The cytokines B lymphocyte stimulator (BLyS) and a proliferation-inducing ligand (APRIL) enhance autoimmune disease by sustaining B cell activation. In RA, B cells contribute to the formation of 3 functionally distinct types of lymphoid microarchitectures in the inflamed synovium: ectopic GCs; T cell–B cell aggregates lacking GC reactions; and unorganized, diffuse infiltrates. We examined 72 tissues representing the 3 types of synovitis for BLyS and APRIL production and for expression of APRIL/BLyS receptors. Biologic effects of BLyS and APRIL were explored by treating human synovium–SCID mouse chimeras with the APRIL and BLyS decay receptor transmembrane activator and CAML interactor:Fc (TACI:Fc). GC+ synovitis had the highest levels of APRIL, produced exclusively by CD83+ DCs. BLyS was present in similar levels in all tissue types and derived exclusively from CD68+ macrophages. In GC+ synovitis, treatment with TACI:Fc resulted in GC destruction and marked inhibition of IFN-γ and Ig transcription. In contrast, inhibition of APRIL and BLyS in aggregate and diffuse synovitis left Ig levels unaffected and enhanced IFN-γ production. These differential immunomodulatory effects correlated with the presence of TACI+ T cells in aggregate and diffuse synovitis and their absence in GC+ synovitis. We propose that BLyS and APRIL regulate B cell as well as T cell function and have pro- and antiinflammatory activities in RA.

Introduction
RA is a systemic autoimmune disease that primarily targets the synovial membrane, cartilage, and bone. Synovial inflammatory infiltrates typically include T cells, B cells, macrophages, and DCs that arrange in defined microarchitectures. The organization of tissue-infiltrating lymphocytes in the synovium is highly complex and recapitulates molecular pathways of lymphoid organogenesis (1, 2). A typical example of ectopic lymphoid neogenesis is the formation of GCs in the synovial sublining layer. Such synovial GCs support B cell selection and affinity maturation (3). In contrast to GC reactions in lymphoid organs, synovial GCs depend upon a unique population of CD8+ T cells positioned in the mantle zone that colocalize with lymphotoxin-β-producing (LT-β-producing) B cells (4). Synovial T cells and B cells can also arrange in clusters, called T cell–B cell aggregates, which lack central follicular DCs (FDCs), B cell proliferation, and mantle zone formation. Recent cohort studies have revealed that GC+ and aggregate+ tissues are present in similar frequencies and account for about 50% of all patients. The other half of patients have diffuse T and B cell infiltrates without distinct topographic arrangements, a pattern called diffuse synovitis (5).

A close correlation exists between the lymphoid architecture and the functional activity of T cells and B cells in the lesion (6). The sharing of T cell receptor sequences between distinct GCs strongly suggests the sharing of antigens (7). However, an antigen-independent pathway may also influence B cell recruitment, organization, and functional activity in rheumatoid synovitis. Recent reports indicate that B cell activation, differentiation, and survival are determined not only by antigen and T cell interaction, but also by cytokines, particularly by members of the TNF ligand superfamily (8).

The TNF superfamily member B lymphocyte stimulator (BLyS), also referred to as BAFF, THANK, TALL-1, TNFSF13b, and zTNF4 (9–12), is known to be a very effective modulator of peripheral B cell homeostasis that promotes B cell survival and differentiation (13). BLyS is expressed by a few stromal cells, T cells, and most myeloid cell lineages—including monocytes, macrophages, DCs, and stimulated neutrophils—and may be the active cytokine by which macrophages and DCs regulate human B cell function (14–18). BLyS transgenic mice show an expansion of the peripheral mature B cell compartment, hypergobulinemia, anti–single-stranded DNA and anti–double-stranded DNA antibodies, and circulating immune complexes. Such mice develop autoimmune-like manifestations reminiscent of human systemic lupus erythematosus (SLE) and Sjögren syndrome (SS) (19, 20).

Consistent with those data, elevated BLyS serum levels have been found in patients with RA, SLE, and SS (21–24). In contrast, in BLyS−/− mice, B cell development in the spleen is impaired beyond the T1 transitional stage and in the peritoneum beyond the B1 stage. There are practically no T2, marginal zone, or follicular B cells in the spleen and lymph nodes or B2 cells in the peritoneum present, indicating that BLyS is fundamental for B cell homeostasis. Along with diminished B cell differentiation, serum Ig levels are profoundly reduced (13).

