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Recent Decrease in Acute GVHD in Children with Leukemia Receiving Unrelated Donor Bone Marrow Transplants

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Abstract

Unrelated donor (URD) BMT is an effective treatment for leukemia in children, but success is limited by graft versus host disease (GVHD) and relapse. In this study we describe the incidence and risk factors for GVHD over time in children receiving URD BMT. We analyzed outcomes of 638 myeloablative URD BMT performed between 1990 and 2003 for treatment of AML, ALL, CML or MDS, using the Center for International Blood & Marrow Transplant Research (CIBMTR) database. Recipients were <18 years of age, and had available high resolution HLA typing for HLA A, B, C and DRB1. Overall, 27% of recipients developed aGVHD grades III-IV and risk was significantly higher in recipients of T-replete compared with T-depleted grafts (OR 3.12, 95%CI 2.02-4.83; p< 0.0001). Acute GVHD significantly reduced risk of relapse in children with ALL (OR 0.34 95% CI 0.13-0.86, p=0.0052) but not AML (OR 0.58, 95% CI 0.22-2.98; p=0.26). Risk of aGVHD was higher in children transplanted in earlier years (1990-1998, n=365) compared to 1999-2003 (OR 1.93 95%CI 1.27-2.91; p=0.002). We conclude that outcomes have changed significantly over time, with reduced risk of aGVHD in more recent transplants.

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Introduction

Bone marrow transplantation (BMT) has been used for over 30 years as a curative therapy to treat malignant and non-malignant diseases in children (1-3). Early success of the procedure in pediatric populations allowed extension of the treatment to adult populations and now the majority of transplants are performed in adults. The distributions of diagnoses, co-morbid conditions and the impact of prior therapies differ significantly between pediatric and adult populations. The majority of studies show superior outcomes in younger patient populations (4-8). However patient numbers are usually too small to analyze in detail the impact of younger age on specific outcomes. The occurrence of graft versus host disease (GVHD) significantly limits the success of unrelated donor (URD) BMT and is a frequent cause of morbidity and mortality. In this study we have used a large CIBMTR database of myeloablative URD BMT to analyze in detail risk factors and clinical impact of acute and chronic GVHD in a pediatric population. The data indicate some significant temporal changes in outcomes of URD BMT for children with leukemia. In particular, risk of aGVHD was higher in children transplanted in earlier years (1990-1998) compared to 1999-2003; in contrast, risk of relapse was lower in children transplanted 1990-1998 compared with 1999-2003.

Patients and Methods

The CIBMTR

The CIBMTR is a research affiliation of the International Bone Marrow Transplant Registry (IBMTR), Autologous Blood and Marrow Transplant Registry (ABMTR) and the National Marrow Donor Program (NMDP) that comprises a voluntary working group of more than 450 transplantation centers worldwide that contribute detailed data on consecutive allogeneic and autologous hematopoietic stem cell transplants to a Statistical Center at the Health Policy Institute of the Medical College of Wisconsin in Milwaukee or the NMDP Coordinating Center in Minneapolis. Participating centers are required to report all transplants consecutively; compliance is monitored by on-site audits. Patients are followed longitudinally, with yearly follow-up. Computerized checks for errors, physicians' review of submitted data and on-site audits of participating centers ensure data quality. The CIBMTR collects data at two levels: Registration and Research. Registration data include disease type, age, sex, pretransplant disease stage and chemotherapy-responsiveness, date of diagnosis, graft type (bone marrow- and/or blood-derived stem cells), high-dose conditioning regimen, post-transplant disease progression and survival, development of a new malignancy and cause of death. Requests for data on progression or death for registered patients are at sixmonth intervals. All CIBMTR teams contribute Registration data. Research data are collected on subset of registered patients selected using a weighted randomization scheme and include detailed disease, and pre- and post-transplant clinical information.

