



The Other Function: Class II-Restricted Antigen Presentation by B Cells

Lital N. Adler^{1,2†}, Wei Jiang^{1,2†}, Kartik Bhamidipati², Matthew Millican³, Claudia Macaubas^{1,2}, Shu-chen Hung^{1,2} and Elizabeth D. Mellins^{1,2*}

¹Department of Pediatrics, Stanford University, Stanford, CA, USA, ²Program in Immunology, Stanford University, Stanford, CA, USA, ³Stanford University, Stanford, CA, USA

Mature B lymphocytes (B cells) recognize antigens using their B cell receptor (BCR) and are activated to become antibody-producing cells. In addition, and integral to the development of a high-affinity antibodies, B cells utilize the specialized major histocompatibility complex class II (MHCII) antigen presentation pathway to process BCR-bound and internalized protein antigens and present selected peptides in complex with MHCII to CD4+ T cells. This interaction influences the fate of both types of lymphocytes and shapes immune outcomes. Specific, effective, and optimally timed antigen presentation by B cells requires well-controlled intracellular machinery, often regulated by the combined effects of several molecular events. Here, we delineate and summarize these events in four steps along the antigen presentation pathway: (1) antigen capture and uptake by B cells; (2) intersection of internalized antigen/BCRs complexes with MHCII in peptide-loading compartments; (3) generation and regulation of MHCII/peptide complexes; and (4) exocytic transport for presentation of MHCII/peptide complexes at the surface of B cells. Finally, we discuss modulation of the MHCII presentation pathway across B cell development and maturation to effector cells, with an emphasis on the shaping of the MHCII/peptide repertoire by two key antigen presentation regulators in B cells: HLA-DM and HLA-DO.

Keywords: B cell, MHC class II, antigen processing, antigen presentation, invariant chain, HLA-DM, HLA-DO, germinal center

INTRODUCTION

In addition to their role as secretors of antibodies, B cells function as professional antigen-presenting cells (APCs) for CD4+ T cells by expressing cell-surface major histocompatibility complex class II (MHCII) molecules with bound peptide, the ligand of the α/β T cell receptor. MHCII-restricted antigen presentation by B cells plays a role in multiple immunological events. For example, recent evidence supports a model of sequential class II-restricted antigen presentation by thymic B cells to CD4+ thymocytes (1). In this scenario, resting IgD+/IgM+ B cells enter the thymus and undergo cognate interactions with CD4+ thymocytes and CD40-mediated B cell activation. This increases MHCII, CD80, and Aire expression, allowing B cell presentation of Aire-dependent antigens and negative selection of self-antigen reactive thymocytes. In another scenario, B cells contribute to the magnitude of primary CD4+ T cell responses (2) and to the development and expansion of Th2 cytokine-secreting CD4+ T effectors (T_{EFF}) (3). B cell division, hypermutation of immunoglobulin genes and engagement of T follicular helper cells (T_{FH}) in the germinal center (GC) are directly

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> *Correspondence: Elizabeth D. Mellins

mellins@stanford.edu

[†]These authors have contributed equally and are listed alphabetically.

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1

proportional to the amount of antigen captured and presented on MHCII (4, 5). Antigen presentation by B cells also is critical to the generation and survival of memory CD4+ T cells (6). Recent work revealed a memory checkpoint when CD4+ T_{EFF} require cognate antigen recognition for further differentiation into memory cells (7); at this checkpoint, B cell APC are required for driving T_{EFF} to T_{FH} (8). In murine models of type 1 diabetes and multiple sclerosis, antigen presentation by B cells contributes to disease pathogenesis (9, 10). Related to this, blockade of B cell antigen presentation is implicated as one mechanism underlying efficacy of anti-CD20 therapy in CD4+ T cell-mediated diseases, including autoimmune hepatitis, rheumatoid arthritis, and multiple sclerosis and mouse models of these diseases [(11–14) and references therein]. In addition, Jackson et al. have proposed that, in certain genetic settings, B cell presentation of self-antigens can initiate a break in CD4+ T cell tolerance (15). In summary, MHCII-restricted antigen presentation by B cells is critical to immune function in health and disease, highlighting the importance of a comprehensive understanding of the associated molecular mechanisms.

Here, we review current information on B cell antigen uptake, processing, and presentation as MHCII/peptide complexes at the cell membrane. We discuss antigen encounter in secondary lymphoid organs (SLOs), the role of the B cell receptor (BCR) in capturing antigen, and B cell polarization, in the case of antigen capture from the surface of other cells. We next cover BCR internalization, the targeting of BCR/antigen complexes to subcellular compartments for interaction with MHCII, the roles of B cell proteases in invariant chain (Ii) degradation and generation of the peptide pool for MHCII loading. We review the shaping of the MHCII/peptide repertoire by HLA-DM and HLA-DO and the MHCII/peptide complex transport to the surface of B cells. Finally, the development and modulation of the MHCII presentation pathway across the stages of B cell differentiation are discussed.

In the mouse, B cells can be separated into B-1 and B-2 subsets, and recent studies argue that these are distinct lineages (16). B-1 cells (which likely have a human equivalent functionally) mostly arise in the fetal liver, self-renew in the periphery, and represent a small population that acts to achieve a rapid, early immune responses independent of T cell help (12, 16, 17). B-2 cells (in mice and humans) are mainly produced in the bone marrow throughout life and predominate in the B cell pool of SLO. B-2 cells are a key component of adaptive immune system, exhibiting antigen specificity and memory and mediating both T cell-independent and T cell-dependent antibody production. This review focuses on the class II antigen presentation pathway in B-2 cells (referred to as B cells in what follows).

