Peripheral B Cell Subsets in Autoimmune Diseases: Clinical Implications and Effects of B Cell-Targeted Therapies

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Antibody-secreting cells (ASCs) play a fundamental role in humoral immunity. The aberrant function of ASCs is related to a number of disease states, including autoimmune diseases and cancer. Recent insights into activated B cell subsets, including naïve B cell to ASC stages and their resultant cellular disturbances, suggest that aberrant ASC differentiation occurs during autoimmune diseases and is closely related to disease severity. However, the mechanisms underlying highly active ASC differentiation and the B cell subsets in autoimmune patients remain undefined. Here, we first review the processes of ASC generation. From the perspective of novel therapeutic target discovery, prediction of disease progression, and current clinical challenges, we further summarize the aberrant activity of B cell subsets including specialized memory CD11chiT-bet+ B cells that participate in the maintenance of autoreactive ASC populations. An improved understanding of subgroups may also enhance the knowledge of antigen-specific B cell differentiation. We further discuss the influence of current B cell therapies on B cell subsets, specifically focusing on systemic lupus erythematosus, rheumatoid arthritis, and myasthenia gravis.

1. Introduction

Antibody-secreting cells (ASCs) refer to short-lived proliferating plasmablasts (PBs) and nonproliferating plasma cells (PCs), with distinct expression profiles, cell morphologies, and a lifespan from B cell lineages [1]. Autoimmune diseases such as systemic lupus erythematosus (SLE) [2], rheumatoid arthritis (RA) [3], and myasthenia gravis (MG) [4] are characterized by T cell hyperactivity and the overproduction of autoantibodies by ASCs, leading to highly activated differentiation to ASCs. For instance, the majority of autoantibodies causing MG are antiacetylcholine receptors (AChR) and AChR+CD21+ B cells in MG patients positively correlate with anti-AChR antibody production by ASCs in the serum [5], suggesting that hyperactivated antigen-specific B cell differentiation to ASCs represents a precursor of autoreactive ASCs. Other antigen-specific B cells, such as ANA+ IgG+ switched cells and IgG+ PBs, are elevated in SLE and further support the highly connected differentiation to ASCs [4]. In SLE patients, next-generation sequencing (NGS) has shown higher naïve to ASC and IgD− memory to ASC connectivity [6].

This highly activated process of differentiation to ASCs is believed to be induced by the disruption of tolerance checkpoints, which promotes survival of autoreactive ASCs with increasing quantities of autoantibodies [7–9]. Through the detection of B cells that recognize nuclear antigens (ANA+B cells) using flow cytometry, the checkpoints between transitional/naïve and naïve/memory cells have been identified in SLE and healthy individuals but naïve ANA+ compartments are defective in SLE [10]. While the numbers of ANA+ IgG PCs have been shown to increase, no changes have been found in ANA+ transitional, naïve, or switched/unswitched memory B cells in SLE [4], the exact tolerance checkpoints limiting the entrance of autoreactive ASCs are unknown. Challenges in this area include aberrant B cell groups with unknown phenotypes and unknown relationships to ASCs following differentiation in autoimmune diseases. Second, PCs such as pre-PCs, early PCs, short-lived PCs, and long-lived PCs fail to provide precise markers [11], increasing the difficulty in clarifying ASC origin and differentiation. Third, the phenotypes of autoreactive B cells with altered B cell receptor (BCR) repertoires [6, 8] are poorly understood, and pathogenic antibodies generated by different clones of
autoreactive B cells may exhibit heterogeneity of effector mechanisms.

Current biological agents targeting B cells including rituximab have been trialed in autoimmune diseases, which to date have shown only limited success, failing to deplete and prevent the replenishment of aberrant ASCs. The reasons for the lack of therapeutic efficacy include memory B cell-mediated relapse [12, 13], some unaffected subsets in peripheral blood [13–17] and in tissue [18, 19], unaffected factors such as BAFF and CD59 [18], and some autoantibody-producing B cell clones protected from rituximab-mediated cytotoxicity [20, 21]. Improving our knowledge of abnormally expanded autoimmune-associated subsets can enhance our understanding of ASC differentiation and explain therapeutic failures. This may reveal more effective targeted therapies and provide potential biomarkers that are appropriate for both diagnostic purposes and prediction of outcome.

We therefore revisited the normal processes of ASCs and conclude possible mechanisms that lead to abnormalities in B cell homeostasis. The existence of specific homing receptors in distinct subpopulations and different activation thresholds amongst the different stages of B cells were used to identify autoimmune-associated subsets [22]. We further summarize the current identified groups and discuss their potential roles as biomarkers for the prediction of organ damage, disease activity, and the influence of current B cell therapy.

2. Generalities during ASC Differentiation

2.1. Immature B Cells. Under normal conditions, immature B cells are generated in the bone marrow (BM), except for B1 cells that are produced in the fetal liver [23]. Those with autoreactive receptors undergo clonal deletion and sufficient receptor editing to enable effective tolerance [24]. Multireactive BCRs exist when leaving the BM, although they remain unresponsive to antigenic stimulation [25].

2.2. Naïve B Cells. Surviving immature/transitional B cells enter the spleen, lymph nodes, or other lymphoid tissues and develop into naïve B cells. Generally, naïve B cells can be divided into B1 cells, marginal zone (MZ) B cells, and follicular (FO) B cells. FO B cells are the most common [26].

