

Distinct cell-specific control of autoimmunity and infection by FcγRIIb

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FcγRIIb is an inhibitory Fc receptor expressed on B cells and myeloid cells. It is important in controlling responses to infection, and reduced expression or function predisposes to autoimmunity. To determine if increased expression of FcγRIIb can modulate these processes, we created transgenic mice overexpressing FcγRIIb on B cells or macrophages. Overexpression of FcγRIIb on B cells reduced the immunoglobulin G component of T-dependent immune responses, led to early resolution of collagen-induced arthritis (CIA), and reduced spontaneous systemic lupus erythematosus (SLE). In contrast, overexpression on macrophages had no effect on immune responses, CIA, or SLE but increased mortality after *Streptococcus pneumoniae* infection. These results help define the role of FcγRIIb in immune responses, demonstrate the contrasting roles played by FcγRIIb on B cells and macrophages in the control of infection and autoimmunity, and emphasize the therapeutic potential for modulation of FcγRIIb expression on B cells in inflammatory and autoimmune disease.

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Abbreviations used: ANA, anti-nuclear antibodies; B-NTG and -TG, B cell nontransgenic and transgenic, respectively; CGG, chicken γ globulin; CIA, collagen-induced arthritis; CII, type II collagen; ERK, extracellular signal-regulated kinase; FDC, follicular dendritic cell; GC, germinal center; M-NTG and -TG, macrophage non-transgenic and transgenic, respectively; NP, 4-hydroxyl-3-nitrophenyl acetyl; PC, phosphorylcholine; SLE, systemic lupus erythematosus.

Fc receptors for IgG are key players in regulating innate and adaptive immunity (1, 2). FcγRIIb, as the only inhibitory Fc receptor for IgG, plays a unique role in regulating immune responses. It is the only Fc receptor expressed on B cells, where it controls the magnitude and persistence of the response to antigen through effects on both mature and memory B cells and plasma cells (3, 4). On myeloid cells, FcγRIIb counteracts activation through activatory Fc receptors (such as FcγRI, III, and IV in the mouse, and FcγRI, IIa, IIIa, and IIIb in the human). The response of the macrophage or dendritic cell to complexed IgG is determined by the specificity of the different low affinity Fc receptors for the isotypes of the IgG involved, as well as the relative ratio of activatory to inhibitory receptor expression on the cell surface (1, 5).

Mice deficient in FcγRIIb develop enhanced inducible immune diseases, such as col-

lagen-induced arthritis (CIA) (6) and inducible alveolitis (7), and on certain genetic backgrounds FcγRIIb deficiency can also contribute to the development of spontaneous systemic lupus erythematosus (SLE) (8). Naturally occurring polymorphisms in the promoter of FcγRIIb reduce expression of the receptor in the mouse and are likely to contribute to spontaneous SLE in several mouse models (9–12). Similar polymorphisms in FcγRIIb may contribute to human SLE (13–15), and a transmembrane mutation associated with SLE is thought to do so by abolishing inhibitory function (16, 17). In addition to this central role in controlling susceptibility to autoimmunity, FcγRIIb controls survival after bacterial infection, balancing bacterial clearance and the risk of septic shock (18), and plays a key role in defense against malaria (19).

Stringent control of FcγRIIb expression is thought to be critical in determining its effect on the immune system. IL-4 enhances FcγRIIb expression and function on human macrophages (20–22), whereas the same cytokine abolishes FcγRIIb function and reduces expression on B cells (23, 24). Expression of FcγRIIb during dendritic cell maturation is also

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modulated by cytokines, as shown by an increase in Fc γ RIIb on mature human dendritic cells after treatment with IL-10 and IL-13 (25). Alteration of Fc γ RIIb expression on plasma cells correlates with the ability of immune complexes to induce their apoptosis (3), and changes in Fc γ RIIb expression are thought to be responsible for the therapeutic effect of intravenous Ig (26). Thus, physiological modulation of Fc γ RIIb, via underlying mechanisms that are incompletely understood, is likely to have important implications for immunity.

It has been shown that restoration of Fc γ RIIb expression using a lentiviral construct could restore tolerance and prevent autoimmune disease in Fc γ RIIb-deficient mice as well as in strains prone to autoimmunity (27, 28). These experiments demonstrated that restoration of Fc γ RIIb expression on only 30% of B cells, together with 20% of immature thymocytes and 10% of macrophages, was enough to prevent disease. This work clearly underlines the potential for restoration of Fc γ RIIb to normal levels, even in a subset of cells, to prevent autoimmunity. Three significant questions remain. The first is the identity of the cell type in which Fc γ RIIb restoration is most important. The second is whether or not a modest overexpression of Fc γ RIIb, rather than restoration of deficient expression, is enough to alter disease progression. This has clear implications for the potential for modulation of Fc γ RIIb to be therapeutically useful. The third question is the relative effect of such a cell-specific increase in Fc γ RIIb on infection as opposed to autoimmunity, which again has therapeutic implications.

To answer these questions, we generated several lines of mice expressing modestly increased levels of Fc γ RIIb on either B cells or macrophages. The relatively small increases in expression that were generated could be demonstrated to enhance the inhibitory function of the receptor on both cell types *in vitro*. Overexpression on the B cell had little effect on T-independent immune responses or on the early IgM response to T-dependent immunization, but it profoundly reduced the IgG response. Overexpression of Fc γ RIIb on the B cell had no effect on the frequency or initial course of CIA but resulted in early resolution, a marked reduction in joint destruction, and reduced levels of anti-collagen IgG antibodies. Spontaneous SLE was also inhibited, but responses to bacterial infection were not. In contrast, overexpression of Fc γ RIIb on macrophages had little effect on autoimmune disease or on immune responses but resulted in reduced bacterial phagocytosis and survival in response to *Streptococcus pneumoniae*. Thus, increasing expression of Fc γ RIIb on B cells and macrophages within a physiologically relevant range has a profound effect on immune responses, as well as therapeutic implications for the treatment of autoimmune, inflammatory, and infectious diseases.

RESULTS

Modest increases in the expression of Fc γ RIIb on B cells enhance its inhibitory effect *in vitro*

A construct was generated in which Fc γ RIIb cDNA was placed under the control of the human μ and V_H promoter

combined with the 3' κ enhancer to achieve B cell-specific expression (29). Three independent mouse lines were generated on the C57BL/6/CBA background, as identified by screening for a V5 tag incorporated in the construct (Fig. 1, A and B). The Fc γ RIIb used was of the Fc γ RIIb1 isoform and of the Ly17.1 allotype, to allow it to be differentiated from Ly17.2, the allotype of the parental strains. Similar expression of the transgene was demonstrated in three independently derived lines, with no effect on expression of the parental Fc γ RIIb allotype being observed (Fig. 1 C). Expression on a second line is shown in Fig. S1 (available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>). The transgene was expressed in all B cell subsets, with highest expression on plasma cells, but was not expressed on other cell types, including follicular dendritic cells (FDCs; Figs. S2 and S3). The major B cell subpopulations in spleen, peripheral blood, and lymph nodes were of normal size in transgenic mice (Fig. 1 D and Table S1). All three founders showed a 2-fold increase in Fc γ RIIb expression over control levels in 80–90% of the B cells, but interestingly, the remaining 10–20% had an approximately 10-fold increase in expression. This subpopulation was stable throughout the lifetime of the mouse, was composed predominantly of follicular B cells evenly distributed throughout the follicles on histology (Fig. S3), and had no phenotypic or functional features to distinguish them from the larger subpopulation, apart from Fc γ RIIb expression (see Supplemental discussion, available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>).

To determine if overexpression of Fc γ RIIb on B cells resulted in increased inhibition, B cells from B cell-transgenic (B-TG) mice were tested by several established assays of Fc γ RIIb inhibition. Purified B cells from B-TG mice stimulated with anti-IgM F(ab')₂, which cannot recruit Fc γ RIIb to the BCR, proliferated equally to controls. As expected, when Fc γ RIIb was cross-linked using an intact anti-IgM antibody, significant suppression was seen in control mice (30). This suppression was enhanced in the B-TG mice (Fig. 2 A). Addition of IL-4 abolished the inhibitory effect of Fc γ RIIb (Fig. 2 A), as previously reported (23). Another experiment with an Fc γ RIIb-deficient control for comparison is shown in Fig. S4 A (available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>). An independent measure of proliferation, the reduction of CFSE with increased cell division, was used to demonstrate the increase in activation threshold induced by expression of the transgene on B cells (Fig. 2 B). Finally, increased Fc γ RIIb expression almost completely abolished the generation of phospho-extracellular signal-regulated kinase (pERK) after BCR cross-linking (Fig. 2 C; quantified in Fig. S4, B and C). The induction of CD86 upon BCR cross-linking was also suppressed in B-TG mice but not in controls (Fig. 2 D). Thus, we generated independent lines of mice that expressed approximately twofold increased Fc γ RIIb on the majority of B cells, which resulted in an easily discernible increase in inhibitory function.

