Chronic graft-versus-host disease in children: incidence, risk factors, and impact on outcome

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Chronic graft-versus-host disease (cGVHD) remains the major cause of late morbidity and mortality after allogeneic hematopoietic stem cell transplantation (HSCT). However, only a few studies specifically focused on children, and little information is available on the antileukemic effect of cGVHD and its impact on diseasefree survival (DFS) in children. We retrospectively analyzed 696 children given allogeneic HSCT for malignant (n = 450) or nonmalignant (n = 246) diseases. The donor was an HLA-identical sibling in 461 cases and an alternative donor in 235. Bone marrow was the stem cell source in 647 cases, peripheral blood in 17, and cord blood (CB) in 32. cGVHD developed in 173 children (25%) at a median of 116 days after HSCT. Three-year cGVHD probability was 27%. In multivariate analysis, variables predicting cGVHD were donor and recipient age, grade II to IV acute GVHD, female donor for male recipient, diagnosis of malignancy, and use of total body irradiation; CB transplants had a very low risk of cGVHD (RR = 0.07, P = .0001). cGVHD occurrence increased transplant-related mortality (P < .05). Nevertheless, in hematologic malignancies, patients with cGVHD had a reduced relapse probability compared with children without cGVHD (16% \pm 3% versus 39% \pm 3%, *P* = .0001) and a better DFS (68% \pm 4% versus 54% \pm 3%, *P* = .01). The antileukemic effect of cGVHD was observed mainly in patients with acute lymphoblastic leukemia (ALL). This study provides novel data on cGVHD in childhood. Use of CB stem cells and preparative regimens without radiotherapy may prevent its development. In patients affected by ALL, cGVHD was associated with a strong graft-versusleukemia effect, improving DFS. (Blood. 2002;100:1192-1200)

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Introduction

Chronic graft-versus-host disease (cGVHD) remains the most frequent late complication after allogeneic hematopoietic stem cell transplantation (HSCT) and it represents the major cause of nonrelapse mortality and morbidity in long-term survivors.¹⁻³ In adults, the reported incidence of cGVHD ranges between 30% and 50% of HLA-identical sibling transplant recipients,¹ with acute GVHD being recognized as the most important factor predicting the development of the chronic form of the disease.⁴⁻⁶ The increased use of matched unrelated volunteers as donors⁷ and of peripheral blood as a stem cell source^{8,9} has led to a further increase of the incidence and severity of this complication.

Chronic GVHD is associated with a graft-versus-leukemia effect (GVL), thus resulting, in the context of transplantation for the treatment of hematologic malignancies, in a decreased incidence of relapse.¹⁰⁻¹² Nevertheless, the reduced relapse rate could be offset by an increase in transplant-related mortality (TRM) and counterbalanced by a severely impaired quality of life for patients experiencing the extensive form of the disease.^{2,3}

Previously published studies documented a lower risk of cGVHD, as well as of acute GVHD, in younger patients.⁴⁻⁶

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However, only a few studies specifically focused on children.¹³⁻¹⁵ For this reason, risk factors for cGVHD in childhood are still poorly defined, and little information is available on the GVL effect of this complication, as well as on its impact on disease-free survival (DFS) and TRM of young patients.

We carried out a retrospective analysis on a large number of children receiving allogeneic HSCT from either a matched or mismatched family donor or an unrelated volunteer, to identify risk factors predicting the development of cGVHD and to assess its impact on clinical outcome.

Patients and methods

Patient characteristics

A total of 696 consecutive patients, undergoing allogeneic HSCT between January 1, 1991, and June 30, 1999, in one of the transplantation centers of the Italian Association for Pediatric Hematology and Oncology (AIEOP) and reported to the AIEOP-BMT Registry were included in this analysis. All new transplantations performed were registered yearly by each center,

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together with the updated follow-up of previously reported transplantations, through disease- and transplant-specific data collection forms.

For the purpose of this study, only patients younger than 18 years of age at the time of HSCT and surviving more than 90 days after the graft were considered for data analysis. Exclusion criteria for the definition of the study population comprised second or subsequent transplantations, the use of a genotypically identical twin as donor, and a diagnosis of solid tumor.

In all donor-recipient pairs, histocompatibility was determined by serology for HLA-A, B, and DR antigens, whereas high-resolution molecular typing of HLA-class II DRB1 was available for all children undergoing HSCT from an unrelated donor. A total of 450 children (65%) had hematologic malignancies; the other 246 children (35%) had a nonmalignant disorder. Children with acute leukemia receiving transplants during first or second complete remission, those with myelodysplastic syndrome (MDS) characterized by a low blast count (ie, refractory anemia and refractory anemia with ringed sideroblasts), as well as patients with chronic myeloid leukemia (CML) in first chronic phase were considered in early phase, whereas all other patients with leukemia or MDS were considered in advanced phase. Of the 696 patients analyzed, 418 were boys and 278 girls. Median age at transplantation was 7 years (range, 0.3-17 years), whereas median donor age was 12 years (range, 0-55 years). The donor was an HLA-identical sibling in 461 cases (66%), a partially matched family donor in 68 cases (10%), and an unrelated volunteer in the remaining 167 patients (24%). The source of stem cells was bone marrow in 647 cases (93%), peripheral blood in 17 (2%), and cord blood (CB) in 32 (5%). GVHD prophylaxis consisted of cyclosporin A (Cs-A) alone in 343 cases (49%) and Cs-A in combination either with methotrexate (MTX) in 127 cases (18%) or with steroids (prednisone [PDN]) in 62 cases (9%). The combination of Cs-A + MTX + in vivo serotherapy (antilymphocyte globulin [ALG] or the monoclonal antibody [mAb] Campath-1G) was adopted for 102 transplants (15%). Nineteen patients (3%) received a T-cell-depleted HSCT.