A proliferation-inducing ligand (APRIL), also known as TALL-2, TRDL-1, and TNFSF13a (25, 26), is a close homolog to BLyS that is expressed by monocytes, macrophages, DCs, T cells, and several types of tumor cells. APRIL is virtually undetectable in normal tissues but is strongly expressed in adenocarcinomas and can accelerate the growth of malignant cells in vitro and in vivo. In contrast to most ligands of the TNF superfamily that are known to be homotrimeric, APRIL and BLyS have also been reported to form heterotrimeric molecules when coexpressed, and those APRIL/BLyS heterotrimers are present in the sera of patients with systemic autoimmune diseases like RA, SLE, and SS (27). Mice

Nonstandard abbreviations used: APRIL, a proliferation-inducing ligand; BAFF-R, BAFF receptor; BCMA, B cell maturation antigen; BlyS, B lymphocyte stimulator; CD21L, CD21 long isoform; FDC, follicular DC; LT, lymphotoxin; SLE, systemic lupus erythematosus; SS, Sjögren syndrome; TACI, transmembrane activator and CAML interactor.

Conflict of interest: The authors have declared that no conflict of interest exists.

BLyS and APRIL in rheumatoid arthritis
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manifested by elevated IgM and IgG serum levels (28). In contrast, TACI share receptors and bind specifically to 2 TNF superfamily receptors, transmembrane activator and CAML interactor (TACI; also known as TNFRSF13b) and B cell maturation antigen (BCMA). The role of the TNF ligands APRIL and BLyS in rheumatoid synovitis, we analyzed a cohort of 72 synovial biopsies for the tissue expression of APRIL/BLyS receptor TACI was expressed on plasma cells, B cells, and T cells. APRIL and BLyS production in rheumatoid synovitis and were absent in GC-containing tissues. The APRIL/BLyS receptor TACI was expressed on plasma cells, B cells, and T cells. However, TACI−/− mice were restricted to aggregate and diffuse synovitis and were absent in GC− synovitis. In vivo blocking of APRIL/BLyS with TACI:Fc essentially abrogated GC− synovitis, suggesting a critical role of the TNF homologs in the formation and maintenance of ectopic GCs. Conversely, treatment with TACI:Fc enhanced T cell function without affecting Ig production in synovial lesions that contained TACI−/− T cells, suggesting that APRIL and BLyS have immunostimulatory as well as immunosuppressive function in RA.

### Results

**APRIL and BLyS production in rheumatoid synovitis.** To determine the role of the TNF ligands APRIL and BLyS in rheumatoid synovitis, we analyzed a cohort of 72 synovial biopsies for the tissue expression of both cytokines. Synovial tissues were classified into 3 categories according to the type of lymphoid microstructure. Sixteen tissues had GC reactions, 30 tissues exhibited GC-negative T cell–B cell aggregates, and 26 tissues showed diffuse synovitis.

![Figure 1](http://www.jci.org)  
**Figure 1** B cell function and tissue expression of APRIL/BLyS in RA synovium. Synovial biopsies from 72 patients were classified according to the lymphoid microarchitectures. Sixteen patients had GC+ synovitis, 30 patients had aggregate synovitis, and 26 patients had diffuse synovitis. CDNA from tissue extracts was adjusted relative to 2 × 10^6 β-actin copies, and specific transcripts were determined by real-time PCR. (A) H&E staining of representative tissues displayed classical GC (left), lymphoid aggregates (Agg) without GC formation (middle), and diffuse (Diff) mononuclear infiltrates without topographical clustering (right). Original magnification, ×100. (B) IgG transcription was highest in GC− synovitis, intermediate in aggregate synovitis, and low in diffuse synovitis. LT-β and CCL19 production correlated closely with the pattern of lymphoid organogenesis in the synovium. (C) Tissue transcripts for APRIL followed the same hierarchy as LT-β and CCL19. BLyS-specific sequences were abundantly found in all tissues with no correlation to synovial lymphoid microstructures. Results are shown as box plots with medians, twenty-fifth and seventy-fifth percentiles as boxes and tenth and ninetieth percentiles as whiskers. P values are indicated where statistically significant.
in GC+ samples (P = 0.008 versus aggregate; P < 0.001 versus diffuse) (Figure 1B). Aggregate synovitis contained intermediate levels of CCL19-specific transcripts in the cohort of 72 tissue extracts. The highest levels of IgG1/2 transcripts were found in tissues with GC reactions. Aggregate-containing tissues produced significantly lower concentrations (P < 0.04), and IgG1/2 transcripts were 100-fold lower in tissue samples with diffuse infiltrates (P < 0.001).

Tissue expression of APRIL mRNA was closely correlated with the production of the DC-derived chemokine CCL19 (Figure 1C). The median concentration of APRIL-specific transcripts in GC+ synovitis was 9,669 copies and was significantly higher (P < 0.001) than that in aggregate synovitis (2,967 copies). In tissue sections with diffuse synovitis, APRIL mRNA concentrations were more than 10-fold lower, with a median of 882 copies (P < 0.001). A different picture emerged for BLyS mRNA which was abundantly detected at indistinguishable levels in all types of synovitis. These findings suggest that APRIL, rather than BLyS, mRNA levels correlate with the variability of tissue B cell function.

