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N-glycan branching is required for development of mature B cells

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Abstract

Galectins have been implicated in inhibiting B cell receptor (BCR) signaling in mature B cells but promoting pre-BCR signaling during early development. Galectins bind to branched N-glycans attached to cell-surface glycoproteins to control the distribution, clustering, endocytosis and signaling of surface glycoproteins. During T cell development, N-glycan branching is required for positive selection of thymocytes, inhibiting both death by neglect and negative selection via enhanced surface retention of the CD4/CD8 co-receptors and limiting TCR clustering/signaling, respectively. The role of N-glycan branching in B cell development is unknown. Here we report that N-glycan branching is absolutely required for development of mature B cells in mice. Elimination of branched N-glycans in developing B cells via targeted deletion of N-acetylglucosaminyl transferase I (Mgat1) markedly reduced cellularity in the bone marrow and/or spleen, and inhibited maturation of pre-, immature and T2 transitional B cells. Branching deficiency markedly reduced surface expression of the pre-BCR/BCR co-receptor CD19 and promoted spontaneous death of pre-B cells and immature B cells *in vitro*. Death was rescued by low dose pre-BCR/BCR stimulation but exacerbated by high dose pre-BCR/BCR stimulation as well as anti-apoptotic Bcl_{xL} over-expression in pre-B cells. Branching deficiency also enhanced Nur77 induction, a marker of negative selection. Together these data suggest that as in T cells, N-glycan branching promotes positive selection of B cells by augmenting pre-BCR/BCR signaling via CD19 surface retention while limiting negative selection from excessive BCR engagement.

INTRODUCTION

T and B cells (or lymphocytes) undergo similar developmental events and checkpoints that involve the dynamic expression of transcription factors, response to extracellular environmental cues such as cytokines, and the somatic rearrangement, expression, and screening of their antigen receptors in order to generate a functionally diverse, self-tolerant lymphocyte repertoire (1–4). Abnormalities in lymphocyte development and selection give rise to congenital immunodeficiencies, leukemias/lymphomas, and autoimmune diseases (5). In the bone marrow, developing B cells rearrange immunoglobulin (Ig) μ heavy (H)-chain and light (L)-chain genes to express surface bound IgM that complexes with

immunoreceptor tyrosine-based activation motif (ITAM) containing Ig α and Ig β (CD79A and CD79B) signaling proteins to form the B cell receptor (BCR) on immature B cells (6–11). BCR binding to antigen directs immature B cell fate – when the BCR is engaged by self-antigen, immature B cells die by clonal deletion (negative selection by apoptosis) or undergo receptor editing (continuous L-chain rearrangement) to reduce/eliminate autoreactivity (12). Immature B cells continue selection through transitional developmental stages (T1 and T2) in the bone marrow and the spleen, but predominantly mature into follicular (FO) or marginal zone (MZ) B cells in the spleen and co-express surface bound IgM and IgD (13). Approximately 25% of developing B cells become mature B cells in the bone marrow (14). Unengaged BCRs induce tonic signaling to drive developmental progression and survival of immature B cells in a CD19 dependent manner (15). CD19 is a transmembrane glycoprotein first expressed in pro-B cells in the bone marrow and augments BCR signaling at multiple points during B cell development, maturation, and differentiation (16–20). Mice with targeted deletion of CD19 have greatly reduced B cell numbers, impaired B cell function, and defective immune responses (21, 22).

The branching of Asparagine (N)-linked glycans with N-acetylglucosamine by the N-acetylglucosaminyl transferases Mgat1, Mgat2, Mgat4a/b and Mgat5 sequentially increase production of ligands for the galectin family of sugar-binding proteins (23, 24). At the cell surface, interactions of galectins with branched N-glycans attached to glycoproteins generates a macromolecular lattice, thereby controlling receptor localization, mobility, clustering, and surface retention to impact cell function/differentiation and disease states (25–33). Our group demonstrated that N-glycan branching regulates thymic positive selection by defining the upper and lower bounds of TCR affinity for peptide-MHC (33). Using a T cell specific (Lck driven cre) knockout of the N-glycan branching enzyme Mgat1, we observed markedly reduced thymic and splenic T cell numbers. Mgat1 deletion enhanced thymocyte death by neglect via decreased CD4/CD8 co-receptor surface retention (i.e., enhanced CD4/CD8 endocytosis) and associated reduced Lck induced Erk signaling, which are important for augmenting low affinity TCR engagement. N-glycan deficient thymocytes simultaneously exhibited increased death by negative selection due to excessive Ca²⁺ flux driven by enhanced TCR clustering. Thus, during T cell development N-glycan branching provides a mechanism for decoupling CD4/CD8 co-receptor and TCR signaling to maintain the appropriate range of TCR signal intensity necessary for thymocyte positive selection and generation of functional circulating CD4⁺ and CD8⁺ T cells.

