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Post Transplant BAFF and B-cell RecoveryPrior to Onset of Chronic GVHD

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Abstract

Excessive levels of B Cell Activating Factor (BAFF) are found in patients with active chronic graft versus host disease (cGVHD). In mice, BAFF has been shown to be essential for B cell recovery after myeloablation. To assess how BAFF levels relate to transplant factors and subsequent cGVHD development, we prospectively monitored 412 patients in the first year after allogeneic peripheral blood or bone marrow (PB/BM) hematopoietic stem cell transplantation (HSCT) and censoreddata at time ofcGVHD onset. In patients who did not develop cGVHD, we affirmeda temporal pattern of gradual BAFF level decreaseas theB cell numbers increase after myeloablative conditioning (MAC). By contrast, after reduced intensity conditioning (RIC), BAFF levels remained high throughoutthe first post-HSCT year, suggesting that the degree of myeloablation resulted in delayedB cell recovery associated with persistence of higher BAFF levels. Since high BAFF/B ratios have been associated with active cGVHD, weexamined differences in early BAFF/B ratios and found significantly different BAFF/B ratios at 3 months post-HSCT only afterMAC in patients who subsequently developed cGVHD. In addition toHSCT conditioning type, use of sirolimus was significantly associated with higher BAFF levels after HSCT and this wasalso potentially related tolower B cell numbers. Together, our results are important for interpretation of BAFF measurements in cGVHD biomarker studies.

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Contribution: C.A.J. designed the research, performed research, and wrote the paper; L.S. and H.T.K. analyzed data and helped write the manuscript; S.M.M., C.G.R, M.I.H. and M.S. performed research; J.K., C.S.C., V.T.H., E.P.A., P.A., B.R.B., R.J.S, J.H.A. provided vital patient samples and clinical information and they helped write the manuscript; J.R. provided vital patient samples, designed the research and wrote the paper; S.S. conceived of the study, analyzed data and wrote the paper.

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Introduction

Bcell activating factor (BAFF) is a key regulator of B-cell homeostasis. BAFF support is required for normal B-cell proliferation and survival and the absence of BAFF or BAFF-R results in profound B lymphopenia.(1, 2) In contrast, excess BAFF has been associated with persistence of autoreactive B cells and a variety of autoimmune diseases.(3) These homeostatic functions of BAFF are exemplified in astudy of patients with common variable immunodeficiency (CVID).(4) In CVID patients soluble BAFF levels wereinversely correlated with peripheralB-cell numbers and the expression of BAFF receptors. The clinical significance of this cytokine pathway is also highlighted by the recent FDA approval of belimumab, an anti-BAFF monoclonal antibody for the treatment of systemic lupus erythematosus (SLE). In SLE, neutralization of high BAFF levels leads to a reduction of pathogenic autoantibodies and clinical improvement of disease.(5, 6)

BAFF alsohas an important role in the reconstitution of B cells and normal B-cell function after stem cell transplantation. In murine models, BAFF is required for reconstitution of the B-cell compartment after myeloablation.(1) Soluble BAFF levels are characteristically high when patients are B-lymphopenic and BAFF levels gradually decrease as the number of circulating B cells return to normal levels.(7-9) After transplantation, increased numbers of bone marrow B-cell precursors and early recovery of circulating B cells have been observed in patients who do not develop cGVHD.(10-13) In addition to supporting recovery of normal B cells after stem cell transplantation, BAFF has also been proposed to contribute to the development of chronic graft versus host disease (cGVHD).(14, 15) Despite having significantly higher BAFF levels, chronic GVHD patients often have low total numbers of circulating B cells compared to those without cGVHD, resulting in relatively high BAFF/B ratios and the presence of circulating activated B cells. While the mechanisms whereby BAFF contributes to the development of cGVHD are not well established, many studies strongly support a role for B cells in this disease. While the cellular sources of BAFF are unknown in cGVHD, inducible expression from neutrophils, monocytes/macrophages, dendritic cell subsets, T-cells and activated B-cells occurs in other inflammatory states.(16, 17)

Previous studies have suggestedthat regulation of B-cell recovery by BAFF might influence the subsequent development of cGVHD.(18) To examine this issue, we prospectively monitored a large cohort of 412patients with serial measurements of BAFF levels and B-cell numbers in the first year after allogeneic HSCTbefore cGVHD development. This study identifies several key variables such as the intensity of transplant conditioning therapy and sirolimus important for interpretation of differences in BAFF levels and B cell numbers in the first HSCT year.