To identify the APRIL- and BLyS-producing cells in synovitis, we utilized 2-color immunohistochemistry of frozen tissue sections. Cells staining positive for APRIL were localized at the outskirts of B cell follicles, typically in the T cell zones. Double immunofluorescence demonstrated that APRIL-producing cells expressed CD83, a marker for activated DCs (37) (Figure 2A). In all types of synovitis, the overlap between CD83+ and APRIL+ cells was complete. BLyS-specific antibodies reacted with a subset of large cells distributed throughout the tissue infiltrates. Double labeling using the macrophage marker CD68 and BLyS confirmed that BLyS-producing cells were tissue macrophages (Figure 2B). CD83+ DCs were negative for BLyS protein.

**APRIL/BLyS receptors in rheumatoid synovitis.** The biologic effects of APRIL and BLyS in rheumatoid synovitis should be dependent on which cell types express the relevant receptors. To assess receptor expression in synovitis, the 72 tissue samples were analyzed for the representation of BAFF-R, BCMA, and TACI-specific sequences by real-time PCR. Transcripts for all 3 receptors were detected in all tissue biopsies (Figure 3). Expressions of BAFF-R and TACI were independent from the synovitis type. Overall, transcripts for BAFF-R were more abundant than those for TACI, but no correlation emerged between the lymphoid microstructure and expressions of these 2 receptors. Conversely, BCMA mRNA was expressed differently, with the highest levels in tissues with organized T cell and B cell clusters. GC+ synovitis and aggregate synovitis both contained significantly higher concentrations of BCMA-specific transcripts than did diffuse synovitis (P < 0.001 and P = 0.006 compared with diffuse synovitis). Quantities of BCMA mRNA were not significantly different in synovial lesions with GC reactions versus aggregate formations (P < 0.1).

**TACI+ T cells in RA tissue lesions.** Whereas BCMA is mostly expressed on B cells, TACI also regulates T cell function (28). Also, TACI−/− mice develop a fatal lymphoproliferative syndrome, suggesting that APRIL and BLyS have immunoregulatory effects beyond supporting B cell differentiation and antibody production. To explore whether APRIL and BLyS have multiple distinct functions in rheumatoid synovitis, we set out to identify TACI+ T cells in the tissue lesions. In all types of rheumatoid synovitis, TACI+ cells were represented in the synovial infiltrates, but the composition of the TACI-expressing population was distinct in GC+ synovitis compared with aggregate and diffuse synovitis. In tissues with typical GC reactions, TACI+ cells were found outside of the follicles and accumulated in 2 distinct areas. They were either lined up in the sublining stroma below the layers of hyperplastic follicles or localized in the interfollicular regions. Dual-color immunohistochemistry of frozen tissue sections demonstrated that these TACI-expressing cells were negative for CD3. A subset of the TACI+ cells in the interfollicular areas stained positive for CD20. The majority of TACI+ cells in GC+ synovitis were identified as plasma cells based on characteristic cellular morphology and expressions of CD138 and VS38c (Figure 4C).

In aggregate synovitis, TACI-expressing cells were localized in the T cell–B cell clusters; in diffuse synovitis, they were distributed throughout the infiltrates (Figure 4A). Double staining with anti-CD3, anti-CD20, and anti-CD138 revealed that T cells, B cells, and plasma cells contributed equally to the TACI+ population. All diffuse and aggregate synovitis tissue sections examined had CD3+ T cells expressing the TACI receptor (Figure 4B). Thus, the representation of TACI+ T cells in rheumatoid lesions correlated closely with the type of lymphoid organization and was typical for synovitis variants with T cell–B cell aggregates and unorganized infiltrates.

To confirm the presence of TACI+ T cells in diffuse and aggregate synovitis and to estimate the frequencies of such T cells in the tissue lesions, we proceeded with multicolor flow cytometric anal-
ysis of cells isolated freshly from tissue biopsies. Synovial tissue samples from 3 patients with aggregate synovitis and 3 patients with diffuse synovitis were digested, and lymphocytes were isolated by density gradient and stained with anti-TACI combined with either anti-CD3, anti-CD20, or anti-CD138. Almost all CD138+ plasma cells were positive for TACI. A subset of about 6–10% of tissue-derived CD3+ T cells expressing TACI on the surface was found regardless of whether the tissues had aggregates or diffuse infiltrates. Representative results obtained from a tissue with classical aggregate synovitis are shown in Figure 5. CD20+ B cells that reacted with anti-TACI antibodies were consistently found, but their frequency varied from patient to patient.

Blockade of APRIL and BlyS in synovitis with GC formation. The production of APRIL and BlyS in all types of rheumatoid synovitis and the presence of tissue-infiltrating TACI+ T cells in selected variants of the rheumatoid lesions provided an experimental system to test the in vivo functions of the 2 TNF homologs in RA. We generated human synovium-SCID mouse chimeras by implanting pieces of synovial tissues collected from patients with active disease into NOD-SCID mice. In this model, full engraftment of the human tissue is reached after 6–7 days. On days 6, 9, and 12, the synovium-SCID chimeras were injected with 200 μg of the soluble decoy receptor TACI:Fc. Control animals received human Fc protein lacking TACI.