Various studies have demonstrated glycosylation to be important in B cell development, selection, and maturation. This includes the sialic acid-binding immunoglobulin-type lectin (Siglec) and B cell inhibitory co-receptor CD22 (34–37), the sialyltransferase ST6Gal1 (38), fucosylation (39, 40), and galectins (41–46). The role of N-glycan branching, however, has not been investigated. Here, we provide evidence that N-glycan branching promotes positive selection of B cells by enhancing low affinity BCR engagement via promoting CD19 surface levels while also reducing high affinity BCR engagement to prevent negative selection.

MATERIALS AND METHODS

Mice

Mgat1^{fl/fl} (006891), *Mgat2^{fl/fl}* (006892), *CD19-cre* (006785), *CD23-Cre* (028197), and *Bim^{-/-}* (004525) mice were obtained from Jackson Laboratory. Eμ-Bcl_{xL} mice were transferred to us by Dr. Brian Iritani from the Department of Comparative Medicine at the University of Washington, Seattle. *Mgat1^{fl/fl}/tetO-cre/ROSA-rtTA* mice were previously described (33). Inter-breeding generated all other mice. Mice used were 5–7 weeks old but otherwise selected randomly for experiments and approved by the Institutional Animal Care and Use Committee of the University of California, Irvine.

Flow cytometry

Flow cytometric analysis was performed as previously described (25, 33, 47). The fluorophore conjugated mouse specific antibodies from Thermo Fisher/eBioscience were CD43 (R2/60), IgM (II/41), B220 (RA3–6B2), CD19 (eBio1D3), CD23 (B3B4), and CD21 (8D9). For flow cytometric analysis of glycan expression, cells were stained with 2 μg/mL L-PHA-FITC or biotinylated L-PHA followed by streptavidin conjugated fluorophore (Vector Labs and Thermo Fisher Scientific, respectively). Fluorophore conjugated Annexin-V was from BD Pharmingen, and fluorophore conjugated Nur77 from Cell Signaling Technologies. Samples were acquired on the Attune NxT (Thermo Fisher Scientific) flow cytometer. Data analysis was performed using FlowJo software.

Cell culture and stimulation

For *in vitro* experiments, isolated bone marrow cells were cultured in RPMI-1640 media (Thermo Fisher Scientific) supplemented with 10% heat inactivated fetal bovine serum (VWR), 2 μM L-glutamine and 100 U/mL penicillin/streptomycin (Gibco), and 50 μM β-mercaptoethanol (Gibco). Cells were cultured at 2.5 × 10⁵ cells per well in a 96-well plate in triplicates without stimulation, or in the presence of functional grade goat anti-mouse IgM F(ab')₂ (polyclonal, Thermo Fisher Scientific) at various concentrations.

Statistical Analysis

Prism 8 software was used for all statistical testing. For comparison of 3 or more groups, we used the non-parametric Kruskal-Wallis test with false discovery rate correction (Benjamini, Krieger and Yekutieli) for multiple comparisons (Fig. 1e,f, 3d–g, 4b,c). The Mann-Whitney test was used when comparing only two groups, with 1-tailed tests in Fig. 2, 3c and 4a as we predicted *a priori* the direction of the effect of *Mgat1* deletion based on our previous T cell data (33) as well as the results of Kruskal-Wallis tests in Figure 1.

RESULTS

N-glycan branching is required for B cell generation

To investigate a role for N-glycan branching in B cell development, we generated B cell specific *Mgat1* and *Mgat2* deficient mice using *CD19-cre*. *Mgat1* deletion completely eliminates while *Mgat2* deletion reduces N-glycan branching (Supplemental Fig 1a). CD19 expression is initiated at the pro-B cell stage (16, 17). The plant lectin L-PHA (*Phaseolus*

