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# The B-cell antigen receptor integrates adaptive and innate immune signals

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B cells respond to antigens by engagement of their B-cell antigen receptor (BCR) and of coreceptors through which signals from helper T cells or pathogen-associated molecular patterns are delivered. We show that the proliferative response of B cells to the latter stimuli is controlled by BCR-dependent activation of phosphoinositidyl 3-kinase (PI-3K) signaling. Glycogen synthase kinase 3 $\beta$  and Foxo1 are two PI-3K-regulated targets that play important roles, but to different extents, depending on the specific mitogen. These results suggest a model for integrating signals from the innate and the adaptive immune systems in the control of the B-cell immune response.

BCR | B-cell proliferation | PI-3K | GSK3β | Foxo1

**E** xpression of a functional B-cell antigen receptor (BCR) is of central importance to nearly every aspect of B-cell biology. During early development, B-lineage cells assemble clonally restricted Ig heavy and light chains that are able to recognize a diverse and unpredictable universe of antigens. The proper assembly and expression of a BCR is a prerequisite for entry into the naive compartment. Furthermore, naive resting B cells remain dependent on the expression of a correctly assembled BCR. If this is prevented by IgH chain ablation or Ig $\alpha$  truncation, the cells rapidly undergo apoptosis. This has been interpreted to reflect a BCR-dependent "tonic" survival signal, which has recently been shown to involve phosphoinositidyl 3-kinase (PI-3K) activity and subsequent inactivation of the Foxo1 transcription factor (1–3).

As a consequence of the large, clonally restricted repertoire generated during development, only a small number of newly generated B cells entering the peripheral mature compartment express a BCR that is able to bind a given antigen. During the early stages of an immune response, antigen-specific B cells proliferate extensively and increase the number of cells that can respond appropriately. The magnitude, duration, and quality of the B-cell response is guided by signals from coreceptors expressed by B cells such as Toll-like receptors (TLRs) and CD40, which provide information to the B cell about the context of the antigen in T-cellindependent and T-cell-dependent responses (4, 5). Given that the raison d'être of B cells is an antigen-specific response, including the production of antibodies, a central role for the BCR in regulating B-cell responses to innate or T-cell-derived signals seems likely. Indeed, early experiments addressing the in vitro response of mouse B cells to bacterial lipopolysaccharide (LPS) have suggested cooperation of LPS stimulation and BCR engagement in the proliferative B-cell response under certain conditions (6, 7). More recent work has uncovered a synergy between TLR and BCR in the control of class-switch recombination in activated B cells (8), and the role of the BCR in the delivery of DNA- or RNA-associated autoantigens to endosomal TLRs in autoimmunity is well established (9). How B cells integrate signals from the BCR and from coreceptors responding to mitogenic stimuli in the induction of the proliferative response is unclear, however.

#### **Results and Discussion**

**Generation of Resting BCR-Negative B Cells.** We generated resting B cells that lack expression of the BCR by transducing splenic B cells

from *bcl-2tg* mice that express a loxP-flanked pre-rearranged variable domain targeted into the *Igh* locus (*B1-8f*) (2) with cell-permeable Cre recombinase, TAT-Cre (10). The transduced cells remained alive throughout the culture because of the ectopic expression of Bcl-2 and lost BCR expression by day 3 in culture with an efficiency of 30–50%, allowing simultaneous analysis of B cells that either express or lack the BCR (BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells, respectively; Fig. 1*A*). Western blot analysis of sorted BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells confirmed the loss of IgH chain expression in the latter (Fig. 1*B*). The proportion of BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells remained constant for at least 2 wk, suggesting the two classes of cells survive equally well in vitro (Fig. S1). Although mature BCR<sup>neg</sup> cells are slightly smaller than their BCR<sup>pos</sup> counterparts and down-regulate some mature B-cell markers (e.g., MHC class II and CD22), they largely maintain the surface phenotype of resting, mature B cells (Fig. 1*C*).

