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# ORIGINAL RESEARCH Anticipation of Relapse and Acute Graft-Versus-Host Disease after Allogeneic Peripheral Blood Stem Cell Transplantation: The Fundamental Role of Antigen-Presenting (Dendritic) Cells

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Background: Dendritic cells (DCs) are antigen-presenting cells. In humans two distinct lineages of DCs exist; DC1 and DC2. Efforts to explore the role of DCs in acute graft-versus-host disease (aGVHD) after allogeneic peripheral blood stem-cell transplantation (PBSCT) are gaining traction. However, further research is needed to identify particular lineages and their values in terms of developing an evidence-based aGVHD- or relapse-prevention strategy. We monitored DC counts and subsets in PBSC grafts while harvesting stem cells in recipients to elucidate their value in anticipating disease relapse or aGVHD.

Methods: We enrolled 29 participants. Using fluorescence-activated cell sorting, total counts/kg of CD34<sup>+</sup>, DCs, and DC subsets were analyzed in 29 PBSC-graft components using CMRF44, CD11c, and CD4 monoclonal antibodies (MoAbs).

**Results:** In the 29 grafts, we detected a significant positive correlation (P < 0.01) between DCs and both DC1 and DC2. Significantly higher counts (P<0.01) of DCs and DC1 in those who had developed aGVHD (nine cases) were also observed. Relapsed cases (two) were also associated with higher counts of DCs and DC2. A significant positive correlation (P < 0.05), was recorded between DCs and DC1 counts and the day of myeloid engraftment, while this was not detected on the day of platelet engraftment. Myeloid engraftment transpired earlier in patients without aGVHD. Increased DC-graft numbers, particularly DC1 measured by CD11c Moabs, were associated with aGVHD. Recipients of higher numbers of CD4<sup>bright</sup> DCs had an increased risk of relapse after allogeneic PBSCT.

**Conclusion:** This study analyzed DCs in PBSC grafts, using novel specific MoAbs and flow cytometry. Our data showed that higher donor DC1 counts were linked to the incidence of aGVHD and DC2 with relapse. We propose a fundamental role for DC-graft monitoring in anticipating aGVHD and disease relapse.

Keywords: dendritic, cell, biomarker, acute GVHD, relapse, allogeneic PBSCT

## Introduction

Dendritic cells (DCs) develop from a precursor in the bone marrow and circulate through the blood to the outermost body tissue, where they participate in antigen scrutiny and T-cell activation.<sup>1</sup> Exact or approximate DC counts in various disease conditions are not well understood. The absence of a standardized method for estimating DC numbers in a certain disease restricts investigations. Some authors have established a technique appropriate for routine checking of DC counts in the blood after short-term culture by fluorescence-activated cell sorting utilizing the CMRF44 monoclonal antibody.<sup>2</sup> CMRF44 antigens represent the phenotype, shape, and functional characteristics of DCs.<sup>3</sup> Measuring CMRF44<sup>+</sup> cells represent DC numbers in peripheral blood stem cell (PBSC) grafts. Because DCs are scarce, culturing mononuclear cells is needed to permit the utilization of DC-motivation antigens for positive recognition of DCs.<sup>2</sup> Aripinati et al suggested that G-CSF– militarized PBSCs, including immunoeffector antigen-presenting cells that stimulate T cells, to yield T<sub>h</sub>2 cytokines. In humans, numerous DC lineages have been identified. Of these, DC1 induces differentiation of T lymphocytes to T<sub>h</sub>1 cells and DC2 augments T-cell differentiation to T<sub>h</sub>2 cells. There are other DC subtypes, such as plasmacytoid, inflammatory, and Langerhans DCs.<sup>4</sup>

The impact of DCs in regulating immunoreconstitution after transplantation likely represents the competing effect of donor DC1 versus DC2 in the graft and donor DCs that are differentiated from CD34<sup>+</sup> cells by the effect of cytokines derived from DC1 or DC2.<sup>5</sup> Recently, Wang et al demonstrated that blocking CD11c on human PB DCs inhibited CD4<sup>+</sup> T-cell proliferation and differentiation into IFN $\gamma$ -generating T<sub>h</sub>1 cells, which are essential in acute graft-versus-host disease (aGVHD) pathogenesis.<sup>6</sup>

Studies have demonstrated reduced DC numbers in the PB of recipients with moderate–severe aGVHD.<sup>7,8</sup> On the contrary, others discovered no link between aGVHD and higher DC counts in the early stages of aGVHD.<sup>9,10</sup> These disparities could be explained in part by the steroid treatment for aGVHD, which causes a rapid drop in circulating DCs. Another report found that CD4<sup>bright</sup> cells had similar phenotypes and counts as CD123<sup>bright</sup> type 2 DC precursors. After short culture, CD4<sup>bright</sup> cells differentiated into typical DCs.<sup>11</sup>

It has been proved that recipients transplanted with higher CD4<sup>bright</sup> DC counts have more reversions. Relapse is the main etiology of disease after autologousPBSC transplantation (aPBSCT).<sup>12</sup> In light of what was previously mentioned, there is still a definite need to undertake studies designed to explore the roles of DCs post-HSCT. The current study was conducted to elucidate the value of DCs in predicting disease relapse and/or aGVHD in aPBSCT recipients.

#### **Methods**

#### Subject and Graft Selection

A total of 29 recipients were prospectively enrolled in the study. They underwent allogeneic PBSCT from 29 HLAmatched siblings. Their age was  $\geq$ 18 years, and women represented >50% of the study sample, as shown in <u>Supplemental</u> <u>Table S1</u>. Willingness to participate in the study was an inclusion criterion.

