Generation and characterization of tabalumab, a human monoclonal antibody that neutralizes both soluble and membrane-bound B-cell activating factor

Abstract: B-cell activating factor (BAFF) is a B-cell survival factor with a key role in B-cell homeostasis and tolerance. Dysregulated BAFF expression may contribute to autoimmune diseases or B-cell malignancies via effects on abnormal B-lymphocyte activation, proliferation, survival, and immunoglobulin secretion. Monoclonal antibodies were generated against human BAFF, characterized for species specificity and affinity, and screened for the ability to neutralize both membrane-bound and soluble BAFF. In addition, studies were undertaken to determine the relative potency of membrane-bound and soluble BAFF. Tabalumab has a high affinity for human, cynomolgus monkey, and rabbit BAFF. No binding to mouse BAFF was detected. Tabalumab was able to neutralize soluble human, cynomolgus monkey, or rabbit BAFF with equal potency. Our data demonstrate that membrane-bound BAFF can be a more potent stimulus for B-cells than soluble BAFF, and tabalumab also neutralized membrane-bound BAFF. Tabalumab prevented BAFF from binding to BAFF receptors and demonstrated pharmacodynamic effects in human BAFF transgenic mice. Tabalumab is a high-affinity human antibody with neutralizing activity against membrane-bound and soluble BAFF. Given our findings that membrane-bound BAFF can have greater in vitro potency than soluble BAFF, neutralization of both forms of BAFF is likely to be important for optimal therapeutic effect.

Keywords: autoimmunity, B-cell malignancies, B-cell survival factor, BAFF

Introduction

B-cell activating factor (BAFF) is a member of the tumor necrosis factor superfamily (TNFSF); it is also known as BLYS, THANK, TALL-1, zTNF4, and TNFSF13b. Like other members of the TNFSF, BAFF is a type II transmembrane protein that naturally forms homotrimers to make a biologically active protein. BAFF may be found in two forms: a membrane-bound form and a soluble form released from the cell surface by an unknown protease. Similar to other family members, an initial report suggested both forms of BAFF are biologically active, but it is unknown whether these two forms stimulate B-cells with equal potency.

Early literature characterizing BAFF expression indicated that it was produced predominantly by monocytes, macrophages, and dendritic cells. However, additional research has found that many other cell types can produce BAFF, including neutrophils, bone marrow stromal cells, follicular dendritic cells, osteoclasts, and some T-cells depending on the physiological conditions or disease state.
BAFF exerts its activity by binding three different cell surface receptors. The expression of BR3 (also known as BAFF-R or TNFRSF13c) is primarily restricted to the B-cell lineage and is first expressed on immature/transitional B-cells, continuing through subsequent B-cell developmental stages. In addition, some activated/memory subsets of T-cells express BR3. TACI (transmembrane activator and calcium modulator and cyclophilin ligand interactor, also known as TNFRSF13b) expression is highest on type 2 transitional (T2) and marginal zone B-cells with modest expression on follicular B-cells and undetectable levels on germinal center B-cells. BCMA (B-cell maturation antigen, also known as TNFRSF17) appears to be almost exclusively expressed on plasma cells. TACI and BCMA also bind a proliferation inducing ligand (APRIL), BAFF’s closest homologue. B-cells may express one or more of these receptors at the same time; how the receptors influence each other’s signaling is still unknown.

BAFF plays a major role in B-cell homeostasis and peripheral tolerance. Mice devoid of BAFF or BR3 are profoundly deficient in mature B-cells demonstrating that BAFF is required for B-cell maturation after the transitional stage. BAFF acts as a survival factor for immature/transitional B-cells as they leave the bone marrow and enter the periphery. Using in vitro assays, BAFF has been shown to provide survival signals to prevent apoptosis of normal B-cells. In addition, BAFF stimulation protects malignant B-cells from apoptosis, including lymphoma, chronic lymphocytic leukemia, and multiple myeloma (MM). Importantly, BAFF stimulation rescues MM cells from dexamethasone-induced growth inhibition and cell death by signaling through both the canonical and non-canonical NF-κB pathway ultimately upregulating anti-apoptotic genes.