**BCR Expression Is Necessary for the Mitogenic Response of B Cells.** B cells express receptors such as TLR-9, TLR-4, and CD40 that induce activation and proliferation when triggered by natural ligands or agonist antibodies (11–13). We tested the proliferative response of BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells to CpG DNA, LPS, or anti-CD40 and found that, in contrast to BCR<sup>pos</sup> B cells, BCR<sup>neg</sup> B cells did not proliferate in response to any of these stimuli (Fig. 24). This defect is not an artifact of protein transduction

#### Significance

Clonal expansion of antigen-specific B cells during an immune response is necessary for effective antibody production. B cells must integrate signals from their clonally restricted B-cell antigen receptor (BCR) and from nonclonal coreceptors that provide contextual cues to the nature of the antigen. How this is accomplished is unclear. We found that B cells require expression of the BCR for mitogenic responses triggered by coreceptors that recognize innate or T-cell-derived signals. The signaling pathway used by the BCR to license coreceptor-induced proliferation is similar to the previously described BCR-dependent survival signal and, thus, couples B-cell survival and mitogenic responses in an unexpected manner that may have implications for B-cell lymphomagenesis.

The authors declare no conflict of interest.

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Fig. 1. In vitro generation of BCR<sup>neg</sup> B cells. (A) Schematic showing Cre-mediated deletion of the loxP-flanked B1-8 allele and subsequent loss of BCR expression. (B) Splenic B cells from B1-8f/+, bcl-2tg mice were treated with recombinant TAT-Cre for 45 min, cultured for 9 d, and sorted by FACS to >98% purity based on IgM expression (data are representative of two biological replicates). Serial dilutions of whole-cell lysates starting at 300,000 cell equivalents were analyzed for Ig $\mu$  and  $\beta$ -actin expression by Western blotting. (C) Splenic B cells from B1-8f/+, bcl-2tg mice were transduced with TAT-Cre for 45 min, cultured for 5 d in medium, and stained for CD19, IgM, and the indicated surface markers (data are representative of three biological replicates). BCR<sup>pos</sup> cells (open histograms) and BCR<sup>neg</sup> cells (closed histograms) were gated as shown.

because *bcl-2tg* BCR<sup>neg</sup> B cells generated by a tamoxifeninducible Cre (14) or in vivo with a mature B-cell-specific Cretransgene (*CD21-Cre*) (1) showed the same proliferation defect (Fig. S2). BCR<sup>neg</sup> B cells are not completely refractory to the mitogenic stimuli, as they activate normal NF- $\kappa$ B DNA binding activity in response to CpG (Fig. S3) and up-regulate CD69 and MHC class II when stimulated with CpG DNA, LPS, or anti-CD40 (Fig. 2*B*). Taken together, these data show that mature

B cells that have lost expression of the BCR are responsive to innate stimuli but are unable to proliferate.

### BCR Expression Licenses B-Cell Proliferation Through PI-3K Activation.

Previous studies demonstrated that constitutive, low-level signaling, a so-called tonic signal, which is dependent on an intact BCR, is necessary for mature B-cell survival (1, 2) and is mediated by the PI-3K/Akt/Foxo1 signaling pathway (3). The requirement of BCR



**Fig. 2.** Expression of the BCR is necessary for B-cell proliferation. BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells were generated by TAT-Cre transduction of B cells from *B1-8fl+*, *bcl-2tg* mice. After 3 d in culture, cells were either (A) labeled with CFSE and stimulated for 3–4 d (data are representative of five biological replicates) or (*B*) stimulated for 24 h with the indicated mitogen and analyzed by FACS (data are representative of two biological replicates). Live cells were gated as ToPro3<sup>-</sup>, CD19<sup>+</sup>.

expression for mitogenic responses of B cells to innate stimuli raised the possibility that the same tonic signal may also regulate B-cell proliferation. BCR<sup>neg</sup> B cells generated in vivo with a mature B-cell-specific Cre-transgene (i.e., CD21-Cre) (1) were used to study Akt status in vivo. Indeed, although intracellular staining for phospho-Akt (Ser473) of CpG-treated BCRpos B cells showed prolonged activation of the PI-3K pathway, induced Akt phosphorylation was not seen in BCR<sup>neg</sup> B cells under these conditions (Fig. 3A). This was confirmed by Western blot analysis of sorted BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells cultured with or without CpG-containing medium for 2 h (Fig. 3B). We conclude that expression of the BCR as mediated by the IgH chain is required for mitogeninduced activation of the PI-3K pathway in response to CpG stimulation. To further test the role of BCR-dependent PI-3K signaling in regulating mitogen-induced proliferation of B cells, we isolated BCR<sup>neg</sup> B cells from B1-8f, CD21-Cre mice that also carried a transgene for an active PI-3K molecule that was induced by Cre activity (B1-8f, CD21-Cre, P110\*) (3). Expression of Cre in mature B cells simultaneously ablates expression of the BCR and induces expression of constitutively active PI-3K, which substitutes for the BCR-dependent survival signal (3). Similar to control BCR<sup>pos</sup> B cells, BCR<sup>neg</sup> B cells expressing constitutively active PI-3K proliferated in response to all the stimuli (Fig. 3C), showing that reconstitution of the PI-3K pathway in BCRneg B cells licenses mitogenic responses to innate stimuli.

