Thymic B cells promote germinal center-like structures and the expansion of follicular helper T cells in lupus-prone mice

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Submitted to Journal: Frontiers in Immunology

Specialty Section: T Cell Biology

Article type: Original Research Article

Manuscript ID: 520837

Received on: 22 Jan 2020

Frontiers website link: www.frontiersin.org
Conflict of interest statement

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Author contribution statement

YH designed the study, performed experiments, analyzed the data, and wrote the manuscript; SN, MJF, FFS, PJS, VM, JD performed experiments and analyzed the data; MRB, DS, SN, MR, EZ, and AMLD designed the study, analyzed the data, and wrote the manuscript. All authors critically read the manuscript.

Keywords

systemic lupus erythematosus, Thymic B cells, Germinal Center, Plasma Cells, follicular helper T cells *Correspondence: Maria Rosa Bono mrbono@uchile.cl, Daniela Sauma dsama@uchile.cl

Abstract

Word count: 200

Systemic lupus erythematosus (SLE) is an autoimmune disease characterized by the activation of autoreactive T and B cells, autoantibody production, and immune complex deposition in various organs. Previous evidence showed abnormal accumulation of B cells in the thymus of lupus-prone mice, but the role of this population in the progression of the disease remains mostly undefined. Here we analyzed the spatial distribution, function, and properties of this thymic B cell population in the BWF1 murine model of SLE. We found that in diseased animals, thymic B cells proliferate and cluster in structures that resemble ectopic germinal centers. Moreover, we detected antibody-secreting cells in the thymus of diseased-BWF1 mice that produce anti-dsDNA IgG autoantibodies. We also found that thymic B cells from diseased-BWF1 mice induced the differentiation of thymocytes to follicular helper T cells (TFH). These data suggest that the accumulation of B cells in the thymus of BWF1 mice results in the formation of germinal center-like structures and the expansion of a TFH population, which may, in turn, activate and differentiate B cells into autoreactive plasma cells. Therefore, the thymus emerges as an important niche that supports the maintenance of the pathogenic humoral response in the development of murine SLE.

Contribution to the field

Systemic lupus erythematosus (SLE) is an autoimmune disease characterized by the activation of autoreactive T and B cells, autoantibody production and immune complex deposition in several organs. Previous evidence has demonstrated the presence of B cells in the thymus of patients with autoimmune diseases and in a murine model of SLE, but the role of this thymic population in the progression of SLE remains largely undefined. Here we analyzed the spatial distribution, function, and properties of this thymic B cell population in the BWF1 murine model of SLE. Our findings suggest that in a prototypic autoimmune disease such as SLE, the thymus shifts its function from promoting negative selection of autoreactive T cells and central tolerance to a function usually associated with the induction of a humoral immune response. We believe this finding will surely contribute to a better understanding of the role of the thymus in the pathogenesis and development of autoimmune diseases.

Funding statement

This work was supported by FONDECYT 1191438 (MRB), FONDECYT 1140431 (MRB), FONDECYT 1180385 (DS), FONDECYT 3170424 (SN), FONDEQUIP/EQM140016 (MRB), CONICYT AFB 170004 (MR), ECOS-CONICYT C14S02 (MRB, AMLD), Doctoral Fellowship CONICYT 21130598 (YH) and Doctoral Fellowship CONICYT 2117036E6 (FFS).
Ethics statements

Studies involving animal subjects
Generated Statement: The animal study was reviewed and approved by Comité de Bioetica de Fundacion Ciencia y Vida CICUA Universidad de Chile.

Studies involving human subjects
Generated Statement: No human studies are presented in this manuscript.

Inclusion of identifiable human data
Generated Statement: No potentially identifiable human images or data is presented in this study.
Data availability statement

Generated Statement: This manuscript contains previously unpublished data. The name of the repository and accession number(s) are not available.
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ABSTRACT
Systemic lupus erythematosus (SLE) is an autoimmune disease characterized by the activation of autoreactive T and B cells, autoantibody production, and immune complex deposition in various organs. Previous evidence showed abnormal accumulation of B cells in the thymus of lupus-prone mice, but the role of this population in the progression of the disease remains mostly undefined. Here we analyzed the spatial distribution, function, and properties of this thymic B cell population in the BWF1 murine model of SLE. We found that in diseased animals, thymic B cells proliferate and cluster in structures that resemble ectopic germinal centers. Moreover, we detected antibody-secreting cells in the thymus of diseased-BWF1 mice that produce anti-dsDNA IgG autoantibodies. We also found that thymic B cells from diseased-BWF1 mice induced the differentiation of thymocytes to follicular helper T cells (TFH). These data suggest that the accumulation of B cells in the thymus of BWF1 mice results in the formation of germinal center-like structures and the expansion of a TFH population, which may, in turn, activate and differentiate B cells into autoreactive plasma cells. Therefore, the thymus emerges as an important niche that supports the maintenance of the pathogenic humoral response in the development of murine SLE.

Running title: Humoral response in the thymus of lupus-prone mice.
Keywords: Systemic lupus erythematosus; thymic B cells; germinal center; plasma cells; follicular helper T cells
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1. Introduction

Systemic lupus erythematosus (SLE) is an autoimmune disease characterized by a set of clinical abnormalities ranging from mild symptoms such as malaise, arthritis or dermatitis to more severe manifestations such as renal disease or compromise of the central nervous system. At the immunological level, SLE patients exhibit hyperactivity of T and B cells against self-antigens that leads to the secretion of autoantibodies against nuclear components such as DNA, RNA, histones, ribonucleoproteins, among others. Autoantibodies bind antigens and form immune complexes that deposit in the skin, joints, kidneys, heart, and central nervous system, generating inflammation and damage.\(^1\)\(^2\)

The thymus is a primary lymphoid organ whose main function is the induction of immune self-tolerance to prevent autoimmunity. This organ is dedicated to T cell generation and maturation, a function that usually declines with age and can be severely compromised in autoimmune diseases.\(^3\) It has been described that patients with autoimmune conditions such as myasthenia gravis exhibit alterations in the thymic structure and its cellular components.\(^4\)\(^5\)\(^6\) One of the important alterations in the thymus of patients with myasthenia gravis is the increase of autoreactive B cells. In the literature, there have been reports of patients with other autoimmune diseases, such as ulcerative colitis and systemic lupus erythematosus that present abnormalities in the structure of the thymus.\(^7\)\(^8\)\(^9\) However, it is unknown if these changes contribute to the disease.

B cells are a scarce population in the thymic medulla of both, healthy humans and mice (approximately 0.3% of total cells) where they are thought to function as antigen-presenting cells for thymic selection.\(^10\)\(^11\)\(^12\)\(^13\) We have previously shown that throughout normal aging, the human thymus accumulates at perivascular spaces memory B cells and plasma cells that generate antibodies with antiviral reactivity.\(^14\) Remarkably, a subset of myasthenia gravis patients has been diagnosed with thymic follicular hyperplasia which encompasses a considerable expansion of B cells and the formation of germinal centers in the thymus.\(^15\)\(^16\) Myasthenia gravis patients often go through thymectomy, a procedure that shows an overall improved clinical outcome, highlighting the contribution of thymic abnormalities to the production of autoantibodies against the acetylcholine receptor and severity of the disease.\(^17\)\(^18\)\(^19\) Recent evidence suggests that infiltration of B cells to the thymus and thymic stroma destruction precedes type 1 diabetes development in NOD mice,\(^20\) supporting the idea that B cell infiltration to the thymus may be a common event in several autoimmune diseases. Alterations in the thymus structure have also been observed in rheumatoid arthritis and SLE, but the contribution of these abnormalities to these diseases are poorly understood.\(^7\)\(^21\)\(^22\) Among patients with SLE, between 1.5 and 2% develop thymomas and undergo thymectomy as treatment. In contrast to myasthenia gravis patients, this procedure on SLE progression has no clear health benefits.\(^5\)\(^23\)\(^9\).