BAFF can also act as a co-stimulatory molecule for B-cells that have been activated through the B-cell receptor. Additionally, BAFF can assist B-cells as they undergo class switching to immunoglobulin G (IgG). Administration of exogenous BAFF to mice in conjunction with T-cell-independent or T-cell-dependent antigens can potentiate antibody responses and increase the number of plasma cells. BAFF can also act as a co-stimulatory molecule for B-cells that have been activated through the B-cell receptor. Additionally, BAFF can assist B-cells as they undergo class switching to immunoglobulin G (IgG). Administration of exogenous BAFF to mice in conjunction with T-cell-independent or T-cell-dependent antigens can potentiate antibody responses and increase the number of plasma cells.

Under physiologic conditions, autoreactive B-cells must compete with other B-cells for survival factors such as BAFF. Overexpression of BAFF, as seen in transgenic (Tg) mice, results in an increase in peripheral B-cells and survival of self-reactive B-cell clones that would normally have been deleted. Depending on the background strain, BAFF Tg mice can develop signs of autoimmune disease with age, such as autoantibodies and immune complex deposition in the kidneys. In addition, autoreactive B-cells have been shown to have an increased dependence on BAFF for continued survival.

Elevated BAFF levels have been observed in patients with autoimmune diseases and B-cell malignancies. In systemic lupus erythematosus (SLE), multiple groups have demonstrated elevated serum BAFF levels, and these have been correlated with autoantibody production and disease activity. Monocytes from SLE patients have increased membrane-bound BAFF expression. In rheumatoid arthritis (RA), high BAFF levels were detected in synovial fluid (SF). In a study with matched serum and SF samples, BAFF levels were higher in SF suggesting a local production.

Therapeutic intervention in the BAFF pathway has been shown to provide clinical benefit in SLE. Belimumab, a human IgG1 antibody that binds and neutralizes BAFF, provided clinical benefit to SLE patients and is approved for the treatment of SLE. Belimumab has been shown to bind soluble BAFF with high affinity, but does not recognize membrane-bound BAFF. Because of the evidence suggesting a role for BAFF in the pathophysiology of multiple autoimmune diseases and B-cell malignancies, monoclonal antibodies were developed that neutralize both membrane-bound and soluble BAFF as a potential therapeutic approach to these diseases.

**Materials and methods**

**Immunizations and hybridomas**

Monoclonal antibodies were generated using the HuMAbMouse™ technology at Medarex (Princeton, NJ, USA) by immunizing mice with soluble human BAFF (amino acids 133–285; Research Diagnostics Inc., Flanders, NJ, USA). Both the HCo7 and HCo12 mice were used. Mice were immunized with 15 µg to 50 µg soluble human BAFF in RIBI adjuvant (RIBI ImmunoChem Research, Inc, Hamilton, MT, USA), Freund’s complete adjuvant or Freund’s incomplete adjuvant. Eight mice producing serum antibody titers to human BAFF were injected intravenously with 10 µg human BAFF in phosphate buffered saline (PBS). The spleen was harvested 3 days later from each mouse and fused with myeloma cells according to the method described by Zola. Hybridomas were tested for specific binding to human BAFF and to make sure they were expressing human immunoglobulin heavy and light chains by enzyme-linked immunosorbent assay (ELISA).

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Microtiter plate wells were coated with either human APRIL, human BAFF, or mouse BAFF overnight at 4°C.
The plates were washed twice with wash buffer and blocked with casein/PBS. Tabalumab and positive control antibodies were diluted in blocking solution, plated and incubated for 1.3 hours at room temperature (RT). The plate was washed three times and then the appropriate anti-IgG (Fc specific)-HRP secondary was dispensed and incubated at RT for 1 hour. The plate was washed three times and developed using an o-Phenylenediamine dihydrochloride (OPD; Sigma-Aldrich Co, St Louis, MO, USA) substrate. The substrate reaction was stopped with 1N HCl and the absorbance 490 was read on a Molecular Devices (Sunnyvale, CA, USA) plate reader.

Here we describe the characteristics of one of these antibodies, tabalumab (ta bal’ue mab) or LY2127399.