Methods

Patient Characteristics

Serial blood samples for analysis of BAFF levels and B-cell reconstitution were obtained from 412 patients who underwent allogeneic HSCT and survived at least 100 days after HSCT at the Dana-Farber Cancer Institute and Brigham and Women's Hospital from 2000-2008. Written informed consent was obtained in accordance with the Declaration of Helsinki and approved by the Human Subjects Protection Committee of the Dana-Farber/ Harvard Cancer Center.

Table 1 summarizes clinical characteristicsof the patient cohort. Umbilical cord blood transplantation(UCBT) patients were excluded from this study since B cell recovery and BAFF levels have previously been shown to be very different in these patients.(19, 20)

Donors were HLA-matched in 86% of transplants, including 42% with related donors. Acute GVHD developed in 24% of patients and chronic GVHD developed in 67% following HSCT. Chronic GVHD was categorized according to documented clinical examination and laboratory studies using both Seattle criteria(21) and National Institutes of Health cGVHD consensus criteria.(22) The median time from HSCT to the development of cGVHD was 222 days (range 71-994 days). Median follow-up for all surviving patients was 72 months (range 30–133 months) after HSCT. Samples obtained after cGVHD development or after disease relapse were excluded from analysis.

Processing of patient plasma

Blood was drawn into standard EDTA-containing collection tubes. Plasma was separated from whole blood cells by centrifugation at 600xg, stored in aliquots at -80°C, and used after first or second thaw for BAFF measurements.

BAFF enzyme-linked immunosorbent assay

Soluble BAFF in patient plasma samples was measured using a commercially available enzyme-linked immunosorbent assay (ELISA) according to the manufacturer's recommended procedures (R&D Systems, Minneapolis, MN).

Flow cytometric analysis of peripheral blood cells

Fluorochrome-conjugated monoclonal antibodies specific for: CD3, CD4, CD19, CD20, CD25, CD27, (BD Biosciences); CD8 (Beckman Coulter, Fullerton, CA) were used to stain fresh, whole blood. After staining, red blood cellswere lysed with BD Pharm Lyse or an automated TQ Prep workstation (Beckman Coulter). Flow cytometry was performed on a FACSCanto II (BD Bioscience) and analyzed using BD FACSDiva software, or on a Beckman Coulter FC500, with Beckman Coulter CXP analysis software. We found no difference betweenB cell numbers using CD19 or CD20 cell surface staining. B-cell percentage was based on enumeration of cells expressing CD20.

Statistical Analysis

Patient baseline and transplant characteristics were reported descriptively, and compared using Fisher's exact test or Wilcoxon rank sum test. BAFF, B cell, BAFF/B-cell ratio data were analyzed descriptively at each time point and compared using the Wilcoxon rank sum test. All p-values are 2-sided at significance level of .05. Multiple comparisons were not considered. The Kaplan-Meier method was used to estimate overall survival (OS) and progression-free survival (PFS). OS was calculated from date of transplantation to date of death. PFS was calculated from date of transplantation to time of relapse/disease progression or death, whichever occurred first. Multivariable linear regression analysis was performed to assess factors that potentially affect elevated BAFF levels. All calculations were performed using SAS 9.2 (SAS Institute, Cary, NC) and R 2.10.1.

Results

BAFF levels and B cell recovery associated with intensity of conditioning regimen and stem cell source

We examined changes in plasma BAFF levelsand reconstitution of B cells in the first year after HSCT in412adult patientswith hematologic malignancies who underwent allogeneic HSCT. This was the first allogeneic HSCT for 96.4% of patients. Patients received granulocyte colony stimulating factor (G-CSF)-mobilized peripheral blood (91%) or bone marrow (9%) stem cells. The 5-year overall and progression-free survival for the entire cohort was 56% and 47%, respectively. Since subsequent treatment would affect both

circulating B cells and BAFF levels, patients were censored at time of cGVHD onset or relapse. None of the patients developed graft failure. Of the 412 patients, 275 (67%) developed cGVHD at a median of 7.4 months after transplant (cGVHD cohort) and 137 patients (33%) did not develop cGVHD during the follow-up period (No-cGVHD cohort). The cGVHD cohort differed from the No-cGVHD cohort in several important variables(Table 1); there were significantly more male patients with female donors, patients with matched unrelated donors (MUD), mobilized PB stem cell grafts and previous history of grade II-IV acute GVHD. All of these variables have previously been associated with increased risk of developing cGVHD.