Treatment with soluble TACI had profound effects in tissues with ectopic GC formation. TACI:Fc efficiently inhibited IgG1/2 production about 13-fold (P < 0.001) and decreased tissue production of IFN-γ by almost 4-fold (P < 0.05) (Figure 6).

To explore how blockade of APRIL and BlyS affected the architecture of synovial follicles, we collected the synovial grafts 2 weeks after TACI:Fc injections and stained tissue sections for the T cell marker CD3 and B cell marker CD20. Treatment with TACI:Fc resulted in the destruction of ectopic GCs, combined with a marked reduction of CD20+ B cells and CD3+ T cells (Figure 7). B cells almost disappeared from the lesions, and T cell densities declined to less than half of the control densities (Figure 7B). To analyze whether the collapse of ectopic GCs was associated with the loss of FDC, cDNAs from the synovial tissues were assayed for the presence of CD21 long isoform (CD21L) transcripts, a marker abrogating CD21L transcription, confirming that FDC networks were no longer maintained (Figure 7C). These findings linked APRIL and BlyS to the process of lymphoid organogenesis with roles in GC formation and regulatory function for B cells and T cells in the rheumatoid lesion.

Blockade of APRIL/BlyS in aggregate and diffuse synovitis. Both aggregate and diffuse synovitis were characterized by the inclusion of TACI+ T cells in the infiltrates. We examined the effects of APRIL and BlyS blockade in tissues from 5 patients with GC− T

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**Figure 3**

Expression of APRIL and BlyS receptors in rheumatoid synovitis. Tissue samples (n = 72) were stratified according to synovitis pattern. cDNAs from tissue extracts were normalized for the concentration of β-actin copies, and BCMA, BAFF-R, and TACI transcripts were analyzed by real-time PCR. All 3 receptors were detected in all tissue samples. Quantities of BAFF-R and TACI transcripts were similar in all 3 types of synovitis. BCMA was produced at low abundance in tissues with diffuse synovitis and in significantly higher amounts in aggregate (P = 0.006) and GC− synovitis (P < 0.001). Results are given as box plots as described in Figure 1.

**Figure 4**

Topography and phenotype of TACI-expressing cells in rheumatoid synovitis. (A) Sections of synovial tissues with GC− (left), aggregate (middle), and diffuse (right) synovitis were stained with anti-TACI Abs. CD3+ T cells and CD20+ B cells forming the GC are marked in the insets. TACI-expressing cells (brown staining) in GC− synovitis were positioned in the interfollicular areas (left). TACI+ cells in aggregate tissues mapped preferentially to the lymphoid clusters (middle). In diffuse synovitis, TACI+ cells were dispersed throughout the tissue (right). (B) In GC− synovitis, TACI+ cells (brown staining) were negative for the CD3 marker (blue staining). In contrast, tissues with aggregate (middle) and diffuse synovitis (right) contained TACI+ cells (blue staining) that expressed CD3 (brown staining). The insets show higher magnifications of representative TACI+ cells. DAB, 3,3′-diaminobenzidine tetrahydrochloride; VB, VectorBlue. (C) The TACI receptor (blue staining) was intensively expressed on a subset of CD138+/VS38c+ plasma cells (brown staining). Immunohistochemistries shown are representative of 15 experiments. Original magnification, ×600 (insets in B and C). Scale bars: 200 μm.

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**Figure 5**

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**Figure 6**

Blockade of APRIL/BlyS in synovitis with GC formation. The production of APRIL and BlyS in all types of rheumatoid synovitis and the presence of tissue-infiltrating TACI+ T cells in selected variants of the rheumatoid lesions provided an experimental system to test the in vivo functions of the 2 TNF homologs in RA. We generated human synovium-SCID mouse chimeras by implanting pieces of synovial tissues collected from patients with active disease into NOD-SCID mice. In this model, full engraftment of the human tissue is reached after 6–7 days. On days 6, 9, and 12,
cell–B cell clusters and in 5 patients with diffuse synovitis. The biologic consequences of inhibiting APRIL and BLyS in these synovitis variants were clearly distinct from the outcome in GC synovitis (Figures 8 and 9). Injection of TACI:Fc had essentially no effects on the transcription of Ig. In both aggregate and diffuse synovitis, transcription of IgG continued despite the disruption of APRIL and BLyS function. Instead, regulation of T cell activity was the primary target of the TNF homologs. In aggregate-containing synovial grafts from TACI:Fc-treated chimeras, mRNA levels of the T lymphocyte activation marker IFN-γ were markedly increased (P < 0.001), and upregulation of IFN-γ mRNA levels increased by approximately 6-fold (P < 0.001) in cases with diffuse synovitis. The quantification of the proinflammatory cytokine TNF-α revealed a significant reduction (P < 0.02) in aggregate synovitis and unchanged levels in tissues with diffuse infiltrates (P = 0.4) (Figures 8 and 9).