**Purification of antibodies**

Cell culture media of subcloned hybridomas was concentrated in Amicon ProFlux M12 tangential filtration systems using an Amicon S3Y30 UF membrane (EMD Millipore, Billerica, MA, USA). The concentrated media was passed over protein-A Sepharose columns (5 to 20 mL column) at a flow rate of 5 mL/min. The columns were washed with buffer A (PBS, pH 7.4) until the absorbance returned to baseline and the bound antibodies were eluted with 50 mM citric acid, pH 3.2. Fractions were immediately neutralized with 1M Tris, pH 8.0. Fractions were then analyzed by SDS-PAGE. Fractions containing antibody were pooled and concentrated using an Ultrafree centrifugal filter unit (EMD Millipore; 10 kDa molecular weight cut-off).

**Surface plasmon resonance**

Antibody affinities to various species of BAFF were determined using a Biacore biosensor 2000 (GE Healthcare Bio-Sciences Corp., Piscataway, NJ, USA). Protein A was coupled via free amine groups to carboxyl groups on Flow cells 1 and 2 of a CM5 biosensor chip (GE Healthcare Bio-Sciences Corp.). The antibody was captured on Flow cell 2, and BAFF was then injected over Flow cells 1 and 2 at increasing concentrations. Affinity for human, cynomolgus monkey, or rabbit BAFF was measured at 25°C using the monomeric mass of BAFF.

**Generation of cells expressing membrane-bound BAFF**

Membrane-bound BAFF is naturally cleaved from the membrane by one or more metalloproteases. To prevent this natural cleavage, an expression construct containing full length human BAFF was generated in which the cleavage site was mutated (KRAV → AAV).³ HEK293 cells were transfected with this non-cleavable BAFF construct (mem-BAFF) or with vector only (negative control). Abrogation of cleavage was confirmed by testing for soluble BAFF using an ELISA (R&D Systems Kit DBLYS0) (data not shown). In order to stimulate B-cells with the transfected cells, cells were removed from the flask using enzyme free cell dissociation media (Chemicon, Bedford, MA), washed with PBS, and either fixed with 1% paraformaldehyde, or membrane preps were prepared by lysing the cells in a hypotonic solution and ultracentrifugation.

**BAFF quantitation assay**

A competition ELISA, using an antibody to BAFF that recognizes a conformationally-dependent epitope, was used to determine the level of BAFF in the native conformation on the memBAFF cells. Recombinant human soluble BAFF was coated on microtiter plates (Greiner Bio-One, Monroe, NC, USA), 3.6 ng/well in PBS, overnight at 4°C. The ELISA plate was then washed (TBS, 0.1% Tween 20) and blocked (Casein/PBS; Thermo Fisher Scientific, Waltham, MA, USA). In a polystrene plate, serially diluted soluble BAFF, cells expressing memBAFF, or vector control cells were incubated with 4.5 ng of anti-BAFF (clone 6B2.1) for 1 hour at RT with shaking. After 2 minutes at RT, the plate was washed and the plate-bound anti-BAFF was developed using donkey, anti-Human IgG (H + L) HRP (Jackson ImmunoResearch Laboratories, Inc., West Grove, PA, USA) followed by OPD substrate. The plate was read on a Molecular Devices plate reader and the data analyzed using SoftMaxPro 3.1.2 software (Molecular Devices, Sunnyvale, CA, USA) using a 4 parameter curve fit. The soluble BAFF titration was used to establish the standard curve to quantify the number of memBAFF molecules on the cells.

**B-cell-line proliferation assay**

T1165.17 is a murine plasmacytoma cell line that requires low levels of interleukin (IL)-1β for survival, yet other cytokines can substitute for IL-1β in short-term proliferation assays.²¹ The cells were washed three times with assay media to remove IL-1β. The BAFF activity assay was performed in Roswell Park Memorial Institute medium (RPMI)-1640 supplemented with 10% fetal bovine serum (FBS), 1 mM sodium pyruvate, 5 × 10⁻⁵ M 2-mercaptoethanol, and antibiotic/antimycotic (Gibco; Thermo Fisher Scientific, Waltham, MA, USA). BAFF in either the soluble or membrane-bound form (or an equivalent amount of vector...
control membrane) was added to a 96-well plate in triplicate. T1165.17 cells were added to each well, and the plate was incubated for 44 hours at 5% CO₂, 37°C. Cell Titer 96 viability substrate (Promega Corporation, Madison, WI, USA) was added and the plate incubated for an additional hour before reading the absorbance at 490 on a Molecular Devices plate reader. To evaluate the inhibition of BAFF by tabalumab, a sub-maximal amount of BAFF was combined with tabalumab or isotype control antibody in the 96-well plate prior to adding T1165.17 cells and then following the protocol above. The data were analyzed using a four-parameter fit and plotted as a Hill-slope curve.