THE BCR AND ANTIGEN ENCOUNTER

A key component of immune surveillance is the circulation of naïve, mature B cells through SLOs. About 45% of naïve B cells migrate to the spleen to sample antigens present in the blood; a roughly equal number move to the lymph nodes to sample antigens drained from tissues, and the remainder move mostly to mucosa-associated lymphoid tissues (MALT), a key interface between the host and its environment (18). Within SLO, the B cells localize to B cell follicles in a chemokine (CXCR13)dependent manner and begin to sample antigen. How antigen traffics to the B cell zone depends on the SLO. In MALT, antigen is passed across the epithelial barrier into B cell zones (19). In the spleen, antigen draining from the splenic artery is intercepted in the marginal zone (MZ) by macrophages and dendritic cells (DCs), as well as by a unique class of MZ B cells with innatelike behavior and limited BCR diversity (20). All three types of APCs are capable of conveying antigen to B cell follicles in the periarteriolar lymphoid sheaths (21), but certain types of antigens-especially repetitive bacterial polysaccharides and other T-independent type 2 antigens-are capable of activating MZ B cells directly (20) in a T-independent process (22, 23). In lymph nodes, antigen enters the subcapsular sinus (SCS) via afferent lymphatics and can reach B cell follicles in soluble form in the case of small antigens (<70 kDa) by movement through a conduit system that permeates the follicles (24, 25), or, for larger antigens and immune complexes, which are typically opsonized by complement components, intercepted by complement receptors on a layer of SCS macrophages (SSMs) lining the follicular (FO) zone, and then passed between complement receptors on various APCs and non-specific B cells. Immune complexes ultimately become tethered to the membrane of a follicular dendritic cell (FDC) (26, 27) for BCR scanning.

The BCR is composed of a membrane-bound immunoglobulin (mIg) for antigen binding and a transmembrane $Ig\alpha/Ig\beta$ heterodimer for signaling (28). The mIg consists of two immunoglobulin light (L) chains and two heavy (H) chains, which have variable numbers hydrophobic amino acid sequence motifs in their cytoplasmic tails, depending on the Ig isotype. Antigen recognition is mediated by the hypervariable regions of mIg $V_{\rm H}$ and V_L segments, which fold to form an antigen-binding site; signaling is mediated by the cytoplasmic immunoreceptor tyrosine activation motifs (ITAMs) of the associated Igα/Igβ heterodimer. The spatial organization of BCRs on resting B cell surfaces and the effect of antigen engagement on this organization are incompletely understood. An early study showed by transmission electron microscopy that almost all plasma membrane-associated proteins, including BCRs, are present in clusters termed "protein islands" (29). Recently, point localization-based, super resolution fluorescence microscopy has provided information on the nanoscale spatial organization of BCRs on B cell surfaces at the level of individual BCRs. The results of three such studies (30–32) are consistent with models in which BCRs exist as monomers and in protein islands, and antigen encounter induces the coalescence of these into active signalosomes (33). By contrast, the results of Maity et al. (34) were interpreted to be consistent with a model in which BCRs exist in clusters on resting B cell surfaces that are disrupted by antigen resulting in the initiation of signaling (35). Clearly much remains to be learned about the nanoscale organization of BCRs that will add to our understanding of the initiation of BCR signaling.

Ultimately, microclusters of BCR with bound antigen and other co-receptors visible by diffraction-limited light microscopy form and encounter the intracellular tyrosine kinase Lyn. Lyn phosphorylates ITAMs on Ig α and Ig β chains in BCR microclusters, providing a docking site for the tyrosine kinase Syk which initiates intracellular signaling cascades that allow the B cell to internalize antigen (36) [see Internalization of BCR and Intersection with MHCII in the MHCII Compartments (MIICs)]. Evidence from high-resolution total internal reflection microscopy in conjunction with fluorescence resonance energy transfer in living B cells argued that newly formed BCR microclusters perturbed the local lipid environment leading to the association of microclusters with a lipid raft probe and that this association facilitated the recruitment of Lyn to the BCR microclusters (37).

Soluble antigens are capable of initiating BCR clustering, but membrane-tethered antigens are more effective at inducing responses *in vivo* (38). This points to a critical role for FDCs and their use of long-term non-degradative compartments to store and recycle immune complexes and serve as an antigen depot (27). SSMs may also play a role in antigen presentation by conveying opsonized antigen directly to B cells after intercepting it in the SCS (38). Cell biological data indicate that APC/B cell interaction involves two major features. First, once stimulated, the B cell exhibits a membrane spreading and contraction response that assists with antigen aggregation and BCR microcluster formation and results in the formation of an immunological synapse with the APC (39, 40). Second, as the B cell separates from the APC, it extracts the target antigen from the surface of its partner cell.

The membrane-spreading response is accomplished within minutes of antigen contact by structural changes in the actin cytoskeleton. These changes involve cofilin-mediated severing of F-actin throughout the cell, followed by actin repolarization in the direction of the APC (41). The severing is thought to increase the mobility of cell-surface BCRs, assisting with formation of BCR/antigen microclusters and their movement into the center of the newly formed synapse (41, 42). The subsequent spreading response involves Arp2/3, resembling cell membrane extension of lamellipodia, whereas contraction requires dyneins and ERM proteins (40, 43). The microtubule cytoskeleton also changes shape in response to stimulation, with the microtubule-organizing center migrating rapidly toward the site of contact, in a process dependent on CDC42 and PKC ζ (44).