vulgaris leukoagglutinin) binds to β 1,6GlcNAc-branched N-glycans generated by the sequential action of the Mgat1, Mgat2 and Mgat5 branching enzymes (Supplemental Fig 1a) (25, 31, 33). The loss of L-PHA binding directly identifies cells deleted for *Mgat1*, *Mgat2* or *Mgat5* (25, 31, 33). Therefore, we utilized L-PHA flow cytometry to assess deletion of *Mgat1* and *Mgat2* in the bone marrow and spleen of B220⁺ B cells (Fig. 1a). In *Mgat2^{fl/fl}/CD19-cre^{+/-}* mice, pro-B cells displayed a small population with reduced L-PHA binding (L-PHA^{int}) that became a larger population of L-PHA^{lo/-} cells while transitioning to the mature B cell stage and entering the periphery (Fig. 1b). In contrast, only a small proportion of L-PHA^{int} pre-B and immature B cells were observed in *Mgat1^{fl/fl}/CD19-cre^{+/-}* mice, with the population largely disappearing at the mature B cell stage and in peripheral splenic B cells (Fig. 1b). The lack of L-PHA^{lo/-} B cells in *Mgat1^{fl/fl}/CD19-cre^{+/-}* mice does not result from L-PHA binding to a glycan structure other than β 1,6GlcNAc-branched N-glycans, as disrupting their biosynthesis via deletion of *Mgat2* (i.e. *Mgat2^{fl/fl}/CD19-cre^{+/-}* mice) results in a large population of L-PHA^{lo/-} B cells. As CD19-cre has a deletion efficiency of only ~75–80% in bone marrow (48), the population of L-PHA⁺ B cells in *Mgat1^{fl/fl}/CD19-cre^{+/-}* mice likely results from inefficient CD19-cre mediated excision of *Mgat1* and continued differentiation/expansion of this L-PHA⁺ cell population (Fig. 1b).

To further investigate these results, we generated *Mgat1^{fl/fl}/CD23-cre⁺* and *Mgat2^{fl/fl}/CD23-cre⁺* mice where CD23-cre will delete *Mgat1* and *Mgat2* in immature transitional 2 (T2) B220⁺ B cells. These cells remain sensitive to BCR regulated selection and apoptosis (49, 50). CD23-cre driven deletion phenocopied the results of CD19-cre, with *Mgat1^{fl/fl}/CD23-cre⁺* displaying L-PHA^{int} but not L-PHA^{lo/-} B220⁺ splenic B cells while ~90% of B220⁺ splenic B cells from *Mgat2^{fl/fl}/CD23-cre⁺* mice were L-PHA^{lo/-} cells (Fig. 1c).

To directly examine whether *Mgat1* deletion impacts mature B cells, we examined doxycycline treated *Mgat1^{fl/fl}/tetO-cre/ROSA-rtTA* adult mice (33). These mice displayed a large population of L-PHA^{lo/-} B220⁺ splenic B cells (Fig. 1d). This confirms that L-PHA detects *Mgat1* deleted B cells and indicates that *Mgat1* deletion specifically impacts survival of developing B cells through to the T2 stage in the periphery, while having little impact on mature peripheral B cell survival.

Consistent with the conclusion that the complete but not partial loss of N-glycan branching leads to death of developing B cells, total cell number in the bone marrow and/or spleen as well as the number of B220⁺ B cells were markedly reduced in *Mgat1^{fl/fl}/CD19-cre^{+/-}* and *Mgat1^{fl/fl}/CD23-cre⁺* but not *Mgat2^{fl/fl}/CD19-cre^{+/-}* or *Mgat2^{fl/fl}/CD23-cre⁺* mice (Fig. 1e,f). Together, these results indicate that partial N-glycan branching levels are permissible for B cell development, while complete elimination prevents production of mature B cells in the bone marrow and spleen without impacting survival of peripheral mature B cell. This parallels our observations in T cells, where *Mgat1* deletion in thymocytes completely blocked development of mature T cells, while deletion in peripheral mature T cells did not impact cell death (33).

N-glycan branching drives bone marrow development and splenic maturation of B cells

Next, we examined at what stage loss of N-glycan branching impacts B cell development and maturation. Cell surface markers and gating strategy used to identify specific B cell

populations were as previously described (16, 51) and is depicted in Fig. 1a. We focused on *Mgat1^{ff}/CD19-cre^{+/-}* and *Mgat1^{ff}/CD23-cre⁺* mice as they displayed significant loss of B cells in the bone marrow and/or spleen. In the bone marrow of *Mgat1^{ff}/CD19-cre^{+/-}* mice, cellularity at each developmental stage revealed little difference at the pro-, pre-, and immature stages, but a drastic loss in the number of mature B cells in (Fig. 2a). In the spleen, transitional T1, transitional T2, follicular (FO) and marginal zone (MZ) B cells are similarly reduced by *Mgat1* deficiency (Fig. 2b). In *Mgat1^{ff}/CD23-cre⁺* mice, the number of transitional T1 B cells was unchanged but there was a marked reduction in transitional T2, follicular (FO) and marginal zone (MZ) B cells (Fig. 2c), the former consistent with CD23 expression beginning at the T2 stage. These data suggest that branching is essential for B cell development, particularly during the transition from immature and T2 to mature B cells.