**Primary human B-cell assay**

Primary human B-cells were isolated from peripheral blood using CD19 positive selection with the MACS magnetic isolation system (Miltenyi Biotec, Auburn, CA, USA). The B-cells were stimulated with 10 µg/mL anti-human IgM (BD Biosciences, San Jose, CA, USA, Clone G20-127; or Jackson ImmunoResearch, 109-005-043), 10 ng/mL IL-4 (R&D Systems) and 2.3-25 ng/mL human soluble BAFF (or 5.5 ng/mL memBAFF) in the presence or absence of various concentrations of tabalumab or isotype control antibody in complete RPMI containing 10% FBS. Three or 5 days later, ³H-thymidine was added and the plate incubated for 6 more hours at 37°C in 5% CO₂. The plate was harvested using a Filtermate cell harvester and radioactivity counted using a TopCount monitor (PerkinElmer Inc., Walther, MA, USA). Incorporation of ³H-thymidine was used as the measure of proliferation.

**BAFF binding to BAFF receptors on transfected cells**

BAFF was biotinylated using Pierce NHS-LC-biotin (Thermo Fisher Scientific); HEK293 cells were stably transfected with human BR3, human TACI, or human BCMA. Expression was confirmed with specific antibody binding and flow cytometry. The cells were removed from the cell culture flask using Enzyme Free Cell Dissociation solution (EMD Millipore), resuspended at 10⁶ cells/mL in FACS buffer (D-PBS, 2% FBS, 0.1% sodium azide) with 1 mg/mL goat IgG added. For cells expressing BCMA, 300 ng/mL biotinylated BAFF + 50 µg/mL tabalumab were pre-incubated for 15 minutes at RT. For BR3, TACI, or vector control cells, 18.8 ng/mL biotinylated BAFF + 5 µg/mL tabalumab were pre-incubated for 15 minutes at RT. The mixture was added to cells and incubated for 20 minutes on ice. The cells were washed with FACS buffer and resuspended with streptavidin-PE (phycoerythrin; Jackson ImmunoResearch) at a 1:150 dilution and incubated on ice for 15 minutes. The cells were washed as above and then resuspended in FACS buffer. The fluorescence was measured using a Guava EasyCyte Plus instrument (EMD Millipore) counting 2,000 cells with the Express Plus protocol. The files were converted to FCS files and the overlays were generated using WinList 5.0 (Verity Software House, Topsham, ME, USA).

**Human BAFF transgenic mice**

For the generation of Tg mice, a DNA fragment encoding the extracellular (soluble) region of human (hu) BAFF was cloned into a liver-specific expression vector consisting of the human Apo E gene promoter and liver enhancer (provided by John Taylor, The J David Gladstone Institutes, San Francisco, CA) modified to include a DNA fragment encoding the mouse preprotrypsin signal peptide upstream of the soluble BAFF coding region. A 6.4 kb DNA fragment encompassing the Apo E promoter-huBAFF fusion gene was isolated and micro-injected into the pronuclei of FVB/N strain zygotes as previously described.²² The presence of the transgene in newborn founder mice and progeny was confirmed by polymerase chain reaction (PCR) analysis of tail or toe tissue using primers specific for the human ApoE sequences. The overexpression of soluble human BAFF in the circulation was confirmed by ELISA (R&D Systems Kit DBL YS0).

