Delayed immune reconstitution after cord blood transplantation is characterized by impaired thymopoiesis and late memory T-cell skewing

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Advances in immune assessment, including the development of T-cell receptor excision circle (TREC) assays of thymopoiesis, cytokine-flow cytometry assays of T-cell function, and higher-order phenotyping of T-cell maturation subsets have improved our understanding of T-cell homeostasis. Limited data exist using these methods to characterize immune recovery in adult cord blood (CB) transplant recipients, in whom infection is a leading cause of mortality. We now report the results of a single-center prospective study of T-cell immune recovery after cord blood transplantation (CBT) in a predominantly adult population. Our primary findings include the following: (1) Prolonged T lymphopenia and compensatory expansion of B and natural killer (NK) cells was evident; (2) CB transplant recipients had impaired functional recovery, although we did observe post-transplantation de novo T-cell responses to cytomegalovirus (CMV) in a subset of patients; (3) Thymopoietic failure characterized post-CBT immune reconstitution, in marked contrast to results in other transplant recipients; and (4) Thymopoietic failure was associated with late memory T-cell skewing. Our data suggest that efforts to improve outcomes in adult CB transplant recipients should be aimed at optimizing T-cell immune recovery. Strategies that improve the engraftment of lymphoid precursors, protect the thymus during pretransplant conditioning, and/or augment the recovery of thymopoiesis may improve outcomes after CBT.

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Introduction

Umbilical cord blood (CB), first demonstrated to have clinical utility by Gluckman et al as a source of hematopoietic stem cells in the setting of Fanconi anemia,1 was later demonstrated to have utility as a source of unrelated donor stem cells for patients lacking matched-sibling donors.2-5 Over the past decade, a large number of studies have demonstrated the clinical utility of CB transplantation (CBT) as a treatment for both malignant and nonmalignant diseases of children and adults.4,6 The establishment of international cord blood banks, advances in supportive care and donor graft selection, and novel clinical approaches aimed at improving engraftment (eg, ex vivo expansion of CB-derived progenitors7,8 and the infusion of pooled unrelated units9) have improved outcomes and led to a dramatic increase in the number of CBTs performed worldwide. CB grafts obtained from matched unrelated donors offer advantages over bone marrow or peripheral blood stem cells (PBSC) such as noninvasive procurement, more rapid availability without the need for the more prolonged process of screening and obtaining stem cells from a matched unrelated donor (MUD), and the apparently greater tolerance for incompletely human leukocyte antigen (HLA)-matched products.10 These advantages are paramount for recipients in historically underrepresented minority groups, for whom the prospect of locating a MUD registry donor remains relatively diminished. At our institution, more than twice the proportion of CB transplant recipients are minorities relative to MUD marrow or PBSC recipients historically undergoing transplantsations. This fact underscores the importance of improving our current approaches to alternative donor transplantation for patients lacking matched-sibling or MUD donors. For these reasons, it remains important for us to clearly define the biologic variables that govern posttransplantation outcomes in these patients.

Of all the clinical challenges that face CBT clinicians, delayed immune reconstitution remains one of the most important causes of morbidity and mortality11-15 (also reviewed in Szabolcs and Niedzwiecki16). Although it is increasingly appreciated that a variety of circulating peripheral blood cell subpopulations may contribute to immune integrity, including B cells, natural killer (NK) cells, peripheral blood monocytes, and dendritic cells, it is also generally accepted that adaptive immune responses mediated by T cells are essential for protective immunity. As is perhaps best illustrated by the HIV/AIDS pandemic, the selective loss of CD4+ T cells is sufficient to trigger profound immunodeficiency that often leads to fatal infection.17 The primary consequence of the loss of CD4+ T-cell help is an attendant loss in the number and/or function of antigen-specific CD8+ T cells, which constitute our primary adaptive response to pathogens, including viruses and fungi.18 A nearly universal characteristic of conditioning regimens used to prepare recipients of unrelated donor grafts is the use of chemotherapeutic agents and/or antibodies that effectively deplete the host of mature T cells. In the setting of CBT, multiple


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chemotherapy drugs are typically combined with polyclonal antithymocyte globulin to decrease the likelihood of donor graft rejection mediated by surviving host T cells. In this setting, T-cell reconstitution after CBT inherently depends on the survival of adoptively transferred T cells from the CB graft or, alternately, the de novo production of T cells in the recipient thymus. Although extrathymic production of T cells has been postulated, no conclusive evidence exists that suggests extrathymic maturation contributes significantly to de novo T-cell production in human stem cell transplantation (SCT) recipients. Furthermore, CB grafts differ from T-cell–replete PBSC grafts in that they contain fewer T cells that are also uniformly naive (eg, antigen-inexperienced).