Inactivation of Glycogen Synthase Kinase 3ß Is Necessary for the Proliferative Mitogenic Response of BCR<sup>neg</sup> B Cells to CpG. Next, we investigated the BCR-dependent signaling pathways downstream of PI-3K as required for mitogenic responses of B cells. PI-3K/Akt activates several downstream signaling pathways including pathways using glycogen synthase kinase  $3\beta$  (GSK3 $\beta$ ) and Foxo1. GSK3B is a constitutively active serine/threonine kinase that inhibits proliferation by phosphorylating and targeting for degradation proteins that are necessary for mitosis, including c-Myc (15) and D-type cyclins (16). Mitogenic signals activate the PI-3K/Akt pathway, leading to phosphorylation and inactivation of GSK3<sup>β</sup> (17), and thereby allowing accumulation of cell cycle-regulated proteins. Consistent with defective Akt phosphorylation in BCRneg B cells, the levels of phospho-GSK3β remained low in BCR<sup>neg</sup> B cells, in contrast to BCR<sup>pos</sup> B cells, after treatment with CpG (Fig. 4A), suggesting that GSK3β remained active in BCR<sup>neg</sup> B cells. To directly assess whether GSK3ß activity prevents the proliferation of mitogen-treated BCRneg B cells, we added the GSK3β inhibitor (2'Z,3'E)-6-bromo-indirubin-3'-oxime (BIO) (18) to CpGstimulated cultures and found that inhibition of GSK3<sup>β</sup> partially restores the ability of bcl-2tg BCR<sup>neg</sup> B cells to proliferate in response to CpG (Fig. 4B). We also found that CpG-induced c-Myc protein levels were much lower in BCR<sup>neg</sup> B cells than those seen in BCR<sup>pos</sup> B cells treated in a similar manner (Fig. 4C). Similar results were seen when LPS or anti-CD40/IL-4 were used as mitogens (Fig. 4C). In contrast, mitogen-induced c-Myc mRNA levels were nearly identical between BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells stimulated with CpG (Fig. S4), suggesting that active GSK3 $\beta$  was destabilizing c-Myc protein in BCR<sup>neg</sup> B cells (15). Unfortunately, pharmacological inhibition of GSK3ß activity using BIO, which partially restores proliferation in response to CpG (Fig. 4B), did not substantially restore mitogen-induced c-Myc protein expression in BCR<sup>neg</sup> B cells (Fig. 4D). Therefore, we conclude that the decreased levels of c-Myc induction in BCR<sup>neg</sup> B cells, although of potential interest, do not explain the defective proliferation of B cells that have lost expression of their BCR. Furthermore, inhibition of GSK3 $\beta$  did not rescue proliferation of BCR^{neg} B cells in response to LPS or anti-CD40/IL-4 treatment (Fig. 4E), suggesting that BCR-dependent PI-3K activation regulates additional signaling pathways required for mitogenic responses, depending on the stimulus.



Fig. 3. Defective activation of PI-3K in BCR<sup>neg</sup> B cells. (A) Splenic B cells from B1-8f/J<sub>H</sub>T, bcl-2tg or B1-8f/J<sub>H</sub>T, CD21-Cre, bcl-2tg mice were stimulated with CpG for the indicated times and stained for CD19 and IgM and intracellular phospho-Akt (Ser473). BCR<sup>pos</sup> (green) cells were gated from B1-8f/J<sub>H</sub>T, bcl-2tg cells, whereas Cre<sup>+</sup> BCR<sup>pos</sup> (red) or Cre<sup>+</sup> BCR<sup>neg</sup> (blue) B cells were gated from B1-8f/J<sub>H</sub>T, CD21-Cre, bcl-2tg cells. Normalized mean fluorescence intensity values for phospho-Akt (Ser473) are shown. Data represent three biological replicates. (B) BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells were sorted from B1-8f/  $J_{HT}$ , bcl-2tg splenic B cells 5 d after transduction with TAT-Cre, placed in culture overnight, and stimulated with CpG for 2 h. Equal amounts of cytoplasmic lysates were blotted for phospho-Akt (Ser473) and Akt. Data are representative of two biological replicates. (C) Splenic  ${\sf BCR}^{\sf pos}$  and  ${\sf GFP}^+$ BCR<sup>neg</sup> B cells were sorted from Rag2<sup> $\Delta/\Delta$ </sup>, c $\gamma^{-/-}$  mice that had been reconstituted with fetal liver from B1-8f/J<sub>H</sub>T, CD21-Cre, P110\* mice, labeled with CFSE and cultured for 3 d in medium (black) or medium plus the indicated stimulus (red). Data are representative of three biological replicates.