Discussion

Ligands of the TNF superfamily are known to critically regulate immune responses and immune-mediated tissue injury. BLyS is now considered to be one of the major factors regulating the size and repertoire of the B cell compartment (39) and thus emerges as a potential driver of B cell hyperplasia and autoantibody production in human autoimmune disease. The contribution of APRIL in B cell autoimmunity is less well understood (40), but both of these TNF homologs have been implicated in breaking tolerance and mediating autoimmunity in mice (18, 19, 41). The data presented here demonstrate that APRIL and BLyS have positive as well as negative regulatory effects in RA, a disease in which B cells appear to play an indispensable role. Absorption of APRIL and BLyS through treatment with the TACI:Fc fusion protein had deleterious effects on the structural and functional integrity of synovial GCs, thereby assigning a role for APRIL and BLyS in the process of inflammation-associated lympho-organogenesis. A complex picture of treatment-related effects was found in tissues that lacked ectopic GCs and possessed TACI+ T cells. Here, preventing APRIL and BLyS signaling through soluble TACI receptors led to increased tissue production of the proinflammatory cytokine IFN-γ.

Thus, BLyS and APRIL regulate synovial inflammation in RA, with negative signals being mediated through TACI+ T cells. TACI+ T cells could possibly be developed into a new therapeutic strategy aimed at downregulating rheumatoid synovitis.

Soon after the discovery of BLyS, it became clear that this pivotal survival factor for B cells derived from myeloid cells such as monocytes, macrophages, and DCs (15). This finding assigned primary control of the B cell compartment size and repertoire to professional APCs. However, recent data suggest that the expression of the relevant receptors on B cells may be regulated through signals mediated by the B cell receptor, thereby implicating antigenic stimulation in this critical aspect of the immune system (42).

The rule that BLyS is produced by myeloid cells was maintained in rheumatoid synovitis. BLyS transcription was highly abundant; however, only a subpopulation of CD68+ macrophages had the capability to produce BLyS protein, and CD83+ DCs were negative. We did not find any distinguishing characteristic features for BLyS-producing macrophages. Their morphology was that of typical tissue macrophages; they did not resemble synovocytes. Their location in the tissue appeared random, and all distinct types of synovitis had BLyS+ macrophages. In this context, it is important to note that synovitis is characterized by the lack of tissue-residing neutrophils, a cell population recently described as being an important producer of BLyS (14). In contrast, neutrophils are often abundantly found in synovial fluid and are possibly the source for the elevated BLyS levels found in the synovial fluid of RA patients (21). It is doubtful that BLyS released in the synovial space would have a major impact on the function of cells deeply embedded in the hyperplastic synovial membrane. BLyS-mediated functions in the synovial infiltrates are likely controlled through macrophage-derived proteins.

A similar selective picture emerged for the cellular origin of APRIL. The protein was exclusively detected in CD83+ DCs. Biologic active heterotrimers of APRIL and BLyS have been previously described (27); however, since these cytokines are supplied from different cellular sources, this mechanism is unlikely to have relevance in synovitis. The density of DCs in synovitis is highest in tissues with ectopic GCs, and DCs in the T cell zones of such GCs were typically positive for APRIL. The profound effects that TACI:Fc had in dismantling such highly organized GCs raises the possibility that DCs, through the release of APRIL, influence lymphoid organogenesis.

The current study identified 2 distinct biologic effects of APRIL and BLyS in RA. Aberrant GCs ceased to exist, B cells and T cells effluxed from the lesions, and Ig production decreased when APRIL and BLyS were blocked with TACI:Fc. BLyS has been implicated in regulating GC function (43). This effect is likely mediated through...
the BLyS receptor, BCMA, which is preferentially expressed on follicular B cells. We did not see expression of TACI on any cell type directly associated with GCs. The essential role of APRIL/BLyS for GC function is well in line with previous reports. The high sensitivity of IgG transcriptions to TACI:Fc treatment may relate to a high frequency of short-lived plasma cells that are highly dependent on a functional GC. It is less likely that TACI:Fc has a direct effect on plasma cell function. The inhibitory action of TACI:Fc on T cell and macrophage function could result from impaired antigen presentation and T cell triggering. Previous adoptive transfer experiments of tissue-derived T cells into SCID chimeras have shown that the functional activity of such T cells depends strictly on B cells, which suggests that B cells in rheumatoid synovitis serve as critical APCs.

The more surprising outcome of the current study relates to the finding that in the majority of rheumatoid synovitis cases, TACI:Fc did not block — but rather enhanced — inflammatory activity. In tissues that did not possess ectopic GCs, such as in aggregate and diffuse synovitis, absorbance of APRIL and BLyS led to enhanced functional activity of such T cells depends strictly on B cells, which suggests that B cells in rheumatoid synovitis serve as critical APCs.