We first examined whether intensity of the conditioning regimen influences B-cell reconstitution following transplantation.(23-27) We found that the patterns of BAFF and B cell recoverydiffered in the MAC and RIC groups. BAFF levels in RIC versus MAC HSCT were significantly different at 3 month and 12 month time points. Specifically, BAFF levels 3 months after MAC HSCT were significantly higher vs RIC at 10.62ng/ml, (range 1.65-40.47) versus 7.93 ng/ml (range 0.79-61-.03), p=0.02; and BAFF levels were significantly higher in RIC at 12 months with MAC at 5.48 (range 2.37-25.3) and RIC at 8.0 (range 2.14-20.69), p=0.02. These increased BAFF levels did not appear to associate with total B cell numbers, since we found no significant difference in B cell numbers or BAFF/B cell ratios at any time point after MAC versus RIC HSCT conditioning (data not shown). To determine whether B cell recovery was related to the level of engraftment with donor cells, we examined available total peripheral blood cell chimerism. No significant correlation between BAFF/B cell and chimerism at Day 30 orDay 100 post transplantation was found (Supplementary Table 1). Since this potential effect of HSCT conditioning on BAFF and B cell values would impact how these values are interpreted in cGVHD studies, we further examined data by stratifying according to subsequent cGVHD development.

BAFF levels and B cell recovery before chronic GVHD development

To identify differences in B cell recovery prior to development of cGVHD, patients were censored after onset of cGVHD, but stratified according to subsequent cGVHD development. First, to further examine the relationship between B cell number and BAFF levels, we compared BAFF and B-cell recovery separately in patients who received either MAC or RIC transplants and either later developed or who never developed cGVHD. Of these412patients,275(67%) developed cGVHD at a median of 7.4 months after transplant (cGVHD cohort) and 137 patients (33%) did not develop cGVHD during the follow-up period (No-cGVHD cohort). BAFF levels and B-cell numbers were measured at 1, 3, 6, 9, and 12 months after HSCT in all available samples from both MAC and RIC cohorts. Figure 1 shows the comparison of patients who received RIC with patients who received MAC before cGVHD development. The general pattern of BAFF and B-cell recovery was similar in patients who do not develop cGVHD in both groups, but these patterns were temporally distinct and appeared to reflect the level and rate of B-cell recovery. Early after transplant, BAFF levels were high in both MAC and RIC groups. B-cell recovery after RIC with support was slightly slower than after MAC transplants. In each group, the decrease in BAFF levels reflected the tempo of recovery of peripheral B cells. This coordinated pattern of BAFF and B-cell number increaseis consistent with previous reports demonstrating a low incidence of cGVHD in patients who have rapid recovery of normal B cells. Taken together and consistent with previous findings, this data suggests a relationship between BAFF levels and degree of B lymphopenia.

We found that patterns of BAFF and B-cell recovery were different in patients who subsequently developed cGVHD. In MAC and RIC patients who subsequently developed cGVHD (Figure 1), BAFF levels remained elevated and were relatively stable while B-cell numbers slowly increased in the first year after transplant. Toascertain whether high BAFF

was related to naïve B lymphopenia, we examined CD27, a marker of antigen-experienced B cells. $CD19^+CD27^+$ B cell percentages were available for only a small subset of these patients (64 out of 412) at various time points. In this small patient subset, thepercentage of CD19+CD27+ B cells was not significantly different between groups at any time point, revealing that the majority of cells were likely naive B cells. Interestingly, the frequency of CD27+ B cells was high at 1 month in both cGVHD and no cGVHD groups after HSCT (data not shown), potentially reflecting increased proportions of antigen-experienced B cells or early bone marrow CD27+ B cells (28, 29). Thus, taken together our data show that cGVHD and No-cGVHD cohorts within each PB/BM transplantation conditioning cohort have variations in the kinetics of BAFF level change with B cell number increase following transplant, but with delayed B cell recovery in those with subsequent cGVHD development.