Here we report the results of a prospective study of T-cell immune reconstitution in recipients of unrelated CB grafts. We conducted a quantitative analysis of T-cell subsets using immunophenotyping and also performed detailed analyses of superantigen-stimulated and virus-specific T cells using cytokine flow cytometry. We also analyzed the recovery of thymopoiesis using a polymerase chain reaction (PCR)–based assessment of T-cell receptor excision circles (TRECs) in CB transplant recipients. Our results suggest that in our studied population, inadequate thymic regeneration after CBT was associated with lymphopenia, delayed functional immune recovery, and skewing of the T-cell compartment away from naive and early memory T cells.

**Patients and methods**

**Patient selection and GVHD management**

A total of 32 patients underwent CBT at the M. D. Anderson Cancer Center were studied. Informed consent was obtained from all patients in accordance with the Declaration of Helsinki for a correlative laboratory study of immune reconstitution that was approved by the M. D. Anderson Cancer Center Institutional Review Board. All patients received graft-versus-host disease (GVHD) prophylaxis consisting of tacrolimus and low-dose methotrexate (5 mg/m² on post-CBT days 1, 3, and 6), in addition to thymoglobulin administered during conditioning. In the event of confirmed or suspected GVHD, initial therapy consisted of methylprednisolone (2 mg/kg per day, with a taper based on clinical response). In the absence of GVHD, tacrolimus was tapered after post-CBT day 100.

**Immunophenotyping**

One million PBMC per sample were stained with monoclonal antibodies against CD4, CD8, CD45RA, and CCR7. Cells were stained in a total of 50 μL staining buffer (PBS + 0.1% BSA + 0.02% sodium azide) and antibodies. Staining was carried out at room temperature for 15 minutes. Cells were then washed with 2 to 3 mL PBS, centrifuged, and resuspended in 200 μL of 1% paraformaldehyde. Cells were acquired on a Cyan flow cytometer (Dako, Fort Collins, CO), and the resulting data were analyzed with FlowJo software (Tree Star, Ashland, OR).

**Assessment of functional T-cell recovery using CFC**

One million PBMC per sample were cocultivated with staphylococcal enterotoxin B (SEB; Sigma, St Louis, MO) and, in cytomegalovirus (CMV)–seropositive patients, a pool of pentadecapeptides spanning CMV pp65 (BD Pharmingen, San Jose, CA) in 96-well V-bottom tissue culture plates in 200 μL of media (RPMI1640; GIBCO Life Technologies, Grand Island, NY) supplemented with 10% fetal calf serum, penicillin, and streptomycin (Sigma). After 1 hour of incubation at 37°C, 5% CO₂, brefeldin A (Sigma) was added at a final concentration of 10μg/mL to enable accumulation of effector cytokines in the cytoplasm. The cells were then incubated an additional 5 hours at 37°C, 5% CO₂, after which time the cells were moved to 4°C overnight. The cells then underwent lysis and permeabilization with FACSLyse and FACSPerm Solution II, respectively (BD Biosciences), the following day. Cytokine flow cytometry (CFC) analyses were performed using fluorescein isothiocyanate-, phycoerythrin-, PerCP-, and allophycocyanin-conjugated monoclonal antibodies (MAB) specific for human CD4, CD8, CD69, and interferon (IFN)γ (BD Biosciences). After staining at 4°C for 20 to 25 minutes, cells were washed, resuspended in PBS with 1% paraformaldehyde, and acquired by 4-color flow cytometry on a FACSCalibur cytometer (BD Biosciences) and analyzed with FlowJo software (Tree Star). For most analyses, at least 30 000 total events were analyzed. Non-specific activation of cells was tested by incubation of paired samples that were unstimulated.

