# **BLyS inhibition eliminates primary B cells but leaves natural and acquired humoral immunity intact**

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**We have used an inhibiting antibody to determine whether preimmune versus antigen-experienced B cells differ in their requisites for BLyS, a cytokine that controls differentiation and survival. Whereas** *in vivo* **BLyS inhibition profoundly reduced naïve B cell numbers and primary immune responses, it had a markedly smaller effect on memory B cells and long-lived plasma cells, as well as secondary immune responses. There was heterogeneity within the memory pools, because IgM-bearing memory cells were sensitive to BLyS depletion whereas IgG-bearing memory cells were not, although both were more resistant than naïve cells. There was also heterogeneity within B1 pools, as splenic but not peritoneal B1 cells were diminished by anti-BLyS treatment, yet the number of natural antibody-secreting cells remained constant. Together, these findings show that memory B cells and natural antibodysecreting cells are BLyS-independent and suggest that these pools can be separately manipulated.**

#### B lymphocyte | BAFF | immune memory | mouse

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Secondary immune responses and natural antibodies are key elements of protective immunity. During primary immune responses, naïve antigen-reactive B cell clones selectively expand, amplifying their frequency and yielding memory cells (reviewed in ref. 1). Although memory populations are a small proportion of total B cells, they turn over more slowly than their naïve precursors and afford protective immunity upon secondary antigen challenge (2, 3). Primary immune responses also generate long-lived plasma cells (LLPC), which persist indefinitely and maintain systemic antibody levels. Another source of protective antibodies are B1 B cells, some of which constitutively generate ''natural'' antibodies against endogenous antigens such as phosphorylcholine (PC) (reviewed in ref. 4).

The B lymphocyte stimulator (BLyS) family of cytokines and receptors plays a central role in B cell homeostasis (reviewed in ref. 5). This family includes the ligands BLyS and APRIL and the receptors BR3, TACI, and BCMA. BLyS can bind all three receptors, whereas APRIL binds only TACI and BCMA. Developing B cells begin to express BR3 as they exit the bone marrow and enter the mature follicular (FO) and marginal zone (MZ) pools (6). BLyS regulates these preimmune B cell pools via survival signals delivered through BR3, such that in the absence of either BR3 or BLyS, FO and MZ cells die rapidly (7, 8).

Memory B cells live longer than their naïve precursors (9), raising the question of whether memory B cells still rely on BLyS-BR3 signaling. Furthermore, LLPC express high levels of BCMA (10), which binds either BLyS or APRIL, raising the possibility that either cytokine might afford survival. Similarly, whether natural antibody-forming B1 cells also rely on BLyS is unclear. For example, peritoneal cavity (PerC) B1 cells require neither BLyS nor APRIL for development (11); but some B1 cells express BLyS, and PerC B1 cells also express APRIL (12).

Direct assessment of whether different B lineage subsets vary in their requirements for BLyS has not been possible, because

there has been no way to selectively manipulate BLyS availability. Herein we ask whether these B cell subsets rely on BLyS by treating mice with an inhibiting anti-BLyS antibody. Whereas this treatment eliminated preimmune B cells and primary antibody responses, PerC B1 and memory B cell numbers were unaffected. Moreover, both secondary antibody responses and natural antibody production were normal. Finally, BLyS inhibition did not affect established IgG antibody titers or LLPC numbers. Together, these findings demonstrate that natural antibody-forming cells, memory B cells, and LLPC are largely BLyS-independent. Moreover, they suggest that BLyS inhibition may be used to target specific B cell populations.

### **Results**

**In Vivo BLyS Inhibition Eliminates Most Primary B Cells.**We generated a hamster monoclonal antibody to murine BLyS (10F4) that effectively inhibited BLyS binding to BR3, TACI, and BCMA [\[supporting information \(SI\) Fig. S1\]](http://www.pnas.org/cgi/data/0807841105/DCSupplemental/Supplemental_PDF#nameddest=SF1) and used it for all work described here. Serum anti-BLyS and BLyS levels, as well as splenic FO B cell numbers, were followed after treatment with 100  $\mu$ g of 10F4 i.p. on days 0 and 5 (Fig. 1 *A* and *B*). The half-life of anti-BLyS *in vivo* was  $\approx$  2 weeks, and serum BLyS levels varied reciprocally with anti-BLyS levels. Control hamster  $IgG_1$  antibody had no effect on lymphocyte numbers or serum BLyS levels (data not shown).