Details on the 696 patients evaluated, as well as on the transplantation procedure, are reported in Table 1.

Diagnosis of acute and chronic GVHD and of chronic GVHD resolution

The diagnosis of acute and cGVHD was made by the site investigators at each institution on the basis of clinical symptoms, laboratory tests, and, whenever possible, on histopathologic findings of skin, oral mucosa, and gastrointestinal tract.¹⁶⁻¹⁸ Acute and cGVHD were classified according to previously described criteria.^{19,20} Limited cGVHD was defined as either or both localized skin involvement and hepatic dysfunction due to chronic GVHD; extensive cGVHD was defined as generalized skin involvement, or localized skin involvement or hepatic dysfunction due to cGVHD plus liver histology showing chronic aggressive hepatitis, bridging necrosis or cirrhosis, or involvement of the eye or minor salivary glands or oral mucosa, or involvement or any other target organ.^{20,21}

Resolution of cGVHD was defined as the disappearance of all active lesions attributable to cGVHD (whenever possible confirmed by histopathologic findings) or normalization of liver function tests, associated with the discontinuation of any form of immunosuppressive treatment.

Treatment of chronic GVHD

The choice among the different options for treatment of cGVHD was based on the policy in use in each single transplantation center and varied over time. Thirty-three children received a single drug treatment (Cs-A alone in 20 cases, PDN alone in 11, ursodeoxycholic acid in 1, and mAbs in 1), whereas the standard combination of Cs-A and PDN was used in 62 children, with the addition of azathioprine in 14. Thalidomide was used, always in combination with Cs-A or steroids, in 14 patients. Fifteen children received 3 or more immunosuppressive drugs. Extracorporeal photochemotherapy was used, as single treatment modality or in combination with other systemic immunosuppressive drugs, in 26 children.^{22,23}

Table 1. Patient characteristics

Characteristics and treatments	Number	Percentages
Number of patients	696	
Sex, M/F	418/278	60/40
Median age at transplantation, y (range)	7 (0.3-17)	
Diagnosis		
Malignant disease	450	65
ALL	268	38
AML	99	14
CML	28	5
MDS	36	5
Lymphoma	19	3
Nonmalignant disease	246	35
Thalassemia	88	12
Severe aplastic anemia	35	5
Fanconi anemia	23	3
Immunodeficiencies	48	7
Inborn errors	52	8
Donor		
HLA-identical family donor	461	66
Partially matched family donor	68	10
HLA-identical unrelated donor	141	20
Partially matched unrelated donor	26	4
Female donor/male recipient pairs	196	28
Median donor age, y (range)	12 (0-55)	
Stem cell source		
Bone marrow	647	93
Peripheral blood	17	2
Cord blood	32	5
Conditioning regimen		
Chemotherapy based	369	53
TBI based	327	47
GVHD prophylaxis		
Cs-A	348	50
Cs-A + MTX	129	19
Cs-A + PDN	63	9
Cs-A + MTX + ALG or mAb	104	15
Cs-A $+$ ALG or mAb \pm PDN	33	5
T-cell depletion	19	2
Acute GVHD		
Grade 0-I	406	58
Grade II-IV	290	42

Finally, 9 children, all with limited cGVHD, did not need any specific immunosuppressive therapy.

Statistical analysis

The occurrence of cGVHD, as well as DFS, TRM, and relapse probability according to cGVHD occurrence and extension, were calculated by the Kaplan-Meier method; comparisons between probabilities in different patient groups were performed using the log-rank test.²⁴ As suggested by Pepe et al,²⁵ cGVHD was presented, besides the Kaplan-Meier curve, also as the prevalence of cGVHD over time after transplantation (Figure 1B). The prevalence function at time *t* was calculated using the following function: (number of patients with cGVHD at time *t*)/(number of patients alive and relapse-free at time *t*).^{25,26}

In DFS analysis, both relapse and death in remission due to any cause were considered events, whereas in relapse probability analysis, only disease relapse was considered as failure. In TRM analysis, all deaths not due to disease recurrence were considered events. Results were expressed as probability (%) and 95% CI.





Figure 1. Cumulative probability of developing cGVHD and its prevalence. (A) Overall cumulative probability to develop cGVHD after HSCT in our study population and probability to develop the limited or the extensive form of the disease. (B) Overall prevalence of cGVHD and prevalence of the limited and extensive form of the disease. The prevalence function at time t was calculated using the following function: (number of patients with cGVHD at time t)/(number of patients alive and relapse-free at time t).