2.3. Activated B Cells. Activated B cells can differentiate in either a T-independent (TI) or a T-dependent (TD) manner.

In TI responses, all B1 cells, MZ B cells, and FO B cells are activated and differentiate into PBs, although these cells show differential responsiveness to antigens, cytokines, or costimulation [27–29]. FO B cells show limited functionality during ASC differentiation in the absence of T cells compared to B1 and MZ B cells [27]. The reasons for these differences include alterations in TLR amongst the subgroups [28, 30] and low Mzb1 (pERp1) expression of FO B cells [31]. Mzb1 is required for ASC formation in TI [31]. This is highly expressed in B1 and MZ B cells, and its silencing impairs ASC differentiation in TI responses [31]. Since long-lived PCs also exist in T cell-deficient mice after immunization with LPS, TI also induces the formation of PCs [32, 33].

In TD responses, both FO B cells and MZ B cells show functionality [34]. Following activation, both undergo somatic mutations of the variable portion of expressed antibodies to alter and improve antigen specificity and affinity through extrafollicular responses and GC formation [1]. The generated PBs lack the ability to form PCs, and many undergo apoptosis. Extrafollicular growth typically occurs in the medullary cords of lymph nodes and in the T zone–red pulp border of the spleen, with low-levels of hypermutations observed [35]. GC reactions are enhanced by activation-induced cytidine deaminase (AID) [35] and result in the formation of plasma and memory B cells. PCs generated via these methods produce high affinity, class-switched immunoglobulins. Memory B cells can be found in both blood and lymph tissue with lower activation thresholds [36] and rapidly differentiate into ASCs. PCs are home to BMs through CXCR4 and continuously produce antibodies in the absence of antigenic stimulation, providing immediate protection [1]. CD19 niches provide external survival signals and are of great importance to PC survival [37] (Figure 1).

3. Potential Mechanism of B Cell Subset Alterations and Failure of Therapy

3.1. Mechanism of Self-Tolerance. In BM stage, immature B cells undergo clonal deletion or receptor editing to complete central tolerance, eliminating 20%–50% of self-reactive clones [38]. Additional peripheral tolerance includes anergy that occurs prior to entering the mature naïve B cell compartment [38, 39]. Specifically, BCR, TLR, and cytokines govern both normal and self-reactive antibody responses to antigens [40]. In autoreactive immature and transitional B cells, the BCR/TLR pathway increases AID to establish tolerance [41]. Further differentiation through GC or extrafollicular stimulation is dependent on initial BCR affinity and antigen density [42]. However, the nature of BCR, TLR, and cytokine interactions remains unclear.

3.2. Relevant Extrinsic and Intrinsic Factors Sustain Alterations in B Cell Subsets. Autoimmune diseases exhibit abnormal central tolerance with unusual BCRs [43]. Unlike central tolerance in the BM stage, the breakdown of peripheral tolerance can be adverse [24] and additional signals are required to overcome regulatory constraints of peripheral tolerance.

Extrinsic factors leading to the disruption of B cell tolerance include the deficient clearance of apoptotic cells by macrophages and neutrophils [44], hyperactivity of Th-cells, alterations in dendritic cells [45–48], extrinsic cytokines such as BAFF, IFNγ, and IL-21 [9, 49–52], TLR stimulation [50, 53, 54], and survival niches for long-lived PCs [37]. Relevant intrinsic factors include changes in major histocompatibility complex (MHC) class II [55], BCR signaling responses [56–58], and TLR responses [59, 60]. These factors in addition to deficient Breg cells [61, 62] and abnormal of extrafollicular germinal centers (GC) formation [6, 63] result in alterations to B cell subsets highly connected to autoreactive ASCs (Figure 2).
3.3. Autoreactive PCs and Memory B Cells in Tissues: Difficult Therapeutic Targets. CXCR3, a chemokine receptor, is associated with migration into bone marrow and/or inflamed tissue, and the majority of B cells in healthy individuals lack CXCR3 expression [64]. However, in disease states, PCs are present in the tissue due to inflammatory factors including...
CXCL10, VCAM-1, and IP-10 [65] and their interaction with CXCR3 [64]. Thymic lymphocytes produce AChR autoantibodies in MG patients either spontaneously or in response to mitogen stimulation [66], suggesting an involvement of autoreactive ASCs in tissues of unknown origin. In addition, TLR4+/CXCR4+ PCs undergo significant infiltration into tissues of SLE patients and correlate with the severity of nephritis [67].

AChR-specific CD27+ memory B cells are also present in the hyperplastic MG thymus, with unknown specificity [68]. Excluding classical memory B cells, unique peripheral memory cells have been identified in tissues. CD11chiT-bet+ B cells are present in nephrotic kidneys, with upregulated chemokine receptors for recruitment to inflamed tissues, such as CCR9 [69]. In addition, circulating CD19+CXCR3hi memory B cells are elevated in SLE and associated with poor clinical outcomes in response to rituximab (RTX) treatment [70], which may also be associated with tissue migration of memory B cells.