Since high BAFF/B ratios in patients with active cGVHD may be pathobiologically relevant(13, 30), we examined whether a difference in BAFF/B cell ratio occurred prior to disease onset. Thus, a more detailed analysis of the data comparing results in the cGVHD and No-cGVHD cohorts is shown in Figure 2. After MAC, BAFF levels were persistently and significantly higher in the cGVHD cohort compared with the No-cGVHD cohort (Figure 2A) (6 months: 9.79 vs 5.71 ng/mL, p=0.025; 9 months: 12.3 vs. 3.99 ng/mL, p=0.004; 12 months: 8.68 vs. 4.3 ng/mL, p=0.02). After MAC,B-cell numberwas significantly lower 3 months after HSCT in patients who subsequently developed cGVHDcompared to those without cGVHD development (10vs. 37cells/uL, p=0.008, Figure2C). This corresponded to ahigherBAFF/B-cell ratio at this time (1.11 vs. o.37, p=0.01,Figure 2E). The only significant difference following RIC was at 6 months when BAFF levels were significantly higher in the No-cGVHD group (Figure 2B)(9.92 vs 7.09 ng/mL in +cGVHD, p=0.05). This peak in BAFF after RIC was followed by a more rapid, albeit not statistically significant, increase in total B-cell numbers at 9 months and 12 months in the No-cGVHD compared with the cGVHD group (Figure 2D). Taken together, our data suggests that BAFF levels in the first year after transplantation are affected by the intensity of the transplant conditioning. The persistence of high BAFF levels associated with the subsequent development of cGVHD is most evident after MAC.

Effects of acute GVHD and corticosteroid therapy on BAFF levels

To examine the role of acute GVHD we compared patients with cGVHD who also had a history of aGVHD with patients with *de novo* cGVHD with respect to BAFF levels, B-cell numbers, and BAFF/B-cell ratios. There were no significant differences in any measure at any time point between the two groups (data not shown). Because high dose corticosteroid therapy is known to be associated with lower BAFF levels (31), each patient was further categorized for use of prednisone, either at any dose or only at doses >30mg/day prednisone (or equivalent) at the time that samples were obtained for BAFF measurement. 216 patients received prednisone at any dose in the first year after HSCT; 100 patients received >30mg/ day prednisone. When samples obtained from patients receiving any dose or only >30mg/ day of prednisone at the time of sample collection were excluded from analysis, we observed a similar temporal pattern of BAFF levels between the two groups as had been observed for all patients: 1 month after HSCT, BAFF levels remained significantly higher in the No-cGVHD cohort (15.21 versus 11.44 ng/ml, $p=0.004$), but decreased with time such that they were significantly lower in the No-cGVHD cohort 12 months after HSCT (5.57 versus 9.57, p=0.009) (data not shown). This analysis suggests that steroid treatment did not affect the relative differences in BAFF levels found in patients who subsequently developed cGVHD versus those who did not develop disease.

Influence of clinical factors and sirolimus on BAFF and B-cell recovery

We attempted to uncover additional factors contributing to high BAFF levels by examining factors that potentially affect BAFF and B-cell levels. Importantly, prior treatment with anti-B cell or lymphocyte antibody therapy was notassociated with cGVHD development in our cohort. None of the patients in this study received ATG in their transplant conditioning regimen. Of the 114 patients with B-cell non-Hodgkin lymphoma or chronic lymphocytic leukemia (CLL), 60 received rituximab or ibritumomab tiuxetan and 6 received alemtuzumab within 6 months prior to transplantation. Importantly, when we excluded the 60 patients who received B-cell specific depletion therapy, the pattern of BAFF levels was the same but B-cell recovery was more robust for the cGVHD cohort and less robust for the "no cGVHD" cohort resulting in no difference between the two groups for B cells or BAFF/ B cell ratios (data not shown). Various clinical factors are known to increase the risk of developing cGVHD, but multivariable linear regression analysis did not detect a significant association between age, donor and patient sex mismatch, related or unrelated HSCT, stem cell source and BAFF levels at any time in the first year post transplant (Table 2). B cell recovery was faster in younger patients than in older patients (median 143.7 vs 27.7, respectively, p=0.049 at 9 month; median 229.4 vs 46, respectively, p=0.004 at 12 month). In multivariable analysis,older recipient age was associated with a significantly lower total B cell number, but only at the 12 monthtime point $(p=0.002)$. In multivariable linear regression