Studies using BWF1 mice, a murine model of SLE, showed an increase in B cell frequency in the thymus of diseased mice compared to those that have not yet developed the disease.\(^24\)\(^25\) In these studies, the authors highlight that the B1/B2 ratio in the thymus is higher than in spleen and blood. They showed that the B1 cell population migrates abnormally to the thymus due to high expression of the CXCR5 chemokine receptor, and aberrant high expression of CXCL13 (B lymphocyte chemoattractant, BLC) by myeloid dendritic cells present in the thymus. However, this study does not address the functional relevance of abnormal B cell numbers within the thymus and their contribution to SLE.
The converging evidence of B cells and plasma cells accumulation in the thymus during aging and particularly in autoimmune diseases prompted us to hypothesize that this lymphoid organ may become a specialized niche for B cells and plasma cells in the context of SLE development. To address this question, we characterized the B cell population of the thymus of BWF1 mice during the autoimmune humoral response. Here we show that upon the onset of the disease, the thymus structure becomes highly disorganized, exhibiting an increasing number of B cells that accumulate into structures that resemble ectopic germinal centers. Accordingly, we observed the presence of antibody-secreting plasma cells, a fraction of which produces anti-dsDNA autoantibodies. Noticeably, we further found that thymic B cells from diseased BWF1 mice induce the activation and differentiation of CD4+ thymocytes to follicular helper T cells. Altogether these data suggest a positive feedback loop, where thymic B cells induce the differentiation of follicular helper T cells that in turn promote the differentiation of autoreactive plasma cells in the thymus.
2. Materials and Methods

2.1 Mice

Female lupus-prone [NZB×NZW]F1 (BWF1) mice were purchased from Jackson Laboratory (Bar Harbor, ME, USA) and maintained at the animal facility of Fundacion Ciencia & Vida. Animal work was carried out under the institutional regulations of the Fundacion Ciencia & Vida and was approved locally by the ethical review committee of the Facultad de Ciencias, Universidad de Chile. Disease incidence and severity was monitored by measuring proteinuria using a standard semi-quantitative Combur Test N (Roche Diagnostics, Germany) and an ELISA to determine antibody titers to double-stranded DNA (dsDNA). To detect early autoimmune disease, proteinuria was measured monthly during the first five months of age and every week after that. In this work, we used 3 and 5 months old BWF1 female mice as young mice which still do not develop autoimmune disease. Diseased mice were 9 months old in average, presented severe proteinuria (i.e., ≥500 mg/dL protein) and high levels of plasmatic antibody titers against double-stranded DNA. In all cases, age-matched [NZW×BALB/c]F1 female mice were used as non-autoimmune controls.

2.2 Flow cytometry and t-SNE

Cell surface staining was performed in ice-cold PBS with 2% fetal bovine serum (FBS) for 30 min in the presence of Fcγ R blocking antibody (CD16/32). Viability dye eFluor 780 reagent (eBioscience) or propidium iodide (PI) were used for live/dead cells discrimination. Monoclonal antibodies (mAbs) against mouse CD8 (53-6.7) FITC, CD138 (281-2) PE or BV421, CD45R/B220 (RA3-6B2) APC or PE-Cy7, CD19 (6D5) FITC, APC or eFluor 780, CD44 (IM7) APC, CD69 (H1.2F3) PE, CD83 (Michel-19) FITC, CD86 (GL1) FITC, IgM (RMM-1) PE-Cy7 or Biotin, purified CD16/32 (93), Ki-67 (11F6) Alexa fluor 488, OX40L (RM134L) Alexa fluor 647, Blimp-1 (5E7) PE and IgG-HRP (polyclonal) were purchased from BioLegend (San Diego, CA, USA). mAbs against mouse IgD (11-26c.2a FITC), CD5 (53-7.3) PE-Cy7, GL7 (GL7) eFluor 660, CD11c (N418) PE, CD62L (MEL-14) FITC, CD25 (PC61.5) APC, CD8 (53-6.7) APC-eFluor 780, and CD279/PD-1 (J43) FITC were purchased from eBioscience (San Diego, CA, USA). mAbs against mouse I-Ad FITC (AMS-32.1), were purchased from BD Pharmingen (San Diego, CA, USA).

Flow cytometry was conducted on a FACSCanto II flow cytometer (BD Biosciences) or FACSaria III (BD Biosciences) and data analysis was performed using the FlowJo software version 8 (Tree Star, Inc., Ashland, OR, USA). The t-SNE analysis was performed in R and FlowJo software version 10.

ELISpot

Millipore® MAIPS4510 96-well plates were activated for 2 min with 50 µl/well of 70% ethanol and washed 5 times with deionized water. Plates were coated with 15 µg/ml of capture antibody anti-mouse IgG or dsDNA at 10 µg/ml and incubated overnight at 4°C. The plates were pre-treated with 10 µg/ml of methyl-BSA for 3 h at 37°C to evaluate reactivity against dsDNA. Subsequently, the plates were washed with PBS and blocked with RPMI medium supplemented with 10% FBS. The number of viable cells was carefully determined
and plated in triplicates. After incubation at 37°C, 5% CO2 for 22 h, the plates were washed 5 times with PBS and added 0.5 µg/ml of biotinylated goat anti-mouse IgG and incubated for 2 h at room temperature. Then plates were washed 5 times with PBS, and avidin-enzyme conjugated to HRP (eBioscience) was added and incubated for 1 h at room temperature. After washing the plates 5 times with PBS, 3-amino-9-ethyl carbazole (AEC) substrate was added and incubated at 37°C for 30 min. The plates were washed with bidistilled water and dry uncovered for 3 h at 37°C. Plates were read using an ELISpot reader AELVIS and the software Eli.Analyse ELISpot Analysis Software V6.0.

2.3 Confocal microscopy

Thymi were extracted from diseased BWF1, and age-matched control animals, thymus lobes were imbibed in RPMI + 10% FBS + 5% low melting point agarose (Invitrogen) solution. Once the agar solidified at room temperature, slices of 400 µm were obtained in a PELCO® 102 vibratome. The slices were fixed with 3.7% paraformaldehyde for 20 min at room temperature and stained for 30 min at 37°C with the following antibodies: CD4 (RM4-5) PE, CD8 (53 6.7) FITC, and CD19 (1D3) APC. After that, slices were washed with PBS, and placed on a slide with ProLong Gold antifade mounting medium (Invitrogen) and covered with a coverslip. Thymic slices were analyzed in the Zeiss LSM 710 confocal microscope, and the images analyzed with ImageJ.

2.4 Immunohistochemistry

Thymi were extracted from diseased BWF1, and age-matched control animals, and frozen at -80°C for 24 h in OCT compound. Six µm cryostat sections were obtained, air dried and fixed in cold acetone for 15 min. Sections were then incubated with a single drop of peroxidase blocker for 7 min at room temperature, washed with PBS and incubated for 1 h with blocking solution (PBS + 1% BSA + 10% goat serum). Then, sections were incubated overnight at 4°C with anti-mouse B220 or anti-mouse cytokeratin 5. After that, the sections were incubated for 1 h with HRP-coupled secondary antibody, washed with PBS and incubated with DAB (3'-Diaminobenzidine) for 4-8 min. Finally, sections were stained with hematoxylin and dehydrated to be mounting and visualized in the Olympus BX51 microscope.

