Toll-Like Receptor Expression and Responsiveness of Distinct Murine Splenic and Mucosal B-Cell Subsets

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Background. Toll-like receptors (TLRs) are pattern recognition receptors that recognize pathogen associated molecular patterns and trigger innate immunity leading to initiation of adaptive immunity. TLR-mediated activation of dendritic cells (DCs) is known to be a critical event in the initiation of cellular and humoral immune responses. Recent work however suggests that B cells also express TLRs, and that they can be activated via TLR ligands. However, whether such B cell activation occurs only on memory B cells, or whether it can also occur on truly naïve B cells remains controversial. Furthermore, the expression and functional relevance of TLRs on distinct subsets of B cells, which are known to play differential roles in humoral responses is not known. Methodology/Principal Findings. In this study, we investigated the expression pattern of different TLRs in distinct subsets of murine B cells (naïve, memory, follicular, marginal zone, B-1 and peyer’s patch). In contrast to the reported restricted expression pattern of TLRs in human peripheral blood naïve B cells, murine splenic naïve B cells express a variety of TLRs with the exception of TLR5 and 8. Consistent with this relatively broad expression pattern, murine naïve B cells proliferate and secrete antibody to a variety of TLR agonists in vitro, in the absence of B-cell receptor cross-linking. In addition, we observed subtle differences in the antibody secretion pattern of follicular, marginal zone, B-1 and peyer’s patch B-cell subsets. Conclusions/Significance. Thus various B cell subsets, including truly naïve B cells, express multiple TLRs, and signaling via such TLRs results in their robust proliferation and antibody secretion, even in the absence of dendritic cell activation, or T-cell help.

INTRODUCTION

TLR-mediated recognition of microbial stimuli often leads to activation of innate immune cells including dendritic cells (DCs) [1,2,3]. TLR signaling promotes activation and maturation of DCs which instruct and support T-cell activation, leading to cell-mediated adaptive immune response [4]. Cognate interaction between activated antigen-specific T cells and naïve B cells promotes B-cell clonal expansion and differentiation leading to a humoral immune response. Differential expression pattern of TLRs have been reported in human and murine DC subsets [5,6]. In humans, myeloid DCs express TLR2 and 4, whereas plasmacytoid DCs express TLR7 and 9 [7].

Recent studies suggests that in addition to TLR signaling in DCs, direct TLR mediated activation of B cells is also required for eliciting humoral immune response [8]. Thus chimeric mice in which only B cells are deficient in TLR signaling, fail to mount antibody responses to protein antigens given with adjuvants [8]. Consistent with this observation, previous work suggests that murine B cells can be stimulated in vitro by TLR4 and TLR9 ligands to proliferate and secrete antibody [9,10]. In contrast to murine B cells, naïve human B cells do not express TLR4 or TLR9 and hence do not respond directly to LPS or CpG [11]. However, it was recently reported that human memory B cells in the blood do express TLR9 and respond to CpG DNA [12], and consistent with this, cross-linking of BCR results in upregulation of TLR9 expression, and responsiveness to TLR9 ligands [12]. In contrast to this, recent studies revealed no differences in TLR expression in naïve versus germinal center versus memory B cells in human tonsils [13,14].

Therefore, TLR activation of B cells likely plays a critical role in the regulation of humoral immunity and memory, and understanding the precise roles played by different TLRs in this regard is a critical first step in exploiting this in the design of vaccines that induce rapid and persistent neutralizing antibody. However, this is complicated by the fact that there are several subsets of B cells that play distinct roles during immune responses. For example, follicular B cells are shown to be important for T-dependent immune responses whereas marginal zone B cells are important for T-independent immune responses [15]. Marginal zone B cells are shown to be in a pre-activated state and respond rapidly to LPS and secrete antibody in vitro [16]. Peyer’s patch B cells in the intestine play critical role in mucosal immune response by secreting IgA that binds to pathogens and prevents enteric infections [17]. B-1 B cells, a subset of B cells in the peritoneal cavity is the source of natural IgM present in the serum and plays an important role in immunity against blood borne pathogens [18]. However, the expression patterns of TLRs on distinct B-cell subsets, and their responsiveness to various TLR ligands is poorly understood.