4. Peripheral B Cell Subset Alterations Associated with ASC Differentiation

Patients with autoimmune diseases show abnormalities during differentiation, with B cell subsets undergoing a wide range of alterations, including transitional B cells, B1, MZ, FO B, and memory B cells giving rise to ASCs, though different stages show preferential responses. In autoimmune diseases, these stages show specific extrinsic and/or intrinsic abnormalities. The clinical significance of these cells is discussed in Table 1.

4.1. B1 Cells. B1 cells localizing to the body-cavity serosa either secrete natural antibody spontaneously (B-1a) or respond to TI antigens (B-1a and B-1b) [91]. B1 cells are present in lymphoid organs and blood [91]. Griffin and colleagues defined the phenotypes of circulating human B1 cells as CD20+CD27+CD43+CD70- [92], although this remains controversial.

ASCs from this group play important roles in the production of protective antibodies, serving as major sources of natural IgM [93]. B1 cells can further differentiate into PCs in BM [94]. The pathophysiological functions of B1 cells in human autoimmune diseases require elucidation. Murine studies have proposed that elevated B1 cells are related to defects in macrophage clearance and represent a source of autoantibodies [95]. CD11b+ B1 cells increase in the peripheral blood of SLE patients, in which higher CD86 expression is observed, and T cell activity is enhanced [96]. This indicates that B cell subsets are activated and promote immunity. Although Murakami and colleagues reported that the elimination of B-1 cells alleviates clinical responses in autoimmune mice [97], its clinical relevance in patients with autoimmune diseases remains unclear.

4.2. Transitional B Cells. Transitional B cells belonging to the immature B2 B cell subset are key players in autoimmune diseases [9]. CD19+IgMhiIgD−CD24hiCD38hi transitional B cells are elevated in SLE but are almost absent in healthy controls [6, 9, 72, 98]. Blair and colleagues reported that the majority of CD19+CD38hiCD24hi cells are IgMhiIgDhiCD5+CD10−CD20+CD27 CD1dhi, which function as regulatory cells [99]. Their regulatory capacity is impaired in SLE [99]. Some B cells using other markers also include transitional B cells [71]. Kosalka and coworkers reported that immature/early transitional B cells (CD27+ IgD+CD21+) are elevated [71]. CD21low subsets (immature and activated B cells) are particularly expanded and correlate with lupus nephritis activity [71].

Studies in SLE patients and animal models show that transitional B cells cause the early loss of B-tolerance since a greater percentage of ANA+ cells in naive or new emigrant/transitional B cells are observed in SLE [4, 10, 100].

Cytokines, including BAFF and IFN, mediate transitional B cell abnormalities in SLE. Elevated BAFF expression contributes to transitional B cell expansion [40, 101], and targeting BAFF can recover the normal function of transitional B cells, promoting negative selection of activated autoreactive B cells [10, 102]. IL-6-producing transitional B cells survive in a type I IFN-dependent manner and positively correlate with disease activity in SLE [72]. Dieudonné and colleagues further emphasized the function of IFN by demonstrating that IFN stimulation combined with CD19 downregulation, and impairment of TLR9 responses disturb transitional B cells, resulting in the expansion of ASCs in SLE [9].

4.3. MZ B Cells. Following the transition of B cells, some remain in the spleen and develop into MZ B cells. MZ B cells become PBs following antigen presentation and rapidly produce high levels of IgM in a TI manner [103]. MZ B cells are present in human peripheral blood [104]. Although it is unclear whether this population is expanded in patients with autoimmune diseases, MZ B cells contribute to autoreactive clones and their numbers correlate with autoantibody production [105]. They can be rescued by BAFF to undergo expansion [40].

4.4. FO B Cells. FO B cells are the largest set of mature B cells following the transitional stage. They circulate freely in the spleen and lymphoid organs, forming an important part of the adaptive immune response, particularly TD responses [26]. TD responses include short-lived PB formation through extracellular responses and GC [1].

CD24-activated naïve B cells with a CD19hiCD21−CD38lowIgMlowCD23− phenotype increase in SLE but are absent in healthy controls [6]. These cells exhibit high clonal lineage with ASC populations [6], suggesting that aberrant subsets preferentially undergo differentiation. Increasing ANA+ and anti-dsDNA+ naïve B cells in SLE patients suggest defective selection at the transitional stage [10], though Suurmond and colleagues reported normal tolerance checkpoints in immature and naïve B cells with increasing total IgG1 PCs [4]. Further studies are now required to explore the exact composition of FO B cells contributing to autoimmune ASC.