Foxo1 and GSK3 $\beta$  Regulate Innate Stimuli-Induced Mitogenic Responses Downstream of the BCR. Inactivation of the transcription factor Foxo1 by BCR-dependent PI-3K/Akt activation is necessary for mature B-cell survival (3). The complete loss of Foxo1 rescues the survival of BCRneg B cells as efficiently as expression of active PI-3K, whereas loss of one Foxo1 allele is sufficient for a partial rescue of BCRneg B cells (3). Because Foxo1 remains active in BCR<sup>neg</sup> B cells (3) and can inhibit proliferation by up-regulating antiproliferative genes (19, 20), we tested the effect of Foxo1 ablation on mitogen-treated BCR<sup>neg</sup> B cells. BCR<sup>neg</sup> B cells that were either heterozygous or homozygous for a Foxo1-null allele were sorted, and their proliferative responses in the absence or presence of BIO to inhibit GSK3ß were compared with mitogen-stimulated BCR<sup>pos</sup> cells (Fig. 5). Although the complete or partial loss of Foxo1 had only minimal effects on the proliferative response of BCR<sup>neg</sup> B cells to CpG, the combination of Foxo1 deletion and inhibition of GSK36 rescued the proliferation of BCRneg B cells in



**Fig. 4.** GSK3β activity prevents the proliferation of CpG-treated BCR<sup>neg</sup> B cells. (A) Splenic B cells from *B1-8fIJ<sub>H</sub>T*, *bcl-2tg* or *B1-8fIJ<sub>H</sub>T*, *CD21-Cre*, *bcl-2tg* mice were stimulated with CpG for the indicated times and stained for CD19 and IgM and intracellular phospho-GSK3α/β (S21/9). BCR<sup>pos</sup> (green) cells were gated from *B1-8fIJ<sub>H</sub>T*, *bcl-2tg* cells, whereas Cre<sup>+</sup> BCR<sup>pos</sup> (red) or Cre<sup>+</sup> BCR<sup>neg</sup> (blue) B cells were gated from *B1-8fIJ<sub>H</sub>T*, *CD21-Cre*, *bcl-2tg* cells. Normalized mean fluorescence interval values for phospho-GSK3α/β (S21/9) are shown. Data represent three biological replicates. (*B*) B cells from *B1-8fIJ<sub>H</sub>T*, *CD21-Cre*, *bcl-2tg* mice or TAT-Cre-transduced B cells from *B1-8fIJ<sub>H</sub>T*, *bcl-2tg* mice were CFSE-labeled and stimulated for 3 d with CpG alone (green) or with Meth-BIO (inactive control inhibitor, red) or BIO (GSK3 inhibitor, blue). Cells were gated on ToPro3<sup>-</sup>, CD19<sup>+</sup>, and IgM<sup>+</sup> or IgM<sup>-</sup>. Data are representative of three biological replicates. (*C*) Splenic B cells from *B1-8fIJ<sub>H</sub>T*, *bcl-2tg* mice were CFSE-labeled and stimulated for 3 d with CpG alone (green) or with Meth-BIO (inactive control inhibitor, red) or BIO (GSK3 inhibitor, blue). Cells were gated on ToPro3<sup>-</sup>, CD19<sup>+</sup>, and IgM<sup>+</sup> or IgM<sup>-</sup>. Data are representative of three biological replicates. (*C*) Splenic B cells from *B1-8fIJ<sub>H</sub>T*, *bcl-2tg* mice were CFSE-labeled and stimulated for 3 d with CpG alone (green) or with Meth-BIO (inactive control inhibitor, red) or BIO (GSK3 inhibitor, blue). Cells were gated on ToPro3<sup>-</sup>, CD19<sup>+</sup>, and IgM<sup>+</sup> or IgM<sup>-</sup>. Data are representative of three biological replicates. (*C*) Splenic B cells from *B1-8fIJ<sub>H</sub>T*, *bcl-2tg* mice were CFSE-labeled and stimulated for 3 d with CpG and BCR<sup>neg</sup> B cells from *B1-8fI*, *bcl-2tg* mice were tore three biological replicates. (*D*) Splenic B cells from *B1-8fI*, *bcl-2tg* mice were tore tore-transduced and cultured for 3 d. Purified BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells were gated in culture overnight and stimulated with