To understand the importance of TLR signaling in B cells and to clarify the differences reported to exist between mouse and human B cells, we determined the expression pattern of TLRs in...
distinct murine B cell subsets and their response to TLR agonists in vitro in the absence of antigenic stimulation (BCR cross-linking). Our data suggest that mouse naive follicular B cells express and undergo polyclonal expansion and differentiation to almost all known TLRs except TLR5 and 8. We demonstrate that different B-cell subsets including follicular, marginal zone, B-1 and peyers patch B cells proliferate and secrete polyclonal antibodies to a variety of TLR agonists in vitro.

**MATERIALS AND METHODS**

**Mice**
C57BL/6 mice were purchased from Jackson Laboratory (Maine). ROSAYFP [19] and germinal center-cre (GCC) [19] mice were genotyped via PCR of genomic DNA derived from tail-tip biopsies. Primers for ROSAYFP PCR and Cre-PCR have been described [19]. All GCC mice were maintained by mating with C57BL/6 and housed under specific pathogen-free conditions at the Emory Vaccine Center. All animal studies were approved by the Institutional Animal Care and Use Committee of Emory University.

**Flow cytometry**
Splenic B cells were bead purified using CD19 microbeads (Miltenyi Biotech). CD19 enriched B cells were stained for Cyochrome anti-CD45R/B220 (RA3-6B2), PE anti-CD23 and FITC anti-CD21 (BD Pharmingen, San Jose, CA). B-cell subsets were FACS sorted using a MoFlo cell sorter (DakoCytoMation, Fort Collins, CO).

**Real-time PCR**
Total RNA was isolated from FACS sorted B cell subsets with the RNA miniprep (Invitrogen, Carlsbad, CA). RNA was quantified by OD260 using a DU® 530 Life Science UV Spectrophotometer (Beckman Coulter Inc, CA) and 2 μg of total RNA was subsequently used to make cDNA using the Superscript II reverse transcriptase (Invitrogen Corp., Carlsbad, CA) according to the manufacturer’s protocol. RT-PCR was performed using Biorad apparatus (Biorad Laboratories, Hercules, CA). Primers for murine TLRs were used as reported previously [5]. The β-actin specific primers were used for loading control (IDT technologies, Coralville, IA).

**Proliferation Assay**
FACS sorted B-cell subsets were incubated at 2×10^5 cells/well in triplicate in 96-well flat-bottom plates in medium consisting of RBMI-1640 supplemented with 10 mM glutamine, 10 mM HEPES, 0.5 mg/ml gentamicin, and 5×10^-5 2-ME (Complete IF-12 media). Stimuli added included (Fab')2 goat anti-IgM (μ-chain specific; ICN Pharmaceuticals, Cappel, ICN Pharmaceuticals, Aurora, OH) LPS (Invivogen), CpG ODN 1826 (Coley Pharmaceuticals, Wellesley, MA), Pam3Cys (EMC microcollections, Germany), MALP-2 (Invivogen, San Diego, CA), and 3M-003 (kind gift from Dr. Sefik Alkan, 3M pharmaceuticals, St. Paul, MN). After 66 h, all cultures were pulsed with 1 μCi of [3H]thymidine and harvested 6 h later onto filter mats using a cell harvester (Packard, Meriden, CT). The levels of radionucleotide incorporation were measured with a Matrix 96 β-radiation counter (Packard, Downers Grove, IL). Results were presented as arithmetic mean of triplicate cultures ±SE.