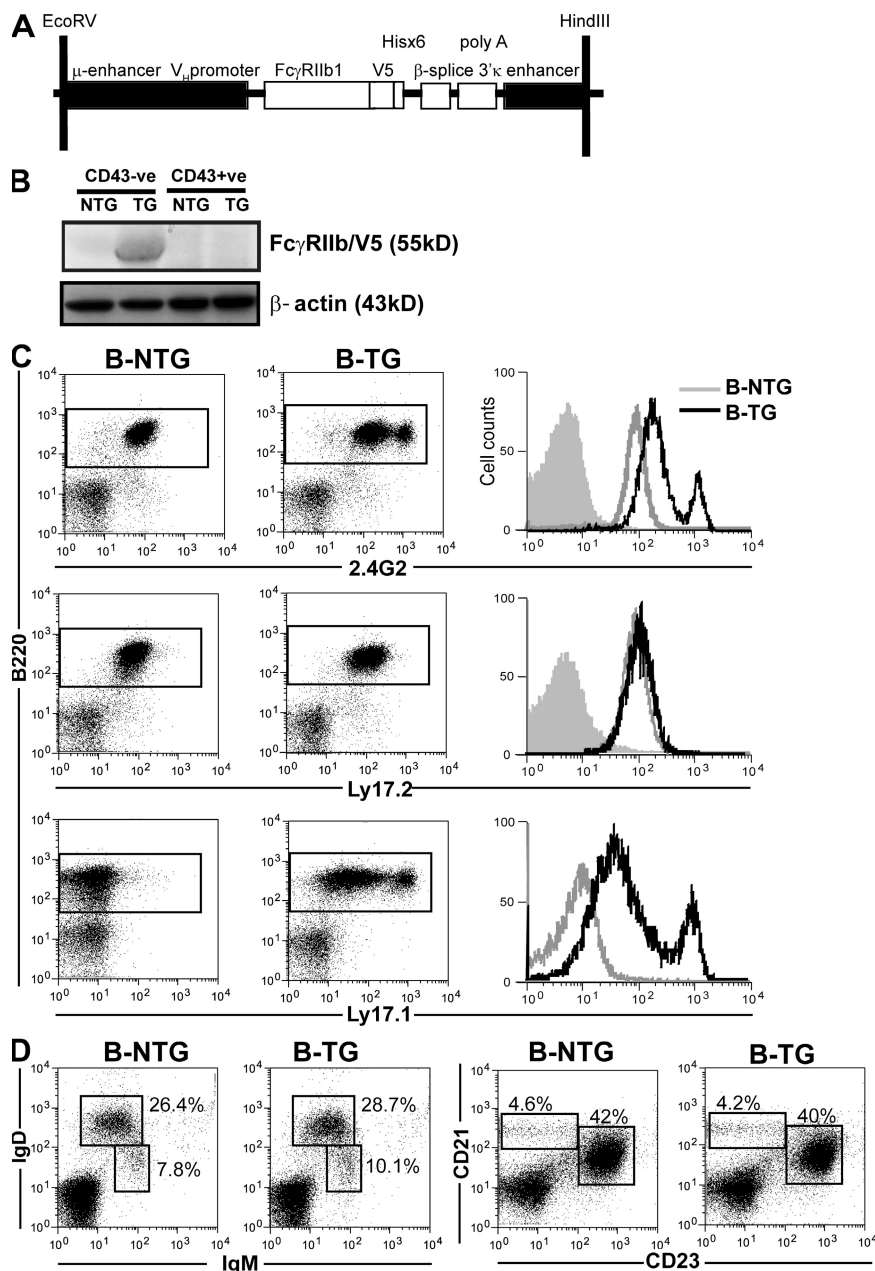


Figure 1. Generation of B-TG mice overexpressing Fc γ RIIb. (A) The construct used to direct B cell-specific expression of transgenic Fc γ RIIb. (B) Western blot, using a V5-specific antibody, of splenic B cell (CD43 $^{-ve}$) and non-B cell (CD43 $^{+ve}$) populations from B-TG and B-NTG mice. (C) Flow cytometric analysis of transgene expression on B220 $^{+}$ splenic cells (as shown in the gated populations on the dot plots) from B-NTG and B-TG mice. Histograms represent either 2.4G2 (which recognizes all Fc γ RIIb on B cells), Ly17.2 (endogenous Fc γ RIIb-specific), or Ly17.1 (transgenic Fc γ RIIb-specific) expression (shaded histogram = isotype control). No detectable transgene expression was demonstrated on non-B cells (Fig. S2). (D) The proportions of B cell subsets analyzed were normal in B-TG mice. Splenocytes were stained with anti-IgM, -IgD, -CD21, and -CD23 and assessed by flow cytometry (percentages of gated populations are shown; see also Table S1). B–D are representative of at least three independent experiments.

The predominant effect of Fc γ RIIb overexpression is to suppress the IgG component of T-dependent immune responses

Titers of total serum IgG, but not IgM, were reduced in B-TG mice (Fig. 3 A). To investigate this, we studied the antibody responses to a range of model antigens. Fc γ RIIb overexpression had surprisingly little effect on T-independent immune

responses (Fig. 3 B). Analysis of T-dependent immune responses showed that the IgM response was largely intact in B-TG mice. Anti-4-hydroxyl-3-nitrophenyl acetyl (NP) IgG titers were normal at day 5 of the response, but there was marked suppression of the IgG response from day 8 onwards (Fig. 3, C and D). Consistent with this, ELISPOT analysis showed little difference in the short-lived IgM and IgG plasmablast