When contraction is complete, the synapse adopts a form with a central supramolecular activation complex (cSMAC), containing a large cluster of antigen-bound BCRs surrounded by a ring of actin and adhesion molecules, the peripheral SMAC (pSMAC) that tethers the B cell to the APC (43, 45). B cell extraction of antigen from the cSMAC is accomplished by mechanical force, in which B cells physically pull on synaptic antigen through the BCR and deform flexible membrane substrates to promote antigen internalization (46, 47). B cells also can extract antigen by release of degradative MHCII-containing vesicles, containing lysosomal proteases and lipases, into the pSMAC (44). Recent evidence suggests that the mechanism of antigen extraction is dependent on the properties of the APC (48). The presence of MHC class II in the synapse along with lysosomal enzymes raises the possibility of peptide loading inside the synapse.

Due to the high affinity of their antigen receptor, B cells expressing specific BCRs require 1,000–10,000× less antigen to stimulate T cells compared to non-specific B cells (49). BCR affinity affects the efficiency of antigen collection from the surface of presenting cells (39), with contractile forces in the presenting cell setting a BCR affinity threshold for successful antigen capture (46). After ligand binding, high-affinity BCR forms larger microclusters for recruitment of signaling machinery (50). Lower affinity BCR also bind and internalize antigens, if a BCR avidity threshold, presumably required for clustering and its consequences, is surpassed (51), as can occur with multivalent antigens, particularly at the surface of presenting cells (52).

Cross-linking of BCR by multivalent ligands initiates B cell activation for antigen presentation. Evidence for activation by monovalent antigen has also emerged (31), with data suggesting a threshold for signaling that must be overcome by monovalent antigen to lead to antigen presentation. Membrane-bound monovalent antigen is apparently more effective than soluble for reaching this threshold [reviewed in Ref. (53)]. For naïve B cells, monovalent antigens can activate IgM-BCR, but immune complexes are required for stimulation of IgD-BCR (35).

INTERNALIZATION OF BCR AND INTERSECTION WITH MHCII IN THE MHCII COMPARTMENTS (MIICs)

After antigen binding, B cells internalize antigens mainly through clathrin-dependent, BCR-mediated endocytosis (46,54). Clathrinmediated endocytosis involves formation of clathrin-coated pits (CCPs), which pinch off plasma membrane to form intracellular vesicles. CCP formation is initiated by the phosphorylation of the adaptor, AP-2, which co-localizes with BCR/antigen clusters (55). The BCR binds AP-2 through its Ig $\alpha\beta$ subunits, the membraneproximal tyrosine of a YxxØ (Ø = bulky hydrophobic) motif on the Ig β being the most important residue for receptor internalization (56). Intracellular vesicle generation requires actin polymerization and myosin activity (57, 58). After vesicle formation, clathrin disassociates, and the vesicle undergoes a cascade of biochemical changes, evolving into an early endosome and then into a late endosome. While most early endosomes are recycled back to the plasma membrane by Rab4, BCR signaling increases the rate of sorting into late endosomes *via* ubiquitination of BCR by c-Cbl and Cbl-b (59). Rab7 shepherds the transition of early endosome to late and appears to be directly modulated by BCR signaling (60). The final stage is delivery of vesicles carrying the antigenbound BCR to peptide-loading compartments.

Reagents that cross-link the BCR and initiate signaling are 10× more efficient in delivering BCR to MHCII peptide-loading compartments than monovalent reagents (61). Lyn kinase activity associated with BCR signaling directly enhances endocytosis *via* phosphorylation of the clathrin heavy chain, which promotes its association with actin-coupling clathrin light chains (62, 63); BCR signaling also results in tyrosine phosphorylation of CCP-associated proteins (epsin15, epsin2, etc.) and downstream proteins (e.g., Vav, Btk) that play roles in both signaling and endocytosis (60, 64). BCR signaling promotes receptor endocytosis, but the exact roles for Iga β motifs remain elusive, as signaling and endocytosis compete for the same motifs (56). Indeed, some evidence supports a model in which distinct BCR/

antigen complexes mediate signaling and antigen internalization (65). Internalization of BCR inhibits signaling, as localization in endosomal compartments reduces surface BCR and abrogates the activity of signaling proteins, such as Lyn, Syk, and BLNK (66).

Endocytosed BCR/antigen complexes intersect with newly synthesized MHCII en route to the cell surface. Nascent MHCII $\alpha\beta$ heterodimers are generated in the endoplasmic reticulum (ER) and rapidly associate with trimers of the type II membrane glycoprotein, Ii (CD74), a dedicated MHCII chaperone. Interaction with Ii facilitates MHCII folding and inhibits ligand binding to class II in the ER. Ii contains a peptide sequence (aa 81–104), referred to as the class II-associated invariant-chain peptide (CLIP) region, which non-covalently binds the MHCII groove. Four Ii isoforms exist: p41 and p43, which are expressed at low levels in B cells compared to other APC, and p33 and p35. p35 associates with clathrin-associated adaptor proteins; however, if all Ii species are p33, the complex can exit the ER as a pentamer (67). Export out of the ER occurs when all the retention motifs present in the trimer are "cancelled" (68). Trafficking of egressed MHCII-Ii complex, first to the trans-Golgi network, is controlled by cytosolic sorting tags. Recognition by AP-2 of two dileucine motifs in the Ii cytosolic domain enables αβIi sorting to clathrin-coated vesicles, and eventual localization to MIICs (69, 70). The exact definition of MIIC has been a matter of debate, but these compartments can be broadly characterized as late endosomal structures enriched in proteolytic enzymes and disulfide reductases and with sufficiently low pH to activate proteases involved in Ii degradation (see below) and antigen processing (54). The compartments exist in a several forms-multivesicular bodies (MVBs) or multilamellar structures—that are generally LAMP-1 positive (71).