**Quantitative real-time PCR for TREC**

**Cell preparation.** One million snap-frozen PBMC were lysed overnight (up to 18 hours) at 56°C in lysis buffer (LB) (0.5% Nonidet P-40, 0.1% Tween-20, and 200 μg/mL of molecular grade [nuclelease-free] Proteinase K in a standard 1× PCR Buffer containing no magnesium [Invitrogen, Carlsbad, CA]). The samples were then heat inactivated at 95°C for 10 minutes. DNA was purified from cell lysates by adding a 1× volume of phenol-chloroform-isoamyl alcohol (25:24:1; Sigma) for extraction using Phase-Lock light gel separator tubes (Eppendorf, Westbury, NY). The aqueous phase was then transferred to a fresh microfuge tube, and ethanol was precipitated using a 0.2× volume of 7.5 M ammonium acetate and a 2× volume of cold 100% ethanol, placement at −20°C for 1 hour, and then spinning at 16 000g for 15 minutes. The supernatant was aspirated and the DNA pellets were dried down and reconstituted in 35 μL of Tris-EDTA (TE) buffer and stored at −20°C until use in the real-time PCR TREC assay.

**Real-time PCR.** Thymic function was assessed using the method of Harris et al.23 Delta-deletion TRECs were amplified and quantified in a Biorad iCycler iQ Real-Time Detection System (BioRad Laboratories, Hercules, CA) using fluorescently labeled oligonucleotides as reporter probes in a 50 μL PCR reaction using 2X iQ Supermix (with additional MgCl₂) to a final 3.5 mM concentration) (Biorad). Primers for the TREC sequence were 5′-CCC TTA CCA CCA TGC TCA CAC-3′ (forward) and 5′-GGG TGC AGG TGC TCA TGA-3′ (reverse), which produced an approximately 80-base pair product detected with the probe 5′-FAM-TCT GGT TTT TGT AAA GGT GCC CAC TCC TG-3′. TREC abundance was normalized to input cell number by a parallel amplification for the β-globin gene. Human β-globin primers were 5′-GAA GAG CCA AGG ACA GGT AGC-3′ (forward) and 5′-CTT GGG ATG AGT TGG GCC AA-3′ (reverse), which produced an 85-base pair product detected with the probe 5′-FAM-CTG TCA TCT AGA CCT CAC CCT GTG-3′. Primers (all from Sigma-Genosys, St Louis, MO) were used at 10 pmol per reaction well, and probes (both from Biosearch Technologies, Novato, CA) were used at 5 pmol per reaction well. Typically, 45 μL of the master mix (appropriate containing either TREC or globin primers and probe) was first added to the 96-well optical grade plate, and then 5 μL per well of standards, controls, and samples were pipetted into the plate and optical-grade tape was applied. All standards were run in duplicate, and all samples were run in triplicate.

**Statistical analyses**

All statistical analyses were performed using Prism software (GraphPad Software, San Diego, CA). Intergroup comparisons were assessed using the Mann-Whitney U test, with significance defined by P less than .05 using a 2-tailed test. Differences in actuarial survival were determined by the method of Kaplan and Meier.24

**Results**

**Clinical characteristics**

A total of 32 CB transplant recipients who underwent transplantation during the study period were assessed, with a median follow-up of 221 days (Table 1).
Clinical outcomes

The majority of patients experienced initial engraftment (28/32), whereas 6 subjects (19%) experienced either primary or secondary graft failure. Neutrophil engraftment occurred at a median of 22.5 days. GVHD occurred after CBT in 20/32 subjects (62.5%). Of the 20 subjects experiencing GVHD, 13 subjects (65%) experienced acute GVHD; in all but 1 of these subjects this was greater than grade 2. Twelve of 20 patients with GVHD experienced chronic GVHD (60%); 7 of these subjects experienced de novo chronic GVHD without antecedent acute GVHD. Five of 12 subjects developed chronic GVHD that was classified as extensive, whereas the remainder had limited disease. Relapse occurred in 15/32 subjects (46%) and was a major cause of death in 12 subjects (37%). Infection was a major cause of death in 10 subjects (31%); in 4 of these subjects, fatal infections occurred in the absence of relapse or GVHD whereas another of these factors contributed to death in the remaining subjects (3 deaths each due to relapse/infection and GVHD/infection). In the 10 patients for whom infection was a major cause of death, the primary infections included the following: suspected or documented fungal pneumonia (4 subjects), presumed infectious pneumonitis without a documented pathogen (4 subjects), CMV colitis with gastrointestinal hemorrhage (1 subject), and polymicrobial infections including herpes simplex virus esophagitis, CMV reactivation, and fungal pneumonia (1 subject).