**Cell culture and flow cytometry**

Normal human monocytes were isolated by CD14 microbead separation from peripheral blood mononuclear cells (PBMCs). PBMCs were isolated from heparinized blood using Ficoll-Paque PLUS (GE Healthcare Bio-Sciences Corp). Positive selection of the monocyte population was performed by magnetic cell sorting (MACS, Miltenyi Biotec) using CD14 microbeads, according to the manufacturer’s protocol. Normal human monocytes (1 × 10⁶ cells/mL) were cultured in RPMI1640 supplemented with 10% fetal calf serum, 100 U/mL penicillin, and 100 µg/mL streptomycin. Cells were maintained at 5% CO₂ at 37°C. Cells were treated with 1 µg/mL of INFγ (R&D Systems) or media alone for 3 days. Cells were stained with the following antibodies: anti-CD14-PE (eBioscience, San Diego, CA, USA), tabalumab-Alexa 488, or isotype control-Alexa 488 (labeled in-house using Alexa488 kit from Thermo Fisher Scientific). Flow cytometric analysis was carried out using FC500 flow cytometer (Beckman Coulter, Indianapolis, IN, USA) and FlowJo software (Tree Star, Ashland, OR, USA).
Splenocytes from human BAFF Tg mice were pelleted and resuspended in cold FACS buffer (Hank’s Balanced Salt Solution [HBSS; Hyclone SH30268.02] containing 2% FBS [Thermo Fisher Scientific] and 0.1% sodium azide) at a density of 3 x 10^7 cells/mL. A 20 µL aliquot of each splenocyte sample was transferred into a 96-well plate and 10 µL of rat anti-mouse CD45R/B220-Cychrome (0553091; BD Biosciences) in FACS buffer were added to the appropriate wells for a final concentration of 1:200. The plates were sealed and incubated at 4°C for 30 minutes. The stained cells were washed with cold FACS buffer, pelleted, and resuspended in 300 µL cold FACS buffer. Data were immediately acquired on a Beckman Coulter FC500 flow cytometer and analyzed using CXP Analysis and Excel software.

Neutralization of human BAFF in transgenic mice

Mice were treated (Day 0) with either human IgG4 isotype control or tabalumab, 500 µg in 200 µL PBS delivered subcutaneously. Animals were taken off study 2, 4, 8, 11, 16 and 23 days after administration of antibody. Splenocytes were processed into a single cell suspension into cold HBSS (14025; Thermo Fisher Scientific), and then put through a 70 µm filter. The cells were pelleted by centrifugation and resuspended in cold PharmLyse (BD Biosciences) for 3 minutes. Ten mL cold HBSS was added to dilute the lysing buffer, and the samples were spun to pellet the cells. The cells were resuspended with 10 mL cold HBSS and counted. ViaCount Reagent (4000-0040; EMD Millipore) was used to determine absolute cell counts and viability of the splenocyte samples using the Guava EasyCyte Plus flow cytometry (EMD Millipore) and analyzed with CytoSoft 5.2 (EMD Millipore) and Excel software.

p52 binding assay

Plates coated with p52 binding DNA (Clontech S3528; Clontech Laboratories, Mountain View, CA, USA) were blocked for at least 15 minutes with 3% blocking solution (Clontech S2794) made in 1x Transfactor buffer (Clontech S2778). Splenocyte lysates from human BAFF Tg mice were diluted to 100 µg/mL final protein concentration with the blocking solution. The blocking solution was removed from the plate, and 50 µL of diluted sample (run in duplicate) was added to the plate and incubated at RT for 1 hour. The plate was washed three times with blocking solution. One hundred µL of a 1:1000 dilution of rabbit anti-p52 (Upstate, Lake Placid, NY, USA) was added to each well and incubated at RT for 1 hour. The plate was washed as before and 100 µL of goat anti-rabbit IgG – HRP (Jackson ImmunoResearch) at 1:2000 dilution was added to each well. The plate was incubated for 40 minutes at RT. The plate was washed three times with 1x Transfactor buffer and then 1 time with PBS. One hundred µL of substrate (Thermo Fisher Scientific) was added and the plate read on a PerkinElmer Victor3 (PerkinElmer Inc.) reader using the Wallac 1420 luminescence software (PerkinElmer Inc.).