Many Ii-mediated events are common to professional MHCII-expressing APC. However, lack of Ii in B cells and DCs in Ii-deficient mice shows some differential impact on MHCII-restricted antigen presentation. Some Ii-deficient DCs undergo normal maturation and present a broad range of peptidic epitopes from antigen internalized by fluid-phase endocytosis (72). By contrast, Ii is required for presentation of BCR-internalized antigen (73), and Ii-deficient B cells failed to present Ag-generated epitopes unless these epitope peptides were exogenously supplied (74), consistent with findings that, in B cells, Ii controls the transport of MHCII to and from MIIC (75) where BCR-delivered antigens are concentrated.

The intersection of antigen-bound BCR and MHCII appears to be actively promoted, as BCR binding by Fab'2 is associated with movement of MHCII to LAMP-1+ MVB vesicles (76) and ligated BCR and Ig- α /Ig- β also localize to these compartments, in a ubiquitin-dependent manner (77). Interaction of internalized BCR/antigen with intracellular MHCII was observed in pulldowns using biotin-labeled antigen (78). Strikingly, a rare (10%) conformation of MHCII seems to be a predominant conformer in complex with the BCR/antigen. This conformer (defined by the Ia.2 epitope) is highly effective for T cell activation, is enriched in lipid rafts, and stimulates intracellular calcium signaling in B cells (79, 80) Selectivity of the BCR/antigen for this MHCII conformer may contribute to the efficiency of BCR-mediated antigen presentation.

GENERATION OF THE MHCII/PEPTIDE REPERTOIRE: PROTEASES AND REGULATION OF PEPTIDE LOADING

Proteolytic events relevant for MHCII antigen presentation in B cells include processing of Ii and processing of endogenous and exogenous antigens, including BCR/antigen (81). Key enzymes implicated in epitope generation for MHCII presentation include lysosomal proteases that can be classified based on their catalytic mechanism into four major families (Table 1): (1) the cysteine cathepsins with a papain-fold (B, C, F, H, K, L, O, S, V, W, and X); (2) the aspartyl cathepsins D and E; (3) the serine cathepsins A and G, and (4) single member family of asparagine endopeptidase (AEP), a cysteine protease grouped with the caspases (82). In addition, gamma-interferon-inducible lysosomal thiol reductase (GILT) cleaves protein disulfide bonds (83). Cathepsins are synthesized as inactive zymogens that get modified by addition of mannose-6-phosphate residues (M6P), allowing recognition by M6P receptors in the trans-Golgi network and delivery to endosomal/lysosomal compartments. Cathepsin activation requires catalysis by other lysosomal proteases or by autocatalysis under certain circumstances, such as low pH or presence of glycosaminoglycans. Lysosomal proteases of the cysteine type have optimal activity at acidic pH, but other proteases are active over a range of pH. Inactivation occurs by protease degradation or by oxidation. Levels of expression and activity of cathepsins are controlled by multiple factors, including mRNA levels, local pH, and other protein factors that influence cathepsin trafficking and activation. For example, endogenous inhibitors like cystatins, stefins, tyropins, and serpins also regulate cathepsin function, although it appears that cystatins do not control antigen presentation (84).

Proteolytic processing of Ii is required for loading of antigen peptides onto MHCII. Ii processing, which starts with formation of a 22 kDa leupeptin-induced protein, can be initiated by AEP (94). The second intermediate, 10 kDa fragment small-leupeptininduced protein (SLIP) is then cleaved to generate the 25aa CLIP fragment. Cat S is the key component for converting the 10 kDa fragment into CLIP. In the human thymus, Cat V or L2 (Cat L in mice) is the main protease converting SLIP to CLIP. Studies of B cells and DCs from animals deficient in Cat S showed accumulation of surface MHCII with the 10 kDa Ii fragment (100), and reduced antigen presentation. In mice, in the absence of Cat S and L, Cat F in macrophages can generate CLIP, indicating cathepsin redundancy (82).

The crystallizable fragment (Fc) and antigen-binding fragment (Fab) domains of immunoglobulin remain largely intact in the MIIC of B cells (101, 102). Notably, however, the two domains can be immunoprecipitated separately, and co-precipitation of antigen with the Fab domain can be detected (101). This finding implies that proteolytic enzymes resident in the MIIC cleave intact immunoglobulin. In bone marrow-derived APC (over 80% macrophages), Cat B appears important for degradation of immunoglobulins that are taken up by Fc receptors (85).