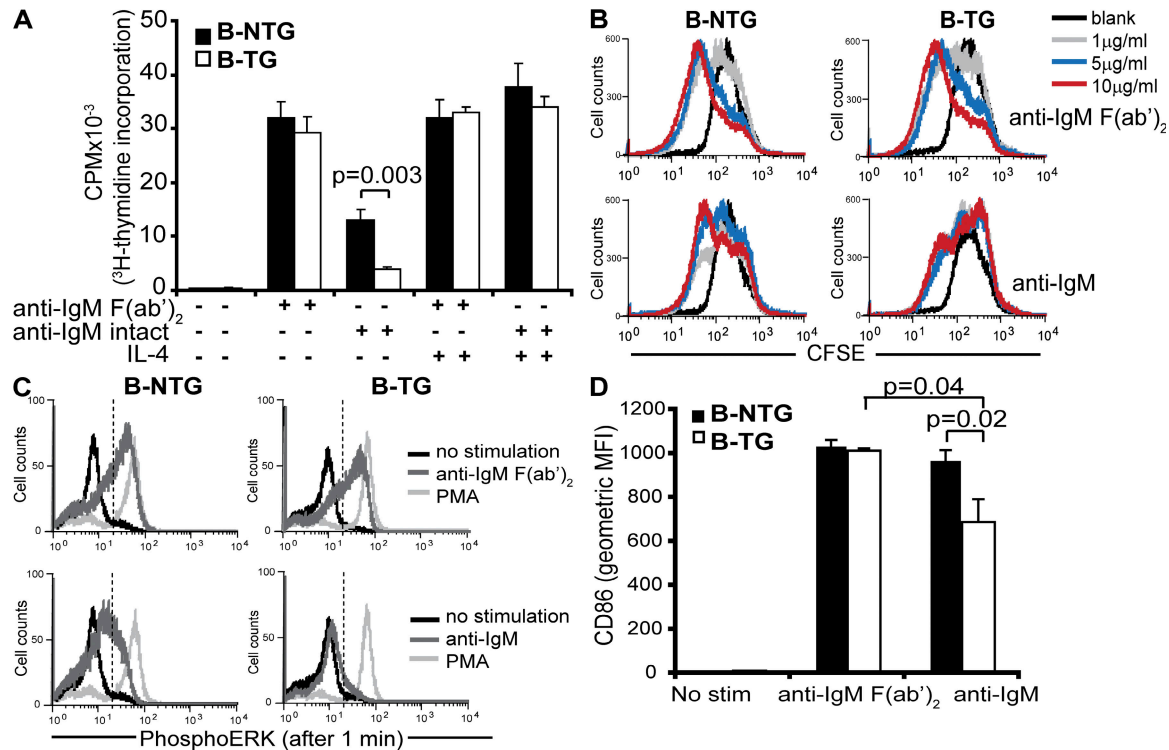


Figure 2. Fc-mediated proliferation, ERK phosphorylation, and CD86 expression are suppressed in B-TG B cells. (A) Proliferation of B cells from B-TG and B-NTG mice after 72 h of activation with either intact anti-IgM or anti-IgM F(ab')₂ antibodies, with and without IL-4, assessed by [³H]thymidine incorporation. (B) CFSE staining of B cells from B-NTG and B-TG mice after activation with varying concentrations of either intact anti-IgM or anti-IgM F(ab')₂ antibodies for 120 h before analysis by flow cytometry. (C) Assessment of ERK phosphorylation by intracellular flow cytometry after B cell activation for 1 min with either anti-IgM intact or anti-IgM F(ab')₂ antibodies (PMA = positive control; dashed lines indicate the gate used to quantify the extent of pERK, as shown in Fig. S4). (D) CD86 expression on B-NTG and B-TG B cells after activation with anti-IgM intact or anti-IgM F(ab')₂ molecules for 48 h, analyzed by flow cytometry. Error bars in A represent SD of triplicate wells from three pooled mice; error bars in D represent SD of three mice. p-values were calculated by the Student's *t* test. A is representative of three independent experiments; B, C, and D are representative of two independent experiments.

response (3). In response to secondary immunization, both IgM and IgG levels were reduced in B-TG mice compared with controls (Fig. 3 C), and we have previously described reduced bone marrow plasma cell accumulation and persistence in these mice (3). These results demonstrate that despite the fact that FcγRIIb cross-linking profoundly suppresses proliferation of IgM⁺ B cells in vitro (Fig. 2, A and B; and Fig. S4 A) (4, 30), IgM production is not influenced in the primary immune response in vivo. Affinity maturation was ascertained by NP₂/NP₁₂ ELISA (31) after secondary immunization. There was a significant increase in the ratio of anti-NP₂ to anti-NP₁₂ IgG in B-TG mice (Fig. 3 E), consistent with more stringent selection of high affinity B cells when FcγRIIb is increased.

Increased macrophage inhibition by transgenic expression of FcγRIIb

Macrophage-specific expression of FcγRIIb was achieved using a construct in which FcγRIIb2 mRNA was placed under the control of the human CD68 promoter (32). This resulted in macrophage-specific expression in two independent

lines, as demonstrated by Western blotting for the V5 tag incorporated in the construct (Fig. 4, A and B). Expression was confirmed on peritoneal macrophages (Fig. 4 C), as well as on macrophages from bone marrow and spleen, and on peritoneal macrophages from an independent strain (Fig. S5, available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>). Expression was not seen on dendritic cells, B cells, FDCs, or other cell types (Figs. S5 and S6). Macrophage numbers in all of these organs were normal (Table S2). To determine if the macrophage transgene was functional, Fc receptor-triggered calcium signaling, known to be inhibited by FcγRIIb (33), was assessed in Fura2-loaded peritoneal macrophages. Although there were no differences in peak intracellular calcium signal after Fc receptor aggregation, plateau calcium levels were significantly reduced in cells from transgenic mice (Fig. 4 D). Phosphorylation of ERK was also measured in peritoneal macrophages as a read out of activation after Fc receptor cross-linking. ERK phosphorylation was suppressed by expression of the transgene, in contrast to controls and to the marked increase seen in FcγRIIb-deficient mice (Fig. 4 E). Thus, a relatively modest increase in

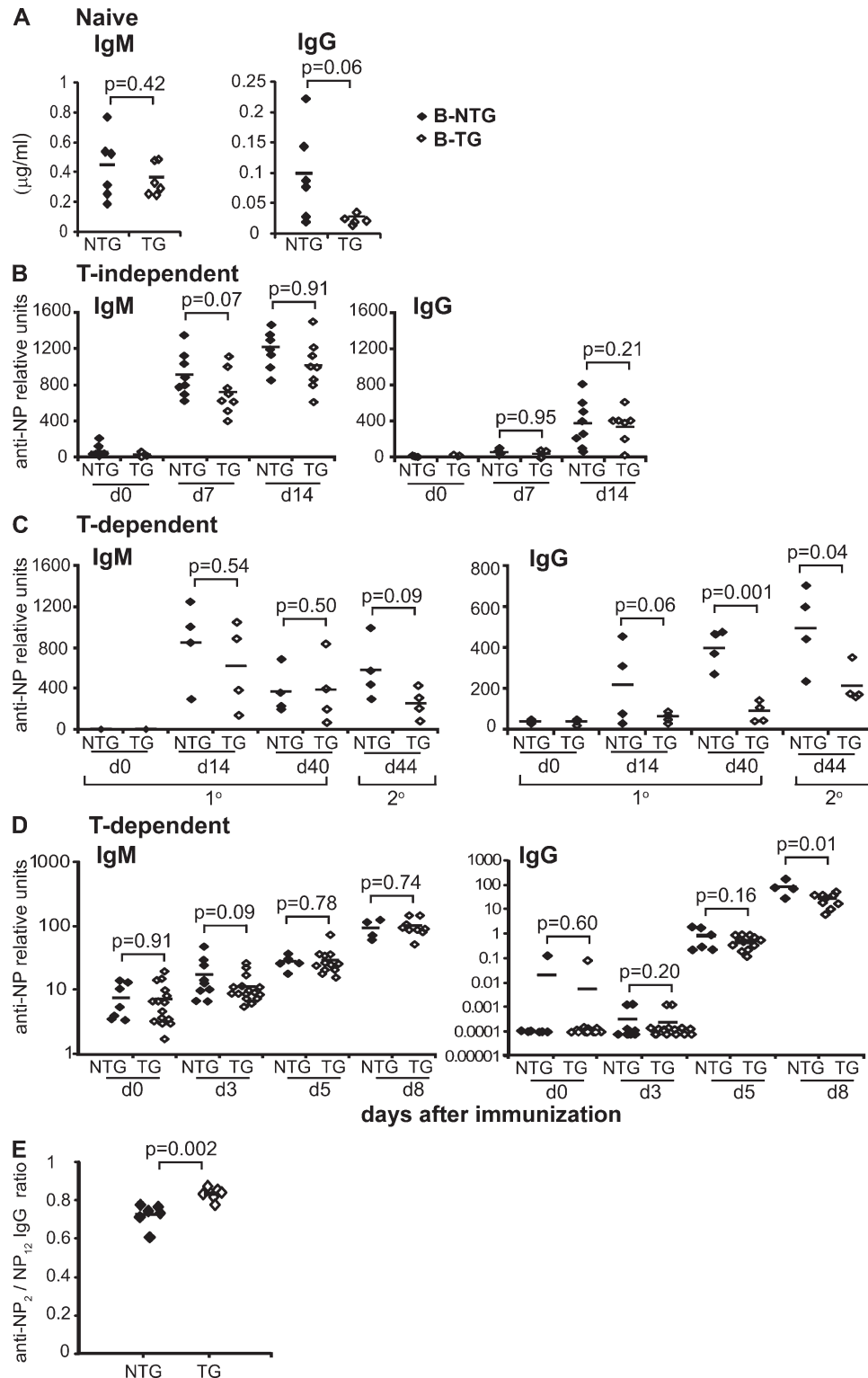


Figure 3. Overexpression of Fc γ RIIb on B cells suppresses the T-dependent IgG immune response. (A) Total IgG and IgM titers were measured in serum from naive B-NTG and B-TG mice. Values were calculated from a standard curve of known Ig concentration. Mice were immunized intraperitoneally with NP-haptenated carriers in alum, and antibody responses were assessed by ELISA. B-TG and B-NTG mice were injected with either T-independent (B, NP-ficoll) or T-dependent (C, NP-CGG) antigens and boosted on day 40 with NP-CGG alone or NP-KLH (D). Serum was then assessed for total anti-NP₁₂ IgM and IgG production at various time points after immunization. (E) The affinity of anti-NP IgG was assessed by calculating the ratio of anti-NP₂ (high affinity)/anti-NP₁₂ (total) IgG (NP₂/NP₁₂) by ELISA from serum obtained from B-NTG and B-TG mice immunized with NP-KLH on day 0, boosted on day 40, and bled on day 49. Each symbol shows data from a single mouse, and the horizontal lines represent the mean. p-values were calculated using the Student's *t* test. A–C are representative of three independent experiments.