Inhibiting programmed cell death fails to rescue B cell death from the loss of N-glycan branching

In T cells, we previously observed that the death of developing thymocytes induced by *Mgat1* deletion was rescued by over-expression of anti-apoptotic Bcl_{xL} as well as targeted deletion of pro-apoptotic Bim (33). Therefore we investigated whether manipulating these programmed cell death pathways (i.e., the mitochondrial apoptotic pathway (52)) could similarly rescue death of N-glycan branching deficient B cells. We generated *Mgat1^{ff}/CD19-cre^{+/-}/Eμ-Bcl_{xL}* to induce B cell specific anti-apoptotic Bcl_{xL} overexpression, as well as *Mgat1^{ff}/CD19-cre^{+/-}/Bim^{-/-}* for deletion of pro-apoptotic Bim. As *Mgat1* deficiency primarily altered the transition from immature to mature B cells, we focused on changes in mature B cells in the bone marrow. Neither *Mgat1^{ff}/CD19-cre^{+/-}/Eμ-Bcl_{xL}* nor *Mgat1^{ff}/CD19-cre^{+/-}/Bim^{-/-}* displayed an increase in the proportion of mature B cells (Fig. 2d). Thus, Bcl_{xL} overexpression, and potentially Bim deletion, fails to overcome the developmental block of N-glycan branching deficient immature B cells into mature B cells.

N-glycan branching promotes CD19 associated tonic signaling to B cell death by neglect

Pre- and immature B cells are subject to positive selection, with CD19 playing an important role in both pre-BCR signaling in pre-B cells (53) and BCR signaling in immature B cells (15). Only intermediate (i.e., tonic) pre-BCR/BCR signaling allows continued development into mature B cells. As N-glycan branching regulates surface retention of cell surface glycoproteins, we assessed whether *Mgat1* deficiency alters positive selection by reducing surface expression of CD19. Indeed, L-PHA^{int}B220⁺ B cells from the bone marrow of *Mgat1^{ff}/CD19-cre^{+/-}* mice displayed a marked reduction in CD19 surface expression (Fig. 3a-c). To assess whether N-glycan branching deficiency drives death of developing B cells, we cultured bone marrow cells for one day in media without stimulation and assessed cell death by Annexin V labeling. Both L-PHA^{int}B220⁺IgM⁻ B cells and L-PHA^{int}B220⁺IgM⁺ B cells from *Mgat1^{ff}/CD19-cre^{+/-}* mice displayed significant increases in Annexin V binding relative to control cells (Fig. 3d,e), consistent with increased sensitivity to death from reduced CD19 dependent pre-BCR/BCR tonic signaling. Bcl_{xL} over-expression in L-PHA^{int}B220⁺IgM⁻ cells from *Mgat1^{ff}/CD19-cre^{+/-}* mice reversed the increase in cell death (Fig. 3f). However, in L-PHA^{int}B220⁺IgM⁺ B cells Bcl_{xL} overexpression displayed only a marginal and non-significant impact on cell death (Fig. 3e). To investigate whether increased death was from reduced pre-BCR/BCR tonic signaling, we assessed whether minimally

enhancing pre-BCR/BCR signaling would reduce the death induced by N-glycan branching deficiency. Indeed, low doses of a polyclonal anti-IgM F(ab')₂ antibody, which should activate both pre-BCR and BCR via the common heavy chain, partially rescued L-PHA^{int}B220⁺IgM⁻ and L-PHA^{int}B220⁺IgM⁺ B cells from death (Fig. 3f,g). Taken together, these data suggest that N-glycan branching promotes CD19 surface expression and associated pre-BCR/BCR tonic signaling to prevent death by neglect of developing B cells.

N-glycan branching inhibits negative selection of developing B cells

Low level stimulation of BCR reduced but did not fully rescue death of N-glycan branching deficient developing B cells. As developing B cells are also subject to death by negative selection via strong BCR signaling from encounter with high affinity self-antigen, this suggests that N-glycan branching may also regulate negative selection in immature B cells. Nur77 is a transcription factor that is expressed at higher levels in B cells subject to excessive antigen encounter and eventual deletion by negative selection (54–56). *Ex vivo* L-PHA^{int}B220⁺IgM⁺ B cells displayed greater Nur77 levels compared to control (Fig 4a). This suggests N-glycan branching deficiency enhances BCR engagement and negative selection *in vivo*. To further support this hypothesis, we examined whether providing high BCR signaling further enhances the death of B cells deficient in N-glycan branching. Indeed, BCR stimulation with high doses of anti-IgM F(ab')₂ enhanced cell death of L-PHA^{int}B220⁺IgM⁺ greater than L-PHA^{int}B220⁺IgM⁻ cells, as evidenced by Annexin V labeling (Fig. 4b,c). These data suggest that N-glycan branching also inhibits negative selection of immature B cells.