Characterization of lysosomal proteases in B cells is still incomplete. Furthermore, one shortcoming of some studies on

Lysosomal protease	Type/activity	Distribution in immune cells	Involved in (reference)
Cathepsin B	Cysteine/endo- and exopeptidase (carboxydipeptidase)	BCLs, DCs, and macrophages	Processing of immune complexes internalized via $Fc\gamma R$ (85)
Cathepsin F	Cysteine/endopeptidase	Macrophages	li processing in absence of Cat S and L (86)
Cathepsin H	Cysteine/endo- and exopeptidase (aminopeptidase)	Ubiquitously expressed	Octapeptid (mini-chain) confers aminopeptidase activity; without the mini-chain function similar to Cat S (87)
Cathepsin S	Cysteine/endopeptidase	BCLs, DCs, macrophages, and epithelial cells	Converting the 10-kDa SLIP fragment from li into CLIP; processing of exogenous and endogenous antigen (88)
Cathepsin V or L2 (human homolog of mouse cathepsin L)	Cysteine/endopeptidase	Cortical human thymic epithelial cells; testis	li proteolysis in thymus (89)
Cathepsin D	Aspartic/endopeptidase	Ubiquitously expressed; also secreted	Destructive processing of myoglobin (90); there is evidence of redundancy among the aspartic cathepsins (Cat D and E)
Cathepsin E	Aspartic/endopeptidase	BCLs, B cells upon activation, monocyte- derived DCs, myeloid DCs, Langerhans cells	Processing ovalbumin; processing of TTC (91); processing of myoglobin (90); there is evidence of redundancy among the aspartic cathepsins (Cat D and E)
Cathepsin A	Serine/exopeptidase (carboxypeptidase)	Widely expressed, including primary B cells, BCLs, myeloid DCs, plasmacytoid DCs (92)	Removes amino acids from proteins or peptides C-terminal; may be important for fine trimming of peptides (92)
Cathepsin G	Serine/endopeptidase	Neutrophils, mast cells, B cells (endogenous/ exogenous source), monocytes, myeloid DCs, plasmacytoid DCs; granule associated	Destructive processing/processing of MBP in primary B cells (93)
Asparagine endopeptidase	Cysteine/endopeptidase	BCLs; low in B cells; thymic CD11c+ DCs; F4/80+ thymic macrophages	Initial processing of li chain (94); <i>in vitro</i> intracellular processing of TTC (95); <i>in vitro</i> destructive processing of MBP (96, 97); not necessary for TTC processing <i>in vivo</i> (98); processing of Cat B, H, and L (99)

BCLs, B cell lymphoblastoid line; DCs, dendritic cells; li, invariant chain; SLIP, small-leupeptin-induced protein; CLIP, class II-associated invariant-chain peptide; TTC, tetanus toxin, C-fragment; MBP, myelin basic protein.

B cell proteases is the use of B lymphoblastoid cell lines (B-LCL), which may not reflect the endo/lysosomal environment in primary B cells. For example, B-LCL degrade myelin basic protein (MBP) by AEP, but human primary B cells have low level of AEP activity and instead use exogenous Cat G (103), a granuleassociated protease that binds the B cell surface via a thrombinlike receptor and is internalized into endosomes (93). In addition, human B-LCL contain active CatS, CatB, and CatD, whereas none of these cathepsins are detected in resting primary human B cells (103). Cat E expression is enhanced in primary tonsillar B cells after activation with Staphylococcus aureus (104); similarly, PMA activation increases expression of Cat E in primary B cells from blood and increases presentation of tetanus toxin C-fragment to T cells (91). Thus, some of the differences between B-LCL and primary B cell lysosomal protease content may be due to activation state. Murine primary B cells express lysosomal GILT, and GILT activity decreases Cat S expression and function (105), suggesting that GILT participates in control of cathepsin activity in B cells.

The cathepsin profile of B cells has similarities to the DC profile (**Table 1**). B cells and DCs have lower levels of lysosomal proteases, with corresponding lower proteolytic activity, compared to macrophages. These differences appear to be specific to the lysosomal proteases and cannot be ascribed to number of lysosomes. Although the rate of antigen internalization is similar between DCs and macrophages, the rate of proteolysis is slower in DCs. For test antigens, protection from degradation is associated with enhanced presentation to T cells (106).

Studies of mice deficient for specific proteases show that proteases can influence the repertoire of peptides presented by MHCII. However, it is largely unknown if these effects are mediated by direct participation of specific proteases in endosomal antigen processing or predominantly influence Ii processing. For example, a series of studies showed that in B cells, as well as in DCs, Cat S does not directly influence the generation of most peptides presented by MHCII, but some peptides are affected [reviewed in Ref. (88)]. An additional challenge is that degradation of a given protein may require a series of proteases. For example, degradation of MBP in vitro is initiated by AEP and Cat S, and intermediates are subsequently degraded by other proteases including Cat S, D, and L (96). Data from an *in vitro* antigen-processing model system using only three cathepsins (Cat B, H, and S) argued that sensitivity to these cathepsins, the structure of proteins, strength of peptide binding to HLA-DR, and HLA-DM susceptibility (see below) were all factors that influence which peptides are presented by MHCII (107). This in vitro system recapitulated generation of naturally processed peptides from test proteins, suggesting its potential value for epitope prediction and for modeling mechanistic steps in antigen processing for MHCII cargo (108). Determination of the in vivo roles for specific proteases remains a subject of on-going work.

Availability of antigen for processing is influenced BCR affinity. Higher affinity binding allows delivery of protected regions to later compartments for peptide generation and binding to nascent MHCII (although highest affinity binding may suppress epitope availability), while lower affinity allows release of antigen in earlier compartments and generation of peptides that can bind recycling MHCII (109). For certain antibodies with highly pH-dependent binding to antigen, lower pH of the later compartments may greatly facilitate epitope generation (110). BCR binding suppresses some and enhances availability of other T cell epitopes, with both types of determinants within the region of the antigen "footprinted" by the antibody (111). BCR-bound antigen can persist in B cells, providing a source of peptide longer than antigen taken up by fluid phase (112).