response to CpG to a level comparable to that observed by CpGstimulated BCR<sup>pos</sup> B cells. In contrast, although heterozygous deletion of Foxo1 was sufficient for some anti-CD40/IL-4-treated BCR<sup>neg</sup> B cells to undergo at least one cell division, homozygous deletion of Foxo1 allowed near-complete rescue of BCR<sup>neg</sup> proliferation to anti-CD40/IL-4. In both cases, inhibition of GSK3 $\beta$  had little additional effect, suggesting that different mitogens require PI-3K to activate different signaling pathways for proliferation in the absence of BCR expression (Fig. 5). To further illustrate this point, BCR<sup>neg</sup> B cells on a Foxo1-null background in

the presence of a GSK3 $\beta$  inhibitor remained unable to proliferate in response to LPS treatment (Fig. 5), suggesting that other or additional pathways downstream of PI-3K/Akt that regulate cell proliferation are involved.

Our experimental model is based on previous evidence that the BCR expressed by resting, mature B cells provides tonic survival signals to the cells (1, 2). The mechanism by which such signals are initiated is controversial, but may involve low-affinity interactions of the BCR with environmental antigens, akin to the positive selection of T cells by self-MHC. It is also possible, however, that loss of the BCR induces a danger signal in the cells from which they can be relieved by PI-3K pathway activation. By extending the analysis to the control of B-cell proliferation, the present data provide the elements of a molecular mechanism by which the BCR controls the proliferative expansion of B cells on the engagement of invariant, "innate" receptors. Such a crosstalk between these two classes of receptors reflects the physiological requirement of integrating, in any meaningful antibody response, input from the BCR regarding specific antigen recognition and information about the nature and context of the immunizing antigen, with the latter being provided to the cells by innate receptors such as TLRs recognizing pathogenassociated molecular patterns presented by microbes and the CD40 coreceptor triggered by T helper cells. The requirement of this regulatory principle is particularly obvious in the case of the somatic antibody mutants generated in the germinal center reaction, whose expansion through T-cell-derived and innate signals should strictly depend on newly acquired BCR specificities or BCR expression in the first place.

Although receptor-proximal signaling events in B cells differ among TLRs and CD40, these receptors share common downstream signaling pathways such as the PI-3K, MAPK, and NF- $\kappa$ B



**Fig. 5.** Foxo1 deletion plus GSK3 $\beta$  inhibition rescues proliferation in CpGtreated BCR<sup>neg</sup> B cells. Splenic B cells were sorted, labeled with CFSE, and cultured either in medium (green) or with the indicated stimuli, with (blue) or without (red) the GSK3 $\beta$  inhibitor BIO. BCR<sup>pos</sup>, Foxo1f/+ and BCR<sup>neg</sup> Foxo1 $\Delta$ /+ cells were sorted from B1-8f/J<sub>H</sub>T, CD21-Cre, Foxo1<sup>f/+</sup> mice, whereas BCR<sup>neg</sup>, Foxo1 $\Delta$ /Δ cells were sorted from B1-8f/J<sub>H</sub>T, CD21-Cre, Foxo1<sup>f/+</sup> mice. Data are representative of three biological replicates.