A univariate analysis of DFS, relapse probability, and TRM according to cGVHD occurrence and severity was performed for the whole study population and, subsequently, stratified for malignant and nonmalignant disorders. Furthermore, a separate analysis on the impact of cGVHD on relapse probability and DFS was performed also for the 4 types of hematologic malignancies including the larger number of patients: acute lymphoblastic leukemia (ALL; 268 patients), acute myelogenous leukemia (AML; 99 patients), CML (28 patients), and MDS (36 patients).

In the univariate analysis of risk factors for cGVHD development, the following variables were evaluated: patient and donor age, sex mismatch, parity of female donors, diagnosis, type of donor, stem cell source, use of total body irradiation (TBI) as part of the conditioning regimen, GVHD prophylaxis, infused cell dose, and development of grade II to IV acute GVHD. For this purpose, continuous variables were categorized as follows: each variable was first divided into 4 categories at approximately the 25th, 50th, and 75th percentiles. If the relative event rates (ratio of the observed number of events to the expected number of events in a category, assuming no variation across categories) in 2 or more adjacent categories (and the mean times-to-event) were not substantially different, these categories were grouped together. If no clear pattern was observed for the primary outcome, the median was taken as the cut point.27

For the multivariate analysis of cGVHD, DFS, and relapse probability, the Cox proportional hazard regression model was used, including in the models all the variables with P < .05 in univariate analysis.28,29

The χ^2 test was used to compare differences in percentages.

All P values were 2-sided and P < .05 was considered statistically significant. P > .1 was reported as not significant (NS), whereas P between .05 and .1 was reported in detail.

The SAS package (SAS Institute, Cary, NC) was used for the analysis of the data.

Results

Data were analyzed as of December 31, 2000. Median follow-up was 57 months (range, 18-119 months) for surviving patients and 9 months (range, 3.4-85 months) for patients not surviving.

Incidence of chronic GVHD and chronic GVHD resolution

Of the 696 patients analyzed, 173 (25%) developed cGVHD at a median of 116 days after transplantation (range, 80-600 days). The limited form of the disease was observed in 109 patients (16%), and extensive cGVHD was diagnosed in 64 cases (9%). As shown in Figure 1A, 2 years after the transplantation, the overall cumulative probability of cGVHD was 27% (95% CI, 23%-30%). The cumulative probabilities of developing either limited or cGVHD were 17% (95% CI, 14%-20%) and 11% (95% CI, 8%-14%), respectively.

For patients with cGVHD, median Karnofsky score at time of last follow-up was 90%, with 72 of the 173 patients (42%) having a Karnofsky score of 100% and only 7 (4%) with a Karnofsky score below 50%.

The skin was the most common target of cGVHD, being involved in 144 cases (83%); the liver was involved in 49 patients (28%) and the gastrointestinal tract in 42 (24%). Oral cGVHD was described in 12 patients (7%); both eye involvement and joint contractures were reported in 6 cases (3%). Finally, 19 children (11%) had lung involvement from cGVHD.

Figure 1B shows the prevalence of cGVHD over time after HSCT. Prevalence was greater during the first year after HSCT, when it reached a value of 18%, and progressively decreased to about 5% at 7 years after transplantation. The prevalence of limited cGVHD was greater than that of extensive cGVHD during the first 20 months after HSCT, whereas the prevalence of extensive cGVHD was higher than that of limited cGVHD from 20 months afterward.

Chronic GVHD resolved in 121 patients (70%), whereas it was still present at the time of last follow-up or at time of death in the remaining 52 children (30%). The median duration of cGVHD, in the 121 patients in whom this complication resolved, was 5 months (range, 1-93 months).

Figure 2 shows the cumulative probability of continuing cGVHD, which was equal to 16% (95% CI, 6%-25%) 8 years after transplantation, and the probability of death with cGVHD over time after development of the complication, which was 12% (95% CI, 3%-18%).

The analysis concerning the impact of cGVHD therapy on the probability of cGVHD resolution and on patient outcome did not show any statistically significant difference among the different treatment modalities (data not shown).

Risk factors for chronic GVHD

A number of demographic and transplant-related characteristics were evaluated as potential risk factors for cGVHD. Tables 2 and 3



Figure 2. Cumulative probability of continuing cGVHD and of death with cGVHD over time after the onset of cGVHD. To calculate the probability of continuing cGVHD, cGVHD resolutions were considered as events, whereas patients whose cGVHD did not resolve were censored at time of last follow-up or of death. To calculate the probability of death with cGVHD, only deaths with active cGVHD were considered events, whereas patients whose cGVHD resolved were censored at time of last follow-up or death.

present the results of univariate and multivariate analysis of risk factors for development of cGVHD.