Like other professional APCs, B cells express a non-classical class II molecule, HLA-DM (DM; H2-M in mice), with highest concentrations in late endocytic compartments. Despite the structural similarity with classical MHCII, DM does not directly bind peptides (113); instead, it modulates the peptidome via a catalytic mechanism (114, 115). DM function is crucial; DM-null APCs are defective in MHCII antigen processing and presentation (116) and DM-deficient mice present a skewed peptide repertoire, impairing selection of CD4+ T cells (117). The generally accepted function of DM is that it stabilizes a peptide-receptive form of MHCII by a transient association (118, 119); and this temporary DM/MHCII interaction facilitates the removal of weak binding peptides, such as CLIP and cryptic epitopes, while promoting the loading of high-affinity peptides that mostly become immunodominant epitopes for MHCII presentation at the surface of B cells (120-123). DM-regulated epitope selection (DM editing) is a pH-dependent process with pH optima (4.5-5.5), recapitulating late endosomal-lysosomal conditions in MIICs (124, 125).

Another non-classical class II molecule, HLA-DO (DO), also plays an essential role in antigen presentation in B cells (126–129), as knockout of the murine DO homolog, H2-O, leads to immunodeficiency and autoimmunity that are directly related to dysfunction of antigen presentation by bone marrow-derived cells (130). Although DO is structurally quite homologous to classical MHCII, there is no evidence that it can directly present peptides. However, DO mimics MHCII in binding to the same site on DM, such that it blocks the catalytic action of DM on MHCII/peptide complexes (113, 131). The requirement for DM inhibition by DO in antigen presentation in a restricted group of APCs (132) implies unique features of the MHCII peptidomes in these cells, especially B cells.

The affinity of DO/DM binding (nanomolar) (125, 133) is 1,000- to 10,000-fold stronger than that of the transient binding of MHCII to DM (micromolar) (131), and DO requires the tight association with DM to exit the ER for translocation to its lysosomal destination (134). The molecular interaction between intact DO and DM is pH insensitive, even at the lysosomal pH (125). This raises questions as to how DO-associated DM is freed to edit MHCII-associated peptide cargo, and what role DO plays to modulate the B cell MHCII peptidome. The stoichiometry of DM:MHCII is a critical factor in the pH-dependence of DM editing (125). DM abundance can compensate for the reduction in its enzymatic activity at pH > 5.5, and this potentially allows DM editing in high DM-expressing APCs, like B cells, to emerge outside late endocytic MIIC. Therefore, DO likely serves as a regulator to control the abundance of free DM and keep DM function focused on acidic MIICs (also see GC B Cells).

Multiple mechanisms have been postulated to explain how DO/DM complexes facilitate peptide loading onto MHCII [details are reviewed in Ref. (129, 135-137)]. Recent evidence argues that DO/DM complexes are inert for peptide exchange catalysis, but that in lysosomal conditions, low pH promotes denaturation of DO within the complex to elevate the level of free, catalytically active DM (125, 133). pH-dependent DO removal is an irreversible process that may be associated with many antigen-triggered events in the B cell, including the acidification of lysosomes, BCR signal-mediated transcriptional modifications of MHCII or the class II transactivator (CIITA), and redistribution of MHCII proteins within a multivesicular MIIC. An alternative view, that DO optimizes presentation of DM-resistant peptides through direct interaction with a peptide-receptive form of MHCII, has been presented (138).

The cytoplasmic tails of both DM and DO contain lysosomal targeting motifs. Interestingly, without DO, DM can reside outside of lysosomes, for instance, on the surface of immature myeloid DCs (139) or on secreted exosomes (140). Although no high-resolution imaging data are available, these studies at least suggest that the sorting motif on DO constrains the intracellular location of DO/DM complexes to lysosomes, where DM is further distributed away from internal membranes (source of exosomes) to the limiting membrane (140, 141). As (usually foreign) antigens associated with BCRs are typically protected from proteolysis at early endosomal pH, DM editing likely is unnecessary at the earliest stages of antigen endocytosis in B cells. Lysosome-oriented trafficking together with functional inhibition conferred by DO would focus DM editing on epitopes derived from BCR/antigen complexes that also localize to MIICs. DO-regulated trafficking may rely on the ubiquitination of cytoplasmic tails of DO heterodimers by, for example, a MARCH family E3 ligase (142).

The ectodomain of DM has reduced activity at pH 4.6 likely due to denaturing of free DM (125). This implies that DO may reciprocally chaperone and stabilize DM through tight association and maintain DM levels in lysosomes; this effect is also suggested by evidence in DO-expressing DCs from transgenic mice (143). Such bi-directional chaperoning would persist until an antigen exposure drives DO removal in MIIC. In agreement with this hypothesis, several studies have shown a concomitant decline of DM and DO during B cell activation (126, 127), although the DM:DO ratio increases in favor of DM editing.

As DO/DM and BCR/antigen complexes meet in MIICs, DM may also function in facilitated antigen hand-off from BCR to MHCII proteins. The recent findings that DM and mIg coprecipitate from cell lysates and that soluble recombinant DM and Fab fragments (but not Fc fragments) co-precipitate *in vitro* raise the possibility of a multimeric class II peptide-loading complex, including membrane Ig, DM, and MHC proteins (144). **Figure 1** summarizes key events described in Section "Generation of the MHCII/Peptide Repertoire: Proteases and Regulation of Peptide Loading" of the review.