Consistent with their lack of BLyS receptor expression (6, 13), developing bone marrow B cell subsets were unaffected by anti-BLyS treatment (data not shown). In contrast, all preimmune splenic B cell subsets were substantially diminished (Fig. 1*B* and [Fig. S2\)](http://www.pnas.org/cgi/data/0807841105/DCSupplemental/Supplemental_PDF#nameddest=SF2). Thus, the TR and FO pools were severely reduced after anti-BLyS treatment, and MZ B cells were eliminated. Autoreconstitution began at days 40–45 and mirrored the kinetics observed in other models (14). Transiently elevated BLyS levels were regularly observed at the onset of reconstitution.

Anti-BLyS treatment ablated splenic but not peritoneal B1 B cells (Fig. 2 *A* and *B*), reducing both B1a and B1b subsets in the spleen. To assess B1 function, we measured the number of spontaneous IgM anti-PC antibody-producing cells, because B1 cells are the main producers of such antibodies (15). The

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**Fig. 1.** BLyS inhibition *in vivo*. (A) C57BL/6 mice were treated with 100  $\mu$ g of anti-BLyS i.p. on days 0 and 5 and subsequently analyzed for serum BLyS (open bars) and anti-BLyS (filled bars). Each bar represents the average  $\pm$  SD for at least three mice per time point. Untreated mice  $(n = 34)$  are shown at day 0. Combined data from three separate experiments are shown. (*B*) Numbers of splenic follicular (FO) B cells  $\pm$  SD for the same mice shown in A.

numbers of these cells were similar in both spleen and PerC in mice treated with anti-BLyS or isotype control antibody, although variance in the spleen was higher in controls (Fig. 2*C*).

**Memory B Cell Compartments and Responses Resist Anti-BLyS Treatment.** Primary immune responses to both T-dependent (TD) and T-independent (TI) antigens were attenuated in anti-BLyS-treated mice [\(Fig. S3\)](http://www.pnas.org/cgi/data/0807841105/DCSupplemental/Supplemental_PDF#nameddest=SF3). To assess the influence of anti-BLyS on NP-specific memory cells and LLPC in wild-type animals, C57BL/6J mice received primary immunizations with NP-CGG, a well characterized TD antigen (Memory System 1) (see refs. 2 and 16 and details in *Materials and Methods*). Some mice were treated with anti-BLyS  $\geq$ 8 weeks later, when memory B cells and standing anti-NP titers had been established, but before secondary (booster) immunization with NP-CGG. Neither splenic NP<sup>+</sup> memory B cell numbers nor high-affinity  $I gG_1$  antibody titers were affected by BLyS inhibition before eliciting a secondary immune response (Fig. 3 *A* and *B*). Moreover, the increases in NP<sup>+</sup> B cells and NP-specific antibody after booster immunizations were similar in anti-BLyS treated and control mice. Finally, established NP-specific BM LLPC were also unaffected (Fig. 3*C*).

To extend these findings, we evaluated how anti-BLyS affected memory B cell survival in two additional memory systems that were based on adoptive transfer of NP-specific B cells from Vh186.2 transgenic (Tg; System 2) or knockin (KI; System 3) mice into recipients that mount poor endogenous responses to NP-CGG (see refs. 2 and 16 and details in *Materials and Methods*). These systems produce large numbers of antigenspecific memory B cells, permitting further resolution of memory subsets. In these systems naïve host B cells can serve as internal controls.

Thirty-three weeks after transfer and immunization, mice from Memory System 2 were treated with anti-BLyS or isotype control given in two doses 5 days apart. After 15 days, naïve recipient B cell numbers were reduced by 5.4-fold  $(P < 0.001)$ ,



**Fig. 2.** Effects of BLyS inhibition on B1 subsets. (*A*) Gating scheme for identification of B2, B1a, and B1b cells in spleen (*Upper*) or peritoneal cavity (Lower). Lymphocyte-gated cells are further identified as B1a (CD5<sup>+</sup> IgM<sup>+</sup>) or B2 (CD5<sup>-</sup> IgM<sup>+</sup>) as shown in left plots and histograms; in addition, B1a cells are  $CD23^-$  (data not shown), whereas B2s are CD23<sup>+</sup> and thus include late TR and FO B cells. B1b cells are CD5<sup>lo/-</sup> CD23<sup>-</sup> IgM<sup>+</sup> Mac1<sup>+</sup> as shown in right plots. (B) Numbers of splenic (*Upper*) and PerC (*Lower*) B2 and B1 cells. Cell numbers SD for three untreated mice (open bars) and three treated mice are shown. n.s., not significant. (*C*) Numbers of PC-specific antibody-producing cells in spleen and PerC of isotype control-treated (open symbols) or anti-BLyStreated (filled symbols) mice at 10 and 21 days after treatment. Each symbol represents one mouse.