DISCUSSION

Here we report that N-glycan branching is required for the development but not survival of mature B cells. Loss of N-glycan branching impacts the survival of pre-B cells, immature B cells and T2 transitional peripheral B cells but not mature B cells, where only the former cells are subject to positive selection. The effects of N-glycan branching during B cell positive selection is similar to what we have observed in T cells (33), with N-glycan branching appearing to be similarly required to maintain the boundaries of pre-BCR/BCR signaling thresholds to permit positive selection by inhibiting both death by neglect and negative selection. Mechanistically, N-glycan branching is required for CD19 surface expression in developing B cells. CD19 is necessary for low affinity pre-BCR/BCR engagement to maintain tonic BCR signaling and prevent death by neglect (15). The mechanism by which N-glycan branching inhibits death by negative selection is less clear, but may be similar to that in T cells, namely limiting antigen induced BCR clustering to prevent excessive signaling and death by negative selection.

CD19 is coupled to both pre-BCR and BCR signaling mediated apoptosis pathways (15, 53). For immature B cells, lack of CD19 results in loss of basal phosphoinositide 3-kinase (PI3K) signaling, continued recombination-activating genes (RAG) expression, and L-chain receptor editing, thus inhibiting positive selection (15). Phosphatase and tensin homologue (PTEN) opposes PI3K activity; the absence of PTEN activity reverses the effects of CD19 loss to promote immature B cell positive selection (57, 58). Like CD19, BCR deletion

Future research in delineating the role of N-glycan branching in B cell tolerance may provide new insight on the role of B cells in diseases such as systemic lupus erythematosus, rheumatoid arthritis, and multiple sclerosis.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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KEY POINTS

1. N-glycan branching is required for generation of mature B cells.
2. N-glycan branching inhibits development of pre-, immature and T2 B cells,
3. N-glycan branching promotes CD19 surface expression to inhibit death by neglect.