4.5. Memory B Cells. Memory B cells reflecting autoimmune-associated reactivation are important as these cells possess...
<table>
<thead>
<tr>
<th>Disease</th>
<th>B subset</th>
<th>Stage</th>
<th>Extrinsic and/or intrinsic mechanism</th>
<th>Relevance to the diseases</th>
<th>References</th>
</tr>
</thead>
<tbody>
<tr>
<td>SLE</td>
<td>CD21\text{low} subsets †</td>
<td>Immature and activated B cells</td>
<td>Type I IFN overactivation with NF-kB activation and reduced Bax</td>
<td>Correlates with lupus nephritis activity</td>
<td>[71]</td>
</tr>
<tr>
<td>SLE</td>
<td>IL-6-producing transitional B cells †</td>
<td>Transitional B cells</td>
<td></td>
<td>Correlates with disease severity</td>
<td>[72]</td>
</tr>
<tr>
<td>SLE</td>
<td>CD19\text{hi}CD21\text{hi}CD38\text{low}IgM\text{low}CD23\text{−} B cells †</td>
<td>Activated naïve B cells</td>
<td>High basal levels of phosphorylated (spleen tyrosine kinase) Syk and ERK1/2</td>
<td>Correlates with disease severity</td>
<td>[6]</td>
</tr>
<tr>
<td>SLE</td>
<td>CD23\text{−}IgD\text{+} CD27\text{−} activated naïve cells †</td>
<td>Activated naïve B cells</td>
<td></td>
<td>Correlates with disease severity</td>
<td>[6]</td>
</tr>
<tr>
<td>SLE</td>
<td>CD19\text{hi}CXCR3\text{hi} B cells †</td>
<td>Naïve B cells, memory B cells, ASCs</td>
<td>Lower CD40 and CD27 expression; increased IL-21R expression; activates IL-21 signaling and drives differentiation</td>
<td>Poor clinical outcomes following RTX treatment</td>
<td>[65, 70]</td>
</tr>
<tr>
<td>SLE</td>
<td>CD11c\text{hi} B cells †</td>
<td>Unique memory B cells</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>SLE</td>
<td>TLR-9 expressing B cells †</td>
<td>Memory and plasma B cells</td>
<td>Activated TLR-9 signaling</td>
<td>Correlates with anti-dsDNA antibodies.</td>
<td>[59]</td>
</tr>
<tr>
<td>SLE</td>
<td>CD27\text{−} IgD\text{−}CD95\text{−} memory B cells †</td>
<td>Memory B cells</td>
<td>Higher levels of CD86, CXCR3, HLA-DR, and CD71</td>
<td>Correlates with disease severity and serological abnormalities</td>
<td>[74]</td>
</tr>
<tr>
<td>SLE</td>
<td>CD27\text{−} memory like B cells with high SYK †</td>
<td>Memory B cells</td>
<td>High expression of p-SYK; enhanced differentiation into CD27++ IgG secreting cells; somatically mutated BCR</td>
<td>Correlates with disease severity; candidate source of plasma cells</td>
<td>[75]</td>
</tr>
<tr>
<td>SLE</td>
<td>IgD\text{−}CD27\text{−} memory B cells †</td>
<td>Memory B cells</td>
<td>Hypermutation in rearranged VH Abs</td>
<td></td>
<td></td>
</tr>
<tr>
<td>SLE</td>
<td>IgD\text{−}CD27\text{−}IgM\text{−} memory B cells †</td>
<td>Memory B cells</td>
<td>BCR signaling abnormalities with Syk and Btk activation</td>
<td>Correlates with disease severity and autoantibodies</td>
<td>[77, 78]</td>
</tr>
<tr>
<td>SLE</td>
<td>IgD\text{−}CD27\text{−} memory B cells †</td>
<td>Memory B cells</td>
<td>Higher CXCR3 and lower CXCR5 expression; Syk, Btk, and JAK</td>
<td>Less susceptible to therapy</td>
<td>[79, 80]</td>
</tr>
<tr>
<td>SLE</td>
<td>TLR4\text{+} CXCR4\text{+} CD27\text{hi}CD38\text{hi}CD138\text{−}B cells †</td>
<td>Plasma cells</td>
<td>TLR4 promotes the secretion of anti-dsDNA IgG</td>
<td>Correlates with disease activity and severe renal damage</td>
<td>[67]</td>
</tr>
<tr>
<td>SLE</td>
<td>HLA-DR\text{hi}CD27\text{hi} plasmablasts †</td>
<td>Plasmablasts</td>
<td>Elevated levels are associated with lupus and anti-dsDNA</td>
<td>Elevated levels associated with disease severity</td>
<td>[81]</td>
</tr>
<tr>
<td>SLE</td>
<td>RP105 B cells †</td>
<td>ASC composition</td>
<td>Preferentially expresses BCMA</td>
<td></td>
<td>[82–84]</td>
</tr>
<tr>
<td>SLE</td>
<td>ARID3a\text{+}B cells †</td>
<td>All stages except for early naïve B cells</td>
<td>Secretes IFNα; not mediated by antibody secretion</td>
<td></td>
<td>[85]</td>
</tr>
<tr>
<td>Disease</td>
<td>B subset</td>
<td>Stage</td>
<td>Extrinsic and/or intrinsic mechanism</td>
<td>Relevance to the diseases</td>
<td>References</td>
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</tr>
<tr>
<td>RA</td>
<td>CD21(^{low}) B cell ↑</td>
<td>Naïve and memory B cells</td>
<td>Increases B cell activation</td>
<td>Correlates with lymph proliferation</td>
<td>[25, 86]</td>
</tr>
<tr>
<td>RA</td>
<td>CD86(^{+}) B cells ↑</td>
<td>Activated B cells</td>
<td>Possible association with ICOS(^{+}) Tfh cells and serum IL-21</td>
<td>Elevated levels associated with disease severity</td>
<td>[87]</td>
</tr>
<tr>
<td>RA</td>
<td>IgD(^{-})CD27(^{+}) memory B cells ↑</td>
<td>Memory B cells</td>
<td>Impaired IgM-production capacity and altered BCR repertoire</td>
<td>Correlates with disease activity and the anticyclic citrullinated protein antibodies</td>
<td>[88, 89]</td>
</tr>
<tr>
<td>RA</td>
<td>IgD(^{-})CD27(^{+}) memory B cells ↓</td>
<td>Memory B cells</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>MG</td>
<td>MuSK-specific CD27(^{hi})/CD38(^{hi}) B cells ↑</td>
<td>Autoreactive ASCs</td>
<td></td>
<td>Present during relapse but not remission</td>
<td>[90]</td>
</tr>
<tr>
<td>MG</td>
<td>AChR(^{-})CD21(^{+}) B cells ↑</td>
<td>Precursors of ASCs?</td>
<td></td>
<td>Elevated levels associated with disease; correlates and anti-AChR antibodies</td>
<td>[5]</td>
</tr>
<tr>
<td>MG</td>
<td>CD19 CD138 ASCs ↑</td>
<td>Plasmablasts</td>
<td>May associate with follicular helper T cells and IL-21</td>
<td>Elevated levels associated with disease severity</td>
<td>[85]</td>
</tr>
</tbody>
</table>