pathways (21, 22). This is supported by the similarity of gene expression profiles of primary B cells stimulated with distinct mitogenic agonists (23). The BCR appeared to specifically control PI-3K signaling in the cells, in that a key PI-3K target, Akt, was not phosphorylated in BCR<sup>neg</sup> cells on CpG stimulation (Fig. 3B), and expression of active PI-3K rescued mitogenic responses in the absence of the BCR (Fig. 3C). The BCR is apparently involved in this process in mitogen-activated B cells, conceivably by controlling the levels, activity, or localization of inhibitors of the PI-3K pathway. This scenario is supported by results showing that mature B cells undergoing Cre-mediated BCR deletion in vivo can be rescued from apoptosis by the expression of constitutively active PI-3K, as well as by the ablation of PTEN (phosphatase and tensin homolog), a lipid phosphatase that directly opposes PI-3K activity (24). Downstream of PI-3K activation, the signaling pathways diverge depending on the specific mitogen. Inactivation of Foxo1 is important for proliferation in response to anti-CD40/IL-4 stimulation, whereas inactivation of GSK3<sup>β</sup> plays a greater role in the mitogenic response to CpG. Interestingly, inactivation of both GSK3β and Foxo1 are not sufficient for LPS-induced proliferation, suggesting that additional BCR-dependent signaling pathways are activated by PI-3K and are necessary for TLR4-mediated proliferation of B cells. The present results indicate that, in addition to its role in the control of B-cell survival, BCR modulation of PI-3K signaling licenses the induction of clonal expansion of B cells by mitogens acting on innate receptors expressed by the cells, a process of crucial importance in the antibody response. In this scenario, BCR engagement by self-antigen or cognate antigen as well as cross-reactivity with mitogens (8) may well play a modulating role. Quite fittingly, the work of Pone et al. has identified an interplay between BCR and TLR signaling in the control of class-switch recombination in the activated B cells. This was mediated by NF-kB signals downstream of both receptors, which, significantly, depended on PI-3K activity in the case of the BCR (8).

The integration of signals delivered by distinct cellular receptors is an obvious requirement for any cellular response in multicellular organisms. An example is the outside-in signaling of integrins bound to extracellular matrix proteins, which mediate the proliferation of adherent epithelial cells in response to growth factors (25). Notably, the PI-3K pathway plays a role in this context as well, and also provides integrin-activated prosurvival signals to the adherent cells (26, 27). The interplay between the antigen receptor and innate immune receptors, which we have described here for B lymphocytes, may thus be just a variation of an ancient principle of receptor crosstalk in higher cells. It will be interesting to see whether this crosstalk also plays a role in the homeostasis of memory cells and to which extent and by what means the BCR licenses the proliferative expansion of B-cell lymphomas (28).

#### **Materials and Methods**

**Mouse Strains.** B1-8<sup>*f*</sup> (2),  $J_HT$  (29), *bcl-2tg* (30), *CD21-Cre* (1), *rosa26-stop<sup>f</sup>-P110\** (3), and CreED-30 (14) mice were housed and maintained under specific pathogen-free conditions. All mouse protocols were approved by Harvard University and the Immune Disease Institute Institutional Animal Care and Use Committees.

**Cell Culture and TAT-Cre Transduction.** TAT-Cre (also known as His-TAT-NLS-Cre or HTNC, expression plasmid available from Addgene) was produced as described (10). Purified B cells were washed three times in serum-free, animal-defined component-free medium (HyClone), prewarmed to 37 °C, and transduced with 50 µg/mL TAT-Cre in the presence of 50 µg/mL polymyxin B sulfate (Sigma-Aldrich) at a density of  $5 \times 10^6$ /mL for 45 min at 37 °C. Cells were then washed in complete medium [DMEM supplemented with 10% (vol/vol) FCS (Biowhittacker), 10 mM Hepes, 2 mM L-glutamine, 1 mM sodium pyruvate, 1× penicillin/streptomycin, 1× nonessential amino acids (Cellgro), and 50 µM β-mercaptoethanol] and cultured at 37 °C for 3–5 d at 2 × 10<sup>6</sup>/mL in complete medium. Mock-transduced cells were treated identically except that TAT-Cre was omitted. B cells were labeled with

carboxyfluorescein succinimidyl ester (CFSE), as indicated, by washing two times in PBS, prewarming for 10 min at 37 °C, and incubating with titrated amounts of CFSE for 10 min at 37 °C. Cells were washed in complete medium and stimulated with 1 µg/mL anti-CD40 (1C10; eBioscience), 20 µg/mL ultrapure LPS (InvivoGen), or 645 ng/mL CpG (ODN1826; InvivoGen) at 2 × 10<sup>5</sup>/mL in complete medium for 3–4 d. The GSK3 inhibitor BIO or the inactive control Meth-BIO (Calbiochem) were added at a concentration of 0.2 µM, as indicated.