Several variables were strongly associated with an increased risk of cGVHD in univariate analysis. In particular, older patient age was significantly associated with cGVHD, with children younger than 5 years having a probability of less than 14% of developing this complication, those aged 5 to 15 years a probability of about 30%, and those older than 15 years a probability of 44% (P = .0001). Also donor age was a significant risk factor for cGVHD occurrence in univariate analysis, with a continuous increase in the incidence of this complication from less than 10% for donors younger than 5 years, to more than 40% for donors older than 40 years of age (P = .0002). Female donor and male recipient combination was another factor predicting cGVHD development (34% versus 23%, P = .01). Furthermore, patients affected by hematologic malignancies had a greater incidence of cGVHD (35% versus 13%, P = .01), as well as those treated with TBI (38% versus 17%, P = .0001) and those with previous grade II to IV acute GVHD (45%) versus 14%, P = .0001). CB transplants had a significantly lower probability of cGVHD (3%) compared with bone marrow or peripheral blood transplants (28%, P = .0073). As reported in Table 2, in our cohort of patients, parity of female donors did not show any effect on the risk of cGVHD development.

In multivariate analysis, the following 6 characteristics were associated with an increased probability of developing cGVHD: patient age 15 years or older (RR = 1.82, P = .03), donor age 5 years or older (RR = 3.36, P < .001), female donor and male recipient (RR = 1.52, P = .01), use of TBI as part of the conditioning regimen (RR = 1.56, P = .02), diagnosis of hematologic malignancy (RR = 1.80, P = .01), and, with the highest relative risk, previous grade II to IV acute GVHD (RR = 2.14, P = .0001). On the contrary, transplants of CB stem cells were characterized by a very low incidence of cGVHD, with only 1 child of 32 presenting this complication (RR = 0.07, P = .001).

Causes of death and TRM

Overall, 191 (27%) of the 696 patients analyzed died; 146 (28%) belonged to the group of 523 subjects without cGVHD, whereas 45 (26%) belonged to the 173 children with cGVHD.

Details on the causes of death for patients with and without cGVHD are reported in Table 4. Disease progression was the main

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Cord blood 32 1 3 $(0-9)$ Stem cell sourceBone marrow + peripheralblood 664 172 28 $(24-31)$ $.0073$ Cord blood 32 1 3 $(0-9)$ Conditioning regimenChemotherapy based 369 60 17 $(13-21)$ $.0001$ TBI based 327 113 38 $(33-44)$ GVHD prophylaxisCs-A 348 82 25 $(20-30)$ $.0052$ Cs-A + MTX 129 32 27 $(19-35)$ Cs-A + PDN 63 13 23 $(12-34)$ Cs-A + MTX + ALG or mAb 104 39 42 $(31-42)$ Cs-A + ALG or mAb \pm PDN 33 4 15 $(1-29)$ T-cell depletion 19 3 16 $(0-32)$ GVHD prophylaxis 592 134 24 $(21-28)$ Cell dose infused $Less than 2 \times 10^8/kg$ 117 37 35 $(26-44)$ Less than $2 \times 10^8/kg$ 117 37 35 $(26-44)$ $.09$ $2 \times 10^8/kg$ or more 556 132 25 $(21-29)$ Unknown 23 4 44 $(11-18)$ $.0001$ Acute GVHD grade 1406 54 14 $(11-18)$ $.0001$	Peripheral blood	17	4	30	(4-55)	
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$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	Conditioning regimen					
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$\begin{array}{c c c c c c c c c c c c c c c c c c c $	TBI based	327	113	38	(33-44)	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	GVHD prophylaxis					
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Cs-A	348	82	25	(20-30)	.0052
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Cs-A + MTX	129	32	27	(19-35)	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Cs-A + PDN	63	13	23	(12-34)	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	Cs-A + MTX + ALG or mAb	104	39	42	(31-42)	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	Cs-A $+$ ALG or mAb \pm PDN	33	4	15	(1-29)	
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	T-cell depletion	19	3	16	(0-32)	
$\begin{array}{c ccccc} Cs-A + MTX + ALG \mbox{ or mAb} & 104 & 39 & 42 & (31-52) & .0001 \\ \hline Other prophylaxis & 592 & 134 & 24 & (21-28) \\ \hline Cell dose infused & & & \\ Less than 2 \times 10^8/kg & 117 & 37 & 35 & (26-44) & .09 \\ 2 \times 10^8/kg \mbox{ or more} & 556 & 132 & 25 & (21-29) \\ \hline Unknown & 23 & 4 & \\ \hline Acute GVHD grade & & \\ \hline Grade \mbox{ 0-l} & 406 & 54 & 14 & (11-18) & .0001 \\ \hline \end{array}$	GVHD prophylaxis					
	Cs-A + MTX + ALG or mAb	104	39	42	(31-52)	.0001
Cell dose infused Less than 2 × 10 ⁸ /kg 117 37 35 (26-44) .09 2 × 10 ⁸ /kg or more 556 132 25 (21-29) Unknown 23 4 Acute GVHD grade Grade 0-1 406 54 14 (11-18) .0001	Other prophylaxis	592	134	24	(21-28)	
Less than 2 × 10 ⁸ /kg 117 37 35 (26-44) .09 2 × 10 ⁸ /kg or more 556 132 25 (21-29) Unknown 23 4 4 Acute GVHD grade Grade 0-1 406 54 14 (11-18) .0001	Cell dose infused					
2 × 10 ⁸ /kg or more 556 132 25 (21-29) Unknown 23 4 4 Acute GVHD grade	Less than 2×10^8 /kg	117	37	35	(26-44)	.09
Unknown 23 4 Acute GVHD grade Grade 0-I 406 54 14 (11-18) .0001	2×10^8 /kg or more	556	132	25	(21-29)	
Acute GVHD grade Grade 0-I 406 54 14 (11-18) .0001	Unknown	23	4			
Grade 0-1 406 54 14 (11-18) .0001	Acute GVHD grade					
	Grade 0-I	406	54	14	(11-18)	.0001
Grade II-IV 290 119 45 (39-51)	Grade II-IV	290	119	45	(39-51)	

*The impact of parity on cGVHD occurrence was evaluated in the 117 donorrecipient pairs with a female donor older than 20 years; data on parity were available for 71 of these 117 donors (61%).