Persistent lymphopenia and compensatory expansion of NK cells and B cells after CBT

Quantification of total numbers of CD4+ T cells, CD8+ T cells, CD16/CD56+ NK cells, and CD19+ B cells was prospectively performed on fresh PBMC samples collected after CBT (Figure 1). Compared with previously published normal values for healthy stem cell transplant donors, baseline CD4+ and CD8+ T-cell counts in this group of CB transplant recipients were reduced; furthermore, the typical ratio of CD4/CD8 T cells was inverted. Of note, the absolute number of B cells at baseline was not reduced relative to that expected for healthy subjects, whereas NK-cell numbers were only slightly below expected values. By post-CBT day 30, CD4+ and CD8+ T-cell numbers plummeted and remained extremely low for at least 6 months after CBT. After this interval, increases in absolute CD4+ and CD8+ T-cell counts were observed in surviving subjects; by 1 year after CBT, the CD4/CD8 T-cell ratio was clearly positive, and absolute CD8+ T-cell numbers plummed and remained extremely low for at least 6 months after CBT. After this interval, increases in absolute CD4+ and CD8+ T-cell counts were observed in surviving subjects; by 1 year after CBT, the CD4/CD8 T-cell ratio was clearly positive, and absolute CD8+ T-cell counts approached normal values (as shown for healthy donors previously studied by Storek et al23). In contrast, absolute numbers of NK cells immediately increased over baseline values and remained increased over expected values in control subjects throughout the study interval, though their numbers declined at 1 year in concert with substantial increases in CD4+ and CD8+ T-cell numbers. B cells, like CD4+ and CD8+ T cells, declined after conditioning and were significantly reduced, relative to that seen at baseline and in healthy controls, by post-CBT day 30. Thereafter, absolute B-cell numbers expanded dramatically, though they also declined slightly by the 1-year time point, at which T-cell counts rebounded. Although these results confirm that quantitative T-cell recovery is delayed...
after CBT, they also suggest that compensatory expansion of non-T lymphoid cells (eg, NK cells and B cells) may accompany T lymphopenia.

**Impaired functional T-cell responses after CBT**

We previously demonstrated that cytokine flow cytometry is a valid measure of T-cell immune reconstitution and that preserved (or reconstituted) responses by CFC are associated with decreased risk of viral reactivation. To examine the extent of functional immune recovery after CBT, we stimulated PBMC from CB transplant recipients with the superantigen SEB and, in the case of recipients at risk for CMV reactivation, with a mixture of pentadecapeptides spanning the dominant CMV capsid antigen pp65. We then assessed the fraction of CD4+ and CD8+ T cells capable of responding functionally, as assessed by intracellular measurement of IFNγ production induced by superantigen or CMV stimulation. Responses in CB transplant recipients were compared with those of other allogeneic SCT recipients assessed at the same time interval (100 days after SC transplantation) using the same assay (Figure 2). Our results demonstrate that the fraction of both CD4+ and CD8+ T cells capable of producing IFNγ after superantigen stimulation was significantly reduced in CB transplant recipients, relative to other SCT patients. Similarly, the proportion of CMV-specific CD4+ T cells that were functionally responsive in the CFC assay was also dramatically reduced relative to other SCT recipients. Although CD8+ CMV-specific T cells were also significantly lower than those of other SCT recipients, the average frequency of CMV-specific CD8+ T cells was greater than 1%, approaching that expected in healthy CMV-seropositive subjects. Overall, a significant proportion of CMV-seropositive recipients assessed by day 100 were found to have circulating CMV-specific CD4+ or CD8+ T cells (4 of 11 with positive CMV-specific CD4+ T-cell responses, 44%; and 6 of 12 with positive CD8+ CMV-specific T cells, 50%) proving definitively that naïve cord blood cells, transferred into CBT recipients, may be effectively initiate a primary immune response leading to CMV-specific memory T-cell generation in the early posttransplantation interval.