2.5 Coculture of follicular helper T cells with B cells

Follicular helper T cells (T\textsubscript{FH}) and B cells of thymus and spleen were isolated by cell sorting from diseased BWF1 mice. T\textsubscript{FH} were stained with antibodies to CD4, CD8, PD-1, CXCR5, and selected as CD4\textsuperscript{+}CD8\textsuperscript{−}PD-1\textsuperscript{−}CXCR5\textsuperscript{−}. B cells were purified as CD19\textsuperscript{+}CD5\textsuperscript{−/int}. CD4\textsuperscript{+} T cells that do not express markers of T\textsubscript{FH} (PD-1\textsuperscript{−}CXCR5\textsuperscript{−}) were used as control. B cells were stained with CellTrace Violet (Invitrogen) according to the manufacturer’s instructions and cocultured with T\textsubscript{FH} at a 5:1 ratio (50,000 B cells and 10,000 T\textsubscript{FH} per well). Cocultures experiments were maintained for five days at 37°C and 5% CO\textsubscript{2}. Subsequently, cells were recovered, and B cells analyzed for GL-7 expression and dilution of CellTrace Violet stain by flow cytometry.

2.6 Coculture of B cells with thymocytes to assess T\textsubscript{FH} differentiation
Thymic B cells from diseased BWF1 and age-matched control mice were isolated by cell sorting (CD19+CD5/intCD11c) while thymocytes were isolated as I-Ad negative cells (to deplete antigen presenting cells). After sorting, thymocytes were stained with CellTrace Violet (Invitrogen) according to the manufacturer’s instructions and cocultured with B cells at a 10:1 ratio (100,000 thymocytes and 10,000 B cells). Coculture was maintained for five days at 37°C and 5% CO2. Subsequently, cells were recovered, and thymocytes analyzed for T FH phenotype (CD4+CD8-PD-1+CXCR5+) by flow cytometry.

2.7 In vitro B cell activation

Total thymic cells were activated at 2x10^6 cells/ml with anti-mouse CD40 at 1.5 µg/ml and anti-mouse IgM at 5 µg/ml for five days at 37°C and 5% CO2 to evaluate OX40L expression on B cells by FACS. On the other hand, total thymic cells were activated at 2x10^6 cells/ml with LPS at 2 µg/ml for three days at 37°C and 5% CO2 to evaluate Blimp-1 expression on B cells after fixation permeabilization with the appropriated buffer.

2.8 RNA-sequencing

RNA extraction from thymic B cells was carried out on 0.6x10^6 thymic B cells isolated by cell sorting (CD19+CD5/intCD11c) recovered directly in 0.5 ml of TRIzol reagent (Life Technologies). Quantification of RNA was performed using specific fluorometry with the Qubit RNA quantification assay (Life Technologies). RNA integrity was assessed using an RNA Quality Measurement Number (RQN) of Fragment analyzer with the High Sensitivity RNA Analysis Kit (Advanced Analytical Technologies), were used RNA samples with RQN values greater than 8.2. Sequencing libraries were prepared using the KAPA Stranded mRNA-Seq kit according to the manufacturer's protocol (Illumina). The length of the libraries was determined by capillary electrophoresis using the Standard Sensitivity NGS Fragment Analysis kit (Advanced Analytical Technologies). Libraries were quantified using the KAPA Library Quantification Kit (Kappa Biosystem) using the Eco PCR system (Illumina), following manufacturer’s protocol. Libraries were sequenced on a Miseq platform (Illumina) using a v3 150 kit with 2x75 bp paired-end. Samples were subsequently analyzed using R/Bioconductor, and the DESeq2 procedure was used to normalize the data. Differentially expressed genes were identified using an adjusted p-value cutoff of 0.05 and a fold change of at least 1.5.

2.9 Apoptosis assay

Cells isolated from the thymus of BWF1 diseased and age-matched control mice were stained with 1 µl of Annexin V FITC (BioLegend) and propidium Iodide at 1 µg/ml in 100 µl of binding buffer (HEPES 10 mM, NaCl 140 mM, CaCl2 2.5 mM) by incubating for 15 min at room temperature. Staining was stopped by adding 200 µl of binding buffer, and the cells were analyzed by flow cytometry.

2.10 Statistical analysis

Statistical analysis was performed with the GraphPad Prism program V6 (GraphPad Software, San Diego, CA, USA). The data were compared using a Student’s t-test after verification of normal distribution. Mann Whitney test was used when the data did not adjust
to a normal distribution. Wilcoxon signed-rank test was used to compare data with hypothetical value. P-values <0.05 were considered significant.

3. Results

3.1 Increased B cell numbers in the thymus of diseased-BWF1 mice correlates with an abnormal thymic structure.

We and others have reported that the frequency of B cells in the thymus increases significantly during normal aging and in several autoimmune diseases. We aimed at studying the dynamics of B cell accumulation in the thymus of BWF1 lupus-prone mice during the development of the autoimmune response. For this, we analyzed the thymus of BWF1 mice at different stages preceding (3 and 5 months old) and after the onset of the disease (9 months old in average) and compared the results to age-matched controls (NZWxBALB/c)F1 mice. We observed a higher than 20-fold increase in B cell frequency and a significant 6-fold increase in absolute B cell number in the thymus of diseased-BWF1 mice compared with age-matched control mice (Fig 1A-B). A modest but significant increase in the frequency of B cells in the thymus of 5 months old BWF1 mice compared to 3 months old BWF1 mice suggest that the frequency of B cells present in the thymus increases before the onset of proteinuria.

Histological examination of the thymus of diseased-BWF1 and age-matched control mice revealed remarkable alterations in the structure of the thymus at the onset of the disease (Fig. 1C). In diseased mice (9 months old on average), there was a reduction of the cortex areas and large numbers of B cells (B220⁺) clustered into structures reminiscent of germinal centers. In contrast, in age-matched control mice, we observed small numbers of B cells disseminated within the medulla, as previously reported. Also, we found that in diseased mice there was an expansion of non-epithelial perivascular spaces (PVS) (cytokeratin-5⁻) where most B220⁺ B cells clustered (Fig 1C and Suppl. Fig. 1). Fluorescent co-staining of CD4, CD8 and CD19 confirmed that the thymus of diseased mice is characterized by the presence of large B cell clusters and the absence of CD4⁺CD8⁺ double-positive (DP) thymocytes (Fig 1C) which are normally found in the cortex. To further evaluate the presence of germinal center-like structures in the thymus of diseased BWF1 mice, we analyzed the expression of the germinal center marker PNA on thymic B cells (Fig. 1D). Although we found no statistical difference in the frequency of PNA⁺ B cells between age-matched control and diseased BWF1 mice, the absolute number of PNA⁺ B cells increase 5-fold in diseased BWF1 mice compared to control mice (Fig. 1E). In agreement with this result, we also found an increase in the absolute number of proliferating B cells (Ki67⁺) in the thymus of diseased BWF1 mice even though the frequency of proliferating B cells do not show statistical differences (Figs. 1F-G). Altogether these results demonstrate that at the onset of SLE, the thymus of BWF1 mice undergoes remarkable changes in terms of structure and B lymphocyte content, with the appearance of ectopic germinal center-like structures.
3.2 The thymus of diseased-BWF1 mice harbors IgG anti-dsDNA antibody-secreting plasma cells.

Present evidence indicates that germinal center formation depends on the activation of antigen-specific B cells by cognate T cells leading to the formation of antibody-secreting plasma cells and memory B cells. The distribution of B cells in germinal center-like structures in the thymus of diseased-BWF1 mice suggests that they may be locally activated and differentiated into memory B cells or plasma cells. We next characterized the thymic B cells by analyzing their expression of differentiation markers. Analysis of isotype switched (IgM-IgD) memory B cells in diseased mice did not show a significant difference compared to control mice, whereas diseased mice present a significant increase in naïve B cells (IgM*IgD*) compared to control mice (Suppl. Fig. 2). These data indicate that a substantial fraction of B cells accumulating in the thymus of diseased mice rather display a naive than a memory phenotype.