Table 3. Multivariate analysis of risk factors for cGVHD development

	Multivariate analysis		
	RR	(95% CI)	Р
Patient age			
Younger than 5 y	1.00		
5-15 y	1.39	(0.93-2.08)	.1
15 y or older	1.82	(1.06-3.12)	.03
Donor age			
Younger than 5 y	1.00		
5-20 y	3.26	(1.57-6.77)	.001
20 y or older	5.73	(2.56-12.8)	.0001
Sex			
Female donor and male recipient vs others	1.52	(1.11-2.07)	.01
Diagnosis			
Malignant vs nonmalignant	1.80	(1.13-2.88)	.01
Donor			
HLA-identical family vs other donors	.69	(0.41-1.17)	NS
Stem cell source			
CB vs bone marrow + peripheral blood	.07	(0.03-0.16)	.0001
Conditioning regimen			
TBI vs chemotherapy	1.56	(1.08-2.26)	.02
GVHD prophylaxis			
Cs-A + MTX + ALG or mAb vs other			
prophylaxis	1.55	(0.92-2.51)	.08
Acute GVHD grade			
Grade II-IV vs grade 0-I	2.14	(1.78-2.59)	.0001

cause of death in 102 patients, whereas 89 children died of transplant-related causes. In patients with cGVHD, we observed a significantly higher percentage of transplant-related deaths (69%) as compared to subjects without cGVHD (40%), and a lower percentage of deaths due to disease progression (31% versus 60%, P = .0006). In the group of children who had cGVHD, this complication was indicated as the primary cause of death in 11 of the 31 patients who died due to transplant-related causes (35%), so that 6% of the 173 patients with cGVHD died because of this complication.

In the 11 cases where cGVHD was reported as the main cause of death, the ultimate cause of death was bronchiolitis obliterans in 4 cases, cytomegalovirus interstitial pneumonia in 1 case, bacterial pneumonia in 1 case, other bacterial infections in 3 cases, hemorrhagic complications in 1 case, and multiorgan failure in 1 case.

For patients treated for nonmalignant disorders, the cumulative probability of TRM was 13% (95% CI, 7%-19%) for children without cGVHD, 5% (95% CI, 0%-15%) for those with the limited form of the disease, and 64% (95% CI, 11%-100%) for those with extensive cGVHD (P = .016). For subjects treated for malignant diseases, the cumulative probability of TRM was 9% (95% CI, 5%-13%) for children without cGVHD, 15% (95% CI, 5%-25%) for those with limited disease, and 24% (95% CI, 12%-36%) for those with extensive cGVHD (P = .036).

Relapse probability

Of the 450 children who underwent HSCT for treatment of hematologic malignancy, 127 (28%) experienced disease relapse. Analyzing the risk of relapse according to cGVHD occurrence, we observed that the 6-year cumulative relapse probability was significantly lower (P = .0001) in children with cGVHD, being 16% (95% CI, 10%-22%) compared with a value of 39% (95% CI, 33%-45%) in patients without cGVHD (Figure 3).

Tables 5 and 6 show the results of univariate and multivariate analysis of the probability of relapse in the 450 children affected by hematologic malignancies. cGVHD confirmed its protective effect

Table 4. Causes of death for patients with and without cGVHD

	No chronic GVHD (523 patients)	Chronic GVHD (173 patients)	Total (696 patients)
Disease progression (%)	88 (60)	14 (31)	102 (53)
Transplant-related causes (%)	58 (40)	31 (69)	89 (47)
Bacterial pneumonia	5	3	
HCMV interstitial pneumonia	2	0	
Bacterial infection	10	5	
Fungal infection	4	4	
Viral infection	0	1	
EBV-PTLD	2	0	
Encephalitis	2	0	
ARDS	5	1	
Idiopathic interstitial pneumonia	5	1	
Autoimmune hemolytic anemia	2	0	
Thrombotic thrombocytopenic			
purpura	2	0	
Liver failure	2	0	
Renal failure	0	1	
Cardiac insufficiency	1	0	
Multiorgan failure	6	2	
Hemorrhage	10	2	
cGVHD	0	11	
Total	146	45	191

 χ^2 = 11.76, *P* = .0006 for the comparison of causes of death (disease progression versus transplant-related cause) in subjects with and without cGVHD.