Multivariable analysis also revealed that sirolimuswasmost significantly associated with higher BAFF levelsat 6months after HSCT (Table 2, p=0.0006). Notably, this significant increase in BAFF level was preceded at the 3 month time point by a significant decrease in B cell number (Table 2, p=0.0005). As shown in Figure 3A, BAFF levels remained high at all times in patients receiving sirolimus and decreased after MAC as B cell number increased and notablydid not decrease asB-cell numbers recovered in those who received sirolimus, suggesting that factors other than total B cell number affect BAFF levels. In contrast, B-cell numbers recovered more rapidly in patients not receiving sirolimus and this was associated with gradual lowering of BAFF levels in these patients(Figure 3B). Of note, GVHD prophylaxis was typically discontinued by 6 months after transplant when patients had no signs of cGVHD. Due to the relatively small number of patients who did not receive sirolimusas GVHD prophylaxis in this study, we were not able to furtheranalyze the effect of sirolimus in the cGVHD and No-cGVHD cohorts.

analysis, we found that conditioning regimen and donor cell source were also significantly

associated with BAFF levels at 3 months after HSCT (p=0.02, Table 2).

Discussion

Chronic GVHD results from a complex series of immune interactionsthat occur as the donor immune system develops in an antigenically disparate host/recipient. Patients with chronic GVHD frequently produce auto-antibodies as well as allo-antibodies, suggesting that the pathogenesis of this disease reflects a critical breakdown in peripheral B-cell tolerance. After allogeneic HSCT, the re-establishment of B-cell homeostasis withoutalloimmunity or autoimmunity may reflectan improved capacity for B-cell recovery early after HSCT.(10) This is similar to what has been demonstrated in mouse models of autoimmunity, wherethe reconstitution of peripheral naïve B cells is critical for the maintenance of B-cell tolerance. Thus, the amount of available soluble BAFF is potentially a pivotal determinant of B cell homeostasis after HSCT(32, 33). Thus, while pre-emptive treatment of cGVHD would likely alter patient outcomes, abrogation of BAFF in the early post transplantation period may not be beneficial. While BAFF has been implicated in cGVHD, whether factors in the early post-transplant period affected patterns of BAFF levels required further examination, since this information impacts interpretation of BAFF levels in ongoing GVHD biomarker studies.

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The current prospective study in 412 patients reveals the dynamic interaction between BAFF and B-cell reconstitution and their relationship to cGVHD. Patients were analyzed serially for up to 1 year prior to the onset of cGVHD to examine the relationship between BAFF and BAFF/B-cell ratios and subsequent development of cGVHD. While induction of lymphopenia after transplant was generally associated withincreased BAFF levels, higher BAFF levels tended to precedemore rapid B-cell reconstitution, and this was particularly evident in patients who subsequently never developed cGVHDafter RIC (Figure 2B and 2D). This finding is consistent with previous studies demonstrating increased numbers of precursor B cells in patients early after HSCT who do not develop cGVHD.(10, 34)Following a more rapid B-cell recovery and concurrent decline in BAFF levels, patients who did not develop cGVHD had a significantly lower BAFF/B-cell ratio 12 months following HSCT. This observation suggests thatonce B cells recover and normal B-cell homeostasis is established, BAFF levels become limiting. This decrease in the amount of BAFF available per B cell (the BAFF/B-cell ratio) potentially promotesB-cell tolerance with preferential survival of non-alloreactive B cells that are less likely to contribute to cGVHD. (32, 33, 35) Taken together, these results suggest that pre-emptive treatment of cGVHD directed at BAFF in the early post-transplantation period may not be beneficial. In contrast, prophylactic rituximab therapy would also delay B-cell reconstitution, but this approach would result in high BAFF levels and allow for a "reset" of BAFF and B-cell kinetics that leads to cGVHD prevention. While, in the current analysis, no patients received prophylactive rituximab, the increased B cell number and decreased BAFF/B ratio found in patients after prophylactic rituximab who did not develop cGVHD affirms this notion.(40)