**Thymic regenerative failure after CBT**

It has been clearly demonstrated that thymopoiesis persists to a variable extent throughout adulthood and that the recovery of thymopoiesis after autologous and allogeneic SC transplantation is probably an important determinant of naive T-cell recovery and T-cell receptor diversity. In prior cross-sectional studies of a wide range of allogeneic SCT recipients receiving both matched-related PBSC grafts as well as matched-unrelated donor marrow grafts, we found that nearly all recipients had detectable thymic function, albeit to a varying extent, even at early time points (eg, within the first 60 days) after allogeneic SC transplantation (L.S.J., K.V.K., unpublished data, June 2002). Although it is conceivable that graft- or conditioning-related factors might influence the degree of thymopoiesis observed after CBT, we initially expected that the relatively younger age of CB transplant recipients in this prospective study (33 years, Table 1) would be associated with relatively robust recovery of thymopoiesis. We expected that this would offset, at least in part, the relatively reduced number of graft T cells, and the inherent lack of memory T cells, transferred from donors in CB transplant recipients versus allogeneic SCT recipients receiving PBSC grafts.

To assess the contribution of thymopoiesis to T-cell immune recovery after CBT, we measured peripheral recent thymic emigrants (TRE) by determining the number of TRECs generated to a variable extent throughout adulthood and that the recovery of thymopoiesis after autologous and allogeneic SC transplantation is probably an important determinant of naive T-cell recovery and T-cell receptor diversity. In prior cross-sectional studies of a wide range of allogeneic SCT recipients receiving both matched-related PBSC grafts as well as matched-unrelated donor marrow grafts, we found that nearly all recipients had detectable thymic function, albeit to a varying extent, even at early time points (eg, within the first 60 days) after allogeneic SC transplantation (L.S.J., K.V.K., unpublished data, June 2002). Although it is conceivable that graft- or conditioning-related factors might influence the degree of thymopoiesis observed after CBT, we initially expected that the relatively younger age of CB transplant recipients in this prospective study (33 years, Table 1) would be associated with relatively robust recovery of thymopoiesis. We expected that this would offset, at least in part, the relatively reduced number of graft T cells, and the inherent lack of memory T cells, transferred from donors in CB transplant recipients versus allogeneic SCT recipients receiving PBSC grafts.
transplantation at the University of Colorado. Individual longitudinal results for all of the CB transplant recipients and median values of TREC levels for all 3 groups are shown (Figure 2A,B, respectively). In addition, we examined a group of healthy donors, matched in age to the CB transplant recipients (n = 18, n = 34) as a reference for the normal values expected in a healthy control population. Our results demonstrate that our heavily pretreated CBT population had markedly impaired thymic function even before transplantation, with only 6 of 21 subjects with samples collected before transplantation having detectable thymopoiesis at baseline. Median baseline TREC values in the 6 subjects with detectable pretransplantation thymopoiesis (1095 TRECs/10⁶ PBMC) were significantly lower than those in healthy donors (7357 TRECs/10⁶ PBMC), suggesting that even those patients had impaired T-cell regenerative potential (Figure 3).

After transplantation, thymopoiesis was undetectable in most CB transplant recipients. Detectable post-CBT thymopoiesis beyond approximately post-SC transplantation day 30 was noted in only 2 subjects; in 1, thymopoiesis recovered and was sustained beyond approximately 8 months after transplantation, and a second subject had only 1 positive post-CBT sample at approximately 6 months posttransplantation. Importantly, average and median TREC values in CB transplant recipients were significantly lower than those seen in autologous and allogeneic SCT recipients at all time points assessed. Notably, the 1 patient with sustained post-CBT thymopoiesis had undetectable TREC.s at baseline, demonstrating no correlation between baseline thymic function and eventual thymic recovery. Because of the nearly universal lack of thymic recovery observed, we were unable to examine donor graft (eg, infused stem-cell dose) or host-related factors governing the recovery of thymopoiesis.

Late memory skewing characterizes T-cell recovery after CBT

T-cell differentiation may be characterized by the expression of surface markers that demarcate naive, early, and late memory T cells. Over the past several years, multiple models, based on the expression of various surface markers (including the reciprocal isoforms CD45RA and CD45RO, the chemokine receptor CCR7, CD27, CD28, and CD57) have been proposed.30-34 Using the combination of CCR7 and CD45RA, it is possible to define subsets of naive cells (CD45RA⁺CCR7⁺), early or central memory cells (CD45RA⁻CCR7⁺), effector memory cells (CD45RA⁻CCR7⁻), and late effector memory cells (CD45RA⁺CCR7⁻).32 At baseline and after transplantation, the CD4⁺ T-cell compartment in CBT transplant recipients was characterized by a relative paucity of naive cells, with large proportions of central and effector memory T cells. The CD8⁺ T-cell compartment was similarly characterized by relatively few naive cells, but with fewer central memory T cells than in the CD4⁺ compartment. The bulk of CD8⁺ T cells expressed markers consistent with effector memory and late effector memory differentiation. To determine how CB transplant recipients compared with subjects not receiving transplants, we assessed the relative frequencies of T-cell differentiation subsets in samples collected from healthy allogeneic SCT donors. Compared with CB transplant recipients (n = 32), healthy donor subjects had significantly greater proportions of CCR7⁺ T cells in the CD4⁺ compartment (wherein CCR7⁺ T cells constituted the majority of CD4⁺ T cells in healthy donors) and in the CD8⁺ compartment (where fewer cells were CCR7⁺ in healthy subjects, though similarly increased, relative to CB transplant recipients; Figure 4).