For the 11 patients with cGVHD reported as the main cause of death, the ultimate cause of death was bronchiolitis obliterans in 4 cases, HCMV interstitial pneumonia in 1 case, bacterial infections in 3 cases, hemorrhagic complications in 1 case, and multiorgan failure in 1 case. HCMV indicates human cytomegalovirus; EBV-PTLD, Epstein-Barr virus-related posttrans-plantation lymphoproliferative disease; ARDS, acute respiratory distress syndrome.

against relapse also in multivariate analysis (RR = 0.32, P = .0001, for patients with cGVHD versus those without cGVHD).

Stratifying the analysis by type of malignancy, we observed that the GVL effect associated with cGVHD was stronger in the 268 patients with ALL. In fact, the relapse probability for children with ALL and cGVHD was 14% (95% CI, 6%-21%) compared with the 47% (95% CI, 39%-54%) observed in children with ALL but without cGVHD (P = .0001). On the contrary, for the 99 patients with AML, the relapse probability was identical (29%) in both groups of patients. Among the 20 children with MDS, development of cGVHD was associated with a lower relapse probability, even

AIEOP BMT REGISTRY Allogeneic BMT in Malignant Diseases Relapse Probability by Chronic GVHD 1.0 cGVHD events = 20 PROBABILITY (95% CI) 70 70 90 800 No cGVHD events = 107 No cGVHD: 39% (33-45) cGVHD: 16% (10-22) Log-rank p = 0.0001 6 YEARS FROM BMT Number of cases at risk 142 90 59 34 cGVHD 58 No cGVHD 308 152 96

Figure 3. Six-year relapse probability after HSCT for children affected by hematologic malignancies, according to cGVHD development. Patients experiencing cGVHD had a significantly lower risk of relapse (P = .0001).

Table 5. Univariate analysis of relapse probability in the 450 patien	ts
with malignant disease	

	Univariate analysis				
	Number of Probability,				
	cases	Events	%	(95% CI)	Р
Patient age					
Younger than 5 y	110	30	28	(20-37)	NS
5-10 y	174	51	34	(26-41)	
10-15 y	125	32	29	(20-38)	
15 y or older	41	14	41	(22-60)	
Donor age				, ,	
Younger than 5 y	81	23	31	(20-42)	NS
5-10 y	95	24	27	(17-36)	
10-15 y	64	18	33	(19-46)	
15-20 y	55	14	28	(15-40)	
20-30 y	69	22	34	(23-46)	
30-40 v	56	17	37	(21-52)	
40 v or older	30	9	44	(15-72)	
Sex				(-)	
Female donor and male					
recipient	103	20	21	(13-29)	.0706
Other	320	90	32	(26-38)	
Diagnosis				(/	
ALL	268	87	35	(29-41)	NS
AML	99	24	31	(19-42)	
MDS	36	8	25	(10-40)	
CMI	28	3	18	(0-39)	
Lymphoma	19	5	28	(7-49)	
Disease phase				()	
Farly	285	66	26	(21-32)	0003
Advanced	165	61	42	(34-51)	
Donor				()	
HLA-identical family	306	85	30	(25-36)	NS
Other donor	144	42	36	(26-46)	
Stem cell source				(/	
Bone marrow + peripheral					
blood	434	121	31	(26-36)	NS
Cord blood	16	6	52	(19-85)	
GVHD prophylaxis				()	
Cs-A + MTX + ALG or mAb	81	25	34	(23-45)	NS
Other prophylaxis	369	102	31	(26-36)	
Conditioning regimen				(/	
Chemotherapy based	148	41	33	(24-43)	NS
TBI based	302	86	31	(26-37)	
Acute GVHD				()	
0-1	223	77	38	(31-45)	.0005
II-IV	227	49	25	(18-31)	
Chronic GVHD				()	
Absent	308	107	39	(33-45)	.0001
Present	142	20	16	(9-22)	
Chronic GVHD		-	-		
Absent	308	107	39	(33-45)	.0001
Limited	89	16	20	(11-28)	
Extensive	53	4	10	(0-19)	

though the advantage was not statistically significant (0% versus 36%, P = .053). Because only 3 of the 28 children with CML experienced a relapse, it was impossible to analyze the role of cGVHD on leukemia recurrence in this group of patients.

Survival and DFS

Overall, 482 of the 696 patients analyzed (69%) are alive and disease free, with a cumulative probability of DFS of 67% (95% CI, 63%-71%).

Figure 4 shows the 6-year cumulative probability of DFS for

Table 6. Multivariate analysis of relapse probability in the 450 patients with malignant disease

	Ν	Multivariate analysis		
	RR	(95% CI)	Р	
Sex				
Female donor and male recipient vs others	0.82	(0.56-1.21)	NS	
Disease phase				
Advanced vs early	2.11	(1.48-3.00)	.0001	
Acute GVHD				
II-IV vs 0-I	0.91	(0.65-1.27)	NS	
cGVHD				
Present vs absent	0.32	(0.20-0.52)	.0001	

Variables with P < .1 in univariate analysis were included in the multivariate analysis.

patients with either nonmalignant disorders (Figure 4A) or hematologic malignancies (Figure 4B), respectively. For children with nonmalignant diseases, no statistically significant difference was observed for DFS between patients without or with cGVHD (85% versus 77%, P = NS), this finding being due to the fact that most of patients with cGVHD had the limited form of the disease. On the





Figure 4. Probability of DFS. Six-year probability of DFS according to the development of cGVHD is shown for patients affected by nonmalignant disorders (A) and for those affected by hematologic malignancies (B). The difference was not statistically significant for nonmalignant disorders, whereas in the group of patients affected by malignant diseases, the advantage for subjects with cGVHD was statistically significant (P = .0013).

contrary, considering children receiving transplants for malignant diseases, patients who developed cGVHD had a significantly better DFS, as compared to children without cGVHD (68% versus 54%, P = .0013).