Patients and Transplant Procedure

The analysis was restricted to recipients less than 18 years of age who had available high resolution typing reported by the transplant center or performed retrospectively by the National Marrow Donor Program (NMDP) (9,10). All patients received unrelated donor bone marrow as the primary stem cell source. Patients receiving peripheral blood stem cell or cord blood grafts were excluded as the natural history of GVHD may differ in these cases.

Study Population

The NMDP retrospectively obtained consent for data submission and study participation from surviving patients or their parent/legal guardian for transplants it facilitated in the U.S. prior to 2002. Thereafter, informed consent was obtained prospectively. The Institutional

Review Board of the NMDP waived consent for patients who had died prior to soliciting consent (transplantations facilitated prior to 2002) and non-consenting surviving patients were excluded. To overcome the bias associated by the inclusion of a proportion of surviving patients (those consenting) but all deceased recipients, and hence their over-representation, a sample of deceased patients was selected using a weighted randomized scheme that adjusts for over-representation of deceased patients in the consented cohort (11). This weighted randomized scheme was developed based on all survivors in the NMDP database. A logistic regression model was fit to identify the factors that predicted whether a patient had consented or not consented to use of data collected by the NMDP. This analysis found the following factors were associated with the likelihood of a patient consenting: age, disease type, race, sex, cytomegalovirus serostatus and country of transplantation (US vs. non US). Using estimated consenting probabilities from this model based on the

non US). Using estimated consenting probabilities from this model based on the characteristics of dead patients, the biased coin method of randomization was performed to determine which of the dead patients likely would have consented to participate had they been alive. Thus this procedure includes the dead patients at the same probability as surviving patients who consented to participate.

The NMDP facilitated 2013 pediatric transplants in patients with acute myelogenous leukemia (AML), acute lymphoblastic leukemia (ALL), chronic myelogenous leukemia (CML) or myelodysplasia disorders (MDS), as their first transplant between 1990 and 2003 using bone marrow as the stem cell source. Of these, 1043 had high resolution HLA typing available. Approximately 12% of dead patients (n=123) were deleted by the weighted randomized method. Patients with more than one antigen mismatch (n=208) or non-myeloablative preparative regimens (n=28) were excluded. An additional 46 patients were subsequently excluded because of presence of > 1 allele mismatch (n= 30) or missing acute GVHD or chronic GVHD information (n=16). The final study population consisted of 638 children.

Patient characteristics are summarized in Table 1. The most frequent diagnosis was acute lymphoblastic leukemia (ALL) and median patient age was 10 years (range <1-17.9 years). Median donor age was 36 years (range 19-60 years).

Patients received a variety of preparative regimens determined by the individual transplant centers. The majority of recipients (90%) received total body irradiation (550cGy-1500cGy); 10% received chemotherapy alone. Chemotherapy drugs administered as part of the preparative regimen included cyclophosphamide, busulfan, cytosine arabinoside, etoposide and thiotepa. Methods for GVHD prophylaxis varied according to the transplant center. The graft was depleted of T-cells ex vivo (TCD) in 37% of cases, who may or may not have received treatment with a calcineurin inhibitor, based on the degree of T-cell depletion; the majority of children (67%) received a non-T depleted graft together with GVHD prophylaxis using the calcineurin inhibitors cyclosporine A or tacrolimus. Risk groups were defined as: Early - includes AML or ALL in CR1; CML in CP1; or MDS with RA or RARS; Intermediate - includes AML or ALL in CR2; CML accelerated phase or CP2 and advanced - includes AML or ALL in relapse or PIF; CML in blast phase or MDS with RAEB-T or IPSS-4.

Donor Selection and Processing of Bone Marrow

The characteristics of the donors are shown in Table 1. High resolution typing at HLA-A, B, C and DRB1 was performed for all donor recipient pairs (12). Forty six percent of recipients were matched with their recipient at all 8 loci; 13% had a single allele mismatch and forty percent had a single antigen mismatch.