Preservation of CCR7⁺ CD4⁺ T cells at day 30 predicts post-CBT survival

In an exploratory analysis, we determined whether the skewing that we observed, toward later memory T-cell subsets, was associated adversely with survival after CBT. Actuarial analysis of survival, stratified by the relative preservation of CCR7⁺ T cells within the CD4⁺ compartment, is illustrated in Figure 5. We found that a higher ratio of CCR7⁺ to CCR7⁻ CD4⁺ T cells (ie, above the median vs below) at day 30 was associated with significantly longer survival (median survival 240 days vs 700 days; P = .013). When we analyzed the association between CCR7⁺ T cells within the CD8⁺ T-cell compartment at day 30 and survival, no similar relationship was seen. We also examined whether the degree of pre-CBT thymic function was associated with survival, given the heterogeneity of baseline thymic function that we observed in this...
In contrast to what has been previously reported in other allogeneic SCT recipients, we found that there was no association between pretransplantation thymic function and survival in this group of CB transplant recipients (data not shown). 

Discussion

Although CB is increasingly used as a stem cell source for adults lacking other suitably matched grafts, delayed T-cell reconstitution remains an important source of morbidity and mortality. As our therapeutic approaches to CBT are continually evolving, it remains essential for us to identify areas in which targeted efforts may improve outcomes. For this reason, we conducted a comprehensive analysis of T-cell immune recovery in 32 adult CB transplant recipients consecutively receiving transplants. Our primary findings include the following: (1) In most recipients, we found impaired baseline thymopoiesis and a relatively diminished naive T-cell repertoire before CBT; (2) Prolonged T lymphopenia was evident after CBT; in contrast, relative and absolute expansion of B cells and NK cells was observed; (3) Although impaired functional responses to superantigens and CMV characterized the group overall, a subset of patients had demonstrable recovery of functional CMV-specific T cells by day 100, indicating the ability of naive cord blood T cells to initiate early primary immune responses to pathogens; (4) Thymopoietic failure characterized post-CBT immune reconstitution, in absolute terms and when CB transplant recipients were compared with other SCT recipients examined prospectively and/or in cross-section; and (5) The failure of thymopoiesis was associated with a relative paucity of CCR7+ naive and central memory CD4+ and CD8+ T cells and greater proportions of effector and late-effector memory (ie, CCR7−) T cells.

The fact that T-cell lymphopenia characterized the post-CBT state was not surprising. In contrast, we also found that relative, and even absolute, increases in circulating B-cell and NK-cell numbers was observed; (3) Although impaired functional responses to superantigens and CMV characterized the group overall, a subset of patients had demonstrable recovery of functional CMV-specific T cells by day 100, indicating the ability of naive cord blood T cells to initiate early primary immune responses to pathogens; (4) Thymopoietic failure characterized post-CBT immune reconstitution, in absolute terms and when CB transplant recipients were compared with other SCT recipients examined prospectively and/or in cross-section; and (5) The failure of thymopoiesis was associated with a relative paucity of CCR7+ naive and central memory cells after CBT, with an apparent association between improved posttransplantation outcome and the preservation of CD4+ CCR7+ T cells in the immediate post-CBT interval.

The fact that T-cell lymphopenia characterized the post-CBT state was not surprising. In contrast, we also found that relative, and even absolute, increases in circulating B-cell and NK-cell numbers characterized post-CBT recovery. Previously, increased numbers of circulating NK cells were found in renal allograft recipients,35 suggesting that T lymphopenia may actually facilitate NK-cell expansion. Antigen-independent homeostatic B-cell proliferation has been postulated and has shown to occur independent of T cells.36 Recently, studies of 2 groups of patients with T lymphopenia (HIV-infected subjects and others with idiopathic CD4+ T lymphopenia) have demonstrated the
expansion of immature/transitional B cells. Taken together, these data suggest that there may be crosstalk between the T-cell, B-cell, and NK-cell compartments in the setting of lymphopenia. The factors that govern this intercompartmental regulation (eg, cytokines including IL-7 or competition for “niches”) deserve further study.