as predicted (8) (Fig. 4 *A* and *B*). In contrast, NP-specific donor-derived IgM<sup>a</sup>-bearing memory B cells were not as affected by anti-BLyS; they showed only a 2-fold depletion that was not significantly different from control treatment. The host cells that do not express the Tg, and thus are positive for the endogenous IgM<sup>b</sup> marker, were also not significantly depleted. Although the exact identity of these cells is uncertain, the most likely explanation is that these rare endogenous cells (17) were expanded by environmental antigens and hence are memory cells.

We next investigated the effects of anti-BLyS in Memory System 3 [\(Fig. S5\)](http://www.pnas.org/cgi/data/0807841105/DCSupplemental/Supplemental_PDF#nameddest=SF5), which enabled comparison of class-switched  $\rm{IgG_1^+}$  vs. unswitched  $\rm{IgG_1^-}$  memory B cells. As expected, naïve



**Fig. 3.** BLyS ablation does not alter NP-reactive memory B cells or LLPC generated in wild-type mice (Memory System 1). (*A* and *B*) Memory B cell analysis of C57BL/6 mice immunized with NP-CGG. Primary immunization was at day 0, anti-BLyS treatment was at day 56, and NP booster immunization was at day 77. Memory B cell numbers and IgG titers were assessed at day 84 for all mice including untreated and primed-only controls. NP-reactive memory B cells were identified according to the FACS gating scheme shown in *A*. After doublet discrimination, DAPI exclusion, and lymphocyte gating, memory B cells were phenotyped as CD4<sup>-</sup>CD8<sup>-</sup>Gr-1<sup>-</sup>B220<sup>+</sup>NP<sup>+</sup>. NT, not treated control. NP-reactive memory B cell numbers and high-affinity anti-NP IgG<sub>1</sub> antibody titers, as assessed by ELISA, are shown in *B*. Each symbol in graphs represents an individual mouse, black lines represent means, and combined results from two separate experiments are shown. Symbols at the bottom indicate treatment given (+) or not given (–). (C) Number of NP-specific LLPC in the bone marrow of control mice or anti-BLyS-treated mice at 21 days after anti-BLyS treatment and 77 days after NP-CGG immunization. Phenotyping strategy for LLPC is presented in [Fig. S4.](http://www.pnas.org/cgi/data/0807841105/DCSupplemental/Supplemental_PDF#nameddest=SF4)

recipient B cells in immunized mice were reduced 8.1-fold (*P* 0.001) after anti-BLyS treatment, as were NP-binding naïve B cells from unimmunized donor mice (Fig. 4 *C*–*F*). In contrast, the total donor-derived memory B cell pool was reduced only 2.6-fold  $(P < 0.01)$ , similar to the extent of reduction seen in System 2 and substantially different from effects on naïve B cells. Among these donor-derived memory B cells, the  $\text{IgG}_1^-$  memory cells, which are nearly entirely unswitched IgM-bearing cells (our unpublished observations), were reduced 3.3-fold  $(P < 0.01)$ . In

contrast, the  $\text{IgG}_1^+$  memory B cells were not significantly depleted. Thus, while all memory B cells were relatively resistant to BLyS depletion compared with their naïve precursors, unswitched IgM-bearing memory cells remain somewhat BLySdependent, and IgG-bearing memory cells do not.

## **Discussion**

Mature B cells in preimmune pools depend on BLyS for survival, as evidenced by their rapid disappearance after *in vivo* BLyS inhibition. In contrast, memory B cells resist BLyS depletion, and recall responses are normal. Furthermore, neither LLPC nor standing antibody titers are impacted by BLyS inhibition. Finally, PerC B1 cells and natural antibody-forming cells are resistant to BLyS depletion. Together, these findings indicate that preimmune and memory B cell pools are governed by distinct survival requisites, suggesting that BLyS inhibition, while eliminating naïve B cells and primary responses, will spare most elements of acquired and natural humoral immunity.