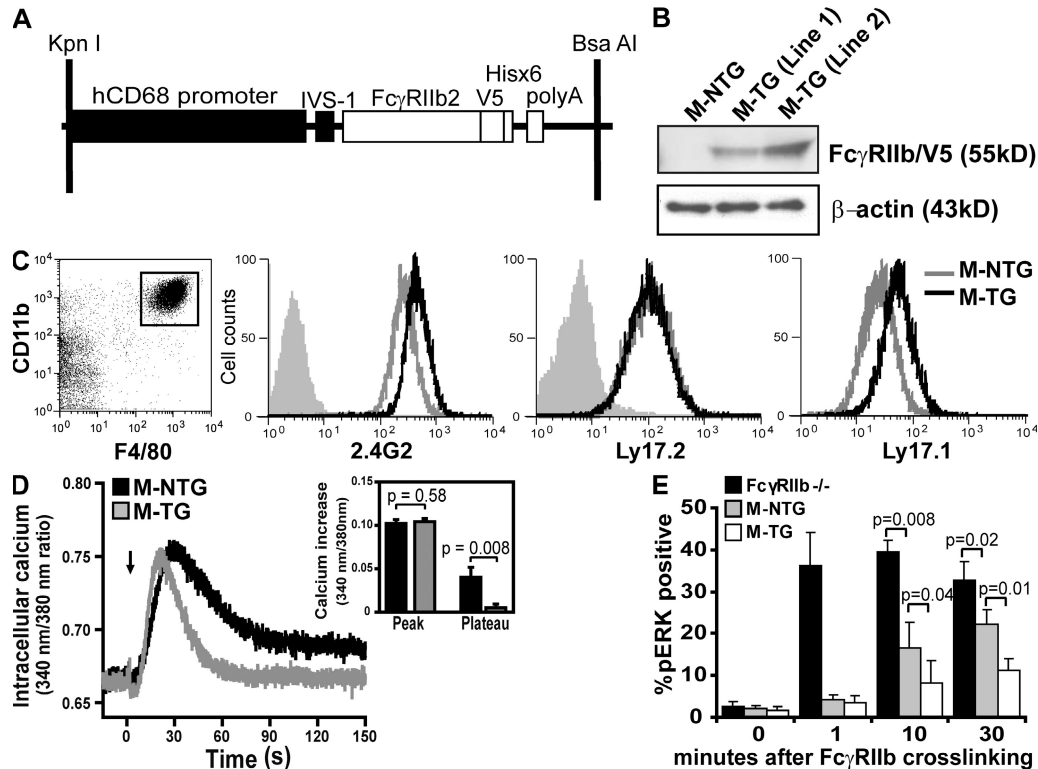


Figure 4. Generation of M-TG mice overexpressing functional Fc γ RIIb. (A) The construct used to direct macrophage-specific expression of transgenic Fc γ RIIb. (B) Western blot, using a V5-specific antibody, of peritoneal macrophages from M-TG (lines 1 and 2) and M-NTG mice to detect the presence of the V5-tagged transgene. All further experiments use line 2. (C) Flow cytometric analysis of transgene expression on F4/80⁺CD11b⁺ peritoneal macrophages (gated population shown on dot plot) from M-NTG and M-TG mice. Histograms represent either 2.4G2 (recognizes all Fc γ RIIb, III, and IV), Ly17.2 (endogenous Fc γ RIIb-specific), or Ly17.1 (transgenic Fc γ RIIb-specific) expression (shaded histogram = isotype control). No detectable transgene expression was demonstrated on other cell types (Figs. S5 and S6). (D) Intracellular calcium measurements in M-NTG and M-TG peritoneal macrophages after Fc γ RIIb cross-linking (arrow). Inset shows mean peak and plateau calcium responses. (E) Measurements of ERK phosphorylation in peritoneal macrophages from M-NTG, M-TG, and Fc γ RIIb^{-/-} mice after Fc γ RIIb cross-linking. The percent increase in pERK-positive cells after stimulation is shown. Error bars represent the SD of triplicate samples from three or four pooled mice. p -values were calculated using the Student's t test. D and E are representative of two and three independent experiments, respectively.

Fc γ RIIb expression on macrophages resulted in a significant increase in inhibitory effect *in vitro*.

Fc γ RIIb overexpression on macrophages does not influence the T-dependent immune response

Macrophage-transgenic (M-TG) mice were immunized with two T-independent antigens (NP-dextran and NP-LPS), and although a possible trend toward a reduction in IgG production was seen at later time points, initial responses to these antigens were not significantly different from control mice (Fig. S7, A and B, available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>). Furthermore, the T-dependent response in M-TG mice was indistinguishable from macrophage-nontransgenic (M-NTG) controls (Fig. S7 C).

Increased Fc γ RIIb expression on B cells, but not macrophages, modifies the course of CIA and spontaneous lupus

CIA is a widely used model of inflammatory joint disease. It can be induced on the C57BL/6 background using a modi-

fied chick type II collagen (CII) immunization protocol (34). B-TG mice showed the same incidence of CIA as controls, and the time of onset and early course of the inflammation were indistinguishable (Fig. 5, A and B). In contrast in the B-TG mice, however, joint inflammation resolved quickly after disease onset, and this resolution was sustained. Consistent with this, there was a reduction in joint damage, as determined by histology at day 74 in B-TG mice compared with controls (Fig. 5 G). IgG anti-collagen antibodies are important in the pathogenesis of CIA (35, 36) and are suppressed in B-TG mice, providing a possible mechanism for the reduction in inflammation (Fig. 5 E). All IgG isotypes were equally suppressed in the B-TG mice (Fig. S8, available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>), but IgM was not affected (Fig. 5 E). In contrast, no substantial difference in incidence, disease course, or antibody production was seen in several experiments in the M-TG mice (Fig. 5, C, D, and F), whereas there was a nonsignificant trend toward increased histological joint damage, perhaps because of a failure of macrophage remodeling and/or immune complex clearance (Fig. 5 H).