SLE: systemic lupus erythematosus; RA: rheumatoid arthritis; MG: myasthenia gravis.
lower activation thresholds for sustaining autoreactive ASCs with variable responses to therapy. Memory B cells exhibit heterogeneity in SLE, and homogeneous groups are difficult to establish. Usually, immunoglobulin isotypes including IgM, IgD, and CD27 are used to discriminate different memory B cells [106] including CD27+IgD+, CD27-IgD−, and CD27-IgD− (DN) types. In disease states, different activation markers or clusters of differentiation markers are observed in a range of disease states. Some CD27−flow memory B cells increase in number, including CD11chI-T-bet+ B cells, CD21+B cells, and spleen tyrosine kinase+t (Syk) memory-like B cells. The phenotypes defined by CD11chI, CD21, and Syk have overlapping populations, and also overlap with the groups defined by CD27 and IgD [75, 107]. These subsets overlap but are unique. CD11chI B cells are often characterized by the expression of CD21−flow in disease states [25, 73, 86], while Golinski e al. revealed that less than 10% of CD11chI B cells were CD21−flow in healthy individuals [108]. CD27 Syk+t memory-like B cells are 64.2±20.9% of CD27 CD21+B cells and 67.4±8.0% of CD27+ IgD−CD95+B cells [75].

4.5.1. CD11chI-T-bet+ B Cells. The number of CD11chI B cells is related to disease activity, anti-dsDNA levels, and ASC frequency in SLE [73]. Nearly all CD11chI cells express T-bet in SLE but with lower T-bet expression in healthy individuals [73]. T-bet is required for the generation of CD11chI cells [109]. In vitro, CD11chI B cells do not spontaneously produce IgG, but are poised to become ASCs and produce the major activation threshold for differentiating in the pathological state partly result from expanded naïve cell compositions. In RA patients, CD21−flow B cells exhibiting differential responses to BCR, CD40, or TLR9 are significantly expanded; the majority of which express autoreactive antibodies. The cells fail to proliferate or activate through BCR and/or CD40 [25]. At the transcriptional level, B cell activation, trafficking, and proliferation decrease, while the expression of integrins, including ITGAX, which encodes CD11c, increases [25].

4.5.3. CD27+IgD+ B Cells and CD27+IgD− B Cells. Several groups have reported that non-switched memory (CD27+IgD+) B cells decrease in SLE [6, 71, 77, 78, 80, 113] and RA [88], with increases in class-switched memory (CD27+IgD−) B cells [6, 71, 78–80].

For nonswitched memory (CD27+IgD+) cells, homogeneous patient cohorts have been assessed in quiescent SLE patients [79]. A decrease in nonswitched memory cells with no increase in other types was observed [79]. The quiescent SLE patients with a low frequency of B cells had lower levels of CD45, which may result from the reduced differentiation to ASCs and tissue homing [113]. IgD+CD27+IgM− memory B cells have a significantly lower association with disease activity and autoantibody concentrations [77, 78]. The cells overexpress CD95, CD80, CD86, CXCR3, and CXCR4 [77], suggesting they contribute to tissue homing. Rodriguez-Bayona and colleagues reported that the phenotype of IgD−CD27+IgM− memory B cells was consistent during both active and remission stages [77]. Their origin remains unclear but may arise from B1 and MZ B cells [78, 104].

The numbers of class-switched memory (CD27+IgD+) B cells increase in SLE [79]. These cells express higher CXCR3 in SLE compared with healthy controls and RA patients, with lower CXCR5 expression [114], which may explain why they are less susceptible to therapy. CXCR5+CXCR3+B cells lead to a B cell class switch through the combined stimulation of BCR and TLR [114], suggesting that class-switched memory B cells originate from CXCR5+ IgM memory B cells. In in vitro studies, lower thresholds were observed due to enhanced CXCR3 expression when stimulated with either CD40L, soluble BAFF, or IL-21, in addition to BCR and IFNγ [79, 80].