Statistical Analysis

Univariate analysis of survival was calculated using the method of Kaplan and Meier and was compared using the log rank statistic (13,14). Rates of acute and chronic GVHD, and relapse were calculated using cumulative incidence, with death acting as a competing risk (15). Cumulative incidences were compared for acute and chronic GVHD and relapse by using a Taylor series linear approximation to estimate the variance (16).

The proportional hazard model was used for the multivariate analysis of survival. (17). Cumulative incidence estimates of acute and chronic GVHD, relapse and treatment-related mortality (TRM) were estimated using the Pseudo-value approach of Klein (18). Additional models were constructed for relapse and TRM treating acute GVHD as a time dependent variable. Variables considered in the analysis included recipient age, gender, performance score, diagnosis and stage of disease at transplantation. Transplant-related variables included donor age, donor-recipient gender match, donor-recipient CMV serology, donor parity and radiation vs non-radiation preparative therapy, GVHD prophylaxis (T-cell depletion vs no T-cell depletion), HLA matching (8 allele match vs 7/8 match, which includes single allele mismatch but antigen match, and single antigen mismatch) and year of transplant (1990-1998 vs 1999-2003).

Results

Acute GVHD

The cumulative incidence (CI) of grades II-IV GVHD was 45% (95% CI 42-49%) and of grades III-IV GVHD was 27%, indicating that most cases of acute GVHD were severe. Multivariate analysis showed similar risk factors for grades II-IV and III-IV acute GVHD . As expected, T-replete grafts were associated with a significantly increased risk of acute GVHD compared with T-depleted grafts (OR 3.12; 95% CI 2.02-4.83; p< 0.0001). In addition, transplant in the earlier era (1990-1998) was associated with significantly increased risk of acute GVHD compared to transplant in recent years (1999-2003) (OR 1.93; 95% CI 1.27-2.91; p=0.002). Notably, HLA mismatch at a single locus did not increase risk of GVHD (OR 1.00; 95% CI 0.68-1.46; p=0.98), and age and diagnosis did not significantly modify risk.

Chronic GVHD

The cumulative incidence of chronic GVHD was 35% (95% CI 31-38%) 3 years after BMT. Multivariate analysis of risk factors for chronic GVHD revealed only a modest non-significant increase in risk with parous female donors (OR 1.45; 95% CI 0.92-2.29; p=0.11), and no effect of HLA-mismatch (OR 1.19; 95% CI 0.78-1.79; p=0.41) or T-cell depletion on risk of chronic GVHD.

Relapse

The cumulative incidence of relapse was 26% (95% CI 22-29%) at 3 years. Multivariate analysis of risk factors for relapse was conducted for the entire group and separately for children with AML and with ALL, as we hypothesized that risk factors might vary in the two categories. Risk of relapse was significantly lower in children with MDS compared with those with acute leukemia, and risk was very markedly increased in children with the most advanced disease (OR 15.37, 95% CI 6.1-38.7; p <0.0001). Risk of relapse declined markedly with increasing recipient age (OR 0.23 for recipients 16-18 years compared with reference group of children less than 5 years old (95% CI 0.10-0.54, p=0.0004)). Risk of relapse was reduced in recipients of marrow from parous female donors (OR 0.43; 95% CI 0.23-0.83; p=0.01), perhaps reflecting the modest increase in chronic GVHD in these children, and was not increased in recipients of T-cell depleted marrow. A perhaps

surprising observation was a decreased risk of relapse in children transplanted in the earlier era compared with more recent transplants (OR 0.47, 95% CI 0.28-0.78; p=0.004).

A multivariate analysis of risk factors for relapse restricted to children with AML (n=156) showed a non significant reduction in incidence of relapse (OR 0.58 comparing relapse incidence in children with GVHD grades III-IV with those without, 95% CI 0.22-1.58; p=0.26) (Table 2). However, relapse risk was significantly higher in the recent era (1999-2003), in children younger than 5 years and in those with advanced disease, as seen in the group overall. In contrast, the occurrence of acute GVHD significantly reduced risk of relapse in children with ALL (OR 0.34 comparing relapse incidence in children with GVHD grades II-IV with those without, 95% CI 0.13-0.86; p=0.0052) (Table 3). Similar to AML, children with ALL older than 5 years and with less advanced disease had a reduced risk of relapse. While there was an increased risk of relapse in the recent era, this was of marginal significance in the children with ALL.