The loss of most naïve B cells after BLyS inhibition is consistent with the lack of FO and MZ B cells seen in BLyS- and BR3-deficient mice (13, 18, 19). Similarly, the attenuation of primary TD and TI responses mirrors prior findings in BLySdeficient mice (20), reflecting the elimination of preimmune subsets. Because the anti-BLyS used herein blocks BLyS binding to its receptors [\(Fig. S1\)](http://www.pnas.org/cgi/data/0807841105/DCSupplemental/Supplemental_PDF#nameddest=SF1), the most likely mechanism involves competition for soluble BLyS that blocks the BR3 signaling required for TR, FO, and MZ B cell survival. Indeed, antigenexperienced subsets also express BLyS binding receptors but were selectively spared, making direct cytotoxic effects unlikely.

BLyS inhibition reduced splenic B1a and B1b pools 2-fold, but peritoneal B1 cells were unaffected, suggesting that they are independently regulated. Because splenic B1 cell numbers and turnover rates are normal in BR3 mutant mice (21), BLyS signaling via TACI or BCMA may influence B1 survival or compartmentalization (20). Alternatively, uncompromised splenic architecture may contribute to splenic B1 maintenance (22). Natural antibody production in anti-BLyS-treated mice is consistent with the idea of functionally distinct B1 subsets in spleen and PerC (23) and suggests that BLyS dependence may distinguish these.

The relative resistance of memory and LLPC to BLyS inhibition suggests that they use alternative survival mechanisms. One possibility is a shift to APRIL dependence. Indeed, activated B cell and plasma cell populations up-regulate TACI and BCMA (but not BR3), both of which can bind APRIL more avidly than BLyS and can promote survival (10, 24, 25). Similarly, memory B cells have increased TACI and decreased BR3 expression but do not appear to express BCMA (26). Memory B cells share the expression of several receptors involved in regulating the renewal of stem cell populations (26), so ligands for these receptors may replace BLyS as key survival factors. TLR stimulation is required for activation of human B cells (27) and leads to TACI up-regulation on murine FO and MZ cells (24), suggesting that TLR ligands play a role.

Concurrent studies by Benson *et al.* (28) have shown that memory B cells require neither BLyS nor APRIL. Our observations confirm these findings and extend them in two ways. First, we have shown that isotype-switched memory cells are more resistant to BLyS depletion than nonswitched IgM-bearing memory cells. This partial dependence of unswitched memory cells is consistent with observations that unswitched human CD27<sup>+</sup> memory cells are intermediate in phenotype between naïve and switched memory cells (29, 30) and supports the notion of heterogeneity among memory B cells (2, 31). We also show heterogeneity within the B1 B cell compartment, because BLyS inhibition spared PerC B1 cells but left a residual B1 subset in the spleen that functionally resembles PerC B1 cells.



**Fig. 4.** Memory B cells generated in adoptive transfer systems are more resistant to BLyS depletion than naïve B cells. Memory B cells were generated from mVh186.2 Tg (Memory System 2; *A* and *B*) or Vh186.2 KI (Memory System 3; *C*, *E*, and *F*) donor B cells after adoptive transfer into AM14/V8R double Tg mice and subsequent immunization i.p. with NP-CGG in alum. Naïve controls were unimmunized Vh186.2 KI donor strain (*D* and *E*) and recipient IgM<sup>a</sup> non-NIP-binding (A–C and E) B cells. Recipient mice >12 weeks after immunization and unimmunized donor strain controls were treated with two 100-μq i.p. doses of anti-BLyS (10F4) or hamster IgG isotype control (IC). Fifteen days later, spleens were harvested and analyzed by FACS. (*A*) Memory System 2. Single, live lymphocytes were identified by FSC and SSC profiles and EMA exclusion. *Upper* and *Lower* represent isotype control (IC) versus anti-BLyS-treated animals, respectively. NIP<sup>+</sup> IgM<sup>a+</sup> B cells were NP-specific donor-derived memory cells. NIP<sup>-</sup> IgM<sup>a+</sup> B cells were recipient AM14/V<sub>K</sub>8R specificity naïve B cells. IgM<sup>a-</sup> IgM<sup>b+</sup> B cells were non-Tg-bearing recipient B cells. This population is expanded in aged recipient mice and may represent endogenous memory B cells. Representative FACS plots are shown. (*B*) Total numbers of B cells in each subgroup were calculated from live splenocyte counts and FACS-based frequencies as shown in *A*. Each point represents an individual animal. Average fold depletion in anti-BLyS-treated (circles) versus isotype control-treated (triangles) animals is indicated. One-way Student's t tests were performed. n.s., not significant. \*\*, *P*  $\leq$  0.01; \*\*\*, *P*  $\lt$  0.001. (C) Memory System 3. Live lymphocytes were identified as in A. NIP<sup>+</sup> k<sup>low</sup> B cells were NP-specific donor-derived memory cells. NIP- B cells were recipient-derived naïve cells. Memory B cells were fractionated into IgG<sub>1</sub>-switched and nonswitched B cells. (*D*) Naïve controls for Memory System 3. Spleens were harvested from unimmunized Vh186.2 KI donor-strain mice. Live lymphocytes were gated as described in *A*. (*E* and *F*) Total numbers of B cells from each System 3 subgroup were calculated from live splenocyte counts and FACS-based frequencies.