#### Analysis of Grafts

Analysis of the allogeneic PBSC grafts (harvests) was done as follows.

#### Mobilization

The matched donors received G-CSF SC injections (10 µg/kg/day) for 5 consecutive days.

• This study was carried out on 29 allogeneic PBSC grafts (harvests) after short-term culture in RPMI1640/10% FBS at 37°C and a 5% carbon dioxide incubator. Heparinized blood samples (5 mL) were collected from harvests and handled within 4 hours. Routine complete blood counts were done using a Coulter STKS. Absolute mononuclear cell counts were determined (mononuclear cells/liter of blood). White blood–cell counts were adjusted between 3,000 and 10,000/mm<sup>3</sup>. The rest of the blood samples were diluted 1:1 with sterile PBS and underlaid with Ficoll–Hypaque before being centrifuged at 520 g for 15 minutes at 37°C. The mononuclear cells were **mended** from the low-density boundary and washed three times, then cultured in RPMI1640/10% FBS for 24 hours at 37°C and the 5% CO<sub>2</sub> incubator. Trypan blue exclusion was used to assess cellular viability after culture.<sup>13</sup>

#### Panel of Monoclonal Antibodies Used

- CD4 FITC/CD34 PE (BD Biosciences)
- CD34 FITC/CD11c PE (BD Biosciences)
- CMRF44 FITC (received as a gift from Professor DNJ Hart (Director, Mater Medical Research Institute, Raymond Terrace, Brisbane, Australia)/CD34 PE (BD Biosciences)
- Isotype controls

#### Flow-Cytometry Analysis

Sample analysis was done using a Coulter Epics-XL-MCL flow cytometer:

- CD34 enumeration ISHAGE protocol,  $CD34^+$  threshold  $\ge 3 \times 10^6$  cells/kg body weight.
- Percentage and absolute CD34<sup>+</sup> cell counts, DC counts, and their subsets were analyzed.

The graft components of all cases were analyzed for enumeration of DC numbers and DC subsets through monitoring of the absolute count/kilogram body weight of the parameters measured by flow cytometry: total nucleated cells, CD34<sup>+</sup> cells, CMRF44<sup>+</sup> DCs, CD11c<sup>+</sup> DCs (DC1), and CD4<sup>bright</sup> DCs (DC2). All these parameters were correlated with one another and the end result of transplantation with regard to complications, especially aGVHD and relapse.<sup>14,15</sup>

#### Treatment, Observation, and Follow-Up

Prospective longitudinal observational follow-up of the recipients was carried out. For conditioning regimens, seven cases (24.1%) underwent minitransplantation (nonmyeloablative SCT), while 22 cases (75.9%) underwent myeloablative transplantation. All patients received primary prophylaxis for aGVHD.

#### **GVHD** Prophylaxis

All patients received GVHD prophylaxis (calcineurin inhibitor, methotrexate, mycophenolic acid, or post-HCT cyclophosphamide). In a majority, cyclosporine (CsA) and short-course methotrexate were given as GVHD prevention. Starting on day –4, CsA was given in an intravenous dose of 2.5 mg/kg twice per day for 4 hours, then switched to an oral dose of 5 mg/kg twice daily once the mucositis was cured. The dose was modified to obtain a target level of 200–400 ng/mL after CsA levels were evaluated. On day 1, the methotrexate dose was 10 mg/m<sup>2</sup>, and on days 3, 6, and 11, the dose was 7 mg/m<sup>2</sup>. Day 6 and day 11 dosages were skipped if the mucositis was severe (grade III/IV) or bilirubin was >20 mg/L.

#### Follow-Up

All patients (recipients) were monitored for 100 days after transplantation for incidence of aGVHD, relapse, and other complications. They were assessed by clinical examination, complete blood count, blood culture, and liver and renal function. aGVHD, relapse, and relapse-free survival were the outcomes of interest. The onset of aGVHD usually occurred during the first 2 months (60 days) following PBSCT, and systemic steroid therapy was the primary line of treatment for aGVHD grade II. Staging and clinical grading for GVHD severity and response criteria were used as previously described<sup>14,15</sup> (as described in Supplementary File Tables S2 and S3).

The presence of any morphological evidence of leukemia at the level of bone marrow or extramedullary sites was used to determine leukemia relapse. This was further confirmed with minimal residual disease monitoring with flow cytometry or PCR. The period from the start of the conditioning regimen to event or final follow-up was used to assess event-free survival (EFS) (aGVHD or relapse or death). The period from the start of the conditioning regimen to death or last follow-up was defined as overall survival (OS). Recipients were then classified into three groups based on clinical outcome: group I (event-free), group II (aGVHD), and group III (disease relapse).

#### Statistical Analysis

Data were collected and analyzed with SPSS 18.0. GraphPad Prism V5 was used also. Descriptive data are presented as means  $\pm$  SD if numeric and percentages if categoric. Correlations and relationships were evaluated using Spearman's

rank-correlation coefficient. Kaplan–Meier survival curves were plotted for estimating OS, EFS, and relapse-free survival. Figures were created using GraphPad Prism 5.02). In all tests, statistical significance was presumed at P<0.05.