Interestingly, the analysis of thymic plasma cells revealed a significant increase in the percentage and absolute number of plasma cells (B220*CD138*) in the thymus of diseased-BWF1 mice compared to age-matched control animals (Fig. 2A-B). These results are consistent with a higher percentage of Blimp-1+ B cells, a transcription factor driving the differentiation of B cells to plasma cells, in diseased mice compared to control mice (Fig. 2C-D). To investigate the presence of functional plasma cells and their specificity, we enumerated antibody-secreting cells (ASC) by ELISpot. These experiments revealed that the thymi from diseased-BWF1 mice contain significantly higher numbers of IgG ASC (>5 times) compared to those from age-matched control mice (Fig. 2E-F). When compared to IgG production from other organs known to harbor ASC such as bone marrow and spleen, we found a comparable number of spots of IgG ASC between the thymus and bone marrow in diseased-BWF1 mice (Fig. 2F).

Of note, the thymus of diseased-BWF1 mice contained few IgM ASC, and we could not detect anti-dsDNA ASC of the IgM class (Suppl. Fig. 3). Thus, the majority of thymic plasma cells in diseased BWF1 mice have gone through isotype switching secreting mainly IgG antibodies, some of which are specific of dsDNA. In summary, our results show that autoimmune BWF1 mice have proliferating B cells in germinal center-like structures within the thymus, which most likely support the differentiation of B cells into anti-dsDNA IgG-secreting plasma cells. Thymic ASC might, therefore, contribute in a significant way to the pool of secreted IgG autoantibodies found in autoimmune mice.

3.3 Thymic B cells from diseased-BWF1 mice express genes associated to cell survival.

Our data show that the B cells found in the thymus of diseased mice are distinct from normal resident thymic B cells in terms of abundance, localization, proliferation, and antibody secretion. To gain insights into the mechanisms that lead to the accumulation of this peculiar B cell population as well as into their function(s), we analyzed their transcriptomic profile using RNAseq. We identified 337 upregulated genes and 492 downregulated genes in thymic B cells from diseased-BWF1 mice compared to age-matched control mice (Fig. 3A and Suppl. tables 1 and 2). Among the upregulated genes, several were related to B cell survival and development including Upf1 and Naip2 (Fig. 3B). Also, CD24a, a negative regulator of early
pre-B cell differentiation in the bone marrow was likewise upregulated in diseased mice. Among the genes that were downregulated in thymic B cells from diseased-BWF1 mice, we found Hif1a, Blk, and Btn2a2, whose low expression has been associated with the induction or development of several autoimmune diseases such as collagen-induced arthritis, experimental autoimmune encephalomyelitis and SLE. The normalized counts of these genes are shown in Figure 3C. To confirm that thymic B cells obtained from diseased BWF1 mice have enhanced survival compared to B cells from age-matched controls, we assessed live B cells through Annexin V/PI assay. As shown in Figure 3D and 3E, the live fraction of thymic B cells was significantly higher in diseased-BWF1 mice, suggesting that the thymus of diseased-BWF1 mice provides a niche that supports the survival of B cells.

3.4 The thymus of diseased-BWF1 mice harbor functional follicular helper T cells.

In addition to the abnormal thymic structure and accumulation of B cells, characterization of the T cell thymic compartment of diseased BWF1 mice showed a significant reduction of CD4⁺CD8⁺ double-positive (DP) thymocytes compared to 3 and 5 month-old BWF1 mice (Suppl. Fig. 4), which is consistent with these animals exhibiting a smaller cortex, as observed by histology (Fig. 1C). Along with the decrease of DP cells, we found an increase in the frequency of CD4⁺CD8⁻ double-negative cells (DN) and both CD4⁺ and CD8⁺ single-positive (SP) thymocytes compared to 3 and 5-month-old BWF1 mice and control mice (Suppl. Fig. 4). These results indicate that during the onset of the disease, the thymus of BWF1 autoimmune mice suffers significant changes in its T cell composition in addition to the accumulation of B cells.

Interestingly, transcriptomic profiling of thymic B cells from diseased BWF1 mice further revealed the increased expression of Id3 and Lgals1 (Fig. 3B-C), two genes known to support the maintenance of germinal center B cells and humoral immune response. Therefore, we investigated whether the frequency and number of follicular helper T (TFH) cells were enhanced in the thymus of diseased-BWF1 mice. tSNE analysis gated on CD4⁺ T cells allowed us to dissect the composition of the naive (CD44lo) and antigen experienced (CD44hi) T cell thymic compartments. As shown in Figure 4A and Suppl. Fig. 5A-B, we observed an abundance of antigen-experienced CD44hi T cells in the thymus from diseased-BWF1 mice compared to age-matched control mice. Among thymic antigen-experienced CD4⁺ T cells found in diseased-BWF1, we detected a variety of different subsets such as memory tissue-resident CD103⁺, some of which also express CD69. The tSNE analysis revealed the appearance in diseased mice of a subset with TFH phenotype co-expressing PD-1 and CXCR5, which was absent in age-matched controls (Fig. 4A). Further analysis revealed that the percentage and absolute numbers of TFH (PD-1⁺CXCR5⁺) increased in the thymus of diseased BWF1 mice compared to the thymus of age-matched control mice (Fig. 4B-C) suggesting that the presence of B cells and GC-like structures in the thymus of diseased BWF1 mice may be associated to the appearance of TFH cells.

Similar to the role of TFH cells in B-cell maturation during normal immune responses, results from animal models of SLE as well as from patients with this disease indicates that TFH cells are required for autoantibody production. To evaluate if TFH present in the thymus of diseased mice could drive B cell activation and proliferation, we performed in vitro co-culture
assays with sorted thymic TFH and B cells from diseased BWF1 mice. Thymic TFH from diseased BWF1 mice induced the proliferation of activated GL7+ B cells to a similar level as splenic TFH obtained from BWF1 mice (Fig. 4D). Of note, non-TFH CD4+ T cells from either thymus or spleen were unable to induce proliferation of activated B cells (Fig. 4D). These results indicate that TFH present in the thymus of autoimmune mice are functional and possibly contribute to the activation and expansion of thymic B cells in diseased BWF1 mice.

3.5 Thymic B cells from diseased BWF1 mice induce the differentiation of follicular helper T cells.

It is known that B cells support T<sub>FH</sub> cell differentiation via OX40L in the spleen. This evidence prompted us to investigate whether thymic B cells could promote the differentiation of thymic T<sub>FH</sub> cells. Accordingly, we found an increase in the frequency of thymic B cells expressing OX40L (2-fold) in diseased mice compared to age-matched controls (Fig. 5A-B). To demonstrate that thymic B cells favor the development of T<sub>FH</sub> cells, we carried out co-culturing experiments of thymus B cells and thymocytes from control animals. We observed that thymic B cells from diseased mice generate a more significant percentage of T<sub>FH</sub> cells (PD-1<sup>-</sup>CXCR5<sup>+</sup>) than thymic B cells from age-matched control mice (5.2% with BWF1 B cells vs. 1.3% with control B cells) (Fig. 5C-D). Of note, thymic B cells from diseased-BWF1 mice induced a higher proliferation of CD4<sup>+</sup>SP thymocytes than B cells from age-matched control mice (Fig. 5E). These results suggest that thymic B cells from diseased mice support the differentiation of thymocytes into T<sub>FH</sub> cells.
4. Discussion and Conclusion

SLE is a chronic autoimmune disease of unknown etiology characterized by the formation of immune complexes, which are deposited in tissues causing inflammation. In SLE, both T and B cells are overactivated and recognize autoantigens related to nuclear proteins. Although the presence of B cells in the thymus in BWF1 mice, a murine model of SLE, has already been demonstrated 24 25, the development of this population during the progression of this autoimmune disease remains mostly unexplored. Using the BWF1 mice, here we report an increase in the number and frequency of B cells, plasma cells, and follicular helper T cells (T<sub>FH</sub>) in the thymus of lupus diseased mice. Moreover, our data provide evidence that these B cells proliferate and cluster in ectopic germinal centers within the perivascular space (PVS) of the thymus. Additionally, cellular suspensions of thymic cells obtained from diseased mice give rise to anti-dsDNA antibody-secreting cells. Finally, we demonstrate that thymic B cells from diseased-BWF1 mice favor the differentiation of T<sub>FH</sub>, which may, in turn, promote the activation and differentiation of B cells into autoreactive plasma cells in the thymus.