We observed profound delays in functional immune reconstitution in CBT when considered as a group. In contrast to other allogeneic SCT recipients, subsets of CD4⁺ and CD8⁺ T cells capable of responding suitably to stimulation with superantigens were markedly reduced. Because the CFC assay measures the proportion of capable responders within a gated T-cell subpopulation, this approach is less vulnerable (eg, relative to conventional lymphoproliferative assays [LPA]) to the effects of lymphopenia, which usually results in diminished LPA responses in patients with CD4⁺ T-cell counts of less than 50 cells/μL. Although it is probable that the transfer of fewer T cells in CBT (vs PBSC grafts) plays a major role in this phenomenon, we cannot discount the effects of nonlymphoid cells (eg, myeloid subsets including monocytes and DC) on these impaired functional responses. When we studied the recovery of CMV-specific CD4⁺ and CD8⁺ T-cell responses by CFC, we found that although responses were reduced in CBT overall, a subset of patients had significant frequencies of functional CMV-specific CD4⁺ and CD8⁺ T cells by approximately 3 months after SC transplantation. These results confirm and extend the results of Cohen et al, who recently demonstrated, using LPA assays measuring herpesvirus-specific T cells, that many CB transplant recipients had at least 1 detectable assay after CBT that demonstrated successful donor-derived, antigen-specific T-cell recovery. Because CFC assays are more quantitative than LPA, our results also provide a more complete window into the quantitative recovery of CMV-specific T cells after CBT.

The most striking finding of our study was the observation that the recovery of thymopoiesis was markedly impaired both at baseline and, in all but 1 subject, at all time points after CBT. Previously, we have conducted cross-sectional studies of more than 140 allogeneic SCT recipients receiving transplants of either marrow or PBSC at our own institution; although thymopoiesis was transiently lower after SC transplantation, we found measurable thymopoiesis in nearly every subject, even at early intervals after SC transplantation. Thymopoiesis peaked approximately 6 to 8 months after SC transplantation and preceded the recovery of naive CD4⁺ and CD8⁺ T cells. Here, we report the results of 2 prospective parallel studies of adult autologous and allogeneic SC transplantation cohorts undergoing transplantation at the University of Colorado. In both groups, consistent with our prior cross-sectional studies, we similarly detected thymic function in nearly all subjects. Thus, the nearly universal lack of thymopoiesis in this group of CB transplant recipients stands out as uniquely different, relative to other SC transplantation subpopulations. Some studies have previously suggested that thymopoiesis can be detected, even at early intervals, in limited numbers of long-term survivors of CBT. Given that other (nonsurviving) patients were not assessed in these studies, it is impossible to know what would have been observed if all subjects in those studies had been examined prospectively. Indeed, our preliminary data suggest that patients who fail to preserve naive cells early after CBT are more likely to experience mortality; furthermore, our longest survivor to date is the only patient to recover significant and sustained thymopoiesis. Clearly, these data suggest that larger prospective studies of thymopoietic recovery are needed, especially in adult recipients who are known to have a diminished (though heterogeneous) capacity to recovery thymopoiesis after SC transplantation.

Phenotypic studies using markers of T-cell maturation and differentiation (eg, CD45RA/RO, CD27, CCR7, and CD28) have demonstrated that multiple subcompartments of the memory T-cell repertoire exist, with marked differences in their distribution, with naive and early memory subsets retaining their ability (via CCR7 expression) to home to lymph nodes where primary immune responses occur. Pantaleo and colleagues have examined the relative subsets of T cells specific for various viral pathogens capable of secreting IFNγ, IL-2, or both cytokines. Their results suggest that the subset of T cells that coproduce both IL-2 and IFNγ are a hallmark of protective immune responses to chronic viral pathogens, although cells producing IFNγ alone are often present in the setting of chronic progressive infection (eg, as seen in HIV-1 infection). In recent preclinical studies, we have demonstrated that the cytokine phenotype of late memory cells is distinct from that seen in earlier memory cells after a variety of stimuli, including phorbol esters (eg, phorbol 12-myristate 13-acetate), antibodies stimulating the CD3/TCR complex (eg, OKT3), and superantigens (eg, SEB). Relative to early- or central memory T cells, late effector memory cells produced were skewed toward MIP-1β and IFNγ production, with few cells producing IL-2. In a murine model of adoptive cellular therapy followed by immunization against melanoma-associated antigens, Restifo and colleagues demonstrated that the transfer of late-effector CD8⁺ T cells (characterized by the production of abundant IFNγ but little IL-2) was associated with progressive tumor growth; in contrast, adoptive transfer of cells with the same specificity but of a central memory differentiation stage resulted in tumor clearance.