# Results

#### Kinetics and Components of the Harvests

First, the kinetics and variable components of the grafts were examined. There were highly significant (P<0.001) positive correlations of CMRF44<sup>+</sup> DC numbers (median 11.88) with both DC1 (median 5.63) and DC2 (median 4.3).Moreover, a significant negative association was noted between DC numbers and CD34<sup>+</sup> cells. Interestingly, graft components showed comparable CMRF44 DC, DC1, and DC2 cell numbers in grafts from male and female donors (<u>Table S4</u>, Figure 1A–D). <u>Supplementary Figure S1</u> shows graft components and their relationship to donor age and sex. Seven cases (24.1%) showed increased CMRF44<sup>+</sup> DC numbers in the grafts. Four of those cases (13.8%) showed increased DC1 numbers, while two cases (6.9%) showed increased DC2 numbers. The last one case had neither increased DC1 nor DC2. Interestingly, all these cases developed complications after allogeneic PBSCT, particularly aGVHD, relapse, and higher fatality rate (Figure 1H–J).

### Characteristics and Indications for Transplantation

As delineated in Figure 1E–G and Tables 1 and <u>S4</u>, the commonest underlying hematologic disorders were acute myeloid leukemia (AML), severe idiopathic aplastic anemia, and acute lymphoblastic leukemia (ALL), with variable cytogenetic abnormalities. A total of 21 cases (72.4%) underwent myeloablative transplantations. Table 1 shows treatments of all patients before allogenetic PBSCT, their cytogenetic profile, disease state at transplantation, and HLA-matched intensity.

# Associations among Graft-DC Counts, Subsets, and Postallogeneic PBSCT Complications

Considering development of complications after allogeneic PBSCT, particularly aGVHD and relapse, recipients were stratified into three groups based on clinical outcomes.

#### Group I

This was the event-free group (18 recipients, 62.1%). These recipients did not develop any complications during followup (0–100 days), either clinically or on laboratory results, and they were doing well.

#### Group II

This group comprised recipients (31%) who suffered from manifestations of aGVHD: grade I (4 recipients), grade II (3 recipients), and grades III and IV (one recipient each). Estimation of DC components in these cases revealed that DC numbers were increased in five cases, and increased DC1 numbers were recorded in four of those five (Figure 1H, Table 2). The numbers of DCs and DC1 reached higher levels with severity of GVHD in the recipient with grade IV aGVHD. This recipient died after 3 weeks. Conversely, the other cases did not show increased numbers of either DCs or DC1 in their grafts, despite clinical evidence of aGVHD. These 4 recipients had low-risk aGVHD (grade I, Table 2). From these results, we found that only group II had significantly (P<0.001) higher DC and DC1 counts (in their grafts) compared than event-free recipients (group I), (Tables 2 and S5, Figure 2A–K). Figure 2L is an example of aGVHD cases that showed increased CMRF44<sup>+</sup> DCs and CD11c<sup>+</sup> DCs (DC1, n=23, grade IV).

#### Group III

This group comprised two recipients (6.9%) who developed relapse after aPBSCT. Higher doses of DCs and DC2 were recorded in the grafts of those cases (Figure 3A–C). Figure 3D and E is an example of relapsed cases that showed increased CMRF44<sup>+</sup> and CD4<sup>bright</sup> DCs (DC2). Analysis of donor-related factors revealed that DC numbers had no significant relationship to age, sex, virological studies, or blood group.



Figure I Graft components and clinical features of the 29 recipients that underwent allogeneic PBSCT. Schematic box plots representing the correlations of CD34<sup>+</sup> cell counts and different types of DC among 29 grafts. (A) Correlation between CD34<sup>+</sup> and CMRF44<sup>+</sup> DC counts; (B) correlation between CMRF44<sup>+</sup> DC and DC1 counts; (C) correlation between CMRF44D<sup>+</sup> and DC2 counts; (D) correlation between DC1 and DC2 counts. Underlying diseases (E), cytogenetic profiles (F), and conditioning regimens (G) following allogeneic PBSCT. (H) Flowchart of PBSC-recipient outcomes. (I) Kaplan–Meier graphs of cumulative survivors free of acute GVHD and relapse 100 days after allogeneic PBSCT among subcohorts with high and low DC counts. (J) Percentage of survivors among subcohorts with high and low DC counts. Abbreviations: DC, dendritic cell; SAA, severe aplastic anemia; β-TM, β-thalassemia major; NHL, non-Hodgkin's lymphoma; MDS, myelodysplastic syndrome; HDL, Hodgkin's lymphoma; ALL, acute lymphoblastic leukemia; AML, acute myeloid leukemia; MM, multiple myeloma; CML, chronic myeloid leukemia; inv(16), chromosome 16 inversion–associated translocations; del, deletion; t, translocation; Ph, Philadelphia chromosome; aGVHD, acute graft-versus-host disease; PBSCT, peripheral blood stem–cell transplantation.

# Associations Between Graft Characteristics and Clinical Outcomes

For graft characteristics that could affect recipients' clinical outcome, those who were transplanted with allografts containing higher CD34<sup>+</sup>cells counts fewer less complications and longer EFS after aPBSCT. Noteworthily, no significant differences were found in study groups I or II in respect if dose of CD34<sup>+</sup>cells/kg, or other variable components