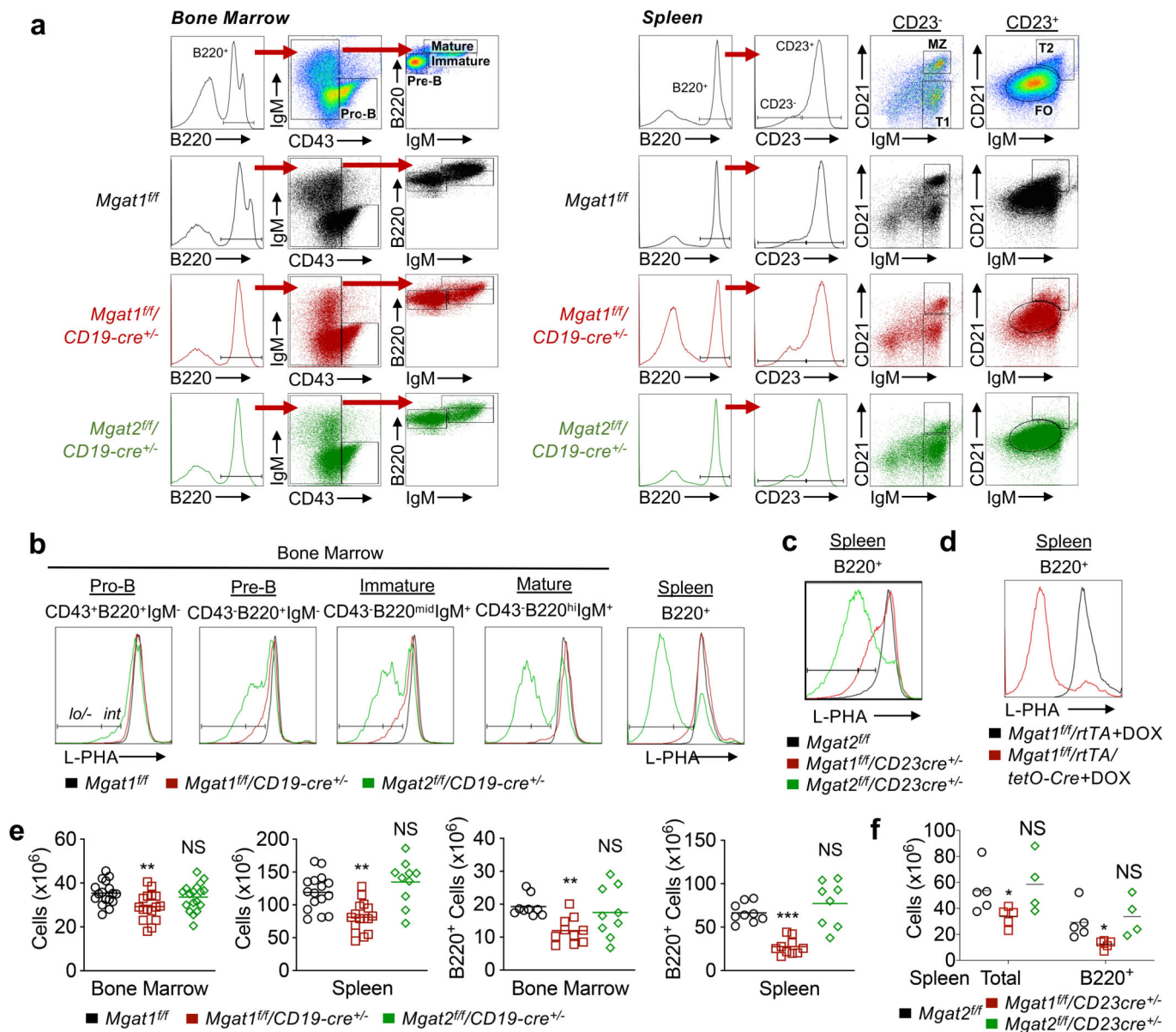


Figure 1. Late B cell development requires N-glycan branching.

(a-d) Flow cytometric analysis with indicated markers on bone marrow B cell subsets and/or splenic B cells from mice of the indicated genotypes. (e,f) Total cellularity or B220⁺ cell numbers (calculated by multiplying percentage of B220⁺ cells acquired by flow cytometry by total cellularity) of bone marrow and/or spleens from mice of the indicated genotypes. Each dot represents one mouse. Kruskal-Wallis test with false discovery rate correction (Benjamini, Krieger and Yekutieli) for multiple comparisons (e,f). Bars indicate mean. NS, not significant;