Previous studies have shown the presence of B cells in the thymus, which has been attributed a role of antigen-presenting cells involved in the negative selection of T cells 10 11 12. However, we demonstrated that the thymus of diseased mice loses the classical structure defined by the functional separation in cortex and medulla, where the processes of selection of the T cells occur. This observation led us to investigate whether these B cells might be involved in different processes independent of antigen presentation and negative selection of T cells. In this line, Pinto and collaborators have reported that thymic B cells produce autoantibodies that attack the thymic stroma, an event that precedes the development of type 1 diabetes 20. Moreover, our previous results demonstrated that the human thymus, as it ages may provide a niche for viral-specific plasma cells 14. The novel data presented here showing the presence of ectopic germinal centers, auto-antibody secreting plasma cells and T<sub>FH</sub> cells strongly argue in favor of the idea that during an autoimmune response, the thymus may acquire a new function as a niche suitable for the development of a humoral immune response.

An important finding presented here is that during the development of the autoimmune response, there is a significant increase in the frequency of B cells present in the thymus of BWF1 mice. This is not only a consequence of enrichment of B cells due to the reduction of double-positive thymocytes since as we report, there is a 6-fold increase in the absolute number of thymic B cells in diseased-BWF1 mice. An unresolved question that arises from this work is the origin of the B cells that accumulate in the thymus of diseased-BWF1 mice. Adoptive transfer experiments with splenic B cells as well as experiments with parabionts have shown that migration of peripheral B cells to the thymus in steady-state conditions does not contribute significantly to the pool of thymic B cells 11 48 49. However, under inflammatory conditions such as systemic LPS treatment, Candida albicans or Trypanosoma cruzi infection, it was demonstrated that mature B and T cells could efficiently migrate to the thymus 50. Thus, an intriguing possibility is that during chronic inflammation in autoimmune BWF1 mice, B cells from the periphery may continuously migrate to the thymus where they survive, proliferate, and differentiate into plasma cells.
The central role of the CXCL13 chemokine (B lymphocyte chemoattractant or BLC) in the recruitment of B cells to the thymus has already been established.\(^{51}\) Using a murine model of myasthenia gravis, Weiss et al. demonstrated that although thymic overexpression of CXCL13 under steady-state condition does not induce B cell recruitment to the thymus, under inflammatory conditions such as after immunization with Poly (I:C), CXCL13 overexpression enhanced B cell migration to this organ.\(^{52}\) In the murine model of SLE, the group of Matsushima demonstrated that dendritic cells in the thymus of BWF1 mice produce CXCL13 which attracts B cells to this organ during the development of the disease.\(^{24}\) The same group further explored this possibility and showed that when B cells are injected intravenously, they can enter the thymic PVS and the medulla of aged BWF1 mice.\(^{25}\) Thus, CXCL13 production in the thymus under inflammatory conditions may be sufficient to drive B cell migration to this organ.

Further evidence of B cell lymphopoiesis within the thymus was previously reported by Perera et al. where the authors use the Rag2-GFP reporter mice and demonstrate that B cells can develop from precursors within the thymus.\(^{11}\) Thus, it may be possible that during the autoimmune response, B cell lymphopoiesis within the thymus might be enhanced or there might be an increase in the survival of B cells in this organ. Whether B cells come from the periphery or are differentiated in situ, our RNAseq data supports the idea that within the autoimmune thymus, B cells might be exposed to an altered environment that effectively boosts their proliferation and survival. Thus, in any possible scenario, the accumulation of B cells may be favored by remarkable changes in the thymic niche during the autoimmune response supporting B cell survival and/or differentiation.

Using a different murine model of SLE, the B6.Sle16 lupus-prone mice, it was reported that B cells support the generation of T\(_{FH}\) through OX40L expression.\(^{46}\) In that report, the authors demonstrate that the ablation of OX40L expression, specifically in B cells, results in the reduction of T\(_{FH}\) and a significant decrease in the autoimmune response in these mice. In agreement with a role of OX40L in the induction of T\(_{FH}\), we also show that in the diseased-BWF1 mice, there is an increase in the frequency of T\(_{FH}\) along with an increase in thymic B cells that express OX40L compared to control mice. Moreover, we demonstrate that only thymic B cells from diseased-BWF1 mice have the capacity to induce the differentiation of thymocytes to T\(_{FH}\). Thus, our results recapitulate the role of OX40/OX40L interactions in the generation of the germinal center reaction observed in secondary lymphoid organs in lupus-prone mice reported by Cortini and collaborators.\(^{46}\) This leads us to propose that the germinal center reaction found in the thymus of lupus-prone mice is directly responsible for the generation of the auto-antibody secreting plasma cells in this organ rather that being the result of plasma cells arriving from the periphery. Activation of autoreactive B cells and differentiation into autoantibody producing plasma cells in germinal centers within the thymus may be favored by the interaction with T\(_{FH}\). Finally, thymic B cells may also induce the differentiation of CD4\(^{+}\) T cells to T\(_{FH}\), generating a positive feedback loop that sustains the humoral immune response within the thymus.

Interestingly, thymic morphological and functional alterations observed in the BWF1 and SLE patients have been described in several other autoimmune diseases including myasthenia gravis, type 1 diabetes, Sjogren’s syndrome and ulcerative colitis.\(^{5,20}\) Despite the distinct pathophysiological features, all these alterations have in common chronic
inflammation, which may be hijacking the normal thymic function of T cell repertoire selection to establish a niche that sustains the humoral immune response. Studies on thymectomized BWF1 mice could give some insight into the role of the thymus as a source of autoantibodies and its real contribution to the development of the disease.
Funding

This work was supported by FONDECYT 1191438 (MRB), FONDECYT 1140431 (MRB), FONDECYT 1180385 (DS), FONDECYT 3170424 (SN), FONDEQUIP/EQM140016 (MRB), CONICYT AFB 170004 (MR), ECOS-CONICYT C14502 (MRB, AMLD), Doctoral Fellowship CONICYT 21130598 (YH) and Doctoral Fellowship CONICYT 2117036E6 (FFS).

Ethics Statement

All mice were kept in the animal facility at Fundacion Ciencia & Vida and maintained according to the “Guide to Care and Use of Experimental Animals, Canadian Council on Animal Care”. Animal work was carried out under institutional regulations of Fundacion Ciencia & Vida and Facultad de Ciencias, Universidad de Chile and was approved by the local ethics review committees.