Diagnosis	n	Cytogenetic	n	Pretransplant therapy	State of disease at transplantation	HLA compatibility
Acute myeloid leukemia (AML), n=8 *MI AML *M4 AML *M5 AML	8 (27.6%) 5 (17.2%) I (3.5%) 2 (6.9%)	Normal t(8;21)(q22;q22) inv(16)(p13q22) t(9;11)(p21–22;q23)	3 (37.5%) 3 (37.5%) I (12.5%) I (12.5%)	Standard 3/7 induction	7 (87.5%) CRI I (12.5%) CRI	Identical (100%) ≥I alleles/ antigens (0)
Acute lymphoblastic leukemia (ALL), n=5 *Pro-B ALL *Common-B ALL *T-cell ALL	5 (17.4%) 2 (6.9%) 1 (3.5%) 2 (6.9%)	Ph <sup>-</sup> Trisomy 8 Normal Normal	5 (100%) 1 (20%) 1 (20%) 2 (40%)	Multiagent chemotherapy	4 (80.0%) CRI I (20.0%) CRI	Identical (100%) ≥1 alleles/ antigens (0)
Chronic myeloid leukemia, n=2	2 (6.9%)	Ph <sup>+</sup> (9;22)(q34;q11)	2 (100%)	Second-line TKIs	Chronic phase	ldentical (100%) ≥1 alleles/ antigens (0)
Multiple myeloma, n=I	I (3.5%)	t(4;14)(p16;q32)	I (100%)	VRD then HDT	Remission	Identical
Non-Hodgkin's lymphoma, n=l	l (3.5%)	Unknown	I (100%)	R-CHOP then high-dose chemotherapy	Remission	Identical
Hodgkin's lymphoma, n=l	I (3.5%)	Unknown	I (100%)	ICE then high- dose chemotherapy	Remission	Identical
Myelodysplastic syndrome	3 (10.3%)	Normal	3 (100%)	Induction chemotherapy	Remission	ldentical (100%) ≥1 alleles/ antigens (0)
Severe aplastic anemia, n=6*	6 (20.7%)	Trisomy 8 7q del Normal	4 (66.6%) I (16.7%) I (16.7%)	CyA+ATG ± Tpo-RA	NR	ldentical (100%) ≥1 alleles/ antigens (0)
β-thalassemia major, n=2	2 (6.9%)	I B-36 del Unknown	I (50.0%) I (50.0%)	Genotyped RBCS transfusions	Transfusion- dependent	ldentical (100%) ≥1 alleles/ antigens (0)
Mean age (years)			19.7	+1.9		
Sex (M/F)			20/9 (	2.2/1)		

Table I Demographic characteristics, ur	derlying hematologic disorders,	, treatment, response, cyt	togenetic profiles and HLA compat-
ibility of the 29 recipients that underwer	nt aPBSCT		

Notes: \*Morphological diagnosis was aplastic anemia in all aplastic anemia patients without cytogenetic abnormalities where cytogenetic studies were carried out at diagnosis. One patient received pediatric-style multiagent chemotherapy.

Abbreviations: inv(16), chromosome 16 inversion-associated translocation; del, deletion; t, translocation; Ph, Philadelphia chromosome; VRD, bortezomib, lenalidomide, and dexamethasone; HDT, high-dose therapy; TKIs, tyrosine-kinase inhibitors; R-CHOP, rituximab, cyclophosphamide, hydroxydaunorubicin hydrochloride (doxorubicin hydrochloride), oncovin, and prednisone; ICE, ifosfamide carboplatin etoposide; CyA, cyclosporin A; ATG, antithymocyte globulin; Tpo-RA, thrombopoietin receptor agonist; CR, complete remission; NR, no response.

of the graft. There was no significant correlation between numbers of DC subsets (DC1 and DC2 and dose of CD34<sup>+</sup> cells), but there was a significant negative correlation between numbers of CMRF44<sup>+</sup> DCs and CD34<sup>+</sup> cells (Table 3).

It was found that a lesser dose of CD34<sup>+</sup> donor cells was linked to aGVHD (two cases) and relapse (one case). A statistically significant association (positive) was found between DC numbers and day of myeloid engraftment, but no significant correlation was detected for day of platelet engraftment (P<0.05, Figure S2). Considering the occurrence of

aGVHD, myeloid engraftment occurred earlier in group I (P<0.05) than group II, while no significant difference was found regarding platelet engraftment (Table 4). Multivariate analysis showed asignificant differences among groups I, II, and III when considered jointly for DCs, DC1, and D2. Wilks's  $\lambda$ =0.094 ( $F_{22,32}$ =3.281, P=0.001, partial  $\eta^2$ =0.693). A separated ANOVA was conducted for each dependent variable, with each evaluated at  $\alpha$ =0.025. There was a significant difference between group I and groups II and III for DCs, ( $F_{2,26}$ =19.5, P<0.001, partial  $\eta^2$ =0.6, with group I (mean 8.62) DC numbers lower than groups I and II (means 20.6 and 23.3, respectively). There was a significant difference between group II and groups I and III for DC1 ( $F_{2,26}$ =12.3, P<0.001, partial  $\eta^2$ =0.495), with group II (mean 13.3) DC1 numbers higher than groups I and III (means 4.6 and 5.3, respectively). There was a significant difference between group III and groups I and III for DC2 ( $F_{2,26}$ =123.67, P<0.001, partial  $\eta^2$ =0.646), with group III (mean 18.4) DC1 numbers higher than groups I and II (means 3.5 and 6.9, respectively). There was no significant differences among the three groups for age or other graft components (Tables S6–S8).

#### Associations Between Hospitalization Days and Graft-DC Counts and Subsets

A statistically significant association was detected among graft-DC numbers and days of hospitalization after aPBSCT, but a significant negative correlation was detected as regards  $CD34^+$  cell numbers and days of hospitalization (Figure 3F–I). EFS was affected by dose of  $CD34^+$  donor cells, and higher DC and lower doses of  $CD34^+$  cells shortened EFS (Figure 3J and K). Figure 4 shows estimated marginal means of DCs and a correlation matrix of variables of graft components and patients' weight.