Survival and Treatment-Related Mortality

Fifty six percent (95% CI 52-59%) of children were surviving at 1 year and 43% (95% CI 39-47%) at 5 years. The occurrence of acute GVHD had an important effect on survival (Figure 1). Survival was notably reduced in children with grades III-IV GVHD, but not in children with grade II GVHD. In a multivariate model of risk factors for mortality, the occurrence of acute GVHD was associated with an almost doubled risk of death (RR 1.96, 95% CI 1.46-2.29, p <0.0001). However, survival was not improved by T-cell depletion, despite significant reduction in GVHD. Mortality was increased in older children, as a consequence of increased treatment related mortality (OR 1.72 in children 16-18 years compared with < 5 years old, 95% CI 1.11-2.67; p = 0.016). Children receiving marrow mismatched at a single locus had significantly reduced survival, despite the absence of a significant effect on GVHD (OR 1.42, 95% CI 1.05-1.92; p=0.02). Treatment related mortality was significantly increased in children with grades III-IV acute GVHD, but not in children with grade II GVHD (Figure 2). Treatment related mortality was higher in the earlier era (1990-1998), OR 1.61 (95% CI 1.03-2.51; p=0.027) compared with more recent transplants (1999-2003). The overall effect of era of transplant and GVHD is shown in Figure 3. Survival is reduced in children with GVHD transplanted in early or later years. The increased risk of relapse in children transplanted in recent years leads to slightly lower DFS in those children, whether or not they have GVHD but is not statistically significantly different (p=0.81).

Discussion

Children constitute an important subset of patients undergoing BMT. Despite differences in diagnoses, organ function and immune reconstitution capability, children are often considered together with adults in studies of GVHD, and reports of the effect of age on incidence of GVHD have been inconsistent (4,5,7,8,19-25). Some reports have described a reduced risk of GVHD in younger patients, while in other reports this has not been seen. The small number of children receiving BMT relative to adults has limited studies focused specifically on children. We examined a large database to examine the incidence, risk factors and outcome of GVHD in children to determine whether children with GVHD have similar risk factors and outcomes as adults, and to examine temporal trends in outcomes of unrelated donor transplants in children.

Our analysis showed that T-cell depletion for GVHD prophylaxis significantly reduced risk of GVHD, as might be expected. In addition, year of transplant was a significant risk factor, with lower risk of GVHD in more recent transplants. Perhaps unexpectedly, risk of GVHD was not increased by an allele or antigen mismatch at a single HLA locus, in contrast to

findings in adult studies (20,21,23,24). This may indicate greater tolerance of modest degrees of HLA disparity in pediatric transplant recipients compared with adults, or greater success of GVHD prophylaxis in controlling alloreactivity. It should be noted, however, that survival was reduced in recipients of marrow with an allele or antigen mismatched bone marrow, perhaps indicating inferior immune reconstitution in this circumstance, despite lack of a measurable excess of GVHD.

Our analysis of risk factors for chronic GVHD failed to identify any important risk factors, aside form a marginal increase in risk with multiparous female donors, as described by others (26-33). Of note, HLA mismatch did not increase risk of chronic GVHD in agreement with our findings in acute GVHD, and risk was not decreased by T-cell depletion of the graft. It should be noted that the overall incidence of chronic GVHD was significant, with approximately one third of children affected, including even the youngest children, and efforts should be made to provide effective treatments for all chronic GVHD for all ages.