4.5.4. CD27+IgD+ B Cells. Double-negative (CD27+IgD+) memory B cells (DN) represent another class of isotype-switched cells. In healthy individuals, their numbers are small with a higher proportion of IgM memory cells [98], suggesting that they are activated in disease states.

DN cells can be expanded and express somatically mutated VH genes [76]. Their frequency correlates with renal involvement, disease activity, and specific autoantibodies, while RA shows no differences [76]. They lack the expression of FcRH4, with higher mutation rates and recirculation in the peripheral blood compared with CD27 (CD27 negative) memory B cells [76, 77, 98, 115]. This suggests
higher activation rates after the GC stage. However, Jacobi et al. observed disease-specific activity and serologic abnormalities with CD27− IgD− CD95+ memory B cell subset, as opposed to CD27+ IgD+ CD95+ memory B cell subset [74]. While increasing numbers of double-negative cells are not consistently correlated with anti-dsDNA [74] and the lack of CD27 expression impairs their binding to T cells, their relationship with ASCs requires additional research. DN cells express higher levels of activation markers (CD86, HLA-DR), chemokine receptor CXCR3, and CD71 [74, 113], suggesting an association with aberrant extrafollicular differentiation. Jenks et al. defined CXCR5+ CD21+ CD11c+ DN cells derived from CXCR5+ CD21− CD11c+ CD19− CD86− CD71− naïve B cells due to additional differentiation to circulating PCs through the extrafollicular activation pathway [107].

4.5.5. CD27 Syk++ Memory-Like B Cells. SYK, a key element of BCR signaling, is critical for B cell antibody TI/TD responses and memory B cell survival [116]. CD27 Syk++ memory-like B cells are expanded in SLE, with CD19++ CD20+ CD38− CD38+ CD19+ CD20− CD38+ phenotypes, primarily CD21+ [75]. The main difference with DN or CD95+ DN B cell subsets is that more than half of CD27+ Syk+ B cells express IgD [75]. In vitro studies, these cells are produced via stimulation with interferon-γ, lipopolysaccharides, or tumor necrosis factor α, showing elevated p-Syk expression and differentiation into CD27++ IgG secreting cells [75]. Thus, these cells represent the precursors of autoimmune ASC in SLE [75].

4.6. ASCs. Autoreactive ASCs produce autoantibodies and can be used to predict disease progression. The increasing number of ASCs not only is responsible for pathogenic autoantibody production but also is associated with accelerated autoimmune disorders [117]. Long-lived PCs in the absence of antigen stimulation represent autoreactive immunological memory cells that secrete pathogenic autoantibodies, but direct studies of autoreactive PCs in humans are challenging since they represent rare and inaccessible cells. The exact origin and pathways of differentiation remain difficult to establish since the markers are unclear [11]. Although it is generally considered that autoreactive ASCs originate from TD responses, elevated MZB1 levels in SLE suggest the contribution of TI responses [118].

In the MuSK MG group, Stathopoulos and coworkers reported that autoreactive ASCs produce MuSK antibodies during relapse [90]. In AChR MG patients, CD19+ CD138+ ASCs significantly increase and strongly correlate with follicular helper T cell frequency in MG patients, while the frequency of follicular helper T cells (FTh) was associated with disease activity [85]. Patients of generalized MG have a higher proportion of CD19+ CD138+ ASCs than clinical forms of ocular MG [85]. In addition, IL-6 and IL-21 which are important to GC activity [119] are increased in the serum [85]. Blocking of IL-21 signaling decreases antibody production [85], suggesting that the function of FTh is in aiding B cell differentiation to CD19+ CD138+ ASCs in an IL-21-dependent manner.

In SLE, ASCs were expanded in the peripheral blood [6, 120]. Using single-cell analysis, ASCs display lower frequencies of SHM and higher mutation frequencies in hypervariable CDR [6]. IgM+ ASCs are elevated in SLE patients; a small proportion of which are derived from newly activated naïve B cells [6], suggesting the importance of antigen-driven selection when differentiating towards ASCs and the importance of precursor identification.

4.6.1. TLR4+ CXCR4+ PCs. TLR4+ CXCR4+ PCs are expanded in the blood and renal tissue of SLE [67]. Their non-Ki-67 expression suggests they are nondividing cells and that their frequency correlates with disease activity and renal damage in SLE [67]. In in vitro studies, TLR4 inhibitors cause decreased anti-dsDNA IgG secretion [67], suggesting the importance of TLR4 signaling and their contribution to autoantibody production.

4.6.2. HLA-DRhiCD27hi PBs. Circulating HLA-DRhiCD27hi PBs are elevated in SLE [81]. Compared with CD27++ CD20+ CD21+ cells, HLA-DRhiCD27hi PBs show a closer correlation with lupus and anti-dsDNA levels [81]. HLA-DRhi PBs are also present in BM and contribute to PC formation in disease states [81]. Further analysis of the chemokines and relevant survival molecules is now required.

4.6.3. RP105+ B Cells. RP105 is a B cell surface molecule of TLRs associated with B cell proliferation and death [121]. Korganow and colleagues reported that RP105+ CD86+ and RP105+ CD38+ cells are persistently elevated in SLE, even in quiescent phases [113]. The frequency of RP105+ B cells correlates with disease activity in SLE [84]. RP105+ B cells cannot be divided into classically categorized B cells and belong to neither GC B cells nor memory B cells but display a highly activated CD95+ CD86+ CD38+ IgD- IgMlo phenotype [83]. In vitro studies in which autoantibodies and polyclonal immunoglobulins have been produced, implying the influence of ASC composition [83, 84].