#### Associations Between Graft-DC Counts, Subsets, and Survival Rates

During the period of this study (0–100 days), the survival rate was 94.4% in the event-free group (group I). This decreased to 77.8% with increased numbers of DCs and DC1 in those who developed aGVHD (group II). High numbers of CD4<sup>bright</sup> DCs were correlated with reduced survival rate following aPBSCT (50%) among relapsed cases (group III, Figure 3L).

#### Discussion

DCs are a group of bone marrow–derived immunoeffector cells that play an important role in presenting antigens and their collaboration with T lymphocytes. As such, they are essential for cell-mediated immunity and starting the chief T-cell immunoreaction.<sup>16</sup> Because of DC scarcity in harvests, their shorter survival in aPBSCT recipients, and the greater number of recipient DCs that are able to present alloantigens to donor T lymphocytes, the role of donor DCs and their subtypes in grafts has been neglected. Moreover, it is unclear if low DC counts after engraftment are linked to underprivileged consequences or if this is an indicator of humble immunological retrieval. Also, it seems unclear whether reduced PB DC numbers after HSCT reflect decreased DC synthesis, treatment effects (steroids) during this period, or enhanced DC migration to tissue impacted by infections and/or aGVHD. Indeed, the role of DC-count assays after HSCT as a predictive value in posttransplantation outcomes is still a mystery.<sup>17</sup> This study focused on exploring DC numbers and subsets in allogeneic PBSCT grafts and investigating their predictive value in incidence of aGVHD and disease relapse. These objectives were examined in a two-phase study: firstly flow-cytometry analyses of grafts and secondly recipient treatment and follow-up.

As far as we know, this is the first time that recognized DC markers have been used for concurrent quantification, recognition, and segregation of two discrete DC subsets in freshly prepared blood samples. Flow-cytometry analysis has been used to enumerate CD11<sup>+</sup> DCs (DC1) and CD4<sup>bright</sup> DCs (DC2) in PBSC grafts.<sup>17</sup> This study reported significant negative correlations between DC counts and CD34<sup>+</sup> cells. This could be explained by a previous report that normal DC counts were detected in patients grafted with CD34<sup>+</sup>-nominated cells, which concluded that CD34<sup>+</sup>-nominated cells had the capacity to repopulate the DC line.<sup>18</sup> Interestingly, this study reported no effect of donor age or sex on CMRF44<sup>+</sup> DC, DC1, or DC2 cell numbers or subsets in grafts.

Our results showed that in recipients with constant allogeneic retrieval, recipients received harvests of lower counts (CMRF44<sup>+</sup> DCs, DC1, and DC2) and had lower incidence of aGVHD and relapse and better survival. Moreover, graft-DC count was found to be an independent effector of severity of aGVHD and fatality rate. Those findings were in

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	Sex	Underlying Diagnosis	Graft-DC components (X10 <sup>6</sup> DC/L blood)			Day engraft	of ment	aGVHD stage		aGVHD grade	aGVHD medication	Response	Outcomes	
			CMR44 <sup>+</sup> DCs	DCI	DC2	Neut	Plt	Skin	Liver	Gut				
Case 19	Μ	SAA	11.7	5.6	6.0	17	18	+	-	+	Ι	Topical	CR	Recovery
Case 20	Μ	SAA	*24.5	*19.6	5.3	27	27	++	+	++	II	S-steroids	CR	Recovery
Case 21	F	HDL	13.1	6.0	7.7	12	12	+	-	-	I	Observation	CR	Recovery
Case 22	F	SAA	*28.0	*17.8	8.8	37	47	+++	++	+++	Ш	S-steroid+	CR	Recovery
Case 23	F	AML (M4)	*31.9	*21.2	10.3	43	0	+++	++++	++++	IV	S-steroid+	Progression	Died
Case 24	Μ	ALL (c-ALL)	*28.0	*23.6	3.9	22	25	++	++	+	II	S-steroid+	PR	Died
Case 25	Μ	ALL (PRO-B)	*18.3	11.2	6.3	13	20	++	+	++	Ш	S-steroids	CR	Recovery
Case 26	М	ALL (T-cell)	15.8	7.6	9.5	16	13	+	_	+	I	Topical	CR	Recovery
Case 27	F	AML (MI)	14.0	8.7	4.7	14	16	+	+	-	I	Topical	CR	Recovery

Table 2 GVHD stage and grade, graft-DC components, medications, and outcomes of those who developed aGVHD

Notes: \*P<0.05. S-steroids: systemic steroids (methylprednisolone) + original immunosuppressive prophylaxis (cyclosporine or tacrolimus). S-steroid+: systemic steroids with additional agent (such as MMF/etanercept/other). Abbreviations: SAA, severe aplastic anaemia; HDL, Hodgkin's lymphoma; AML, acute myeloid leukemia; ALL, acute lymphoblastic leukemia; ALL (c-ALL), common-B ALL; DC, dendritic cell; Plt, platelet count >20×10<sup>9</sup>/L; Neut, neutrophil count >0.5×10<sup>9</sup>/L; aGVHD, acute graft-versus-host disease; PR, partial response; CR, complete response.



