCT, which is a particular concern in patients with MECOM-associated syndrome [10]. The risk of malignancy in MECOM-related disorders has not been evaluated since the responsible gene was only identified in 2015 [4]. Approximately 44% of patients with familial platelet disorders that are predisposed to hematologic malignancies caused by autosomal dominant RUNX1 mutations progressed to acute myeloid leukemia caused by second-hit mutations in CDC25C or other genes [16–18]. Moreover, alterations in EVI1 are involved in dysplastic hematopoiesis and acute leukemia of the megakaryocytic lineage in both humans and mice [19-23]. Therefore, patients with MECOM mutations may be at high risk of developing malignant diseases because of the long-term natural history of the disease or treatment with low-dose irradiation.

In conclusion, RIC regimens were feasible, and all infantile patients had perfect overall survival. In addition, they achieved stable, complete chimera of the donor type. Based on this retrospective study, we propose the RIC regimen comprised FLU, alkylating agents at appropriate doses, and low-dose rATG instead of low-dose irradiation if the patient is not at high risk of rejection to prevent the risks of short stature and secondary malignancy. Nevertheless, further investigations that include a larger number of infantile patients should be conducted to assess the optimal doses of alkylating agents and rATG in the RIC regimen followed by allogeneic HSCT in *MECOM*-associated syndrome.

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Data availability statement All data generated or analysed during this study are available on reasonable request.

Declarations

Conflict of interest The authors have no competing financial interests related to the research or publication of this study.

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