Whereas BCMAs display specificity for B cells, BAFF-R and TACI can also be expressed by T cells (32, 45). In circulating T cells, TACI is infrequently seen, and polyclonal stimulation is necessary to bring the receptor to the surface of some T cells. TACI+ T cells in the tissue lesions had a preference for the T cell–B cell aggregates and were distinctly absent in those tissues that had formed organized GCs.

Figure 7
APRIL/BLyS blockade results in collapse of ectopic GCs. GC+ synovium was implanted into SCID mice, and the chimeras were treated with TACI:Fc or control Fc as described in Figure 6. Explanted synovial grafts were embedded, and frozen tissue sections were stained for CD3 and CD20. Tissue extracts were assayed by PCR for the presence of CD21L transcripts, a marker for FDCs exclusively expressed in synovitis with intact GCs. (A) Treatment of GC+ synovium with TACI:Fc resulted in the destruction of ectopic GCs. Scale bars: 50 µm. (B) In synovial grafts treated with TACI:Fc, few CD20+ B cells remained, and CD3+ T cells were markedly reduced. HPF, high-powered field. (C) TACI:Fc injection caused loss of CD21L, confirming that FDC networks were no longer sustained. First lane, size markers. +, positive control; –, negative control.
In mice, the depletion of TACI has been associated with massive cells infiltrating diffusely in organs, lymphoma development, autoabnormalities in the B cell compartment, including hyperplastic B downregulatory functions of APRIL and BLyS dominate over their of such negative signals is T cells or B cells has not yet been explored activity for T cells (46), quite opposite to the in vivo effects seen here. It has been proposed that APRIL and BLyS may have costimulatory activity for T cells (46), quite opposite to the in vivo effects seen here. For patients in whom the demasking this inhibitory function of APRIL/BLyS through block role in sustaining B cells and their organization in the synovium, have shown that the patterning of synovitis is stable over time and sets of producer cells in this synovitis type, such that APRIL or BLyS contributed to TNF-α induction while suppressing IFN-γ through targeting TACI- T cells. Results presented here extend the role of the TNF homologs BLyS and APRIL in human autoimmune disease. BLyS serum levels are significantly increased in patients with RA (21), SLE (47), or SS (22). A recent report found elevated serum levels of APRIL in SLE patients compared with normal controls and patients with RA (48). A contributory role of BLyS and APRIL in promoting B cell survival, hyperplasia, and autoantibody production can be easily imagined. To conceptualize on this model, clinical trials aimed at disrupting BLyS activity in patients with SLE and RA have been started. Data on clinical efficacy have not yet been published, but preliminary results suggest that autoantibody levels are only partially affected. Based on the results of blocking APRIL and BLyS in RA tissue lesions, we propose that the 2 mediators may have additional pathways through which they interfere with the risk for and the severity of autoimmune disease. By triggering TACI- T cells, APRIL/BLyS seem to provide a negative signal, suppressing production of the key T cell cytokine IFN-γ. The suspicious absence of TACI- T cells in all cases in which tissue-invading T cells and B cells organized in GC reactions suggests that TACI- T cells may be capable of stopping certain pathways. Specifically, they may disrupt the formation of very stable extralymphoid architectures, which facilitate breakdown of tissue tolerance through optimizing antigen capture, storage, and recognition and generating memory responses. Our study also emphasizes the complexity of B cell function in rheumatoid synovitis. The clinical efficacy of B cell depletion via treatment with anti-CD20 antibodies has identified B cells as excellent targets in immunomodulatory therapy. The difference in therapeutic outcomes when GC- and GC- synovitis was treated with TACI-Fc supports the notion that the regulation of B cell function in RA is complex and will require choosing therapeutic targets and interventions that are individualized for each patient.

**Figure 8**
Therapeutic effects of TACI:Fc in aggregate synovitis. Human synovium–SCID mouse chimeras implanted with synovial tissues with aggregates were treated with TACI:Fc or control Fc and then analyzed as described in Figure 6. Results from experiments with tissues from 5 patients are shown. Blocking APRIL/BLyS left production of IgG1/2 transcripts totally unaffected. Following treatment with TACI:Fc, IFN-γ production markedly increased. DC function assessed through CCL19 remained unchanged. Transcription of TNF-α was significantly diminished. Results are given as box plots, as described in Figure 1.

**Figure 9**
Treatment with TACI:Fc increases T cell activation in diffuse synovitis. Synovial tissues from 5 patients with diffuse synovitis were implanted into SCID mice. Chimeric mice were treated with TACI:Fc and control Fc, and explanted tissue was analyzed as described in Figure 6. Neutralization of APRIL/BLyS in diffuse synovitis markedly increased IFN-γ transcripts, whereas productions of IgG1/2, CCL19, and TNF-α mRNA were unaffected. Results are given as box plots, as described in Figure 1.
Table 1
Clinical characteristics of the study population

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<th>Patients (n)</th>
<th>Synovitis type</th>
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<th>Aggregate</th>
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<td>Sex (% female)</td>
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<td>16</td>
<td>30</td>
<td>26</td>
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<tr>
<td>RA duration (years; mean ± SD)</td>
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<td>Age at disease onset (years; mean ± SD)</td>
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<tr>
<td>RF (% positive)</td>
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<td>84</td>
<td>81</td>
<td></td>
</tr>
<tr>
<td>No. joint surgeries</td>
<td>6.1 ± 3.5</td>
<td>4.2 ± 3.3</td>
<td>5.2 ± 3.7</td>
<td></td>
</tr>
</tbody>
</table>

Extra-articular disease
Nodules (%) | 47 | 35 | 32 |
Major organ disease (%) | 7 | 17 | 12 |

RF, rheumatoid factor.