Figure 2 Comparisons between variables of graft components in group II (cases with acute GVHD) versus group I (event-free group). Total leukocyte count (**A**), volume of PBSC (**B**), total nucleated cells (**C**), percentage of gated cells (**D**), percentage (**E**) and count (**F**) of CD34<sup>+</sup> cells, CMRF44<sup>+</sup> DC count (**G**), DC1 count (**H**), DC2 count (**I**), and recipient weight (**J**) between group I and group II. (**K**) Comparison of CMRF44D<sup>+</sup> DC and DC1 counts among patients with different grades of aGVHD. (**L**) Flow-cytometry analysis of DC counts in PBSC harvests for recipients (n=23) who suffered grade IV aGVHD after allogeneic PBSCT: right, increased DC counts represented by CMRF44<sup>+</sup> DC FITC; left, increased DC1 counts represented by CD11c<sup>+</sup> DC PE. **Notes**: \*\*\*P<0.001: \*\*\*P<0.01.

Abbreviations: TLC, total leukocyte count; TNC, total nucleated cells; DC, dendritic cell; aGVHD, acute graft-versus-host disease; PBSC, peripheral blood stem cell; NS, not significant; FITC, fluorescein isothiocyanate; PE, phycoerythrin.

accordance with Rajasekar et al, who concluded that a significant increase in DCs, particularly DC1, was linked with higher incidence of aGVHD.<sup>19–21</sup> On the other hand, the current findings contradict with Reddy et al.<sup>22</sup>

This study proved that in allogeneic PBSCT recipients of grafts with high DC1 or DC2 counts are linked to a higher incidence of aGVHD or relapse, respectively. This suggests that assessment of DC numbers in the grafts could help identify patients who are at greater risk of aGVHD and losses, as well as allowing for a future evaluation plan and potential early therapeutic strategies or the development of an aGVHD-control strategy. These results support the findings of Waller et al, who found that the amount of donor DC2 was linked to a reduced graft-versus-leukemia effect and increased incidence of relapse following PBSCT, denoting a pivotal role for donor DCs in influencing immunoreactions after allogeneic PBSCT.<sup>19</sup> In a previous study, recipients with fewer DCs following HSCT had a greater risk of



Figure 3 (A–C) Comparisons between variables of graft components in group III (cases with relapse) versus group I (event-free group). CMRF44<sup>+</sup> DC (A), DC1 (B), and DC2 counts (C) in group I and group III. (D and E) Flow-cytometry analysis of DC counts in PBSC harvests for recipients (n=28) who suffered relapse after allogeneic PBSCT: (D) increased DCs represented by CMRF44<sup>+</sup> DC FITC; (E) increased DC2 counts represented by CD4<sup>bright</sup> DC FITC. (F–I) Correlations of CD34<sup>+</sup>-cell and DC counts and days of hospitalization after PBSCT: CD34<sup>+</sup> cell count (F), CMRF44<sup>+</sup> DC count (G), DC1 count (H), DC2 count (I), and days of hospitalization after PBSCT. CD34<sup>+</sup> cell (J) and CMRF44<sup>+</sup> DC counts (K) in group of patients without events and complications group. (L) Percentage of survivors among group I, II, and III subcohorts. Notes: \*\*\*P<0.001; \*\*P<0.01; \*\*P<0.05.

Abbreviations: DC, dendritic cell; PBSCT, peripheral blood stem-cell transplantation; NS, not significant; FITC, fluorescein isothiocyanate; PE, phycoerythrin.

aGVHD and disease relapse, despite the fact that their original grafts had more DCs than grafts administered to patients without aGVHD.<sup>20</sup> Our results are consistent with earlier research showing that high CD4<sup>bright</sup> DC numbers in the marrow of donors<sup>12</sup> or high pDC content in the PBSC graft<sup>21</sup> are linked to higher relapse rate.

Fast and robust engraftment is the instantaneous apprehension after transplantation. It was found that higher quantities of CD34<sup>+</sup> are constantly linked with swift engraftment; on the opponent little doses are recognized to yield late engraftment. A significant clarification can be obtained from the ISHAGE study. It showed that recognized swift engraftment occurred in all recipients who received  $>10-15\times10^6$ /kg total CD34<sup>+</sup> and sluggish engraftment in those received a dose of  $<2\times10^6$ /kg body weight.<sup>22</sup> The current study reported, delayed myeloid engraftment in recipients of

	Group I (n=18)	Group II (n=9)	Р
Prepharesis TLC (×10 <sup>9</sup> /L)			
•Mean + SD	48.9±15.2	51.9±18.1	0.681
●95% CI	41.3-56.5	37.9–65.8	
●Median	47.6	50.4	
Volume (mL <sup>3</sup> )			
•Mean + SD	192.9±36.7	207±29.8	0.280
•95% CI	174.7-211.2	184.3-230.1	0.200
•Median	195	200	
Total nucleated cells (X10 <sup>6</sup> /I)			
•Moan + SD	411 5+206 8	459 3+233 9	0 777
•95% CI	308 7 514 4	2795 6 397	0.777
•Median	385	400	
	505	400	
Weight (kg)			
●Mean + SD	55.9±22.4	53.4±20.4	0.662
●95% CI	44.9–67.0	37.8–69.1	
●Median	61.5	56	
Gated cells (%)			
•Mean + SD	30.3±10.8	24.6±11.8	0.111
●95% CI	24.9-35.6	24.6-33.7	
●Median	34.1	28	
CD34 cells (%)			
•Mean + SD	2.5±1.3	2.5±1.6	0.837
●95% CI	1.8-3.2	1.3–3.8	
●Median	2.2	2.2	
•Mean + SD	145+115	92+73	0117
•95% CI	88-202	36-148	0.117
•Median	9.6	4.6	
	7.0	1.0	
DC count (absolute)			
●Mean + SD (×10 <sup>6</sup> /L)	8.6±3.7	20.6±7.6	<0.001***
●95% CI	6.8–10.5	14.6–26.4	
●Median	9.9	18.3	
DCI count (absolute)			
•Mean + SD (×10 <sup>6</sup> /L)	4.6±2.4	13.5±7.1	<0.001***
●95% CI	3.4–5.8	8.1–18.9	
●Median	4.1	11.2	
DC2 count (absolute)			
•Mean + SD (×10 <sup>6</sup> /L)	3.5±3.3	6.9±2.2	0.009**
•95% CI	1.9–5.2	5.2-8.7	
●Median	3.2	6.3	
		1	