Compared with healthy individuals, RP105+ B cells demonstrate increasingly higher relative ratios of BCMA/BAFF-R expression in SLE [82], suggesting that RP105+ B cells are dependent on BAFF/APRIL when differentiating into ASCs. The exact mechanisms of these pathways now require further elucidation (Figure 3).

5. Influence of B Cell Targeting Therapy

Currently, belimumab therapy for SLE and rituximab for RA are approved by the US Food and Drug Administration [122, 123]. Rituximab is recommended for clinical use for severe or MuSK MG [124, 125] and SLE in refractory- and corticosteroid-dependent forms of kidney or central nervous system involvement or severe autoimmune thrombocytopenia [126]. Here, we discuss their effects on different B cell subgroups and their association with clinical efficacy (Table 2).

5.1. Anti-CD20 Monoclonal Antibodies. CD20 is a 33 kD protein expressed by all mature B cells, except for plasma B cells. Rituximab (RTX), ofatumumab, and ocrelizumab are monoclonal therapeutic anti-CD20 antibodies considered treatments for autoimmune diseases [127].
RTX is recommended for patients who are refractory to standard therapy in RA [128], severe or MuSK MG [124, 125], and SLE [126]. The depletion of precursor cells that differentiate into autoimmune ASCs is considered the cause of effective treatment [129]. Except for the effect on antibody secretion, RTX mediates tropocytosis of human B cells by producing and releasing IL-6 in vitro and has no effect on tumor necrosis factor α, IL-1β, interferon-γ, or IL-10 [130]. In RA, patients have higher levels of IL-6 after 6 months of therapy and high IL-6 levels are also good predictors for RTX response [131, 132]. Third, the few remaining and/or regenerating B cells exhibit incomplete deficiency in costimulatory molecule expression, thus having impaired antigen presentation function [133, 134]. Following their depletion, immature and transitional B cells can be detected for several months [135] and B cell reconstitution is observed. The efficacy of rituximab in autoimmune diseases is mediated by decreased rates of PC synthesis and improved selection for autoreactivity by receptor revision. Numbers of PBs and PCs decrease indirectly after RTX therapy in SLE [136] and 16 months of RTX therapy in RA [137]. Good and shorter clinical responses after RTX therapy are associated with a sustained IL-6 and TNF-α levels [138]. RTX alone may not be sufficient to delete autoreactive clone.

5.2. Anti-CD22 Monoclonal Antibodies. CD22 is a transmembrane protein that regulates adhesion and inhibits BCR signaling [146]. CD 22 is expressed on the majority of developing B cells except for plasmablasts and plasma cells [147]. Epratuzumab is a humanized antibody directed against CD22 [148], inhibiting B cell proliferation and maturation and reducing production of proinflammatory cytokines including IL-6 and TNF-α [16, 147, 149]. Although phase Ib studies have shown improvements [150], recent phase III data reveal no differences compared with standard therapy without epratuzumab in SLE [151, 152]. Post hoc analysis of open trials of SLE patients with primary SS demonstrates that anti-SSA levels were consistently reduced after epratuzumab treatment [153]. Additional research is required to fully explore responsive clinical subgroups and relevant mechanisms.

Immature B cells, transitional B cells, naïve B cells, and limited memory B cells are affected [16, 154]. CD27+ memory B cells are less affected, partly due to low CD22 expression [16] and lower binding with epratuzumab [147]. In an in vitro study, epratuzumab binding leads to the expression of CD62L, decreased β7 integrin, and increased β1 integrin, and the primary effect is observed on CD27+ B cells [147]. The unaffected CD27+ memory B cells may contribute to the failure of the therapy.

Figure 3: Differentiation of aberrant ASCs and the involved subgroups. From immature B cells to ASCs, B cell subgroups show expression changes in autoimmune diseases. Unique autoimmune-memory phenotypes include CD11c+ B cells.

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Table 2: Summary of the effects of B cell targeting therapies on circulating B cell subsets.

<table>
<thead>
<tr>
<th>B cell targeting therapy</th>
<th>Mechanism of action</th>
<th>B subsets affected</th>
<th>B subsets not or less affected</th>
<th>Relevance to clinical relapse</th>
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<td>Targeting B cell surface antigens</td>
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<tr>
<td>Anti-CD20</td>
<td>General B cell depletion mediates complement-dependent cytotoxicity</td>
<td>All mature B cells except plasma B cells</td>
<td>Stem cells, PCs, and PBs</td>
<td>IgD−CD27+ and IgD−CD27+ memory B cells</td>
<td>[12, 138]</td>
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<td></td>
<td>Moderate B cell depletion impairs B cell signaling</td>
<td>CD27+ transitional and naïve B cells</td>
<td>CD27+ memory B cells</td>
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<td>[16, 147, 149, 154]</td>
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<td>Targeting ASCs with proteasome inhibitors</td>
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<tr>
<td>Bortezomib</td>
<td>ASC depletion induces proapoptotic unfolded protein response components and inhibits of NF-κB signaling</td>
<td>Short-lived and long-lived PCs</td>
<td>Precursor B cells</td>
<td>Precursor B cells</td>
<td>[17]</td>
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<td>Targeting B cell survival factors</td>
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<tr>
<td>Belimumab</td>
<td>Impaired B cell survival promotes negative selection of activated autoreactive B cells</td>
<td>Transitional naïve B cells, CD11c−CD21− B cells, and double-negative memory B cells</td>
<td>PBs and switched memory B cells</td>
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<td>[13–15, 102, 166]</td>
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5.3. Targeting ASCs with Proteasome Inhibitors. CD19+ ASCs in the BM or spleen and CD19- BM ASCs in BM have a similar capacity to contribute to immunological memory [155]. Both contribute to the failure of current therapeutic approaches.