Our study showed a marked reduction in relapse in children with ALL who developed grades III/IV acute GVHD. These data are perhaps surprising as clinical experience of quite limited responses to donor lymphocyte infusions in ALL suggest that any graft-versusleukemia effect in ALL might be modest (reviewed in 34). However, data from previous non-pediatric registry studies of sibling and URD BMT support the observation of a reduced relapse rate in cases with ALL who develop GVHD (35,36). Horowitz et al analyzed 2,254 sibling donor transplants performed in children and adults with early stage leukemia (ALL or AML in CR1 and CML in first chronic phase) reported to the International Bone Marrow Transplant Registry (IBMTR) (35). This study included 439 ALL cases and showed an almost 3-fold reduction in risk of relapse associated with acute GVHD, but no effect of chronic GVHD. Similarly, a low relapse rate, attributed to graft-versus-leukemia, has been described in a report of 127 adults receiving URD BMT for high risk ALL, facilitated by the NMDP (36). It is unclear, and perhaps equally surprising that a reduction of relapse was not seen in children with AML. It is possible that competing mortality from severe GVHD has obscured any benefit from GVHD, or that small sample size has rendered a biologically meaningful effect statistically insignificant. Future studies will address this question again as additional transplant data are accrued.

The data in our study show a significant effect of grades III/IV acute GVHD on mortality, with an almost doubled risk of death in children with acute GVHD, demonstrating that current treatments for acute GVHD are inadequate. In contrast, grade II GVHD had only a minor effect on survival. Performance of randomized controlled trials to optimize treatment of acute GVHD is an essential step in improving survival for children receiving URD BMT.

Our analysis has identified some striking temporal changes in outcome of URD BMT for children with leukemia. The risk of acute GVHD is reduced in more recent transplants (1999-2003) compared with those performed prior to 1999, even when T-cell depletion and HLA-match are included in the multivariate model. These data may indicate greater success in optimizing use of calcineurin inhibitors, and or methotrexate as GVHD prophylaxis in recent years. In contrast to this improvement in outcome, risk of relapse is increased in children transplanted in the more recent era and the increased risk is most notable in children with AML. This may reflect the use of increasingly intense chemotherapy in the primary treatment of children with AML. Of note, 35% of children with AML were transplanted with advanced disease compared with 11% of children with ALL. In addition, criteria for selection of children with leukemia for transplantation change frequently, based on improvements in identification of genetic characteristics that alter risk of relapse, for example and improvements in the effectiveness of chemotherapy. Notably, the era of transplant was not a significant risk factor for overall survival, perhaps indicating that recent

reductions in GVHD are counterbalanced by the increase in relapse, illustrating the dynamic nature of leukemia patient populations as chemotherapy outcomes continue to evolve.

This study is a registry analysis and so is limited by the heterogeneity of treatment regimes and clinical practices at different institutions. The strength of the study is the size of the population, allowing analyses of subsets that could not be examined in single institution studies.

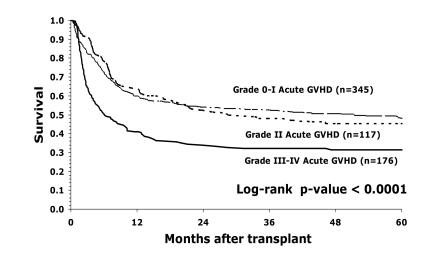
In summary, this study is the largest single analysis of GVHD in URD BMT in children to date. We have identified important temporal changes in clinical outcomes that will merit continued observation as chemotherapy and transplant strategies both continue to improve. In addition we have shown a significant effect of GVHD in reducing relapse in children with ALL, but not AML. However, overall, acute GVHD has a significant negative effect on survival, demonstrating the need for further improvements in both GVHD prophylaxis and the treatment of acute GVHD.