One important caveat is that the patients studied here consisted of a relatively heavily pretreated group of patients. This fact is reflected by the fact that half of the acute myeloid leukemia (AML) recipients received transplants with active disease and overall had received an average of 3 prior treatment regimens and is also demonstrated by the fact that thymopoiesis at baseline, which has previously been demonstrated to correlate with post-CBT immune recovery in 1 pediatric study, was absent in most subjects and markedly reduced (relative to age-matched healthy controls) even in the minority of patients with detectable thymopoiesis before transplantation. Thus, we cannot necessarily generalize these results to other groups of CB transplant recipients who are less heavily pretreated. It should also be noted that all of our recipients received regimens containing antithymocyte globulin (ATG). Although the biologic half-life of ATG is prolonged and is likely to have influenced both central (ie, thymic) and peripheral T cells infused within CB grafts, this intervention is a standard component of most CB conditioning approaches and appears to be necessary to decrease the otherwise significant risk of graft rejection. It is also probable that stem cell dose, which is reduced in adults relative to children, may also be an important factor in the recovery of lymphoid precursors and thymopoietic function. Given the modest number of patients studied to date, it will be important to conduct larger, multi-institutional studies examining a greater number of immune recovery endpoints before we can draw firm conclusions regarding the broader relevance of the findings of this single-institution study. Ultimately, we need to understand how prior therapies, age, conditioning regimen, and graft characteristics influence thymic regeneration and functional T-cell recovery.
Overall, our data support the generally accepted notion that attempts to improve outcome in the setting of CBT for adults will depend on the success of these approaches in improving T-cell immune reconstitution. Already, encouraging progress has been made in the development of clinical strategies aimed at increasing stem-cell dose, including ex vivo expansion of progenitor cells,44,45 and the use of pooled CB units.9,46 It will be important to prospectively assess the influence of these various approaches on thymopoietic recovery and to consider immune recovery endpoints as a primary measure of the success of these trials, in addition to the typical primary endpoints (eg, date of neutrophil and platelet engraftment). Among approaches being considered, the use of Notch ligand-based approaches that may preferentially expand lymphoid progenitors deserves special consideration given our findings.7,48 Additional strategies, already demonstrated to have efficacy in murine models and in human trials, include the use of thymoprotective agents (eg, keratinocyte growth factor or related family members)49-51 and the use of thymopoietic factors (eg, androgen ablation using leuprolide, IL-7 administration, and growth hormone administration).52,53 Finally, the ex vivo expansion of naïve cord blood cells in the presence of viral antigens may offer a solution to the elusive goal of adoptively transferring antigen-specific memory to CB recipients.54,55 Although such strategies are attractive, our data also suggest that attention should be paid to the activation status, differentiation stage, and cytokine production signatures of expanded cell populations, because it is possible that in vitro stimulation might favor the production of late-stage memory cells that may be functionally ineffective or proliferatively impaired or even increase the otherwise manageable incidence of GVHD, which remains one of the principal advantages of CBT versus the use of other unrelated donor grafts.

References


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Authorship

Contribution: K.V.K. designed the study, analyzed the data, and wrote the manuscript; L.S.J. performed the primary research and analyzed the data; M.d.l. and R.E.C. designed the parent clinical trials; J.M. helped to design the parent clinical trials and contributed clinical data; S.R. and I.M. contributed to the study of control subjects; S.G.B., I.K., and S.M. performed the research; E.D.W. contributed valuable flow cytometry expertise; L.W., D.P., and L.J.N.C contributed to the parent clinical trials; J.J.M. assisted with data analysis and interpretation; and E.J.S. designed the parent clinical trials and contributed to writing the manuscript.

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