Author Contributions

YH designed the study, performed experiments, analyzed the data, and wrote the manuscript; SN, MJF, FFS, PJS, VM, JD performed experiments and analyzed the data; MRB, DS, SN, MR, EZ, and AMLD designed the study, analyzed the data, and wrote the manuscript. All authors critically read the manuscript.
Figure legends

**Figure 1.** Diseased-BWF1 mice present proliferating thymic B cells that cluster in germinal center-like structures. (A) Representative FACS analysis of CD19+ cells in the thymus of diseased-BWF1 and age-matched-control mice. CD5 was used to exclude T cells. (B) Thymic B cell frequency (left) and absolute number (right) in BWF1 mice at different ages prior (3 and 5 months old) and after the onset of the disease and age-matched control mice. B cells were analyzed as CD19+CD11c−CD5/int cells. Each dot represents one mouse (n=6-20 mice per group). Student’s t-test, *p≤0.05; **p≤0.01; ***p≤0.001. (C) Representative light microscopy images of B220 (left panel) and cytokeratin 5 (middle panel) staining of thymic tissue from diseased-BWF1 and age-matched-control mice. Scale bar: 200 µm. PVS: perivascular spaces. GC-like: germinal center-like structures. The right panel shows representative confocal microscopy images of CD4+ T cells (red), CD8+ T cells (green) and CD19+ B cells (blue) in thymic tissue of BWF1-disease and age-matched control mice. Scale bar: 50 µm. (D) Flow cytometry plots of PNA expression in thymic B cells (CD19+CD11c−CD5/int gate) from diseased-BWF1 and age-matched-control mice. (E) Frequency and absolute number of PNA+ B cells in the thymus of diseased-BWF1 and age-matched-control mice. Mann-Whitney test, ps0.05. (F) Flow cytometry plots of Ki-67 expression in thymic B cells from diseased-BWF1 and age-matched-control mice (G) Frequency and absolute number of Ki-67+ B cells (CD19+CD11c−CD5/int gate) in the thymus of diseased-BWF1 and age-matched-control mice. Student’s t test, p≤0.05. Data represent 3-4 independent experiments.

**Figure 2.** The thymus of diseased-BWF1 mice harbors IgG anti-dsDNA antibody-secreting plasma cells. (A) Flow cytometry plots of B220intCD138+ plasma cells gated on live thymic cells of diseased-BWF1 and age-matched-control mice. (B) Frequency (left) and absolute number (right) of plasma cells (B220intCD138+) in BWF1 mice and age-matched controls at different stages prior and after the onset of the disease. Each dot represents one mouse (n=3-14 mice). (C) Flow cytometry plots and (D) frequency of Blimp-1+ thymic B cells from diseased-BWF1 and age-matched control mice after activation with LPS. Each dot represents one mouse (n=4-6 mice). Mann-Whitney’s, **p≤0.01. (E) Representative ELISPOT of total and anti-dsDNA IgG antibody-secreting cells (ASC) present in the thymus of diseased-BWF1 and age-matched control mice. (F) Quantification of total and anti-dsDNA IgG antibody-secreting cells in the thymus, bone marrow and spleen from diseased-BWF1 and age-matched control mice. Each dot represents one mouse (n=5 mice). Student’s t test, *p≤0.05; **p≤0.01; ***p≤0.001.

**Figure 3.** Thymic B cells from diseased-BWF1 mice present a transcriptional profile associated to cell survival. (A) RNAseq transcriptome analysis of thymic B cells from diseased-BWF1 and age-matched-control mice. Heatmap shows genes that are upregulated (red) and downregulated (blue) with at least 1.5-fold change and adjusted p-value < 0.05 (detailed list of genes is provided in supplementary table 1 and 2). (B) Volcano plot of selected genes related to apoptosis and B cell development. (C) RNA-seq normalized counts for selected genes. White bars: age-matched control mice; gray bars: diseased BWF1 mice. (D) Flow cytometric analysis of live and apoptotic B cells as assessed by Annexin V and PI in cells from thymus of diseased-BWF1 mice and age-matched control mice. (E) Frequency of live thymic B cells (Annexin V/PI-) from diseased-BWF1 mice and age-matched control mice. Mann-Whitney’s t test, *p≤0.05.
Figure 4. Functional follicular helper T cells increase in the thymus of diseased-BWF1 mice. (A) tSNE analysis of CD4SP (CD4+CD8- gate) cells of diseased-BWF1 and age-matched-control mice. Bottom panel shows populations colored by manual gating. (B) Flow cytometry plots of PD-1 and CXCR5 expression in CD4SP cells from the thymus of age-matched-control and diseased-BWF1 mice. (C) Frequency and absolute number of follicular T helper cells (PD-1+CXCR5+) in diseased-BWF1 and age-matched control mice. Each dot represents one mouse (n=4-16 mice). Student’s t-test, *p≤0.05; ***p≤0.001. (D) Percentage of proliferating thymic GL7+ B cells (CD19+) after coculture with either PD-1+CXCR5+ thymic follicular helper T cells (B+T FH) or thymic PD-1-CXCR5- non-follicular CD4+ T cells (B+T) isolated from diseased-BWF1 mice. The same experiments were performed with cells from spleen. Each dot represents one mouse (n=4-5 mice). Student’s t-test, *p≤0.05.

Figure 5. Thymic B cells favor the expansion of follicular helper T cells in diseased-BWF1 mice. (A) Flow cytometry plots of OX40L expression in thymic B cells from diseased-BWF1 and age-matched-control mice following 5-day activation with anti-IgM and anti-CD40. (B) Frequency of thymic OX40L+ B cells from diseased-BWF1 relative to OX40L+ B cells from age-matched-control mice. Data represent 3 independent experiments. Wilcoxon signed rank’s test, ***p≤0.001. (C) Flow cytometry plots and (D) frequency of PD-1+CXCR5+ follicular helper T cells (in a CD4+CD8- gate) after co-culture of thymocytes (Thy) with thymic B cells from diseased-BWF1 (Thy+B BWF1) or age-matched control mice (Thy+B control). Data shows representative results of 3 independent experiments. (E) Proliferation of CD4SP thymocytes assessed by cell-trace violet dilution after co-culture of control thymocytes (Thy) with thymic B cells from diseased-BWF1 (Thy+B BWF1) or age-matched control mice (Thy+B control). Data represent the results of 3 independent experiments.
Supplementary Figure Legends

**Supplementary Figure 1.** Thymic B cells from diseased-BWF1 mice are localized in perivascular spaces. Representative images of thymic tissue from diseased-BWF1 stained with B220 (left panel) and cytokeratin 5 (right panel).

**Supplementary Figure 2.** Subpopulations of B cells are altered in diseased-BWF1 mice. (A) IgD and IgM expression in thymic B cells (CD19^+GL7^-) from diseased-BWF1 and age-matched-control mice. (B) Summary of the frequency (up) and absolute number (down) of the IgM^+IgD^+, IgM^+IgD^- and IgM^+IgD^- B cells populations.

**Supplementary Figure 3.** IgM and IgM anti-dsDNA antibodies in thymus of diseased-BWF1 mice. ELISPOT analysis was performed with total cells obtained from the thymus of diseased-BWF1 and age-matched control mice. Representative experiment of two independent experiments.

**Supplementary Figure 4.** Subpopulations of thymocytes are altered in diseased-BWF1 mice. (A) Representative FACS analysis of thymic DP, CD4SP, CD8SP and DN populations in BWF1 mice at different ages prior and after the onset of the disease and age-matched control mice. The cells were analyzed in a I-Ad^+ gate (n=4 mice per group). (B) Summary of the frequency (up) and the absolute number of thymocyte populations (down).