MIC. Although li processing begins as MHCII/li complexes traverse earlier endocytic compartments, exchange of the final li remnant peptides (CLIP) bound to the MHCII peptide binding groove with antigenic peptides or self-peptides is inefficient before MHCII enters MIIC. Mechanisms limiting peptide schange until this stage include (1) DO significantly inhibits DM catalysis of peptide loading and (2) Fab-associated antigen fragments are largely protected from proteolysis to peptides. At late stages of these pathways, lysosomes merge with MIIC. This generates the highly acidic condition that favors denaturation and degradation of DO and Fab, and DM-mediated peptide exchange. Most derivatives of self-proteins internalized without mlg cannot survive endosomal/lysosomal proteolysis. Arrows indicate protein trafficking through different intracellular compartments (indicated by colors) along each pathway. Dotted lines indicate unfavorable processes. ECF, extracellular fluid; MHCII, major histocompatibility complex class II proteins; MIIC, MHCII compartments; CLIP, class II invariant-chain-derived peptides.

MHCII/PEPTIDE COMPLEX TRANSPORT TO THE SURFACE

The molecular details of transport of newly loaded MHCII/peptide complexes from the MIIC to the cell surface are not delineated in mechanistic detail for any APC, although clearly CLIP replacement with other peptides is not a requirement for egress from MIICs. More is known about the regulation of MHCII surface expression and its relationship to recycling of newly synthesized MHCII in complex with peptide. Available data argue that B cell activation does not influence the endocytosis of surface MHCII, yet it clearly alters the level of MHCII expression. Cho et al. demonstrated that internalized MHCII efficiently recycles in activated B cells, but not in resting B cells. The lack of MHCII recycling in these cells is a consequence of oligoubiquitination of a single conserved lysine (K225) in the C-terminal cytosolic tail of the MHCII β chain by the E3 ubiquitin ligase MARCH1. Further, MHCII ubiquitination promotes sorting to MVB and turnover of MHCII in lysosomes. Upon antigen ligation of BCRs, MARCH1 expression and MHCII ubiquitination is halted, enabling MHCII escape from degradation, rapid recycling, and increased surface expression. The activation induced boost in MHCII also promotes the ability of B cells to present the activating antigens (145).

MHCII ANTIGEN PRESENTATION OVER THE SPAN OF B CELL DEVELOPMENT

Early B Cell Development

Early stages of B cell development in the bone marrow (pro- and pre-B cells) are defined by stepwise rearrangement of V, D, and J gene segments at the Ig heavy and light chain loci [(146); see **Figure 2**]. The first BCR complex, expressed at the pre-B cell stage, is composed of the heavy-chain gene product with a surrogate light chain (λ 5-Vpre-B), together with the Ig α /Ig β heterodimer (147, 148). Interestingly, antigen-presenting machinery is already present at the pro- and pre-B stages, as CIITA is expressed and drives expression of MHCII, HLA-DM, and Ii (149, 150). The presence of most MHCII pathway components (except DO β) in the absence of mature BCR raises the question of their potential function in B cell development. Strikingly, the Ii N-terminal domain (aa 1–82) has an alternate activity in B cells that does not relate to its role in antigen presentation. Ii-deficient mice



show an arrest in B cell development with an enlarged population of immature IgD^{low} CD23^{low} transitional B cells. When the Ii N-terminal domain alone is expressed in transgenic mice, MHCIi trimer formation and egress from the ER is recovered, but MHCII/peptide formation and surface expression is not restored (151). Rather, this Ii domain initiates a signaling cascade, resulting in NFkB activation and B cell maintenance (152). The intramembrane protease SPPL2a (presenilin homolog signalpeptide-peptidase-like 2a) catalyzes the generation of this Ii fragment in mice and human B cells (153) Ii has third role for B cells as the receptor for macrophage migration inhibitory factor and interaction leads to signaling cascade that promotes B cell chemotaxis and survival (154).

In contrast to effects of Ii knock out, B cell development is only subtly altered in MHCII-deficient mice. In these mice, a mature B cell population is established, although the overall number of mature B cells and their capacity for antigen presentation is impaired (155, 156). B cells also reach the mature stage in bare lymphocyte syndrome patients, who lack MHCII expression due to mutation of CIITA (157). In DM-deficient mice, B cells mature, but display a reduced response to antigen (117). In light of these results, Chen and Jensen (156) proposed that the presence of the class II antigen presentation machinery in pre-, pro-, and immature BM B cells serves as a mechanism to induce peripheral CD4+ T cell tolerance through presentation of peptides from selfantigens (156). Low or no MHCII/CLIP complexes are detected on these B cells, implying that MHCII undergoes at least one peptide exchange cycle to remove CLIP and bind self-peptides. This possibility is consistent with expression of DM without DO at these early stages. As CD40 is not expressed on immature B cells, recognition by autoreactive T cells would likely result in anergy or deletion. Additional work is needed to confirm this hypothesis.

Immature Transitional and Naïve B Cells

Following BCR maturation, IgM is expressed on the surface of immature B cells that exit the BM and migrate to the spleen as transitional B cells (12). Close to 85% of the original B cell repertoire is eliminated by negative selection of cells expressing autoreactive BCR. Transitional B cells further mature to naïve/ pre-immune B cells of the FO B subset in blood and SLO or to MZ B cells of the splenic white pulp (158, 159). Expression of DO emerges in naïve (IgM+/IgD+/CD40+) B cells (142). DM levels are relatively stable throughout B cell development, leading to a high DO:DM ratio at this stage. This ratio inhibits DM activity, particularly in early, higher pH endosomal compartments, which are accessed by fluid-phase endocytosis of self-proteins. Reduced DM peptide exchange activity results in high levels of MHCII/ CLIP complexes on naïve B cells. DO activity may thus limit the MHCII/self-peptide repertoire in naïve circulating B cells, reducing the possibility of recognition by autoreactive T cells (128, 160).