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Probability of overall survival by acute GVHD grade in children with leukemia receiving unrelated donor bone marrow transplants

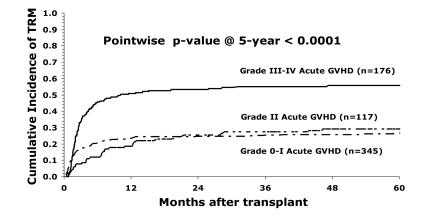
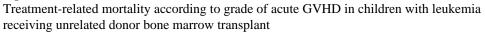
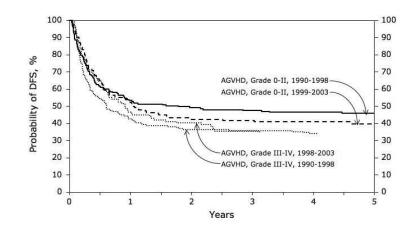


Figure 2.





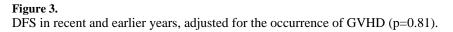


Table 1

Characteristics of pediatric patients (age < 18 years) who underwent an unrelated bone marrow donor transplant receiving a myeloblative conditioning regimen, reported to CIBMTR between 1990 and 2003.

Number of patients 638 Age at transplant, median (range), years $10 (<1-17.9)$ Age at transplant, years <5 <5 $151 (24)$ $6-10$ $183 (29)$ $11-15$ $167 (26)$ > 15 $137 (21)$ Male sex $365 (57)$ Performance score <90 <90 $111 (17)$ $90-100$ $513 (80)$ Unknown $14 (2)$ Disease AML ALL $329 (52)$ CML $75 (12)$ MDS $66 (10)$ Disease status at transplant ^d Early Early $175 (27)$ Intermediate $310 (49)$ Advanced $128 (20)$ Other b $25 (4)$ Donor age, median (range), years $36 (19-60)$ Donor age, years $18-30$ $169 (26)$ $31-40$ $240 (38)$ $41-60$ $229 (36)$ Donor-recipient sex match Male -> Male $132 (21)$ Female -> F	Characteristics	N (%)
Age at transplant, years <5 151 (24) $6-10$ 183 (29) $11-15$ 167 (26) >15 137 (21)Male sex365 (57)Performance score <90 <90 111 (17) $90-100$ 513 (80)Unknown14 (2)DiseaseAMLAML168 (26)ALL329 (52)CML75 (12)MDS66 (10)Disease status at transplant [#] Early175 (27)Intermediate310 (49)Advanced128 (20)Other b25 (4)Donor age, median (range), years36 (19-60)Donor age, years169 (26) $31-40$ 240 (38) $41-60$ 229 (36)Donor-recipient sex matchMale -> MaleMale -> Male135 (21)Female -> Male135 (21)Female -> Female141 (22)Donor-recipient CMV match125 (20)D(-)/R(+)125 (20)D(+)/R(-)125 (20)D(+)/R(+)88 (14)	Number of patients	638
<5	Age at transplant, median (range), years	10 (<1-17.9)
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>15 137 (21) Male sex 365 (57) Performance score <90	6-10	183 (29)
Male sex $365 (57)$ Performance score <90	11-15	167 (26)
Performance score <90 111 (17) 90-100 513 (80) Unknown 14 (2) Disease AML 168 (26) ALL 329 (52) CML 75 (12) MDS 66 (10) Disease status at transplant ^{<i>A</i>} Early 175 (27) Intermediate 310 (49) Advanced 128 (20) Other ^{<i>b</i>} 25 (4) Donor age, median (range), years 36 (19-60) Donor age, years 18-30 169 (26) 31-40 240 (38) 41-60 229 (36) Donor-recipient sex match Male -> Male 135 (21) Female -> Female 132 (21) Female -> Female 141 (22) Donor-recipient CMV match D(-)/R(-) 277 (43) D(-)/R(+) 136 (21) D(+)/R(-) 125 (20) D(+)/R(+) 88 (14)	>15	137 (21)
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D(-)/R(+) 136 (21) D(+)/R(-) 125 (20) D(+)/R(+) 88 (14)	Donor-recipient CMV match	
D(+)/R(-) 125 (20) D(+)/R(+) 88 (14)	D(-)/R(-)	277 (43)
D(+)/R(+) 88 (14)	D(-)/R(+)	136 (21)
	D(+)/R(-)	125 (20)
Unknown 12 (2)	D(+)/R(+)	88 (14)
	Unknown	12 (2)