Tables 7 and 8 summarize the results of univariate and multivariate analysis of DFS probability for the 450 children affected by malignant diseases. As for relapse, the positive impact of cGVHD on DFS was confirmed also in multivariate analysis, where this variable was associated with a relative risk of DFS of 2.07 (P = .0001).

Table 7. Univariate analysis of DFS probability in the 450 patients with malignant disease

		Univariate analysis				
	No. of		Probability,			
	cases	Events	%	(95% CI)	P	
Patient age						
Younger than 5 y	110	35	66	(56-76)	NS	
5-10 y	174	72	57	(49-64)		
10-15 у	125	52	57	(48-66)		
15 y or older	41	17	54	(35-72)		
Donor age						
Younger than 5 y	81	31	61	(50-72)	.09	
5-0 y	95	31	65	(55-76)		
10-15 у	64	22	63	(49-76)		
15-20 у	55	18	65	(52-78)		
20-30 у	69	32	52	(40-64)		
30-40 y	56	24	48	(34-62)		
40 y or older	30	14	45	(21-70)		
Sex						
Female donor and male						
recipient	103	53	59	(50-67)	NS	
Other	320	123	59	(53-65)		
Diagnosis						
ALL	268	116	56	(50-62)	NS	
AML	99	30	64	(53-76)		
MDS	36	15	54	(35-73)		
CML	28	8	66	(45-87)		
Lymphoma	19	7	63	(41-85)		
Disease phase				,		
Early	285	93	65	(59-71)	.0001	
Advanced	165	83	48	(39-56)		
Donor				()		
HLA-identical family	306	103	65	(59-70)	.0003	
Other donors	144	73	45	(34-55)		
Stem cell source				()		
Bone marrow + peripheral						
blood	434	167	59	(54-64)	.1032	
Cord blood	16	9	38	(9-66)		
GVHD prophylaxis		-		(****)		
Cs-A + MTX + ALG or mAb	81	39	51	(39-62)	.0533	
Other prophylaxis	369	167	61	(55-66)		
Conditioning regimen	000		01	(00 00)		
Chemotherapy based	148	52	59	(50-69)	NS	
TBI based	302	124	58	(52-64)		
	002	121	00	(02 04)		
0-1	223	80	58	(51-65)	NS	
11-11/	223	87	60	(53-67)	140	
	221	07	00	(55-67)		
Abcont	200	124	55	(40.60)	0012	
Absent	140	134	55	(49-00)	.0013	
- LESELIL	142	42	00	(00-70)		
Abcont	200	104	FF	(40.60)	0055	
ADSENT	308	134	55	(49-60) (50-70)	.0055	
Limited	89	26	68	(58-79)		
Extensive	53	16	68	(54-81)		

Table 8. Multivariate analysis of DFS probability in the 450 patients with malignant disease

	Multivariate analysis		
	RR	(95% CI)	Р
Donor age			
Younger than 20 y vs 20 y or older	1.23	(0.79-1.90)	NS
Disease phase			
Advanced vs early	0.57	(0.42-0.78)	.0004
Donor			
HLA-identical family vs other donors	1.52	(0.95-2.43)	.08
GVHD prophylaxis			
Cs-A + MTX + ALG or mAb vs other prophylaxis	.97	(0.58-1.64)	NS
cGVHD			
Present vs absent	2.04	(1.43-2.91)	.0001

Variables with P < .1 in univariate analysis were included in the multivariate analysis.

As already observed for relapse probability, also for DFS the favorable influence of cGVHD was observed only for children with ALL, where the cumulative probability of DFS was 74% (95% CI, 64%-83%) for patients developing cGVHD and 47% (95% CI, 38%-53%) for children without cGVHD (P = .0001), whereas the effect on other hematologic malignancies was nonsignificant. In fact, in the 99 patients with AML, the DFS probability was 59% (95% CI, 36%-82%) for children with cGVHD and 68% for those without cGVHD (P = NS), and in the 28 children with CML it was 64% (95% CI, 35%-92%) and 71% (95% CI, 44%-97%) for subjects with and without cGVHD, respectively (P = NS). Finally, also considering the 20 children with MDS, the outcome of patients with and without cGVHD was similar, DFS being 67% (95% CI, 36%-97%) and 64% (95% CI, 35%-92%), respectively (P = NS).

Discussion

Chronic GVHD remains a substantial problem of allogeneic HSCT. Several studies, published in recent years, introduced the notion that recipient age is a major risk factor for the development of cGVHD and that children have a lower probability of presenting this complication.⁴⁻⁶ However, this information, as well as our clinical knowledge of cGVHD and of its risk factors, are based primarily on results of analyses performed in adults.