Methods

Study population. Synovial tissue specimens were collected from a cohort of 72 patients with RA who were undergoing synovial biopsy, synovectomy, or total joint replacement surgery. All patients were diagnosed according to the 1987 revised criteria of the American College of Rheumatology, were seropositive for rheumatoid factor, and had unequivocal destructive RA, and had active disease at the time of surgery. Most of the patients were on methotrexate as a disease-remitting agent; none of the patients were on anti-TNF antagonists. Demographic data for the study population is given in Table 1. The protocol was approved by the Mayo Clinic Institutional Review Board and the Emory University Institutional Review Board.

Histopathologic evaluation of tissue specimens. Hematoxylin-stained sections of the synovial tissue were analyzed for the cellular organization of inflammatory infiltrates, with particular attention paid to the lymphoid microstructure and the arrangement of T cells, B cells, and macrophages. All analyses of tissue specimens were performed by one hematopathologist (P.J. Kurtin) who was unaware of the clinical course or any laboratory findings. Synovial tissue specimens were grouped according to the following criteria: (a) T cell and B cell follicles with GC formations, (b) T cell and B cell aggregates without GCs, and (c) diffuse infiltration of T cells and B cells with the absence of a distinct lymphoid microstructure. GC formation was confirmed by standard histological criteria (49) and PCR analysis for the CR2/CD21L expressed on FDCs (38). This histological classification has proven useful for us in pathogenetic studies. However, it has not yet been validated in clinical studies. In particular, its importance for clinical outcome and treatment responses is unclear.

Immunohistochemistry. Frozen human synovial tissues embedded in OCT were cut into 5-μm sections, fixed in acetone for 10 minutes, air dried, and fixed in 1% paraformaldehyde/EDTA (pH 7.2) for 3 minutes. Endogenous peroxidase activity was blocked with 0.3% H2O2 in 0.1% sodium azide, and endogenous alkaline phosphatase activity was inhibited by the addition of levamisole (Vector Laboratories). Nonspecific binding was blocked with 5% normal blocking serum from the appropriate species for 15 minutes. The following primary Abs were used for immunohistochemistry: anti-human CD3 mAb (1:100 dilution), anti-human CD20 mAb (1:100 dilution), anti-CD138 mAb, and anti-human plasma cell p63 Ab (all from DakoCytomation); anti-human CD3 mAb (1:100 dilution); anti-human CD20 mAb (1:100 dilution); anti-human p63 Ab (all from DakoCytomation); and anti-human TACI mAb (1:5 dilution; R&D Systems). Primary antibodies were incubated either for 60 minutes at room temperature or overnight at 4°C. After incubation and washing, secondary species-specific Abs (all at 1:400 dilution; DakoCytomation) were applied for 30 minutes at room temperature followed by detection with either the streptavidin-biotin complex immunoperoxidase (VECTASTAIN Elite ABC Kit; Vector Laboratories) or alkaline phosphatase technique (VECTASTAIN ABC-AP Kit; Vector Laboratories). Then the slides were developed with 3,3′-diaminobenzidine tetrahydrochloride (DAB; DakoCytomation) substrate for 3 minutes or alkaline phosphatase substrate kit III (VectorBlue; Vector Laboratories) for 30 minutes and counterstained with hematoxylin for 10 seconds. For double labeling, the slides were incubated with a second primary Ab for 1 hour and developed using Vector ABC-AP reagents (Vector Laboratories) and alkaline phosphatase kit I (VectorBlue; Vector Laboratories) for digital fluorescent visualization. After the final washes, the slides were mounted in VectaMount permanent mounting medium and examined under a light microscope. Control sections were treated in parallel with nonimmune isotype-matched Ig substituting for the primary antibody.

Flow cytometry of tissue-derived cells. Lymphocyte isolation from synovial tissue has been described previously (50). Fresh synovial tissues were digested with 0.05 M HEPES buffer, 3 mg/ml type I collagenase, 1 mg/ml hyaluronidase, and 0.1 mg/ml type IV deoxyribonuclease I at 37°C for 30–45 minutes. Lymphocytes were isolated by Ficoll-Hypaque centrifugation and stained with PerCP-labeled anti-CD3 mAb, FITC-labeled anti-CD20 mAb, and APC-labeled CD138 mAb (BD). For TACI staining, synovial lymphocytes were incubated with mouse anti-TACI mAb (32); PE-labeled goat anti-mouse IgG was used as secondary Abs. For all staining experiments, control stains with isotype-matched Abs were included. Stained cells were analyzed on a FACSCalibur flow cytometer (BD), and WinMDI was used for analysis (Scripps Research Institute).