Bortezomib, a proteasome inhibitor, has been used in the treatment of autoantibody-mediated autoimmunity, including SLE [156–158], RA [159], and MuSK MG [160]. Further studies are required to ensure safety, since murine studies showed higher mortality rates after drug use [161]. The drug effectively depletes both short-lived and long-lived PCs in the peripheral blood and bone marrow by ~50% including the CD19 phenotype by inducing apoptosis [17] with decreasing levels of pathogenic autoantibodies [156]. In addition, fluctuations in anti-dsDNA antibody levels have been observed in relation to each bortezomib cycle [17], suggesting the dynamic depletion of autoreactive ASCs. However, precursor B cells remain unaffected, resulting in the rapid repopulation of ASCs in the absence of bortezomib [17]. Bortezomib withdrawal is accompanied by rapid repopulation of short-lived PCs with increasing autoantibody levels [17].

5.4. Targeting B Cell Survival Factors. Blocking targets include BAFF, its homolog APRIL, and their receptors including BAFF-R, BCMA, and TACI. BAFF-R is expressed on the surface of human peripheral B cell subsets excluding PCs and centroblasts located in the dark zone of GCs [162] while BCMA is expressed constitutively by long-lived plasma cells and is important for their survival [163]. TACI is expressed on activated B cells, MZ B cells, switched memory B cells, and PCs [164]. BAFF-R is the major receptor molecule for BAFF-dependent response in the peripheral blood [165], and the interaction of BAFF and BAFF-R is required for the survival and late transition of MZ and mature naïve B cells [105]. Relevant agents include belimumab, tabalumab, atacicept, and blisibimod, which primarily block BAFF.

Belimumab, an inhibitor of BAFF, is recommended for those with active or flaring extrarenal disease in SLE [126]. Belimumab functions rapidly in the early developmental stages of B cells, especially naïve B cells and transitional stages [14]. Malkiel and colleagues elaborated that belimumab treatment restored the censoring of ANA+ transitional B cells through anergy [10]. In SLE, earlier longitudinal studies demonstrated that inhibiting BAFF using belimumab selectively reduced the total number of transitional and naïve B cells with no effects on memory B cells [15]. CD11c+CD21+ B cells [14] and double-negative memory B cells [13, 14] continuously declined with stable PBs [14, 15] and switched memory B cells [13, 14], suggesting the importance of unique memory B cells. PBs slowly decrease after 532 days [13] or 18 months [14]. The level of anti-dsDNA autoantibodies also decreased but only at an early stage [14]. Switch and nonswitch memory B cells and PCs exhibit resistance to belimumab therapy. While the lack of BAFF-R on PCs may be one reason, it cannot be explained why switched memory B cells can survive. These delayed clinical effects require a longer therapeutic regimen.

6. Conclusions and Future Perspectives

Autoimmune diseases result from B cell hyperactivity and a disturbance of ASC homeostasis. The development of multi-chromatic flow cytometry has improved the identification of aberrant B cells and promotes the expression of extrinsic and/or intrinsic molecules. In this review, we have discussed the normal progression towards ASCs (Figure 1), the potential mechanisms of imbalance (Figure 2), and potential B cell subgroups that mediate autoimmune diseases (Figure 3). Aberrantly activated B cells may further contribute to ASCs and their extrinsic and/or intrinsic alterations (Table 1).

The limited success of current B cell therapies coupled with the depletion of precursor B cells is key to the identification of phenotypes of these heterogeneous pathological groups. Technically, more effective identification methods are required. Recombinant antibodies and mass cytometry may aid the discrimination of subgroups.

Moreover, the relationship with autoreactive ASCs, their differentiation, and their sensitivity to chemokine and homing molecules requires further understanding for the generation of long-lived PCs in tissues. Single-cell RNA sequencing and serum proteomics to identify autoantibodies can provide new insight into autoreactive ASC differentiation and identify the landscape of B cells in autoimmune diseases, including additional peripheral tolerance checkpoints, distinct distributions, gene expression analysis, the association amongst subsets, and their underlying mechanisms of differentiation.

In addition, the assessment of self-antigen reactivity of the expanded groups coupled with the analysis of the differential response to therapy may provide more effective targets for refractory groups.

Conflicts of Interest

The authors declare that they have no competing interests.

Authors’ Contributions

Wanlin Jin wrote the manuscript and Zhaohui Luo and Huan Yang revised the manuscript.

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