Our results suggest that targeting the BLyS/BR3 axis (32, 33) will spare memory, LLPC, and natural antibody-producing B cell populations. Because many human B cell lymphomas are likely derived from postgerminal center cells (34) and autocrine BLyS attenuates apoptosis of non-Hodgkin's lymphoma B cells (35), these data provide an important caveat for BLyS/BR3 targeting in B cell malignancies. Similarly, if autoimmune pathology stems from memory or LLPCs (36), BLyS-targeted treatments may prove ineffective. However, in some autoimmune situations self-reactive antibodies may be replenished by activation of extrafollicular foci and short-lived antibody-forming cells (37). In these scenarios, such directed therapies might deplete pathogenic naïve B cells but spare subsets that maintain induced and natural immunity.

#### **Materials and Methods**

**Mice.** Female C57BL/6 mice (age 6–12 weeks) were obtained from The Jackson Laboratory. The mVh186.2 Tg, Vh186.2 KI (B1.8), and AM14 Vk8R CB17 strains have been described (38-40). mVh186.2 Tg and Vh186.2 KI mice were maintained on the Jh knockout (KO) strain (41) and crossed to J $\kappa$  KO mice (42) to increase the frequency of NP-reactive B cells. All procedures were approved by the University of Pennsylvania or the Yale University Institutional Animal Care and Use Committees.

**BLyS and Anti-BLyS Assays.** Serum BLyS was measured by ELISA using mBAFFR-Fc (Alexis) as a capture reagent and an anti-murine BLyS monoclonal antibody (16D7; Human Genome Sciences) as a detector. Samples were diluted to a final concentration of 10% matrix on the assay plate. The limit of detection was 0.8 ng/ml. Serum anti-BLyS (10F4) levels were measured by ELISA; the limit of quantification was 0.075  $\mu$ g/ml, and the limit of detection was 0.039  $\mu$ g/ml.

**In Vivo Inhibition of BLyS.** Monoclonal IgG<sub>1</sub> hamster anti-mouse BLyS (clone 10F4; Human Genome Sciences) was diluted in PBS and injected i.p. Purified Armenian hamster IgG<sub>1</sub> (clone G235-2356; BD Pharmingen) was the isotype control.

**Anti-NP Antibody ELISAs.** Plates were coated with 10  $\mu$ g/ml NP<sub>3</sub>BSA in 100 mM bicarbonate buffer and blocked with 2% BSA in PBS, and serum dilutions were incubated for 1 h. NP-specific  $\log_1$  standard was a gift of G. Kelsoe (Duke University, Durham, NC). HRP-conjugated goat anti-mouse IgG<sub>1</sub> (Southern Biotechnology Associates) was used for detection with a TMB substrate kit (BD Biosciences). Washes were performed by using PBS plus 0.1% Tween 20.

**Anti-PC Antibody ELISPOTS.** Immobilon-P plates (Millipore) were coated with 10  $\mu$ g/ml PC16-BSA (Biosearch Technologies) and blocked with 2% BSA in PBS. After erythrocyte lysis, splenic or PerC cell suspensions were enumerated, and cell suspensions were diluted 2-fold, then incubated for 4 h. Plates were developed with biotin-conjugated anti-mouse IgM (Southern Biotechnology Associates) followed by ExtrAvidin-Alkaline Phosphatase using NBT/BCIP substrate (Sigma); color development was terminated with 1 M NaH<sub>2</sub>PO<sub>4</sub>. Spots were enumerated on CTL-ImmunoSpot reader (Cellular Technologies). The number of background spots obtained with BSA-coated wells was subtracted from the number of spots counted on PC16-coated wells for each mouse.