Our study of a large number of transplant patients allowed for an investigation of the impact of various HSCT-related factors on BAFF levels and B cell recovery and the subsequent development of cGVHD. We found that transplant conditioning (MAC versus RIC) significantly impacted the kinetics of BAFF and B-cell numbers following transplant. Older age of HSCT recipients was significantly associated with decreased B cell number at the 12 month time point. An unexpected finding in our study was the highly significant effect of sirolimus on BAFF levels and B-cell reconstitution. Overall, 64% of patients in this study received sirolimus as part of their GVHD prophylaxis and this was associated with persistence of high BAFF levels as well as delayed B-cell reconstitution. Although not statistically significant, and not found in previously published reports at our center [rev. in(36)], these patients also experienced a higher incidence of cGVHD than patients who did not receive a sirolimus-containing GVHD prophylaxis regimen (67% vs 59% respectively, p=0.1) in the recently presented Clinical Trials Network study.(37) Our analysis did not allow us to decipher the mechanisms responsible for elevated BAFF associated withsirolimus administration. The higher BAFF levels that were associated with cGVHD development in the sirolimus cohort may have been independent of sirolimus use. Since sirolimus was associated with significantly lower B cell numbers prior to the significant increase in BAFF levels, it is tempting to speculate that sirolimus itself may result in higher BAFF levels because of B lymphopenia induction. This is plausible because sirolimus in mice is known to hinder B cell development.(38) Previous studies havealso suggested that mTor inhibition may contribute to increased BAFF production(39) or elimination of activated B cells.(40, 41) Taken together,our data reveal the effects of HSCT conditioning, age and sirolimus onB cell recovery kinetics and BAFF levels after HSCT.

Understanding a potential role for early high BAFF levels in relation to B-cell reconstitution and attainment of B-cell tolerance versus whether high BAFF relates to the promotion of autoreactive B cells will be critical if we are to understand cGVHD pathophysiology.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Patients who received either myelablative (MAC) or reduced intensity conditioning (RIC) are shown separately according to subsequent cGVHD development. **A)** BAFF levels (squares) and B cell numbers (triangles) after MAC in patients who never developed cGVHD. **B)** BAFF levels (squares) and B cell numbers (triangles) after MAC in patients who developed cGVHD. **C)** BAFF levels (squares) and B cell numbers (triangles) after RIC in patients who never developed cGVHD. **D)** BAFF levels (squares) and B cell numbers (triangles) after RIC in patients who subsequently cGVHD.

*Indicates differences that are statistically significant $(p<0.05)$.

Note: Blood samples after the clinical onset of cGVHD were excluded from analysis.

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Figure 2. Patterns of BAFF levels, B-cells and BAFF/B-cell ratios after stem cell transplantation before cGVHD onset

Serial values during the first year after transplant for patients who received eitherMAC (three left panels, **A, C**, and **E**) or RIC (three right panels, **B, D**, and **F**) prior to HSCT. Patients who later developed cGVHD (triangles) are compared to patients who did not develop cGVHD (squares).

*Indicates differences that are statistically significant ($p<0.05$).

Median values and ranges (in parentheses) are listed for those that were significantly different after MAC are as follows: 6 months: yes cGVHD 9.79 (0.62-58.43) vs no cGVHD 5.71 (0.08-34.49) ng/mL, p=0.025; 9 months: yes cGVHD 12.3 (1.75-37.38) vs. no cGVHD 3.99 (0.83-31.82) ng/mL, p=0.004; 12 months: yes cGVHD 8.68 (2.37-22.17) vs. no cGVHD 4.3 (3.48-25.3) ng/mL, $p=0.02$); or after RIC are as follows: 6 months: yes cGVHD 7.09 (0-26.01] vs no cGVHD 9.92 (0-80.49) ng/mL, p=0.05). Significant differences in total B cell numbers (median values with range in parentheses) are listed as follows: 3 months after MAC: yes cGVHD 10 (0- 631.41) versus no cGVHD 37 (0-434.64) cells/uL, p=0.008; BAFF/B-cell ratios were calculatedas previously described and are depicted on a log scale. Significant difference was determined at the 3 month timepoint after MAC with median values and ranges (in parentheses) as follows: yes cGVHD 1.11 (0.05-20.07)vs No-cGVHD 0.37 (0.03-10.75), p=0.01.