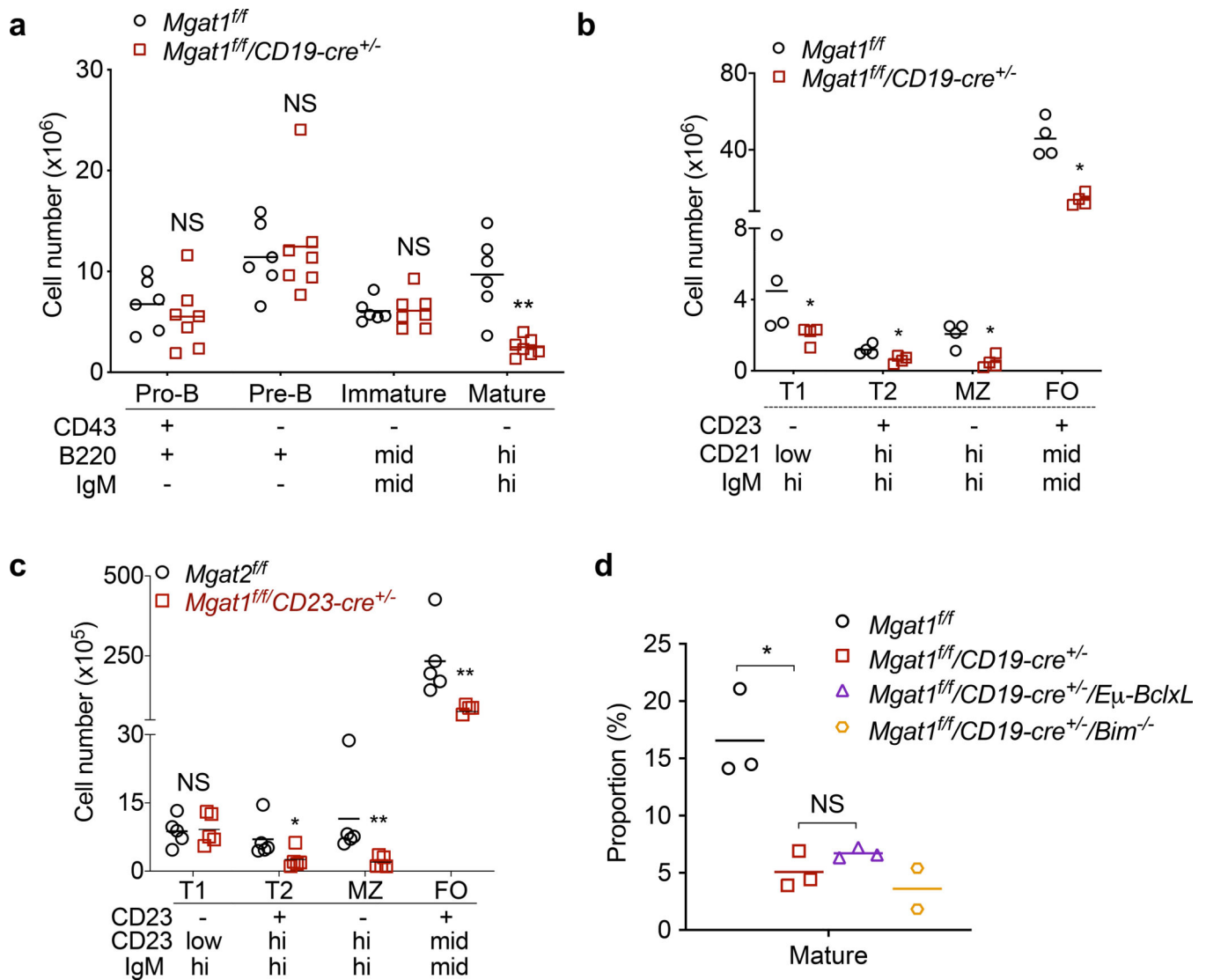


Figure 2. N-glycan branching promotes transition of immature to mature B cells in bone marrow and B cell maturation in the spleen.

(a-d) Flow cytometric analysis of total number of B cell subtypes in the bone marrow (a) and spleen (b,c) using indicated markers, and proportion of mature B cells in spleen (d) from mice of the indicated genotypes. Each dot represents one mouse, unpaired one-tailed Mann-Whitney *t*-test. Bars indicate mean. NS, not significant; ***P* < 0.01; **P* < 0.05.

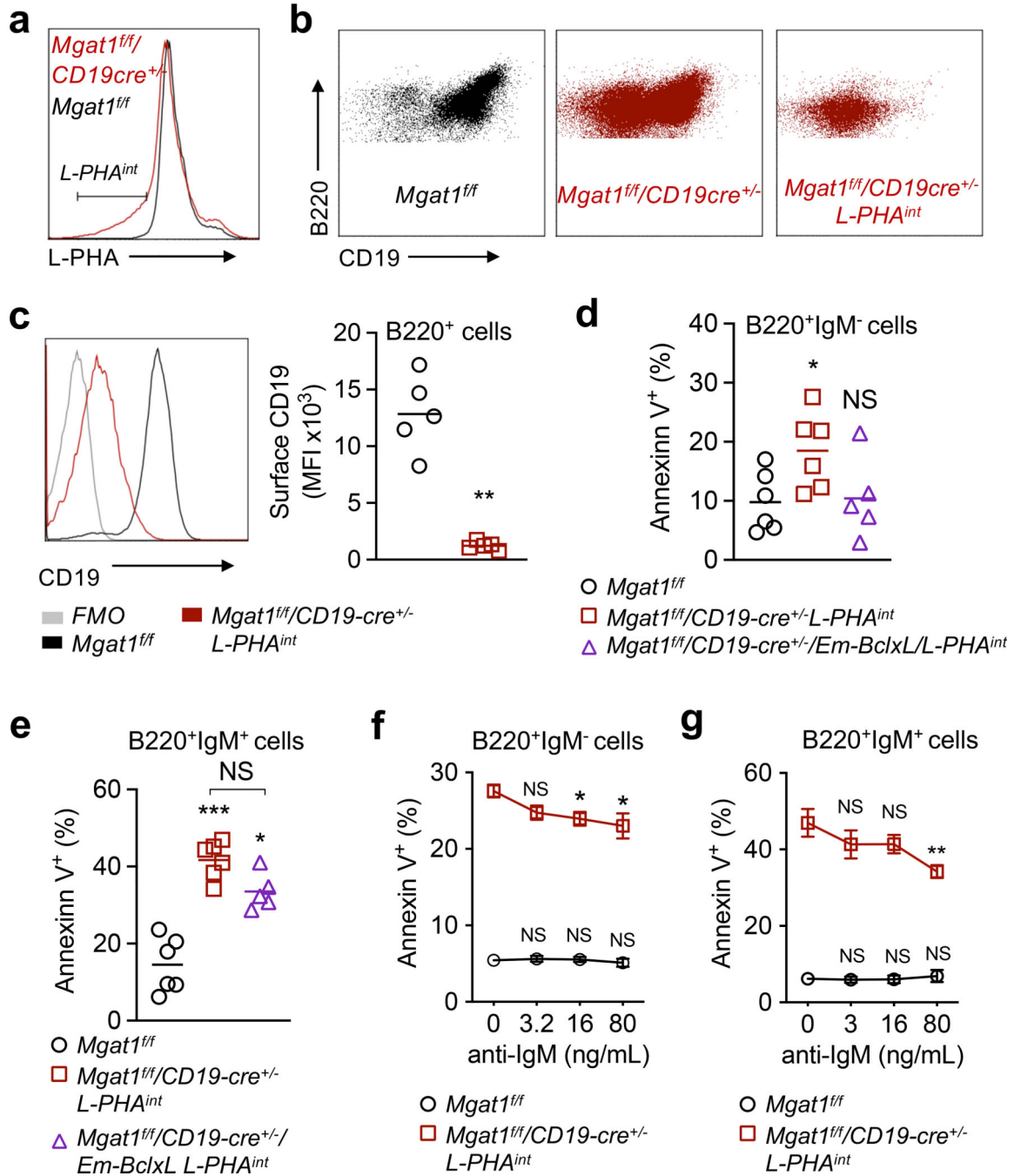


Figure 3. N-glycan branching promotes immature B cell positive selection via CD19 surface expression.