Results

Binding and affinity of tabalumab

Tabalumab was obtained by immunization of human immunoglobulin Tg mice with soluble human BAFF resulting in many hybridomas of multiple isotypes. Initially, hybridomas were tested for binding to human BAFF and to make sure they were expressing human immunoglobulin heavy and light chains. There was no binding to APRIL, the family member with the highest homology to BAFF. Affinity measurements using surface plasmon resonance indicated that tabalumab had a high affinity for human (126 pM ± 37 standard deviation [SD]), cynomolgus monkey (101 pM ± 78 SD), and rabbit (81 pM ± 24 SD) soluble BAFF. Binding to mouse BAFF was not detected by ELISA (data not shown).

The hybridoma producing the antibody that eventually became tabalumab was a human IgG1 isotype. Since our goal was complete neutralization of BAFF without depletion of BAFF-expressing cells, we engineered the isotype to human IgG4. This should reduce interactions with Fc receptors and eliminate antibody-dependent cellular cytotoxicity and complement-dependent cytotoxicity. It is known that human IgG4 can form two half antibodies (half-mer) each with a single heavy and light chain and these can recombine in vivo. To prevent half-mer formation, a specific serine-to-proline mutation has been engineered into the hinge region of human IgG4 connecting the CH1 and CH2 of the heavy chain. This specific mutation makes the hinge more similar to that of the human IgG1 subclass, and has been shown to minimize the antibody-molecule heterogeneity. To prevent half-mer formation, a specific serine-to-proline mutation has been engineered into the hinge region of human IgG4 connecting the CH1 and CH2 of the heavy chain. This specific mutation makes the hinge more similar to that of the human IgG1 subclass, and has been shown to minimize the antibody-molecule heterogeneity. To prevent half-mer formation, a specific serine-to-proline mutation has been engineered into the hinge region of human IgG4 connecting the CH1 and CH2 of the heavy chain. This specific mutation makes the hinge more similar to that of the human IgG1 subclass, and has been shown to minimize the antibody-molecule heterogeneity.

Tabalumab was able to bind to a HEK293 cell line stably expressing non-cleavable human BAFF, or membrane-bound BAFF (Figure 1A). To confirm binding to cells naturally expressing membrane-bound BAFF, human monocytes were activated with IFN-γ, which is known to induce BAFF expression. Figure 1B shows that tabalumab binds membrane-bound BAFF on activated human monocytes.
Biological activity of BAFF

Similar to other TNFSF members, both the soluble and membrane-bound form of BAFF could be biologically active. To confirm this in vitro, a B-cell line or primary human B-cells were stimulated with either soluble human BAFF or cells expressing a non-cleavable form of membrane-bound BAFF. Data in Figure 2A show that membrane-bound BAFF is a more potent stimulus of T1165.17 cells, a murine plasma cytoma cell line, compared with soluble BAFF. In contrast, primary human B-cells responded slightly better to soluble BAFF (Figure 2B). The differences in response could be due to species’ differences or in receptor expression. T1165.17 cells strongly express TACI whereas primary human B-cells predominantly express BR3. The results demonstrate that soluble and membrane-bound BAFF were found to be biologically active.

Tabalumab neutralizes soluble and membrane-bound BAFF

Neutralization of BAFF was tested using T1165.17 cells and primary human B-cells. Using T1165.17 cells, the IC₅₀ (half maximal inhibitory concentration) for tabalumab neutralization of soluble human, cynomolgus monkey, or rabbit BAFF was 104 pM, 143 pM, and 176 pM, respectively (Figure 3). Tabalumab was tested for the ability to inhibit human B-cell proliferation induced by soluble or membrane-bound human BAFF in the presence of IL-4 and anti-human IgM. Figure 4A illustrates the ability of a dose titration of tabalumab to neutralize the effect of soluble BAFF on anti-IgM/IL-4-induced B-cell proliferation. Figure 4B demonstrates the ability of tabalumab to neutralize the membrane-bound form of BAFF compared with the soluble form of BAFF. Neutralization of cynomolgus monkey membrane-bound BAFF was also observed (data not shown).

Tabalumab inhibits BAFF binding to BR3, TACI, and BCMA

The ability of tabalumab to interfere with BAFF binding to each of the three different receptors for BAFF was investigated. HEK293 cells were transfected with either BR3, TACI, or BCMA, then incubated with BAFF in the presence of tabalumab or isotype control. As shown in Figure 5, tabalumab prevents the binding of BAFF to BR3, TACI, or BCMA on the cell surface. In contrast, once BAFF has docked into a receptor, tabalumab is unable to recognize it (data not shown). Tabalumab neutralizes the biological activity of BAFF.
Tabalumab neutralizes soluble or membrane-bound human BAFF co-stimulation of B-cells.