**Supplementary Figure 5.** Diseased-BWF1 mice present an increase in the frequency of antigen-experienced CD44^hi T cells in the thymus. Analysis of the thymic antigen-experienced T cell population in BWF1 mice at different ages prior and after the onset of the disease and age-matched control mice (A) Representative example of CD44 and CD62L expression in thymocytes from diseased-BWF1 and age-matched control mice. Analysis was carried out in a CD4SP (CD4^+CD8^-) gate. (B) Frequency of thymic antigen-experienced T cell populations (CD44^hi) in BWF1 mice at different ages prior and after the onset of the disease and age-matched control mice.
**Supplementary table 1.** RNAseq. List of genes upregulated in thymic B cells from diseased-BWF1 compared to thymic B cells from age-matched control mice. The genes in the list were selected with at least 1.5-fold change and p-value < 0.05.

### Upregulated genes

| Gene1 | Gene2 | Gene3 | Gene4 | Gene5 | Gene6 | Gene7 | Gene8 | Gene9 | Gene10 | Gene11 | Gene12 | Gene13 | Gene14 | Gene15 | Gene16 | Gene17 | Gene18 | Gene19 | Gene20 | Gene21 |
|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|
| Gstm4 | Gasm3 | Sle6a15 | 1600002K03Rk | Argge7 | Rffl | Sdf121 | Pd4b1 | Cens3 | Tamra1 | Hf11 | Kiz12 | Dhx8 | Fsa4 | Pit3 | EdEd1 | Lmo1 | Klk1b16 | Spc25 | Vgpa39 | Fgfl2 |
| Agata4 | Ly6a | Tpg1 | Cd3d | Binc2 | Mi84gd | Ptara1 | Syng3 | Arhgef10 | Hecw2 | Co10a1 | Zfp68 | Gm12789 | Scafl8 | Gga1 | Zfand2b | Ly6h | Alms1 | Tbc1d15 | Bpfla1 | Add3 |
| Ampd2 | Cpq | N4b3p | Il1f6 | Rgs9bp | Zfp790 | Stap2 | Tnnd2 | Arhgef3 | Dcn | Htb1 | Tbc1 | Tmem94 | Scarf2 | Guk1 | Hist12bc | Pole | Gm4858 | Tmem262 | Dphl | Il17b |
| Kfl2 | Avpr1a | S1x4 | Mrqf35 | Sccar3 | Pop7 | Papp2s | Hrasl5 | Cd79a | Aboed8 | Rhoa | Tmem127 | Mycbpap | Ifgfas | Mypt55 | Gapdh6 | Vipr2 | Bcd48507 | Bg1-ps2 | Cox5a | Scd2 |
| Nfplc | Gm732 | Tcgf | Tmem206 | Mjdx | Agxl1 | Cacfl1 | Edc2 | Tor1b | Prsc27 | Ppo | Cacfl1 | Gascam | Crap2 | Snn2 | Sgce | Myoc | Ncf4 | Tegpr2 | Jmpd4 |
| Sbpl | Chka | Tpds21l | Ptnp3 | Tnrc6a | Il10rb | Muc13 | Dc3f3 | Ythdf3 | Gm45253 | Tmem83 | Lipt1 | Spag11b | Acer2 | Dcplb | Rabac1 | Arhgap18 | Mapk8ip1 | Sck39a5 |
| Rad17 | Rpsi6a1 | S1c52a12 | Kad2b | Coq8a | Zgpat | Toppa | Wdr47 | Ppp1r18 | Gpelp1 | Cstal1 | Selp | Hs3t3b1 | Rasd1 | Ppla | Adgre5 | Sld8b1 | Gm4070 | Rplp1 |
| Cytol | Stm4 | Tbp | Tlbs1 | Krt25 | Acer2 | Pdcd9 | Gabarap | Pld3 | Bcl9l | Mup2 | Sk25a29 | Lgal1s | Scd2 | Stat5a | Serpina3f | Ankar | Inf2 | Smp32 | Shi3p65 | Dmnt1 | Sin4 | Ppp1r3d |
| Sk26a1 | Ranbp9 | Zfat | Tmem108 | Cx47b | Mad2l2 | 433405020Rik | Hmmpu1l | Reg3a | Mx2 | Opl1 | Adgb1 | Pel2 | Gdpgpl | Ube2a | Sit14 | Srec1 | Cnlfln |
| Lipo | Gdpgpl | Ube2a | Sit14 | Srec1 | Cnlfln | Epb41 | Pst2 | Zfp281 | Anap5 | Taf10 | Srebf2 |
| Gln3a | Rumx1 | Gm15922 | Aldh9a1 | Fafl | Id3 | Far1 | Ube2zn | Pira2 | Cascl1 | Perme3 | Rbbp9 | Sk9a3 | Zbbx | Serpina3n | Fam114a2 | Yhdfl3 | Trex1 |
| Cd24a | Kcnal6 | Wipfl | Cul7 | Rora | Wfcl8 | Igsfl1 | Lpact1 | Parp11 | Hcn1 | Zmynd8 | Ppaa | Notch4 | Ube2j2 | Slc35b3 | Gaint10 | Sox10 | Ztb2b3 | Wasi | Nas16 | Lca5l | 1700001J03Rk | Hfl20b2 | Pdcm3 |
| Klf5a | Kkk12 | Zfp219 | Slan1 | Nap2 | Tspan32 | Crl32a2 | Adk | Eid2 | Etfbkmt | Upl1 | Dsc | Vegfa | Rg9 | Cica3a1 | Eao8 | Tmem165 | Cant1 | Tbk22 | Era1 | Bc7b | Mfsd14a | 1700017805Rk | Apmap |
| Lpc | Snrpa | Mn1 | Spata1 | Ncapg | Ctnnb1 | Cdkn1a | Cdkn2aip | Ncs1 | Lrnf1 | Mnt | Cser1 | Pbx1 | Stk17b | Lce3f | Mex3 | Cit | Rip2 |
**Supplementary table 2.** RNA-seq. Genes downregulated in thymic B cells from diseased-BWF1 compared to thymic B cells from age-matched control mice. The genes in the list were selected with at least 1.5-fold change and p value < 0.05.