**Generation of TD Primary and Memory Responses in Wild-Type Mice.** For primary responses, female C57BL/6J mice were immunized i.p. with 50-100  $\mu$ g of NP16-CGG in alum (2, 16). To generate memory B cells and LLPC (Memory System 1), mice were rested  $\geq$ 8 weeks after primary immunization, then rechallenged with 50  $\mu$ g of i.p. NP<sub>16</sub>-CGG. Memory B cells, LLPC, and serum NP<sup>+</sup> antibodies were analyzed 7 days after the boost dates.

**Generation of Naïve and TD Memory B Cells in Adoptive Transfer Systems.** ''Memory System 2'' has been described in detail previously (2, 26). Briefly, splenocytes containing  $1 \times 10^6$  NP-specific naïve B cells from mVh186.2 Ig heavy chain transgenic (Tg) Jh KO mice (39) were adoptively transferred into recipient mice carrying rearranged Ig Tgs of irrelevant specificities (AM14 IgH and V<sub>K</sub>8R, which produce anti-IgG<sub>2a</sub><sup>a</sup> antibody), and which therefore mount poor endogenous responses to NP. Six hours after transfer, recipient mice were immunized i.p. with 50  $\mu$ g of NP<sub>25</sub>-CGG in alum. More than 12 weeks after transfer, when recently activated germinal center B cells (3, 39, 43) were

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undetectable, there was a large and stable donor-derived NP-specific memory B pool, identified as NP-binding  $IgM^{a+}$  B cells. This population was virtually (99%) free of ''contaminating'' donor-derived naïve B cells, because few donor B cells survived in the absence of immunization. The vast majority of recipient-derived B cells were AM14/V<sub>K</sub>8R Ig-bearing naïve cells. There was also a small population of recipient-derived non-Tg-bearing B cells, but, because recipient mice were on the CB.17 background, these were readily distinguished from BALB/c donor B cells by Ig allotype (Igb versus Ig<sup>a</sup>, respectively). A further advantage of this system is that mVh186.2 Tg-derived B cells do not secrete NP-specific antibody (16), which can confound memory cell identification based on NP specificity alone (44, 45). Additionally, because the Vh186.2 Tg prevents class switch, this system models IgM B cell memory.

Memory System 3 [\(Fig. S5\)](http://www.pnas.org/cgi/data/0807841105/DCSupplemental/Supplemental_PDF#nameddest=SF5) was identical to System 2, except donor B cells were isolated from Vh186.2 KI (B1.8) mice (38). An advantage of this model is that B cells can undergo isotype class switching. More than 12 weeks after transfer and immunization, NP-specific donor-derived B cells averaged 1.1 million per spleen in comparison to 25,000 per spleen in adjuvant-only-treated animals. Thus, in immunized animals, an average of 2.3% of NP-specific B cells were naïve whereas 97.7% were memory. In System 3, donor-derived memory B cells were identified as NP-binding  $\kappa^{\text{low}}$  IgG<sub>1</sub><sup>+</sup> or IgG<sub>1</sub><sup>-</sup> (virtually all of which are IgM-bearing) cells.

**Antibodies and Flow Cytometry.** Splenocytes and bone marrow were harvested and stained as described (ref. 50 and [Fig. S2\)](http://www.pnas.org/cgi/data/0807841105/DCSupplemental/Supplemental_PDF#nameddest=SF2). Splenic and PerC B1 B cells were analyzed by using the following antibodies: FITC-anti-CD5, PE-anti-Mac-1 (CD11b), biotin-anti-CD23, and APC-anti-IgM (all from BD Biosciences). NPbinding B cell analysis in Memory System 1 used the following antibodies: DAPI vital dye (Invitrogen), PE-Cy5-anti-CD4 and anti-CD8, PE-anti-Syndecan-1, FITC-anti-GL7 (BD), PE-Cy5-anti-Gr-1 (eBioscience), biotin-anti-IgD, FITC-antilambda, FITC-anti-Igß (Southern Biotechnology Associates), Streptavidin-Pacific Blue (Invitrogen), PE-Cy7-anti-B220 (eBioscience) and APC-NP (M.P.C. laboratory). Analyses of Systems 2 and 3 used APC-Cy7-anti-B220 (RA3-6B2; BD), FITC-anti-IgG<sub>1</sub> (A85-1; BD), and AI488-anti-IgM<sup>a</sup> (RS3.1), biotinylated anti-IgM<sup>b</sup> (AF6-78), Pacific Blue anti- $\kappa$  (187.1), and APC-NIP, all produced in the M.J.S. laboratory. Data were collected on a BD LSR II flow cytometer and analyzed with FlowJo software (Tree Star).

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