**ELISA**
FACS sorted B-cell subsets were incubated at 2×10^5 cells/well in triplicate cultures in 96-well flat-bottom plates for 5 days. Day 5

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**Figure 1. TLR expression profile of murine B-cell subsets.** Real-time PCR profile of TLR expression in FACS sorted follicular B-cells (CD19^+ B220^+CD23^-CD21^-, panel A), marginal Zone B (CD19^+ B220^-CD23^-CD21^+, panel B), peritoneal B-1 (CD19^-B220^-CD23^+, panel C) and peyer’s patch B cells (CD19^-B220^, panel D) as indicated with β-actin as loading control. CD11c^+ dendritic cells were used as a positive control (panel E). Values represent the ratio of the TLR to β-actin.

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supernatants were analyzed for antibody isotypes using ELISA kit (BD Biosciences, San Jose, CA).

**Immunization**

C57BL/6 mice were immunized with sheep RBC (SRBC) 10% v/v intraperitoneally. On day 12, mice were sacrificed and B220⁺ PNA⁻ IgG⁺ splenic memory B cells FACS sorted using a MoFlo cell sorter (DakoCytomation, Fort Collins, CO) as described previously [20].

**RESULTS AND DISCUSSION**

**Expression profiles of TLRs in different B-cell subsets**

We determined the expression pattern of TLRs1-9 in the following murine B-cell subsets i) splenic follicular ii) splenic marginal zone, iii) peritoneal B-1 and iv) mucosal peyer’s patch B cells. In initial experiments, splenic follicular, marginal zone, peritoneal B-1 and peyer’s patch B-cells were screened for expression of 9 known TLRs. Total B cells were isolated from murine spleen using CD19 microbeads and stained with antibodies to B220⁺ PNA⁻ IgG⁺ splenic memory B cells FACS sorted using a MoFlo cell sorter (DakoCytomation, Fort Collins, CO) as described previously [20].

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**Follicular, marginal zone, B-1 and peyer’s patch B cells proliferate and secrete polyclonal antibodies to various TLR agonists**

First, we studied the functional relevance of TLR expression by various B-cell subsets. We found that murine B-cell subsets proliferate in response to 1 μg/ml of Pam3Cys (TLR2 agonist), LPS (TLR4 agonist), TLR2/6 (MALP-2), 3M-003 (TLR7/8 agonist) and CpG (TLR9 agonist) (Figures 2A-D) with varying degrees. Follicular B cells proliferate robustly in response to most of the TLR agonists except for TLR3 and TLR6, where the response is weak or absent and for TLR5 where the response is absent. Interestingly, marginal zone B cells proliferate robustly in response to TLR2 (Pam3Cys) and TLR6 (MALP-2) agonists but weakly to TLR4, 7 and 9 agonists. B-1 B cells respond weakly to most of the TLR agonists except TLRs 4, 7 and 9. Robust proliferation of peyer’s patch B cells was observed in response to TLR2, 6 and 7 ligation but not other TLRs. Alternately, CFSE
dilution of follicular B-cells in response to various TLR agonists was measured and B-cells appear to undergo multiple cell divisions in vitro (data not shown). The dose we used to stimulate B cells is optimal since dendritic cells stimulated with 1 μg/ml of TLR ligands secrete cytokines like IL-6, IL-12p40 and IL-10 depending on the stimulus [21,22]. Follicular and marginal zone B cells not only clonally expand in response to TLR ligation but also secrete antibodies of all isotypes except IgA which was abundantly secreted by B-1 and peyer’s patch B cells (Figures 3A-D). Consistent with their pre-activated phenotype, B-1 B cells have higher background IgM response and Peyer’s patch B cells have higher IgA response in unstimulated conditions [18,23]. The isotype profile appears to be different depending on the subset studied and the ligand used. Thus, our studies demonstrate that not only do different B-cell subsets express TLRs, but that they can also be stimulated directly by TLR ligands.