Notes: (A) Primary human CD19+ B-cells were co-stimulated with 25 ng/mL soluble BAFF plus anti-IgM/IL-4 in the presence of a dose response of tabalumab or isotype control. Proliferation was measured using ³H-thymidine incorporation at day 5. (B) Primary human CD19+ B-cells were co-stimulated with 2.3 ng/mL soluble BAFF or 5.5 ng/mL membrane-bound BAFF plus anti-IgM/IL-4. Gray bars represent cells treated with 1 μg/mL tabalumab, black bars represent cells treated with 1 μg/mL isotype control. Stimulation with anti-IgM/IL-4 alone produced a relatively low proliferative signal of 3640 ± 592 CPM. Proliferation was measured using ³H-thymidine incorporation at day 3. Data are presented as mean ± SD.

Abbreviations: BAFF, B-cell activating factor; CPM, counts per minute; Ig, immunoglobulin; IL, interleukin; mem, membrane-bound; SD, standard deviation.
Discussion

BAFF plays a critical role in the development and maintenance of normal B-cell populations and in numerous B-cell functions. Several lines of evidence also suggest that BAFF contributes to pathophysiology in multiple autoimmune diseases and B-cell malignancies.4

Similar to TNF, both soluble and membrane-bound BAFF are biologically active, but it is not clear whether there are distinct roles for membrane-bound BAFF compared with soluble BAFF. Despite years of research and several unique pharmacologic agents for TNF neutralization, the distinct roles for each form of TNF still remain elusive.26,27 Our data show that depending on the context, membrane-bound BAFF can be a more potent stimulus than cleaved soluble BAFF. This is consistent with a report by Bossen et al28 demonstrating that membrane-bound BAFF is at least 50-fold more active on cells expressing BCMA. In addition, we have confirmed reports that T-cells can be co-stimulated with BAFF through BR3, but only if the BAFF is immobilized rather than soluble.5 This suggests that T-cells require membrane-bound BAFF expressed by an antigen-presenting cell for co-stimulation. The fact that membrane-bound BAFF is tethered to a cell suggests that it will play a role in direct cell-to-cell communication.

The majority of soluble BAFF appears to be produced by radiation-resistant stromal cells, as lethally irradiated mice...
reconstituted with bone marrow from BAFF-deficient mice have normal levels of serum-soluble BAFF. Soluble BAFF appears to be sufficient for normal B-cell maturation. BAFF-deficient mice administered exogenous soluble BAFF can restore normal B-cell development and T-independent antibody response.

Recent data suggest that local BAFF production by T follicular helper (T<sub>FH</sub>) cells is required to promote and maintain germinal center B-cells that achieve high affinity B-cell receptor maturation. The data suggested that T<sub>FH</sub> cells express membrane-bound BAFF locally in the germinal center, although cleavage from the cell surface to produce soluble BAFF in the microenvironment cannot be ruled out.

An attempt to create mice expressing only membrane-bound BAFF was reported by Bossen et al using a transgene containing full-length mouse BAFF with two mutations to prevent cleavage, overexpressed in BAFF-deficient mice. The resulting phenotype was similar to BAFF-knockout mice, and they concluded that membrane-bound BAFF was unable to support normal B-cell development. Unfortunately, the authors were unable to demonstrate the expression of membrane-bound BAFF on any cells or tissues from these mice, a key piece of evidence required for proper interpretation of the data.

Membrane-bound BAFF expressed by fibroblast-like synoviocytes cultured from patients with RA appears to be critical for the local activation of B-cells resulting in re-expression of enzymes involved in immunoglobulin gene rearrangement, RAG1 and RAG2. The effect could be inhibited in vitro with anti-BAFF or shRNA gene interference to downregulate membrane-bound BAFF expression, however, soluble BAFF could not recapitulate B-cell induction of RAG expression. The reactivation of B-cells in the synovium, outside the context of a normal germinal center, may lead to loss of self-tolerance and further promote autoreactivity.