(a-c) Flow cytometric analysis of *ex vivo* B220⁺ bone marrow B cells for the L-PHA^{int} gate (a) and CD19 surface levels (b,c). Each dot represents one mouse, unpaired one-tailed Mann-Whitney *t*-test (c). ***P* < 0.01. (d-g) Annexin V binding to bone marrow B cells cultured for 1 day without BCR stimulation (d,e) and with low doses of anti-IgM F(ab')₂ (f,g), gated on B220⁺IgM⁻ (d,f) and B220⁺IgM⁺ (e,g) cells. Cells from control mice are gated on all B cells, while all others are gated on the L-PHA^{int} population as indicated below

each panel. n = 3. Each dot represents one mouse (d,e). Kruskal-Wallis test with false discovery rate correction (Benjamini, Krieger and Yekutieli) for multiple comparisons (d-g), with comparison to zero anti-IgM (f,g). Error bars indicate mean \pm s.e.m. *** $P < 0.001$; * $P < 0.05$.

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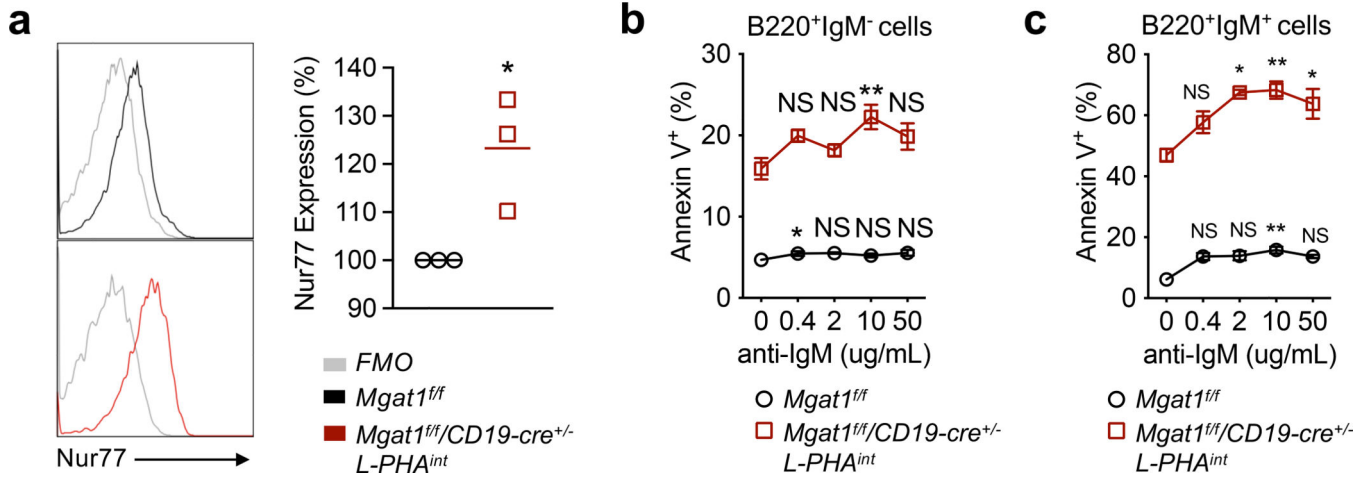


Figure 4. N-glycan branching limits immature B cell negative selection by limiting BCR engagement.

(a) Flow cytometric analysis of *ex vivo* B220⁺IgM⁺ bone marrow B cells for intracellular staining of Nur77. Each dot represents one mouse of the indicated genotypes; unpaired one-tailed Mann-Whitney *t*-test. **P* = 0.05. (b-c) Annexin V binding to bone marrow B cells following a 1-day culture with high doses of anti-IgM F(ab')₂, gated on B220⁺IgM⁻ (b) and B220⁺IgM⁺ (c) cells. n = 3. Kruskal-Wallis test with false discovery rate correction (Benjamini, Krieger and Yekutieli method) for multiple comparisons, with comparison to zero anti-IgM for each genotype. Error bars indicate mean ± s.e.m. **P* < 0.05.