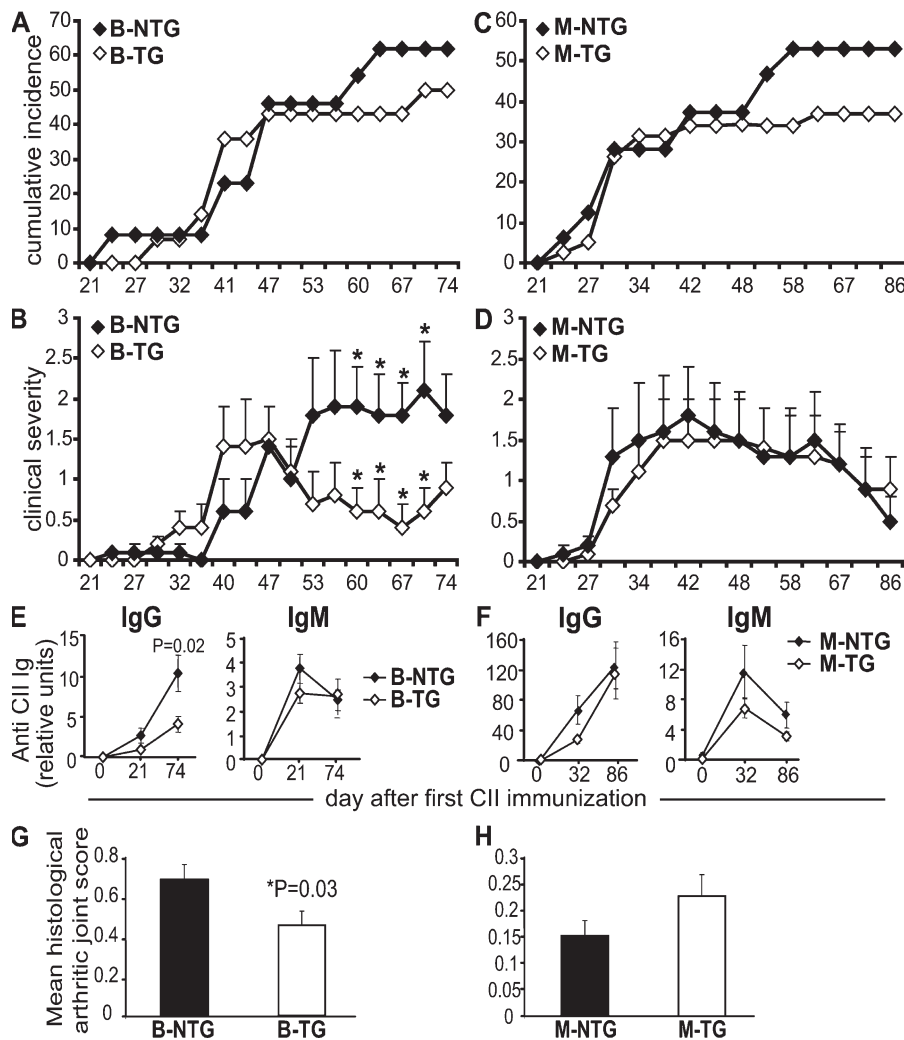


Figure 5. Suppressed severity of CIA in B-TG but not M-TG mice. (A) Cumulative arthritis incidence and (B) mean clinical scores (\pm SEM) for B-TG ($n = 14$) compared with B-NTG ($n = 13$) mice over 74 d. (C) Cumulative arthritis incidence and (D) mean clinical scores (\pm SEM) for M-TG ($n = 32$) compared with M-NTG ($n = 38$) mice (data represent three independent experiments combined). p-values for clinical scores were calculated using the unpaired Student's *t* test. *, $P < 0.05$; no significant differences were detected in the incidence by χ^2 analysis. (E and F) Circulating levels of CII-specific antibody were determined in individual sera by ELISA. Results show mean total IgG and IgM levels (\pm SEM) as arbitrary units per milliliter (levels of IgG isotypes are shown in Fig. S8). (G and H) At the end of the study, paws were processed for histology and scored blinded for signs of arthritis. Data represent mean \pm SEM.

MRL/*lpr* mice develop a spontaneous autoimmune disease resembling human SLE. The disease is characterized by the development of antinuclear antibodies (ANA), glomerulonephritis, and inflammatory skin lesions (37). Although other cells play their role, the importance of B cells is highlighted by complete abrogation of disease in B cell-deficient MRL/*lpr* mice (38). B-TG and M-TG mice were crossed onto the MRL/*lpr* background (Table S3, available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>) and assessed for the development of autoimmunity. There was a striking reduction in mortality in the B-TG MRL/*lpr* mice. All B cell-nontransgenic (B-NTG) MRL/*lpr* mice were moribund by 34 wk of age, compared with only 14% of B-TG MRL/*lpr* mice (Fig. 6 A). Consistent with this protection from disease, B-TG mice had significantly lower levels of protein-

uria and autoantibodies (Fig. 6, C, E, and F). In contrast, no significant differences in disease progression were observed in the M-TG mice (Fig. 6, B, D, and G).

Fc γ RIIb overexpression on macrophages, but not B cells, suppresses responses to bacterial infection

In view of the effect of Fc γ RIIb on bacterial infection in Fc γ RIIb-deficient mice (18), we examined the response of the transgenic mice to pneumococcal infection. In the B-TG mice, "natural" IgM antibody titers against pneumococcus were normal, and IgG was only slightly reduced (Fig. S9, available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>). In addition, phagocytosis of opsonized bacteria in vitro by peritoneal macrophages was also normal (unpublished data). Consistent with these findings, no effect

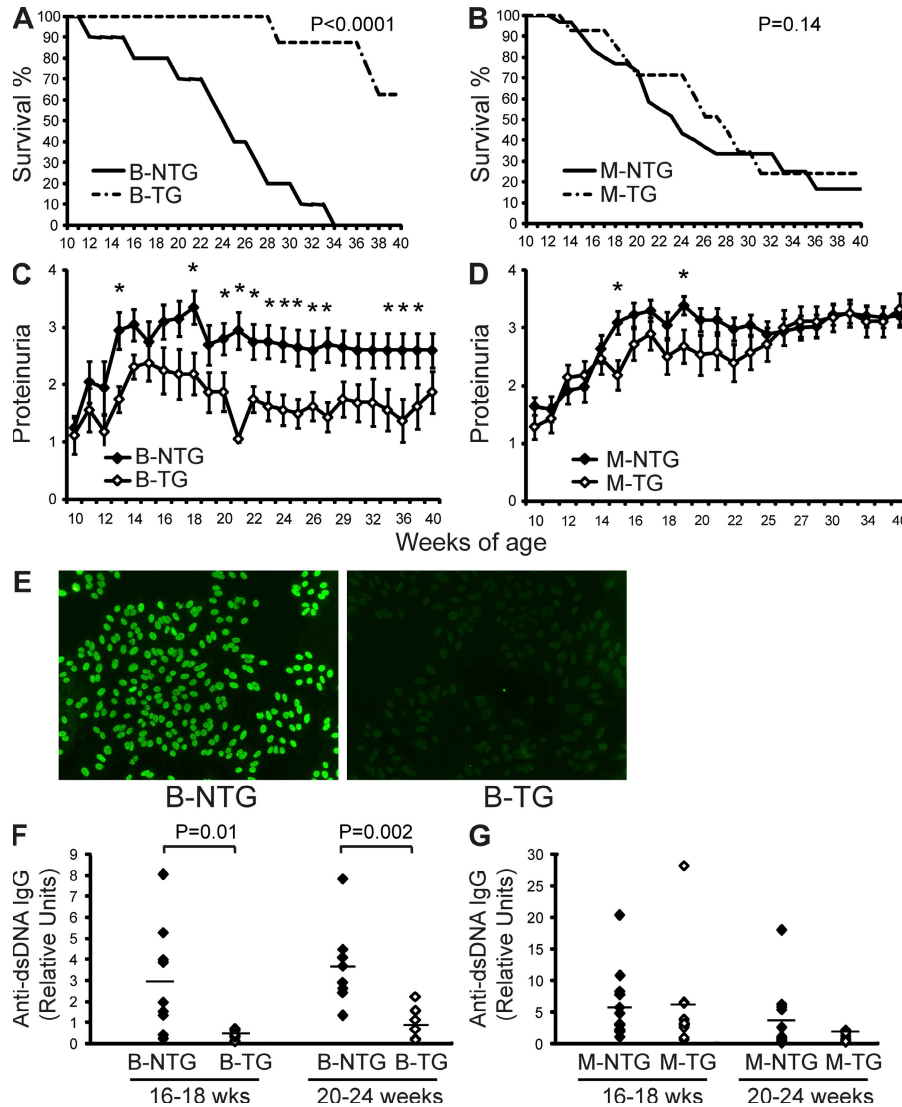


Figure 6. Protection from lupus in B-TG mice. B-TG and M-TG mice were crossed with MRL/lpr mice and spontaneous lupus was monitored for up to 40 wk of age. (A and B) Survival rates are shown for groups of B-TG ($n = 8$), B-NTG ($n = 10$), M-TG ($n = 12$), and M-NTG ($n = 21$) mice. (C and D) Proteinuria levels were recorded weekly in all mice throughout the study. Data are presented as mean \pm SEM. p-values were calculated using the unpaired Student's t test. *, $P < 0.05$. (E) All mice were screened for serum levels of ANA at 20–24 wk of age using Hep-2 slides, and representative results for B-NTG and B-TG mice are shown. (F and G) Serum from each mouse was analyzed for anti-dsDNA IgG. Each symbol represents an individual mouse. Error bars represent SEM. p-values were calculated using the χ^2 test (A and B) and the Student's t test (all other values).

on mortality was seen in the B-TG mice when they were challenged with pneumococcus (Fig. 7 A). In contrast, although natural antibody levels were normal (Fig. S9), M-TG mice showed a marked reduction in their ability to phagocytose opsonized pneumococci in vitro (Fig. 7 B). In keeping with this, there was a modest but consistent and significant reduction in mortality in M-TG mice after challenge with pneumococcus given either intraperitoneally or intranasally (Fig. 7, C–E).

DISCUSSION

Enhancing the expression of Fc γ RIIb on both B cells and macrophages was shown to increase the inhibitory effect of

this receptor on both cell types. The degree of expression difference seen in both transgenic mice was less than the difference that can be induced by cytokine treatment, for example (20–24), and was thus likely to fall within the physiological range of expression of the receptor. Nonetheless there was a profound increase in inhibitory effect, as measured by several independent in vitro assays. This implies that the physiological variation in Fc γ RIIb expression seen in humans and mice with Fc γ RIIb promoter polymorphisms, and the changes in expression that can be induced by cytokines, are very likely to have substantial effects on Fc γ RIIb function.