Both naïve and memory murine B cells respond to TLR agonists in vitro in the absence of BCR cross-linking

In humans, memory but not naïve B cells express and respond to TLR9 ligands in the absence of BCR triggering [12,24]. However, in mice previous studies suggest that so-called naïve splenic B cells do respond directly to TLR stimulation [9,10]. In our preliminary experiments, we sorted B220⁺CD23⁺CD21⁺IgG⁺, putative naïve B cells and stimulated them in vitro with TLR ligands and observed proliferation and antibody secretion (data not shown). However it was formally possible that putative IgM⁺ memory B cells might have been present in the B220⁺CD23⁺CD21⁺IgG⁺ fraction. To definitively address the issue of whether bona fide naïve B cells express TLRs and are responsive to TLR ligands, we utilized the recently described transgenic mouse system where memory B cells are marked based on YFP expression and can be tracked in vivo [19]. This transgenic system utilized the I-Eαa promoter and gene expression from this promoter was restricted in B lineage cells that had entered the germinal center (GC) pathway of B cell differentiation. The transgenic mice were subsequently bred to the ROSA-YFP cre-reporter strain, in which constitutive expression of YFP occurs upon cre-mediated recombination of the ROSA locus [19]. We FACS sorted naïve follicular (B220⁺CD23⁺YFP⁺IgM⁺) B cells from these transgenic mice thereby effectively excluding the contaminating memory B-cell population (Figure 4A). In parallel, we FACS sorted memory (B220⁺CD23⁺IgG⁺) B cells from sheep RBC (SRBC) immunized mice, and studied the expression and functional role for TLRs.
Interestingly, the TLR expression profile of naive and memory B cells was almost identical (Figure 4B). Unlike human peripheral blood naive B cells, murine splenic naive B cells respond to TLR ligation (proliferation and antibody secretion) equally well compared to unsorted B cells, ruling out potential memory B cell contamination in the subsets studied (Figure 4C & D).

In summary, we report here the expression pattern and function of TLRs in different murine B-cell subsets including follicular, marginal zone, B-1 and Peyer’s patch. B-cell subsets express all known TLRs except 5 and 8. Consistent with their expression pattern, FACS sorted purified B-cell subsets respond to TLR ligands in vitro (Figure 5). Robust proliferation and antibody secretion of follicular B cells was observed in response to various TLR ligands, particularly ligands to TLR2/1, TLR2/6, TLRs 4, 7 and 9 (Figure 5). Marginal zone B cells have a similar pattern of TLR expression and response (Figure 5). However, B-1B cells do not respond strongly to TLR2 ligands. In contrast, Peyer’s patch B cells respond robustly to TLR2, 6 and 7 ligation, but poorly to ligands for TLR9 and 4. Our results are consistent with a very recent report that B-cell subsets that participate in T-independent immune response express and respond to TLR agonists in vitro [25]. But in contrast to their observations, we show that follicular B cells that participate in T-dependent immune response do express and secrete antibodies of IgG3 and IgM isotypes in response to a variety of TLR agonists in vitro [25]. The observed differences between our study and the previous study could be due to different CpG motifs that were used (CpG 1826 versus CpG 1668). It is possible that the CpG ODN we used (CpG 1826) was more potent than the one used in the previous study (CpG 1668) [25,26].

Furthermore, we addressed the issue of TLR expression and function in naive versus memory B cells and found that mouse...
naive follicular B cells express multiple TLRs and respond to a variety of TLR agonists in vitro. The extent to which TLR signaling in B cells modulate antibody responses to vaccines needs to be examined. Surprisingly, a recent report from Gavin et al demonstrated that certain commonly used adjuvants including CFA and Ribi do not require TLRs for eliciting antibody responses [27]. In contrast, it was shown that VSV induced IgM neutralizing antibody was partially dependent on TLR7 [28]. Furthermore, our recent study suggests that the yellow fever vaccine-17D, one of the most effective vaccines stimulates T cell responses by activating multiple TLRs [29], although to what extent TLR signaling influences humoral immune responses need to be determined.

In conclusion, the data presented here highlight the importance of TLR expression and function in different murine B-cell subsets which contribute to T-dependent and T-independent responses and signifies the potential fundamental differences that exist between human and mouse naive B cells.

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Author Contributions
Conceived and designed the experiments: BP MG. Performed the experiments: MG. Analyzed the data: BP MG. Contributed reagents/materials/analysis tools: JJ. Wrote the paper: BP MG.
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