**FACS.** Single-cell suspensions of ex vivo splenocytes or cultured B cells were stained with antibodies to IgM (goat Fab; Jackson ImmunoResearch), IgD (1.3–5), Ig<sub>K</sub> (R33-18.10), Ig<sub>β</sub> (HM79), CD19 (1D3), B220 (RA3-6B2), CD22 (CY34), CD21 (7G6), CD23 (B3B4), CD24 (M1/69), CD95 (Jo2), CD93 (AA4.1), I-A<sup>b</sup> (AF6-120), or CD69 (H1.2F3). Dead cells were excluded by addition of ToPro3 (Invitrogen). For intracellular staining, cells were fixed and permeabilized with Perm/Fix and Perm/Wash solutions (BD Bioscience) according to manufacturer's instructions, stained with anti-phospho-Akt (S473; Cell Signaling) or anti-phospho-GSK3 $\alpha/\beta$  (37F11; Cell Signaling), and detected with anti-rabbit-FITC. Data were collected on a FACSCalibur (BD Bioscience) and analyzed with CellQuest (BD Bioscience) or FlowJo (TreeStar) software.

**Cell Separation.** Red blood cells were lysed by hypotonic shock from splenocyte suspensions. B cells were purified by CD43 depletion using magnetic-activated cell separation (MACS; Miltenyi). For some experiments, BCR<sup>neg</sup> B cells were isolated by MACS depletion of IgM<sup>+</sup> B cells from TAT-Cre-transduced cultures using biotinylated anti-IgM (331-12) and streptavidin-beads (Miltenyi). Purity was >98%, as determined by IgD staining. Alternatively, B cells were stained with Fab anti-IgM and anti-B220 before FACS sorting on an Aria2 or FACS-Vantage (BD Bioscience).

**Immunoblot Analysis.** Whole-cell lysates were prepared in lysis buffer [40 mM Tris at pH 7.6, 150 mM NaCl, 2 mM EDTA, 1% Nonidet P-40, 0.25% sodium deoxycholate,  $1 \times$  EDTA-free protease inhibitor mixture (Roche), 1 mM PMSF, 5 mM NaF, 0.5  $\mu$ M, 1 mM Na<sub>3</sub>VO<sub>4</sub>]. The protein concentration was determined by detergent-compatible protein assay kit (Bio-Rad). Equal amounts of protein were separated by SDS/PAGE and transferred to PVDF membranes (Millipore).

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The membranes were immunoblotted, as indicated, with phospho-Akt (S473), Akt (Cell Signaling), c-Myc (Epitomics),  $\beta$ -actin (Sigma-Aldrich), or biotinylated goat Fab anti-mouse IgM (Jackson ImmunoResearch).

**NF-κB EMSA.** Splenic B cells from B1-8f/J<sub>H</sub>T, bcl-2tg mice were TAT-Cretransduced and placed in culture for 3 d. Purified BCR<sup>pos</sup> and BCR<sup>neg</sup> B cells were then cultured overnight in 1% FCS-containing medium before the addition of CpG for the indicated amounts of time. Cells were lysed in buffer A (5 mM Hepes at pH 7.9, 1.5 mM MgCl<sub>2</sub>, 10 mM KCl, 0.5 mM DTT, and protease inhibitors) to prepare cytosolic extracts and nuclei. Nuclear extracts were subsequently prepared using buffer C [20 mM Hepes at pH 7.9, 25% (vol/vol) glycerol, 0.5 M NaCl, 1.5 mM MgCl<sub>2</sub>, 0.5 mM EDTA, 0.5 mM DTT, and protease inhibitors]. Nuclear extracts were incubated with competitor DNA (pdldC; Sigma) and a <sup>32</sup>P labeled-kB consensus probe (Promega). Protein–DNA complexes were then resolved by electrophoresis on a nondenaturing polyacrylamide gel.

**Quantitative RT-PCR.** RNA was purified using mirVana RNA isolation kit (Ambion) and subjected to cDNA synthesis, using random hexamer primers (Invitrogen). Quantitative RT-PCR analysis was performed using *c-myc-* and *hprt*-specific primers on a StepOne Plus (Applied Biosystems) machine. Primers used were: c-MycF1, 5'-gagccctagtgctgcatgag-3'; c-MycR1, 5'-ttgcctcttctccacagacacc-3'; HPRTF1, 5'-ctataagttctttgctgacctgctggattaca-3'; HPRTR1, 5'-ccagttaaagttg-agagatcatctcccacaataact-3'.

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