The effect of Fc γ RIIb on B cell proliferation has traditionally been measured in vitro by cross-linking surface IgM

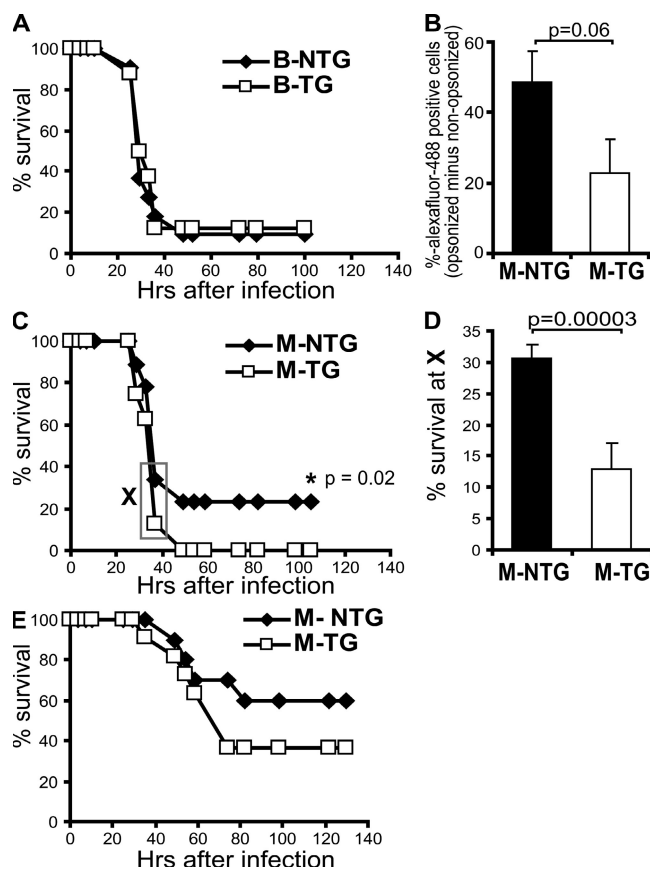


Figure 7. M-TG mice are more susceptible to pneumococcal infection. Survival after *S. pneumoniae* infection. (A) Groups of B-TG ($n = 8$) and B-NTG ($n = 11$) mice were inoculated intraperitoneally with 2×10^4 CFU *S. pneumoniae* type 2 (D39). (B) Peritoneal macrophages were incubated with Alexa Fluor 488-labeled *S. pneumoniae* either unopsonized or opsonized with heat-inactivated immune serum, followed by flow cytometric analysis. Antibody-dependent phagocytosis is expressed as the percentage of Alexa Fluor 488⁺ cells relative to the nonopsonized sample. Error bars represent SD of three mice per group and are representative of four independent experiments. (C) Groups of M-TG ($n = 8$) and M-NTG ($n = 9$) mice were inoculated intraperitoneally with 10^2 CFU *S. pneumoniae* type 2 (D39). A significant reduction in survival was observed for the M-TG mice ($P = 0.02$). (D) Graphical representation of data pooled from five separate infection experiments (including C) using various doses of *S. pneumoniae* (3×10^1 , 10^2 , 10^4 , 10^6 , and 10^7). Data represent mean \pm SEM. The kinetics differ depending on dose; to normalize for this, data are shown as survival at the time point before 100% mortality of the M-TG mice (indicated by the gray box in C). (E) Intranasal infection of M-NTG ($n = 11$) and M-TG ($n = 10$) mice with 2×10^6 CFU *S. pneumoniae* type 2 (D39). p -values were calculated using the Student's t test (B and D) and the log-rank test (C).

with or without ligation of Fc γ RIIb. This has profound effects on the activation threshold of B cells, as shown in several previous studies (4, 30), as well as in Fig. 2. It is therefore interesting that the overexpression of Fc γ RIIb on B cells has little effect on T-independent responses or on IgM responses to T-dependent antigens. This, together with the fact that IgG was only markedly affected from day 8 after immunization,

suggests two possible explanations. The first is that Fc γ RIIb-mediated suppression of B cell responses requires the production of an adequate titer of antigen-specific IgG that can cross-link Fc γ RIIb. Such cross-linking may not be achieved in T-independent responses, as they are dominated by IgG3 (39), which has extremely low affinity for Fc γ RIIb (5). The relative lack of effect on the T-dependent IgM response could reflect the fact that most IgM is made within the first week of immunization, before IgG rises. This model is also consistent with the initial normal production of anti-NP IgG at day 5, with mild differences beginning to be seen at day 8, presumably as anti-NP IgG titers have risen sufficiently to allow Fc γ RIIb cross-linking to occur. An alternative or additional explanation is that Fc γ RIIb-mediated suppression in vivo requires substantial cross-linking only achieved on the surface of the FDC. This model would predict that suppression would only occur once the germinal center (GC) reaction had become important, thereby accounting for the lack of effect on T-independent responses, where GCs play at most a minor role. In T-dependent responses, it would explain both the lack of effect on IgM (most early IgM is made by plasmablasts of the extrafollicular foci, and GCs predominantly contain isotype-switched B cells) and on early IgG, made before GC formation has become important (usually estimated to be from 6 d onwards) (40). The low affinity plasmablast response (both IgM and IgG) is controlled independently of Fc γ RIIb, as we have previously shown using Fc γ RIIb^{-/-} mice (3). Thus, overexpression of Fc γ RIIb has helped define the precise stage of the immune response at which Fc γ RIIb-mediated suppression is important, and suggests the need for further work to elucidate the mechanism underlying this. In addition, we have presented data that suggest increased stringency of affinity maturation in B-TG mice. This suggests that an increased B cell activation threshold controls affinity maturation. Currently, we are testing this at the single-cell level, as described previously (31).

The Fc γ RIIb-deficient mouse was made using a C57BL/6/129 embryonic stem cell line (4). When other genes in the same region of distal chromosome 1 have been deleted, they have also shown an autoimmune phenotype (e.g., complement receptor 1/2 [41], decay accelerating factor CD55 [42], and serum amyloid P component [43]), much of which has subsequently been found to be caused by the adjacent region of 129 origin rather than to the gene deletion itself (44, 45). The effect of our transgene in suppressing autoimmunity is clearly independent of this 129 region. This confirms that Fc γ RIIb plays a role in controlling autoimmunity and suggests that the autoimmune phenotype of the Fc γ RIIb-deficient mouse is not an artifact of 129 contamination.

Antibodies have long been known to be important in the pathogenesis of CIA. Mice deficient in B cells are totally resistant to CIA induction (34), and it is possible to induce arthritis using anti-CII antibodies (36, 46). Despite B-TG mice having decreased anti-collagen antibodies, the incidence and initial course of CIA was unchanged. Rather, the suppression of anti-collagen antibodies by transgenic Fc γ RIIb results in

early disease resolution and an improved long-term outcome. This shows that relatively modest modulation of Fc γ RIIb expression can have a critical effect on inflammatory arthritis despite not affecting onset. If it was possible to achieve such a change in expression pharmacologically, it would have clear therapeutic implications, particularly in humans, where preemptive therapy is rarely possible and patients present with established disease. The fact that overexpression of Fc γ RIIb on B cells also markedly reduces spontaneous SLE provides further encouragement that this avenue may be therapeutically useful. The effect of Fc γ RIIb overexpression on the macrophage, in contrast, had little effect on either CIA or spontaneous SLE. Overall, these results indicate that the B cell, rather than the macrophage, should be the primary target of therapeutic modulation of Fc γ RIIb in inflammatory disease.

The effect of overexpressing Fc γ RIIb on autoimmunity will be influenced by the fact that different Fc receptors selectively engage IgG subclasses with varying affinities (1, 47–49). Fc γ RI binds IgG2a with high affinity and Fc γ RIV binds IgG2a and IgG2b with intermediate affinity, whereas Fc γ RIIb and Fc γ RIII bind IgG2a, IgG2b, and IgG1. Moreover, the affinity with which Fc γ RIV binds IgG2a or IgG2b is at least twofold higher than Fc γ RIIb and between 20 and 40 times higher than Fc γ RIII (50). Thus, IgG2a and IgG2b immune complexes would preferentially bind to Fc γ RIV, and little effect of Fc γ RIIb co-cross-linking may be seen. In contrast, immune complexes containing IgG1 would more likely bind Fc γ RIIb and, thus, induce its inhibitory effect. CIA is predominantly associated with IgG2a anti-collagen antibodies (Fig. S5) (51), and in SLE, autoantibodies of the IgG2a subclass are also dominant (52). Thus, in the M-TG mouse, immune complexes are likely to engage Fc γ RIV in preference to Fc γ RIIb, which might explain the lack of effect of Fc γ RIIb overexpression on macrophages in both CIA and SLE.