Characteristics	N (%)
Donor pregnancy	
Male donor	362 (57)
Female, no pregnancy	117 (18)
1 or more pregnancies	153 (24)
Unknown	6(1)
Conditioning regimen	
Cy+TBI+/-other	543 (85)
TBI+other	29 (5)
Bu+Cy+/-other	57 (9)
Bu+/other	9(1)
GVHD prophylaxis	
T-cell depletion	234 (37)
CSA+MTX+/-other	338 (53)
CSA+/-other	18(3)
FK506+/-other	46 (7)
MTX+/-other	1 (<1)
Missing	1 (<1)
HLA match	
Matched	296 (46)
Single allele mismatch	84 (13)
Single antigen mismatch	258 (40)
Median follow-up of survivors, months	87 (3-195)

^aRisk groups are defined as: Early - includes AML or ALL in CR1; CML in CP1; or MDS with RA or RARS; Intermediate - includes AML or ALL in CR2; CML accelerated phase or CP2 and advanced - includes AML or ALL in relapse or PIF; CML in blast phase or MDS with RAEB-T or IPSS-4.

^bOther includes (N=25): 1) ALL, disease status prior to transplant is missing (n=1) 2) MDS, NOS (n=3) 3) CMML Chronic myelomonocytic leukemia (n=3) 4) PNH Paroxysmal nocturnal hemoglobinuria (n=1)

Table 2

Multivariate analysis for relapse in children who were diagnosed with AML (N=156) and received unrelated donor BMT.

Variables	N	Odds Ratio (95% Confidence Interval)	P-value
HLA matching			
(1) Matched	69	1.00 ^a	
(2) 7/8 matched	87	0.63 (0.26, 1.55)	0.36
Other significant covariates:			
Grade III-IV aGVHD			
(1) No	89	1.00 ^a	
(2) Yes	67	0.58 (0.22, 1.58)	0.26
Disease stage			< 0.0001
(1) Early	33	1.00 ^a	
(2) Intermediate	69	0.90 (0.23, 3.54)	0.88
(3) Advanced	54	12.59 (3.09, 51.32)	0.0004
Recipient age categories			0.0059
(1) < 5 years	46	1.00 ^a	
(2) 6-10 years	32	0.40 (0.13, 1.28)	0.12
(3) 11-15 years	47	0.12 (0.04, 0.39)	0.0005
(4) 16-17.9 years	31	0.12 (0.02, 0.65)	0.014
Transplant year group			
1999 - 2003	64	1.00 ^a	
1990 - 1998	92	0.14 (0.13, 0.73)	0.0075

^aReference group.

Acute GVHD III-IV is treated as a time-dependent variable.

Table 3

Multivariate analysis for relapse in children who were diagnosed with ALL (N=311) and received unrelated donor BMT.

Variables	N	Odds Ratio (95% Confidence Interval)	P-value
HLA matching			
Matched	147	1.00	
7/8 matched	164	0.72 (0.37, 1.43)	0.45
Grade III-IV aGVHD			
No	168	1.00	
Yes	143	0.34 (0.13, 0.86)	0.0052
Disease stage			0.0012
Early	62	1.00	
Intermediate	214	1.96 (0.74, 5.17)	0.17
Advanced	35	12.77 (3.83, 42.55)	< 0.0001
Recipient age categories			0.05
< 5 years	67	1.00	
6-10 years	114	0.38 (0.17, 0.89)	0.025
11-15 years	75	0.22 (0.07, 0.66)	0.0069
16-17.9 years	55	0.31 (0.11, 0.87)	0.025
Transplant year group			
1999 - 2003	113	1.00 ^a	
1990 - 1998	198	0.51 (0.26, 1.01)	0.054