| Downregulated genes |
|----------------------|
| Tdh1 | Jsrp2 | Stk32b | Tce3 | Iil2g | Arhgef42 | Clec2d | Tmff2 | Cgc26 | Washc6 |
| C77080 | Ctl | Pyf1a | Pip | Adss | Dhh16 | Pprz2d | H2-M4 | Agbi3 | Smx22 | Anat3 |
| Plek2 | Rasgef1b | Rgs6 | Papd5 | Bax1b | Rfp56 | Dcn1d6 | Agbi3 | Smx22 | Faim | Capsp2 |
| Myc | Emp2 | Ptpd4 | Stk32b | Rfp56 | Dcn2d | H2-M4 | Agbi3 | Smx22 | Faim | Capsp2 |
| Letm2 | Rnf127 | Dgkh | Eh3 | Smad1 | Klg3 | Smx22 | Faim | Trmt112 | Pmp | Ccdd167 |
| Gypa | BC051143 | Znrd1 | 11100121l19Rk | Ptf1a | Dctn7 | Cacng4 | Mf22 | Tagln4 |
| Anxa7 | Snx16 | Arpc5l | Prpg3 | Dctn7 | Coro2a | Mf22 | Csrp4 | Trns1bp2 |
| Gpc2 | Rgs5 | Etv6 | Ndufc3 | Coro2a | Ahrge41 | Csrp4 | Slc25a14 | Gfap |
| Jsp1 | 1810037117Rk | Tcte2 | Mars | Arhgef41 | Nans | Slc25a14 | Mfps5 | Igfb2 |
| Ctsb | Fsp2 | Pip | Lnc58 | Nans | Pmp5 | Fsd3 | Cct2 |
| Rasgef1b | Id2 | Ptpd4 | Stk32b | Rfp56 | Dcn2d | H2-M4 | Agbi3 | Smx22 | Faim | Capsp2 |
| Emp2 | C77082 | Dhx15 | H2-M3 | Klg2 | Fgo9 | Fbo2d | Ints9 |
| Ref126 | Plek4 | Ehd2 | Atpt6v1b2 | Klg2 | Fanc | Fbo2d | Rasal1 | Supo4 |
| Ef23s3x | Mxc | Atpt6v1b2 | Setd7 | Fanc | Glrl4 | Spb | Cebpe | Gsk3b |
| Zeb2s3 | Letm3 | Setd7 | 11100121l19Rk | Glrlx | Hal | Cdx2 | Drmt3b | Pdeo4d |
| BC051142 | 4933415F23Rk | 11100121l19Rk | Png2 | Hal | Pah | Lopn3 | Slc25a16 | Dova3 |
| Smx15 | Tran5 | Ppng2 | Ndufc2 | Pah | Fam126a | Aagalt | Sumo2 | Epo |
| Rgs4 | Gypa | Ndufc2 | Mars | Fam126a | Ptnpb | Jaz2 | Srr | Rnf215 |
| 1810037117Rk | Anxa9 | Mars | Lnc57 | Ptnpb | Pnf7a2 | Cyp3a26 | Osp1g | Sptint2 |
| Fos1 | Id2 | Lnc57 | Stk32b | Pnf7a2 | Iil2g | Tekt6 | Cnp | Rpdf3 |
| Id2 | C77081 | Stk32b | Pfpfa2 | Iil2g | Xirp2 | Clec2d | Map1a | Cgo26 |
| C77081 | Plek3 | Pfpfa2 | Rgs7 | Spa87 | Ptkg | Tmff2 | Ints8 | Washc6 |
| Id2 | Myc | Rgs7 | Adss | Bax1b | Ncomp | Dcn1d6 | Sumo3 | Mecox |
| Myc | Letm3 | Adss | Dgkh | Pprz2d | Rasa3 | Agbi3 | Gsk3b | Anat3 |
| Letm3 | 4933415F23Rk | Dgkh | Ddh3 | Smad2 | Ddah2 | Smx22 | Pdeo4d | Capsp2 |
| 4933415F23Rk | Traf4 | Ddah3 | Kel | Bk | Vps13a | Faim | Ocva2 | Pmp |
| Traf4 | Gypa | Kel | Znrd2 | Rgpg2 | Abcd1b | Trmt113 | Epo | Ced167 |
| Gypa | Anxa8 | Znrd2 | Arpc5l | Ptf1a | Ml6p | Cacng4 | Rnf214 | Tagln4 |
| Anxa8 | Gyc | Arpc5l | Abc9 | Dctn7 | Cyp3a25 | Mf22 | Sptint1 | Trns1bp2 |
| Gjc3 | Jsrp2 | Etv7 | Tcm | Coro2a | Tekt5 | Csrp4 | Rpdf2 | Gfap |
| Jsp1 | Ctsb | Tcte3 | Gdb1 | Arhgef41 | Clec2d | Slc25a14 | Cc25 | Igb2 |
| Ctsb | Rasgef1b | Tptd2 | Nans | Tmff1 | Mrp5 | Washc5 | Cct2 |
| Rasgef1b | Emp2 | Papd5 | Hsp1a | Rfp56 | Dcn1d4 | Fsd3 | Mecox | Map1a |
| Emp2 | Rnf127 | Dhx16 | Trnn2 | H2-M3 | Agbi3 | Fgo9 | Anat2 | Ints10 |
| Ref127 | Ef23s3x | Ehd2 | C9 | Klg3 | Smx21 | Fbo2d | Capsp1 | Supo6 |
| Ef23s3x | Zeb2s3 | Atpt6v1b2 | Smad2 | Fanc | Faim | Spb | Pmp | Gsk3b |
| Zeb2s3 | Tmem176b | Setd7 | Rgo1 | Glrlx | Trmt112 | Cdx2 | Ccdd166 | Pdeo4d |
| BC051143 | Atqg9a | 11100121l19Rk | Ptf1a | Hal | Cacng3 | Lopn3 | Tagln3 | Ovca4 |
| Smx16 | St17 | Ppng2 | Dctn6 | Pah | Mf11 | Aagalt | Trns1bp1 | Epo |
| Rgs5 | Selonep | Ndufc2 | Coro2a | Fam126a | Csrp3 | Jaz2 | Gfap | Rnf216 |
| 1810037117Rk | Tgfb2 | Mars | Arhgef40 | Ptnpb | Slc25a13 | Cyp3a27 | Igb1 | Sptint3 |
| Fos2 | Tftp | Lnc57 | Nans | Ptf7a2 | Mrp4 | Tekt7 | Cts2 | Map1a |
| Id2 | Chmb2 | Stk32b | Rfp4 | Iil2g | Fsd2 | Clec2d | Map1a | Ints9 |
| C77081 | Fdo4 | Pfpfa2 | H2-M2 | Spa87 | Fgo8 | Tmff3 | Ints9 | Supo4 |
| Plek3 | Ehd1 | Rgs7 | Klg1 | Bax1b | Fbo41 | Dcn1d6 | Sumo4 | Gsk3b |
| Myc | Atpt6v1b1 | Adss | Fanc | Pprz2d | Spb | Agbi4 | Gsk3b | Pdeo4d |
| Letm3 | Setd6 | Dgkh | Smad2 | Cdx2 | Smx23 | Pdeo4d | Ovca3 |
| 4933415F23Rk | 11100121l19Rk | Ddh3 | Hal | Bk | Lopn2 | Faim | Ocva3 | Epo |
| Traf4 | Prng1 | Kel | Pah | Rgpg3 | Aagalt | Trmt114 | Epo | Rnf215 |
| Gypa | Ndufc1 | Znrd2 | Fam126a | Ptf1a | Jazf1 | Cyp3a26 | Rnf215 | Sptint2 |
| Anxa8 | Mars | Arpc5l | Ptnpb | Dctn8 | Cyp3a26 | Tekt6 | Sptint2 | Rpdf3 |
| Gjc3 | Lnc56 | Etv7 | Ptf7a1 | Coro2a | Tekt6 | Clec2d | Rpdf3 | Cc26 |
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Figure 2.

A. Age-matched control vs. Diseased-BWF1

B. Plasma cells (%)

C. CD19 vs. Blimp-1

D. Blimp-1+ cells (%)

E. Total IgG and IgG a-dsDNA

F. IgG-ASCs/10^6 cells

In review.
**Figure 4**

A.

| Age-matched control | CD44 | PD-1 | CXCR5 | CD103 | CD69 | CD62L | CD25 |
|---------------------|------|------|-------|-------|------|-------|------|
| BWF1-diseased       |      |      |       |       |      |       |      |

![tsNE2 tSNE1 plot](image)

- **CD44** T cells
- **CD44** T cells

**Legend**

- Follicular T helper cells (PD1^−CXCR5^+)
- Tissue-resident memory T cells (CD103^−CD69^+)
- Tissue-resident memory T cells (CD103^+CD69^+)
- CD25^+ T cells
- PD-1^+ T cells
- Mature naive T cells (CD62L^+CD69^+)
- Immature naive T cells (CD62L^−CD69^+)

B.

| Age-matched control | Disease-BWF1 |
|---------------------|--------------|
| PD-1 | CXCR5 |

C.

![Bar charts](image)

D.

![Bar charts](image)

- Proliferating GL7^+ B cells (%)
- Thymus
- Spleen