GC B Cells

Presentation of MHCII/peptide by FO B cells in the lymph node mediates cognate interaction with CD4+ T cells, co-stimulated by CD40-CD40L interaction. Initial T cell contact drives B cell differentiation into centroblasts and centrocytes organized in GCs, with B cell-rich dark zones, where centroblasts undergo somatic hypermutation, yielding clonal variants of GC B cells with modified antigen affinity and specificity (161). Centroblasts further differentiate into centrocytes and enter the GC light zone, where encounters with FDC reintroduces antigen to the modified BCR. Antigen uptake and presentation permits cognate interaction with T_{FH} cells, in proportion MHCII/peptide levels (4). Centrocytes with higher affinity for antigen are selectively preserved, whereas those with reduced BCR affinity are deleted. Additionally, a subset of centrocytes undergoes class-switch recombination for expression of IgG, IgA, or IgE (161–163). Reduced DO expression marks GC entry of B cells and is primarily driven by post transcriptional events, such as ubiquitination by MARCH9 (126, 127, 142). Centrocytes likely benefit from maximum DM editing both kinetically and stoichiometrically to achieve high levels of stable MHCII/peptide complexes and garner T cell help. Such a high level of DM activity cannot simply stem from transcriptional upregulation; indeed, GC B cells express slightly lower levels of DM protein than naïve or memory B cells (127). This suggests that the boost in DM editing originates from pre-existing inactive DM that was most likely associated with DO.

Major histocompatibility complex class II/peptide complexes on the surface of GC centroblasts are ubiquitinated and targeted for degradation, whereas centrocytes express long-lived MHCII/ peptide complexes (164). The MHCII turnover in these cells is controlled by oligoubiquitination of the MHCII β chain cytosolic tail by MARCH1. MHCII ubiquitination prevents the accumulation of older MHCII/peptide complexes in GC centroblast B cells and allows antigen captured *via* affinity modulated BCR to be presented on newly expressed MHCII, thus reporting on the SHM process (165). Once cells move to the light zone as centrocytes, MHCII/peptide complexes on the cell surface are stabilized, facilitating productive interactions with CD4+ T cells during selection for high-affinity clones.

Post GC Memory B and Plasma Cells (PC)

Centroblasts and centrocytes circulate between the light and dark zone until an optimal level of BCR affinity is achieved, and the selected B cells can further differentiate into long-lived memory B cells or antibody-secreting PC (161). Memory B cells express DO and DM and reestablish a higher DO:DM ratio compared to GC B cells. The model of irreversible DO removal (see Generation of the MHCII/Peptide Repertoire: Proteases and Regulation of Peptide Loading) need not conflict with the observation that memory B cells leaving the GC have upregulated DO levels (126, 127), given the rapid differentiation and cell division of phenotypically polarized GC B cells (166). Asymmetric segregation of proteins relevant for T cell interaction could allow polarized GC B cells to give rise to daughter cells with distinct fates: an effector-prone cell that bears most of free DM capable of antigen presentation to T cells, and a memory-prone cell that inherits most of the DO/DM complex and exits GC with high DO:DM ratio. Surface MHCII/CLIP levels on memory B cells are lower than levels on naïve cells (167); this finding could result from stability of MHCII/peptide complexes acquired during the GC stage.

Studies of mice lacking MHCII expression in memory cells showed impaired ability of these cells to differentiate into PC upon antigen restimulation (168). During the B memory response, B/T cell cognate interaction again promotes selective activation of affinity matured memory B cells that will yield PC secreting high-affinity antibodies. After B cell differentiation into PC, expression of CIITA, which controls most components of the class II processing pathway, is repressed, yielding marked reductions in the levels of DM, DO α , Ii, and MHCII (167). Though DO β expression together with DO requirement for DM to egress from the ER results in loss of functional DO.

CONCLUSION

Many essential molecular mechanisms modulating the protein dynamics of MHCII-restricted antigen presentation by B cells have come to light in recent years, revealing the elegance of this system for carefully regulated functions across the span of the B cell life cycle. In parallel (though not the focus of this review), emerging evidence indicates a critical role for B cell antigen presentation function in various immune and autoimmune reactions. Nonetheless, key questions remain to be answered. What initiates and regulates fluid-phase endocytosis in B cells and what is the role of this process in early B cells that lack mature BCR? What molecular events mediate independent and interdependent signaling and antigen internalization downstream of antigen binding to the BCR? How does antigen binding influence the architecture of the endocytic pathway and the composition and protease content of intracellular compartments? What factors in addition to pH influence the DM:DO ratios in endocytic compartments? What are the mechanisms involved transport to the surface of MHCII/peptide complexes? Are there unique features of the MHCII pathway in memory B cells? Undoubtedly, B cell-mediated MHCII antigen presentation is an exciting and important field to watch in the coming years.

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LA, S-cH, MM, KB, CM, WJ, and EM contributed sections to this review. EM edited the full review and generated the final draft. LA, WJ, and CM contributed the tables and figures.

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Adler et al

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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.

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