Table 2
Characteristics of PCR primers

<table>
<thead>
<tr>
<th>Target</th>
<th>Sequence (5′−→3′)</th>
<th>Annealing temperature (°C)</th>
</tr>
</thead>
<tbody>
<tr>
<td>APRIL</td>
<td>5′-AGGGTGAGGAGGTCACAGGC-3′</td>
<td>55</td>
</tr>
<tr>
<td>Bls-S</td>
<td>5′-GGAGCTGAGGATCAAATTCTC-3′</td>
<td>55</td>
</tr>
<tr>
<td>LTβ</td>
<td>5′-CTGGAGAAAAACAAAATGAGT-3′</td>
<td>60</td>
</tr>
<tr>
<td>IgG1z</td>
<td>5′-ACTCCAGGACCCTTCCTTGTTCTC-3′</td>
<td>55</td>
</tr>
<tr>
<td>BCMA</td>
<td>5′-GCTTCTCGGAGTATGGTGTT-3′</td>
<td>55</td>
</tr>
<tr>
<td>TACI</td>
<td>5′-CTAGGCTCTGTCCTGTGTTCT-3′</td>
<td>55</td>
</tr>
<tr>
<td>BAF-R</td>
<td>5′-AGACAAAGGAGGGCCAGACGCCC-3′</td>
<td>55</td>
</tr>
<tr>
<td>IFN-γ</td>
<td>5′-ACCTCAAGAATTATTTTATGTC-3′</td>
<td>55</td>
</tr>
<tr>
<td>TNF-α</td>
<td>5′-CTTTGGGATCTGCTGGTTGTC-3′</td>
<td>60</td>
</tr>
<tr>
<td>CCL19</td>
<td>5′-ACCAATGACTGCTGGTAC-3′</td>
<td>55</td>
</tr>
<tr>
<td>CD21L</td>
<td>5′-GGGGTGTATGGTGTTGTC-3′</td>
<td>55</td>
</tr>
<tr>
<td>β-Actin</td>
<td>5′-ATGCCGACGCTCTCCACGACG-3′</td>
<td>55</td>
</tr>
<tr>
<td>CD138</td>
<td>5′-CATGGTGTCGCGGCGAAGCAC-3′</td>
<td>55</td>
</tr>
</tbody>
</table>
PCR analysis of tissue transcripts. Total RNA was extracted from synovial tissue specimens using TRIzol reagent (Invitrogen Corp.), and cDNA was synthesized. The cDNA was amplified by multiple real-time quantitative PCR (Mx4000; Stratagene) in the presence of SYBR green dye under nonsaturating conditions with cytokine-specific primers. Primer sequences of analyzed cytokines are shown in Table 2. PCR reactions were validated using gene-specific sequences with known copy numbers. Amplification cycles included initial denaturation at 95°C for 30 seconds, annealing at primer-specific temperature for 1 minute, and polymerization at 72°C for 1 minute, with 5 minutes’ initial denaturation at 96°C and a final 1-minute extension at 95°C. Serial dilutions of known amounts of gene-specific sequences were amplified in parallel. The number of cytokine-specific sequences was determined by interpolation on a standard curve and expressed as the number of cytokine transcripts per 2 × 10⁴ copies of β-actin to adjust for equal amounts of cDNA.

Treatment of human synovium–SCID mouse chimeras with TACI:Fc mAbs. NOD.CB17-PrkdcscidJ mice were used at the ages of 6–9 weeks (Jackson Laboratory). The protocol was approved by the Animal Care and Use Committee of the Mayo Clinic. For tissue implantation, the mice were anesthetized with i.p. 50 mg/kg pentobarbital (Abbott Laboratories) and methoxyflurane (Medical Developments International Ltd.) inhalation. Pieces of human synovium–SCID mouse chimeras were injected i.p. with either a single 200-μg dose of TACI:Fc or 200 μg of IgG on days 6, 9, and 12. Fourteen days after implantation, the mice were sacrificed, and the synovial tissue graft was harvested and either embedded in OCT compound (Tissue-Tek; Sakura Finetek Co.) for immunohistochemical analysis or shock frozen in liquid nitrogen (−160°C) for RNA extraction and cytokine transcript analysis.

Statistics. Analyses, plotting, and data graphing were performed using SigmaStat version 3.0 and SigmaPlot version 8.0 software (SPSS Inc.). Experimental groups were compared by nonparametric Mann-Whitney rank sum tests. P values less than 0.05 were considered to be significant.

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ing TACI as an inhibitory BLyS receptor. *Immunity.*