Natural antibody levels against pneumococcus were similar to controls in both B-TG and M-TG mice, consistent with the fact that they were also normal in Fc γ RIIb-deficient mice (18). Given that pneumococcal challenge results in very rapid onset of illness before an IgG response can be initiated, it is thus not surprising that survival was normal in the B-TG mice. In M-TG mice, however, where reduced phagocytosis of pneumococcus could be demonstrated *in vitro*, there was a significant reduction in survival. Although of a similar magnitude, this effect is the converse of that seen with the Fc γ RIIb-deficient mouse (18), thus underlining the likelihood that the latter observation was caused by Fc γ RIIb deficiency on the macrophage. This highlights the importance of Fc γ RIIb in controlling initial bacterial clearance in infection. It focuses attention on the potential for manipulating expression of the receptor on macrophages to either be useful in therapy, perhaps in the treatment of septic shock (9, 18), or to be responsible for adverse effects if expression is increased in the treatment of inflammatory disease.

Cell-specific overexpression of Fc γ RIIb on B cells and macrophages highlights the role that therapeutic modulation of

receptor expression on B cells might play in inflammatory and autoimmune disease. It has also defined the role of Fc γ RIIb in suppressing the T-dependent immune response, and has shown that increased expression on macrophages reduces survival after bacterial infection, demonstrating the importance of macrophage Fc γ RIIb in the control of infection and emphasizing the importance of careful cell-specific targeting in therapeutic modulation.

MATERIALS AND METHODS

Mice

C57BL/6 Fc γ RIIb^{-/-} mice (4) were provided by J. Ravetch and S. Bolland (The Rockefeller University, New York, NY). MRL/*lpr* mice were purchased from Harlan. All other mice were obtained from Charles River Laboratories. All experiments were performed according to the regulations of the UK Home Office Scientific Procedures Act (1986). The animal experiments were approved by the UK Home Office.

Generation of B-TG and M-TG mice overexpressing Fc γ RIIb. To direct B cell-specific expression of Fc γ RIIb, we used a construct containing a V_H promoter, the IgH intron enhancer, and the Ig κ 3' enhancer, which was a kind gift from T. Tsubata (Tokyo Medical and Dental University, Tokyo, Japan) (29). To direct macrophage-specific expression of Fc γ RIIb, we used a construct containing the human CD68 (hCD68) promoter, as previously described (32). We introduced the mouse Fc γ RIIb.1 (Ly17.1 allotype) tagged with both the V5 and Hisx6 epitopes into these constructs. Transgenic mice were established by injecting the DNA fragment containing the tagged Fc γ RIIb cDNA and the respective promoters, or enhancers in the case of the B cell construct, into CBA-fertilized eggs. The presence of transgene was identified by tail DNA PCR assays (B-TG primers: 5'–3', 5'–TTCTCGGCTCTGTGAATGACA–3', and 3'–5', 5'–CAGCCCTCTCTTGAAAGGAG–3'; M-TG primers: 5'–3', 5'–CATAGCTGGAGGAACAACTACTGAAC–3', and 3'–5', 5'–GTAGAATCG AGACCGAGGAGAGG–3'). Transgenic offspring were backcrossed with C57BL/6 mice for at least seven generations. Transgenic mice were crossed to MRL/*lpr* mice for at least five generations using a marker-selected backcrossing method (Table S3) predicted to result in >99% MRL genetic background by N5 (53).

B cell purification and stimulation

Splenic B cells were negatively selected by magnetic cell purification using anti-CD43 beads (Miltenyi Biotec) to >90% purity. 2×10^6 cells/ml were usually cultured in 24-well flat-bottom plates (Corning Inc.) in RPMI 1640 (Sigma-Aldrich) and 10% FCS (Invitrogen), and were stimulated with either goat anti-mouse IgM μ chain-specific F(ab')₂ or intact IgG at 10 μ g/ml, unless otherwise stated in the figures (Jackson ImmunoResearch Laboratories). IL-4 added to these cultures was used at 10 ng/ml (PeproTech). PMA was used at 1 ng/ml.

Peritoneal macrophages, splenic dendritic cells, and bone marrow-derived macrophages (BMDMs) and dendritic cells (BMDCs)

Resident peritoneal macrophages were harvested from mice by peritoneal lavage using 5% BSA in PBS. Splenic dendritic cells were isolated from spleens that were gently disrupted between glass slides and cut up into 5 ml of 2% BSA/PBS containing 1 mg/ml collagenase (Worthington) and 0.1 mg/ml DNase I (Roche). Suspensions were kept at room temperature for 30 min with regular pipetting before RBC lysis using ammonium chloride. Bone marrow cells were flushed with cold RPMI 1640 containing 10% FCS from the dissected tibiae and femurs of NTG and TG mice. RBCs were lysed with ammonium chloride, and the remaining cells were washed three times. 2×10^6 bone marrow cells per well were seeded into six-well tissue culture plates. Cells were cultured for 7 d in the presence of 4 ng/ml of recombinant M-CSF (PeproTech) to generate BMDMs, or in the presence of 20 ng/ml of recombinant GM-CSF (PeproTech) to generate BMDCs.

Flow cytometric analysis

Cells were stained with combinations of anti-FcγRII/III-FITC and -biotin (2.4G2), anti-FcγRII (Ly17.1-FITC and Ly17.2-biotin; Cedarlane), CD19-FITC (1D3), CD22-FITC (Cy34.1), B220-allophycocyanin (APC), B220-biotin, B220-PE (RA3-6B2), IgM-FITC (R6-60.2), I-A^b-FITC (25-9-17), IgD-PE (11-26; SouthernBiotech), CD21/CD35-FITC (7G6), CD23-PE (B3B4), CD86-PE (GL1), CD3-PE (17A2), CD4-PerCP (GK1.5), CD8-biotin (RA3-6B2), CD43-biotin (S7), F4/80-RPE (MCA497PE; Serotech), CD11b-APC (M1/70), CD11c-APC, CD11c-PE (HL3), or isotype controls (all antibodies were purchased from BD Biosciences unless otherwise stated). Biotinylated antibodies were cross-linked with streptavidin conjugated to FITC, PerCP, or APC (BD Biosciences). 1% normal rat serum was used to block nonspecific antibody binding. 7-amino-actinomycin D (Invitrogen) was used to exclude dead cells. Cells were analyzed using a flow cytometer (FACSCalibur; BD Biosciences) and FCS Press software (provided by R. Hicks, University of Cambridge, Cambridge, UK).

Immunofluorescence microscopy

Spleens from B-NTG and B-TG mice were sectioned 7 μm thick using a cryostat and left to dry overnight. Sections were fixed in cold acetone for 5 min and blocked for 1 h with 5% BSA/PBS and 5% normal goat or rat serum before staining. Sections were stained in blocking medium at room temperature with 2.4G2-FITC and B220-biotin for 1 h, and then stained with streptavidin-Alexa Fluor 546 for 1 h. For FDC staining, sections were initially stained with rat anti-mouse FDCM-1 (BD Biosciences) and detected with goat anti-rat IgG Alexa Fluor 555 (absorbed against mouse Ig; Invitrogen). Sections were then stained with 2.4G2-FITC, B220-biotin, and goat anti-mouse IgG Alexa Fluor 633 (highly cross absorbed; Invitrogen), followed by streptavidin-Pacific blue (Invitrogen). Alternatively, spleens from M-NTG and M-TG mice were sectioned and stained with FDCM-1, as described, and then stained with 2.4G2-FITC and B220-Alexa Fluor 647 (BioLegend). Vectorshield hardset was used to mount sections before analysis by confocal microscopy.

Intracellular signaling measurements

Purified B cells were stimulated with either goat anti-mouse IgM μ chain-specific F(ab')₂ or intact IgG for between 1 and 30 min. Peritoneal macrophages were stimulated by incubating cells with 2.4G2-biotin before cross-linking with streptavidin for between 1 and 30 min (Dako). To determine the extent of ERK phosphorylation, stimulated cells were fixed in 4% formaldehyde for 10 min at 37°C and permeabilized by adding 100% methanol for 30 min at 4°C. Samples were incubated with phospho-p44/42 Map kinase (Thr202/Tyr204) antibody (Cell Signaling Technology) for 1 h at room temperature, washed, and stained with an Alexa Fluor 488 F(ab')₂ fragment of goat anti-rabbit IgG (H+L; Invitrogen). Cells were analyzed by flow cytometry. To determine intracellular calcium flux, macrophages were loaded with Fura2-AM ester before activation and measured by cuvette fluorimetry, as previously described (54).