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Figure 3. Changes in BAFF levels and B-cell recoveryin patients receiving sirolimus for GVHD prophylaxis after stem cell transplantation

Serial median BAFF levels (left y-axis, squares) are compared to B cell numbers (right yaxis, triangles) in patients at 1, 3, 6, 9, and 12 months post-transplantation. **A)** Patients who received sirolimus after transplant. **B)** Patients who did not receive sirolimus after transplant. Note: All patients studied after HSCT, before cGVHDdevelopment, were combined in this analysis.

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Table 1

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Clinical characteristics of study population Clinical characteristics of study population

HLA = human leukocyte antigen; GVHD = graft-versus-host disease; NHL = non-Hodgkin lymphoma; CLL = chronic lymphocytic leukemia; HD = Hodgkin disease; AML = acute myelogenous leukemia; HLA = human leukocyte antigen; GVHD = graft-versus-host disease; NHL = non-Hodgkin lymphoma; CLL = chronic lymphocytic leukemia; HD = Hodgkin disease; AML = acute myelogenous leukemia; CML = chronic myelogenous leukemia; MDS/MPD = myelodysplastic syndrome/myeloproliferative disorder; ALL = acute lymphoblastic leukemia; MM = multiple myeloma; aGVHD = acute GVHD; CML = chronic myelogenous leukemia; MDS/MPD = myelodysplastic syndrome/myeloproliferative disorder; ALL = acute lymphoblastic leukemia; MM = multiple myeloma; aGVHD = acute GVHD; $\mathrm{cGVHD} = \mathrm{chronic}\ \mathrm{GVHD}$ cGVHD = chronic GVHD

Details of conditioning regimens and GVHD prophylaxis: Details of conditioning regimens and GVHD prophylaxis:

Myeloablative Conditioning (MAC): TBI d-4 to -2; cyclophosphamide 1800mg/m2 IV QD d-6,-5 GVHD: MTX 15mg/m2 IV QD d1 then 10mg/m2 IV QD d3,6,11; Tacrolimus 0.02mg/kg IVCI starting Myeloablative Conditioning (MAC): TBI d-4 to -2; cyclophosphamide 1800mg/m2 IV QD d-6,-5 GVHD: MTX 15mg/m2 IV QD d1 then 10mg/m2 IV QD d3,6,11; Tacrolimus 0.02mg/kg IVCI starting day -3 and titrated to level day -3 and titrated to level

Reduced Intensity Conditioning (RIC): Fludarabine 30mg/m2 IV QD d-5 to -2; Busulfan 0.8mg/kg IV QD d-5 to -2 Reduced Intensity Conditioning (RIC): Fludarabine 30mg/m2 IV QD d-5 to -2; Busulfan 0.8mg/kg IV QD d-5 to -2

GVHD Prophylaxis: MTX 5mg/m2 IV QD d1,3,6 (day 11 is optional); Tacrolimus 0.05mg/kg BID starting day -3 and titrated to level; Sirolimus 12mg po QD d-3, then 4mg PO QD from day-2 on and GVHD Prophylaxis: MTX 5mg/m2 IV QD d1,3,6 (day 11 is optional); Tacrolimus 0.05mg/kg BID starting day -3 and titrated to level; Sirolimus 12mg po QD d-3, then 4mg PO QD from day-2 on and GCD from day-2 on and titrated to level titrated to level

Table 2

Multivariable linear regression models on Log (BAFF) and Log10 (B cell) at 1,3,6,9, and 12 months after HSCT. Multivariable linear regression models on Log (BAFF) and Log10 (B cell) at 1,3,6,9, and 12 months after HSCT.

 a , older' is defined as age $>=50$ in MAC and age
 $>=60$ in RIC a : 'older' is defined as age >=50 in MAC and age>=60 in RIC