 Table 3 Comparison between different variables of graft components in group II (cases with acute GVHD) versus group I (event-free group)

**Notes**: \*\*\*P<0.001; \*\*P<0.01.

Abbreviations: TLC, total leukocyte count; DC, dendritic cell.

grafts with high DC counts, without significant effect on platelet engraftment. In the current study recipients of allografts with higher DC counts had longer in hospital days, post PBSCT, on the contrary higher CD34+ cells was associated with shorter in hospital stay.

	Time to engra	lftment (days)		Р	
	Group I	Group II	aGVHD	DCs	DCI
	Mean ± SD	Mean ± SD			
Myeloid <u>(</u> ≥0.5×10 <sup>9</sup> /L)	13.94±1.12	21.88±3.7	0.014*	0.02*	0.04*
Platelets (≥ 20×10 <sup>9</sup> /L)	4.38±1.13	17.55±4.8	0.4	0.88	0.66

**Table 4** Associations between DC numbers and day of myeloid and platelet engraftments in allogeneic

 PBSCT recipients

Note: \*P<0.05.

Abbreviations: DCs, dendritic cells; aGVHD, acute graft-versus-host disease.

This study showed that recipients of allografts with higher CD34+ cells had lower incidence of complications and prolonged EFS. This was in accordance with others who concluded that patients of greater number of CD34<sup>+</sup> cells had longer EFS; whereas those with less CD34<sup>+</sup> cells had overdue hematopoietic engraftment and augmented disease from infection.<sup>12,22</sup> The findings of this study provided strong scientific evidence that donor DCs could have a role in pathogenesis of aGVHD, and the applicability of this in advances in SC therapy.

Now the question is how donor DCs participate in aGVHD. The answer could be extracted from the current study and previous studies. DCs could eventually present antigens in either a tolerogenic or immunogenic way based on the degree of their maturation, which in turn is dependent on the circumstances in which they faced antigens. Therefore, DCs either generate a suitable immunoreaction to pathogens or avert autoimmunity.<sup>23</sup> The net influence of DCs in modifying immunoreconstruction after transplantation possibly signifies the opposing effects of graft- donor DC1 versus DC2 and CD34<sup>+</sup>-differentiated donor DCs cells that are augmented by DC1 or DC2 cytokines.<sup>5</sup>

	0										
25 20- 15- 10- 5-	Graft Components	Preph. TLC (X10 <sup>9</sup> /L	Vol. (ml <sup>3</sup> )	TNC (X10%/L)	Weight (kg)	GC %	CD34 %	CMRF-44+DC (X106/L)	DC-1(X10 <sup>6</sup> /L)	DC-2 (X10 <sup>6</sup> /L)	1.
0- Group I Group II Group III	Preph. TLC (X10%/L)	1.000	0.207 N.S.	0.251 N.S	-0.068 N.S	-0.143 N.S.	0.043 N.S.	0.227 N.S.	0.470 *	-0.239 N.S.	0
14	Vol. (ml <sup>3</sup> )	0.207 N.S.	1.000	0.079 N.S	0.640 ***	-0.563 **	0.196 N.S.	0.103 N.S.	0.150 N.S.	-0.053 N.S.	
12-	TNC (X10 <sup>6</sup> /L)	0.251 N.S	0.079 N.S	1.000	0.098 N.S.	0.121 N.S.	0.118 N.S.	0.128 N.S	0.117 N.S.	0.113 N.S.	
8-	Weight (kg)	-0.068 N.S	0.640 ***	0.098 N.S.	1.00	-0.310 NS	0.086 N.S.	-0.015 N.S.	0.002 N.S.	-0.041 N.S.	0
6- 4-	GC %	-0.143 N.S.	-0.563 **	0.121 N.S.	-0.310 N.S.	1.00	-0.394 *	-0.074 N.S.	- 0.140 N.S.	0.074 N.S.	
Group I Group II Group III	CD34 %	0.043 N.S.	0.196 N.S.	0.118 N.S.	0.086 N.S.	-0.394 *	1.000	-0.264 N.S.	-0.192 N.S.	-0.266 N.S.	
15-	CMRF-44+DC (X10 <sup>6</sup> /L)	0.227 N.S.	-0.103 N.S.	0.128 N.S	-0.015 N.S.	-0.074 N.S.	-0.293 N.S.	1.000	0.840 ***	0.640 ***	-0
10-	DC-1 (X10 <sup>6</sup> /L)	0.417 *	-0.150 N.S.	0.117 N.S.	0.002 N.S.	-0.147 N.S.	-0.192 N.S.	0.810 ***	1.00	0.171 N.S.	
5	DC-2 (X10 <sup>6</sup> /L)	-0.239 N.S.	-0.053 N.S.	0.113 N.S.	-0.047 N.S.	0.089 N.S.	-0.254 N.S.	0.640 ***	0.171 N.S.	1.00	-1
	25 20 15 10 5 0 Group I Group II Group III 4 20 6 4 Group I Group II Group III 20 5 6	25 20 15 10 5 0 Group I Group II Group III 14 12 14 12 6 4 Group I Group II Group III 14 12 14 12 10 8 6 4 Group I Group II Group III 14 12 10 10 10 10 10 10 10 10 10 10	25 20 15 10 5 0 Group I Group II Group II 4 20 6 4 Group I Group II Group II Croup I Group II Group III 20 5 10 5 10 5 10 10 10 10 10 10 10 10 10 10	25       20       100       100	25       0       Graft       0 <td>25       0</td> <td>25       0</td> <td>25       0</td> <td>25       20       30       <td< td=""><td>20       0</td><td>20       30       Graft       1000       1000</td></td<></td>	25       0	25       0	25       0	25       20       30 <td< td=""><td>20       0</td><td>20       30       Graft       1000       1000</td></td<>	20       0	20       30       Graft       1000       1000