In our study, conducted on the largest pediatric population reported so far, we observed a cumulative probability of cGVHD of 27%, which is similar to probabilities ranging from 22% to 29% reported by 3 other analyses conducted on smaller cohorts of pediatric patients in recent years.¹³⁻¹⁵ This probability is nearly half of the estimated probability of 40% to 50% described in adults.⁴⁻⁶ In our group of patients younger than 18 years, older patient and donor age were predictive of a higher risk of cGVHD both in univariate and in multivariate analysis. On the other hand, the type of donor and of GVHD prophylaxis was not predictive of the development of this complication. In multivariate analysis, other factors, such as the female donor-male recipient combination, a diagnosis of malignant disease, the use of TBI, and, as expected, previous grade II to IV acute GVHD, were characterized by a significantly higher risk of cGVHD. Furthermore, in agreement with the observation of Rocha et al,30 use of CB progenitor cells was confirmed to be characterized by a very low probability of cGVHD. In contrast, we could not confirm the observation of an increased risk of cGVHD in subjects receiving peripheral blood stem cells.^{8,9} This was possibly due to the limited number of patients receiving peripheral blood stem cells.

Our study confirms previously published reports documenting that male patients given a transplant from a female donor experience more GVHD,⁶ an observation that has been biologically explained hypothesizing that male antigens (ie, H-Y) not shared by females are recognized and attacked by the donor's immune system.^{31,32} However, our data are particularly intriguing because most female donors were young and, thus, not sensitized toward male antigens by previous pregnancies. Furthermore, even if evaluated on a limited number of subjects, parity of female donors was not a risk factor for cGVHD development; this suggests that even a primary immune response toward these molecules may produce cGVHD.

Children with nonmalignant disorders had a reduced risk of developing cGVHD. Several factors may be considered to explain this observation. Because children with these diseases do not benefit at all from GVHD, it is reasonable to hypothesize that the most effective pharmacologic strategies for both GVHD prevention and therapy of acute GVHD were used in these patients. Moreover, it has been already reported that these patients can have a mixed chimerism,³³⁻³⁵ and this condition is associated with reduced susceptibility to GVHD, probably through mechanisms of central tolerance with negative selection of both host-reactive and donor-reactive T cells.36 Finally, some of the children with nonmalignant disorders (ie, those with aplastic anemia or with congenital immunodeficiencies) were given less intensive preparative regimens and it has been hypothesized that the cytokine storm, which is dependent on the intensity of the conditioning regimen, triggers development of GVHD.³⁷ In this regard, the finding of an increased risk of cGVHD in children given TBI as part of the preparative regimen is also of particular interest. Moreover, because it has been suggested as a relevant role of recipient's thymus in promoting tolerance of donor cells, we speculate that the higher risk of cGVHD in children given TBI can be partly explained by a radiation-induced damage to thymic epithelial cells.³⁸⁻⁴⁰

We also found an increased risk of cGVHD in patients receiving a transplant from an older donor. The detrimental effect of increased donor age on cGVHD has been recently reported in a cohort of patients given HSCT from unrelated volunteers.⁴¹ The biologic mechanisms accounting for this effect of donor age remain unclear and deserve further investigation.

Considerable experimental and clinical data suggest a role of the immune system in controlling cancer and, particularly, leukemia, through the GVL effect.^{12,42-45} In our study, we document a strong protective effect of cGVHD against relapse in children with ALL, whereas the advantage was lower and not statistically significant for patients with MDS and absent for those with AML. Moreover, in children with ALL, the increased TRM associated with cGVHD was largely offset by the impressive reduction in relapse probability, thus resulting in a significantly better survival and DFS.

Despite a beneficial GVL effect against relapse, cGVHD remains a major determinant of poor long-term outcome and of impaired quality of life after allogeneic HSCT.² For this reason, several efforts have been made to identify factors having an unfavorable prognostic significance, to select subjects with higher risk of either death or invalidating sequelae (as severe joint contractures requiring physical therapy, impairment of lung function, keratoconjunctivitis sicca with corneal lesions, malnutrition due to gastrointestinal involvement), who need to be treated promptly and more intensively.⁴⁶⁻⁴⁹ Over the last few years, new drugs and different treatment modalities have been developed for patients with cGVHD,⁵⁰ even though the optimal treatment is still controversial and no single therapy has been demonstrated to be superior and most effective.⁵¹

In our study population, subjects developing cGVHD had a significantly higher TRM, as compared to those without the disease, even though only 6% of children with cGVHD died directly or indirectly because of this complication. The negative effect of cGVHD was particularly evident in patients with the extensive form of the disease and for those affected by nonmalignant disorders. These observations emphasize the need for a more effective prophylaxis for patients undergoing transplantation with the aim of curing nonmalignant disorders and a more effective treatment for those with extensive GVHD or with poor prognostic clinical features.

Our data document that children have a lower incidence of cGVHD compared with adults. Use of CB stem cells and of conditioning regimens without TBI, as well as better strategies for preventing acute GVHD occurrence, could further decrease the risk of this complication. We provide evidence for a strong antileukemia effect of cGVHD for ALL, but not for other hematologic malignancies. Despite an increased TRM, the probability of DFS in ALL was not impaired by the occurrence of this complication.

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