The most reproducible cell type found to express membrane-bound BAFF is the activated monocyte. We observed good staining of activated monocytes either using tabalumab as an intact antibody or a Fab fragment of tabalumab, precluding the possibility for Fc binding (unpublished observation). Interestingly, we were unable to see binding to resting monocytes in culture or in whole blood from healthy volunteers. Despite a few reports that B-cells express BAFF, we were not able to observe tabalumab binding to B-cells in whole blood from healthy donors, even at concentrations up to 100 μg/mL (unpublished observation). This is consistent with our data that B-cells do not express BAFF mRNA, even following activation, and that once membrane-bound BAFF expression, however, soluble BAFF could not recapitulate B-cell induction of RAG expression. The reactivation of B-cells in the synovium, outside the context of a normal germinal center, may lead to loss of self-tolerance and further promote autoreactivity.

Because tabalumab does not recognize mouse BAFF, it is not feasible to evaluate tabalumab in preclinical disease models. Tg mice expressing soluble human BAFF were used
to confirm in vivo neutralization of BAFF by tabalumab demonstrating a reduction in NF-κB signaling and a reduction in B-cells in the spleen. The increased splenocyte number in the BAFF Tg mice was a result of increased numbers of B-cells due to the proliferative/survival effect of human BAFF on murine B-cells, as demonstrated by our studies with tabalumab. The only preclinical disease model that tabalumab was tested in was an MM xenograft model that also included a human bone graft as a source of human BAFF. Treatment of the mice with tabalumab demonstrated a significant reduction in tumor burden, prolongation of survival, a decrease in osteoclast recruitment and activation resulting in less lytic lesions in the bone.

Several different BAFF antagonists with distinct mechanisms of action, and therefore potentially differing effects on patient outcomes, are being investigated in clinical trials. Belimumab is a human IgG1 monoclonal antibody that neutralizes only soluble BAFF, and it was recently approved for the treatment of SLE. The rationale for selecting an antibody to neutralize only soluble BAFF is unclear in light of the data showing expression on activated monocytes, upregulation on monocytes from SLE patients, and in vitro biological activity. In contrast, tabalumab is a human IgG4-variant monoclonal antibody that neutralizes both soluble and membrane-bound BAFF. Using a binding assay for Fc receptors and C1q, tabalumab does not bind to any of the Fcγ receptors tested (CD16a, CD32a, and CD64) nor to the complement component C1q (data not shown). These results suggest that it is highly unlikely that tabalumab will be able to induce Fc-mediated biological effects in vivo, an important consideration since our goal was complete neutralization of BAFF without depletion of BAFF-expressing cells.

Tabalumab has demonstrated the expected pharmacokinetic and pharmacodynamic responses in human clinical trials based on preclinical studies in cynomolgus monkeys. At doses ranging from 0.01 mg/kg to 8.0 mg/kg, a single intravenous dose of tabalumab exhibited nonlinear pharmacokinetics suggesting target-mediated clearance at low doses due to binding of membrane-bound BAFF. In patients with RA treated with multiple infusions of tabalumab, there was a transient increase in circulating B-cells followed by a reduction. The most sensitive population measured was naïve B-cells (CD20+ IgD+ CD27-), with an approximate 60% reduction in circulating B-cells over the 24 week study. Curiously, there was an increase in circulating memory B-cells (CD20+ IgD-CD27+) for at least 16 weeks, however, they returned to baseline by the end of the study. The mechanism responsible for the increase in circulating memory B-cells is unknown but may be related to changes in homing ability of the cells. These pharmacodynamic observations are consistent with other BAFF antagonists in clinical development, however, the clinical relevance is unknown.

Conclusion

Both membrane-bound BAFF and soluble BAFF are biologically active, suggesting that a greater clinical response might be achieved by inhibiting both forms of BAFF instead of either BAFF form alone. However, to date it has been difficult to determine if the different BAFF forms have unique, non-overlapping biological effects, so this hypothesis remains speculative. Tabalumab is a human IgG4 antibody that neutralizes both soluble and membrane-bound BAFF. We are awaiting the outcome of several clinical trials, evaluating the efficacy of tabalumab for treatment of autoimmune disease and MM.

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Disclosure

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References