Figure 4 (A) Estimated marginal means of DCs, DCI, and DC2. (B) Matrix of correlations showing associations between variables of graft components and patient weight. Values in cells are Spearman correlation coefficients.

**Notes**: \*\*\*P<0.001; \*\*P<0.01; \*P<0.05.

Abbreviations: Preph. TLC, peripheral total leukocyte count; Vol, volume of the graft; TNC, total nucleated cells; GC, gated cells; DC, dendritic cell; NS, not significant.

The role of APCs (DCs) in pathogenic donor T-cell development and differentiation is beginning to emerge, although the identification of DC types responsible for this abnormal T-cell differentiation remains puzzling. Gonçalves et al reported that greater pDC levels in the graft were linked to a higher risk of aGVHD, but pDC counts in the PB following transplantation had the reverse impact.<sup>24</sup> Furthermore, Lee et al demonstrated that transplanting bone marrow deficient in MyD88 led in a slanted DC differentiation toward CD11c<sup>+</sup> DCs that encompassed a majority of graft cells in GVHD hosts and resulted in GVHD and deterioration.<sup>25</sup>

It has been stated that G-CSF has a starring role in the differentiation of DCs. G-CSF injections of donors with 10–16  $\mu$ g/kg/day for 5 days amplified PB DC2 numbers, whereas those of DC1 did not changed. G-CSF might increase DC2 numbers in the circulation by enhancing their production in the bone marrow, prolonging their survival, persuading mobilization, or reducing their movement out of the vascular enrollment into lymphoid organs.<sup>26</sup> It is plausible that aPBSCT of G-CSF–mobilized PBSC does not result in devastating aGVHD, as the graft has mostly T<sub>h</sub>2-persuading DCs. Adoptive transmission of filtered DCs may bring immunoaberration after transplantation of HSCs or organ allografts. The significance of such reports has been highlighted by recent findings on progression of therapeutic approaches against cancers and transmittable agents using DCs. Some studies have proved that this method holds potential in preventing cancer progression or even causing cancer regression.<sup>25,26</sup>

### Conclusion

This study aimed to monitor DC numbers and subsets during SC harvesting in PBSCT and their possible roles in prediction of incidence of aGVHD after PBSCT, represented by increased DC1 numbers in the grafts (harvests), and testing the hypothesis that larger doses of CD4<sup>bright</sup> DC (DC2) in PBSC grafts are associated with higher relapse rates after PBSCT. The results revealed that donor progenitor DCs, particularly DC1, can have a significant impact on engraftment associated with an increased incidence of aGVHD, while higher DC2 was linked to disease relapse. These results provide strong evidence that DCs can have an important role in the pathogenesis of aGVHD. Furthermore, we speculated on graft-DC monitoring as a predictive biomarker of aGVHD and/or relapse. These findings together with others suggest it is necessary to create a technique for accurate DC monitoring that can measure their numbers and subsets. We established a new technique that might facilitate quantitative measurements of DCs and subsets, and accordingly could simplify its usage as a biomarker in PBSCT allografts (harvests). The technique relies on exploiting modern flow-cytometry procedures and instruments using CD4 FITC/CD34 PE, CD34 FITC/CD11c PE, and CMRF44 FITC antibody panels. The specificity of the technique was primarily reliant on the specificity of the anti-DC antibody, whereas precision was founded on flow-cytometry capability in providing exact measurements.

# **Data Sharing**

All data connected with and supporting the study results are within the manuscript and the Supplementary File.

# Ethics

The research protocol, objectives, and methods were in accordance with the WMA Declaration of Helsinki. The ethics committee of the National Cancer Institute (NCI), Cairo University, Egypt granted approval for this study. The study had no impact on patients' treatment, and no extra procedures were undertaken.

# **Patient Consent**

Before inclusion in the study, written informed consent was obtained from all PBSCT recipients and their matched sibling donors who participated in the study voluntarily. Objectives, possible risks, and withdrawal rights were clarified to all participants before the study. Anonymity was addressed.

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# **Author Contributions**

All authors made a significant contribution to the work reported, whether in conception, study design, execution, acquisition of data, analysis and interpretation, or all these areas, took part in drafting, revising, or critically reviewing the article, gave final approval to the version to be published, have agreed on the journal to which the article has been submitted, and agree to be accountable for all aspects of the work.

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### Disclosure

The authors declare no conflicts of interest in relation to this work.

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