Proliferation assays

2 × 10⁶ purified B cells/ml were stimulated with anti-IgM μ chain-specific F(ab')₂ or intact antibody for 72 h. 0.5 μCi tritiated [³H]thymidine (GE Healthcare) was added for the last 8 h of culture, and incorporation was determined using a scintillation counter (PerkinElmer). For CFSE labeling, 5 × 10⁷ purified B cells/ml were loaded with 5 μM CFSE (Invitrogen) before culturing 2 × 10⁶ cells/ml with anti-IgM and anti-IgM F(ab')₂ molecules. The extent of CFSE labeling was analyzed by flow cytometry.

Western blot analysis

Standard Western blot analysis was performed using anti-V5-horseradish peroxidase (HRP) conjugate (Invitrogen) and chemiluminescent detection. Blots were stripped and reprobed with a β-actin-specific antibody (Abcam) as a loading control.

T-independent and T-dependent immunization

Samples of preimmune sera were taken before the initial immunization. Groups of mice were immunized intraperitoneally with 100 μg NP-ficol, NP-dextran, NP-LPS, NP-chicken γ globulin (CGG), or NP-KLH (Biosearch Technologies) emulsified in alum (Alu-Gel-S; Serva Electrophoresis). Samples of blood from tail veins were taken at various time points after immunization. For secondary responses, mice received a further 100 μg NP-CGG or -KLH/alum on day 28 or 40. Tail bleeds were taken on day 44.

ELISA

For analysis of serum anti-NP antibodies, plates (Nunc Maxisorp; Thermo Fisher Scientific) were coated with 5 μg/ml NP₁₂- or NP₂-BSA (Biosearch Technologies) overnight (the NP₂/NP₁₂ ratio was used to assess affinity maturation). Plates were blocked by the addition of PBS/1% BSA for 1 h. Serial dilutions of sera were added. To detect specific antibody isotypes, HRP-conjugated secondary antibodies were used (SouthernBiotech). EC₅₀ values were calculated by nonlinear regression analysis using Prism software (Microsoft). Standard curves were constructed from the pooled sera of NP-hyperimmunized mice.

Serum IgM, IgG, IgG2a, IgG2b, IgG1, and IgG3 were assayed using paired capture and HRP-conjugated antibodies (SouthernBiotech), and a standard curve of known concentration of the same mouse Ig isotype.

Anti-dsDNA antibody levels were measured by ELISA, as previously described (55). In brief, 96-well ELISA plates were coated with 5 μg/ml of calf thymus dsDNA (Sigma-Aldrich) in sodium salt citrate buffer at 37°C overnight. Plates were washed with PBS/0.05% Tween 20 and blocked with 1% BSA/PBS, and sera were added in serial dilutions starting at 1:100 and incubated for at least 1 h at room temperature. After a further wash, HRP-conjugated goat anti-mouse IgG was added. Pooled MRL/lpr serum was used as a standard curve.

Levels of antiphosphorylcholine (anti-PC) antibodies were measured by ELISA, as previously described (18). In brief, 96-well Maxisorp microtiter plates were coated with 20 μg/ml PC conjugated to BSA (Biosearch Technologies) overnight at 4°C. Plates were blocked with PBS/10% FCS and washed, and dilutions of serum in PBS/0.05% Tween 20 were added for 2 h. A positive control (pooled serum from mice immunization with Pneumovax II; Aventis-Pasteur MSD) was included on each plate. Plates were revealed with HRP-conjugated goat anti-mouse IgM and IgG.

CIA

The protocol followed to induce CIA in C57BL/6 mice was reported by Campbell et al. (34). CFA was prepared by grinding 100 mg heat-killed *Mycobacterium tuberculosis* (H37Ra; Difco Laboratories) in 20 ml of incomplete Freund's adjuvant (Sigma-Aldrich). An emulsion was formed by dissolving 2 mg/ml of chick CII (Sigma-Aldrich) overnight at 4°C in 10 mM of acetic acid and combining it with an equal volume of CFA. Mice were injected with 100 μl CII/CFA intradermally into the base of the tail. The same injection was repeated at day 21. Animals were assessed for redness and swelling of limbs, and a clinical score was allocated for each mouse two to three times per week for up to 86 d. The maximum score per mouse was 12. At the end of the experiment, the paws of the mice were removed, fixed, decalcified, and paraffin embedded. 5-μm sections were stained with hematoxylin and eosin and examined for histological changes of inflammation, pannus formation, and cartilage and bone damage. ELISA for antibody to CII was performed as previously described (34). HRP-conjugated goat anti-mouse IgM, IgG, IgG1, IgG2a, IgG2b, or IgG3 (SouthernBiotech) detection antibodies were used. Standard curves were constructed from the pooled sera of CII-hyperimmunized DBA/1 mice.

Proteinuria measurements

Urine protein levels were measured with Multistix 10 SG (Bayer) and scored as follows: 0 = negative; 0.5 = trace; 1 = 30 mg/dl; 2 = 100 mg/dl; 3 = 300 mg/dl; and 4 = 2,000 mg/dl or more.

ANA test

ANAs were examined using a commercial indirect fluorescent antibody assay (ANA Hep-2 slide; Cambridge Life Sciences) according to the manufacturer's

instructions. Mouse serum, starting at 1:40 in PBS, was titrated across the slide. Slides were incubated for 30 min at room temperature and washed twice with PBS. Alexa Fluor 488–conjugated rabbit anti–mouse IgG antibody was added to the wells at 1:150 in PBS for 30 min at room temperature. After two more washes, slides were mounted on coverslips for blinded fluorescence microscopic analysis.

Phagocytosis and infection studies with *S. pneumoniae*

For phagocytosis assays, *S. pneumoniae* type 2 (D39; provided by J. Brown, University College London, London, UK) was cultured to log phase in Todd–Hewitt broth with 0.5% yeast extract (Oxoid Ltd.), heat inactivated at 60°C for 1 h, and labeled with Alexa Fluor 488 (Invitrogen) (18, 56). Alexa Fluor 488–labeled *S. pneumoniae* were incubated in PBS or heat-inactivated immune serum at 37°C for 1 h before washing. Immune serum for opsonization was obtained from mice immunized with pneumovax (Aventis–Pasteur MSD). Serum-opsonized and nonopsonized Alexa Fluor 488–labeled pneumococci were added to plastic-adhered peritoneal macrophages at 37°C (and at 4°C as a control). Adhered macrophages were washed, harvested, and analyzed by flow cytometry. Peritoneal macrophages were identified by scatter characteristics and CD11b⁺ F4/80⁺ staining. The percentage and geometric mean fluorescence of Alexa Fluor 488⁺ macrophages were used as measures of phagocytosis. For infection studies, groups of 7–14 B–TG or M–TG mice and littermate NTG controls were inoculated intraperitoneally with 200 μ l PBS containing between 3×10^1 and 10^7 CFU *S. pneumoniae*, or intranasally with 50 μ l PBS containing 2×10^6 CFU *S. pneumoniae*. The development of disease was monitored as previously described (57). During observation, mice were scored by a blinded observer for the presence or absence of physical signs of progressive sepsis. Mice that became moribund were considered to have reached the end point of the experiment and were killed. Survival data were analyzed using Kaplan–Meier graphs and log-rank tests.

Online supplemental material

Fig. S1 shows transgene expression in two independent strains of B–TG mice. Fig. S2 demonstrates no transgene expression on CD11b⁺, CD11c⁺, or CD3⁺ cells in the B–TG line. Fig. S3 shows transgenic Fc γ RIIb expression in spleens of B–TG mice by immunohistology. Fig. S4 demonstrates decreased Fc-mediated proliferation in B–TG B cells, and quantification of pERK measurements shown in Fig. 2 C. Fig. S5 shows that transgene expression is restricted to macrophages in two strains of M–TG mice. Fig. S6 shows no transgene expression on dendritic cells and FDCs in M–TG mice. Fig. S7 shows T-independent and T-dependent immune responses in M–TG mice. Fig. S8 shows levels of anti-CII IgG isotypes during CIA. Fig. S9 shows levels of natural anti-PC antibody. Table S1 shows absolute cell numbers of various cell populations in B–TG and B–NTG mice. Table S2 shows absolute cell numbers of various cell populations in M–TG and M–NTG mice. Table S3 lists the primers used for the MRL/lpr backcross. Supplemental discussion discusses the transgenic expression profile in B–TG mice. Online supplemental material is available at <http://www.jem.org/cgi/content/full/jem.20072565/DC1>.

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