

Supplemental material

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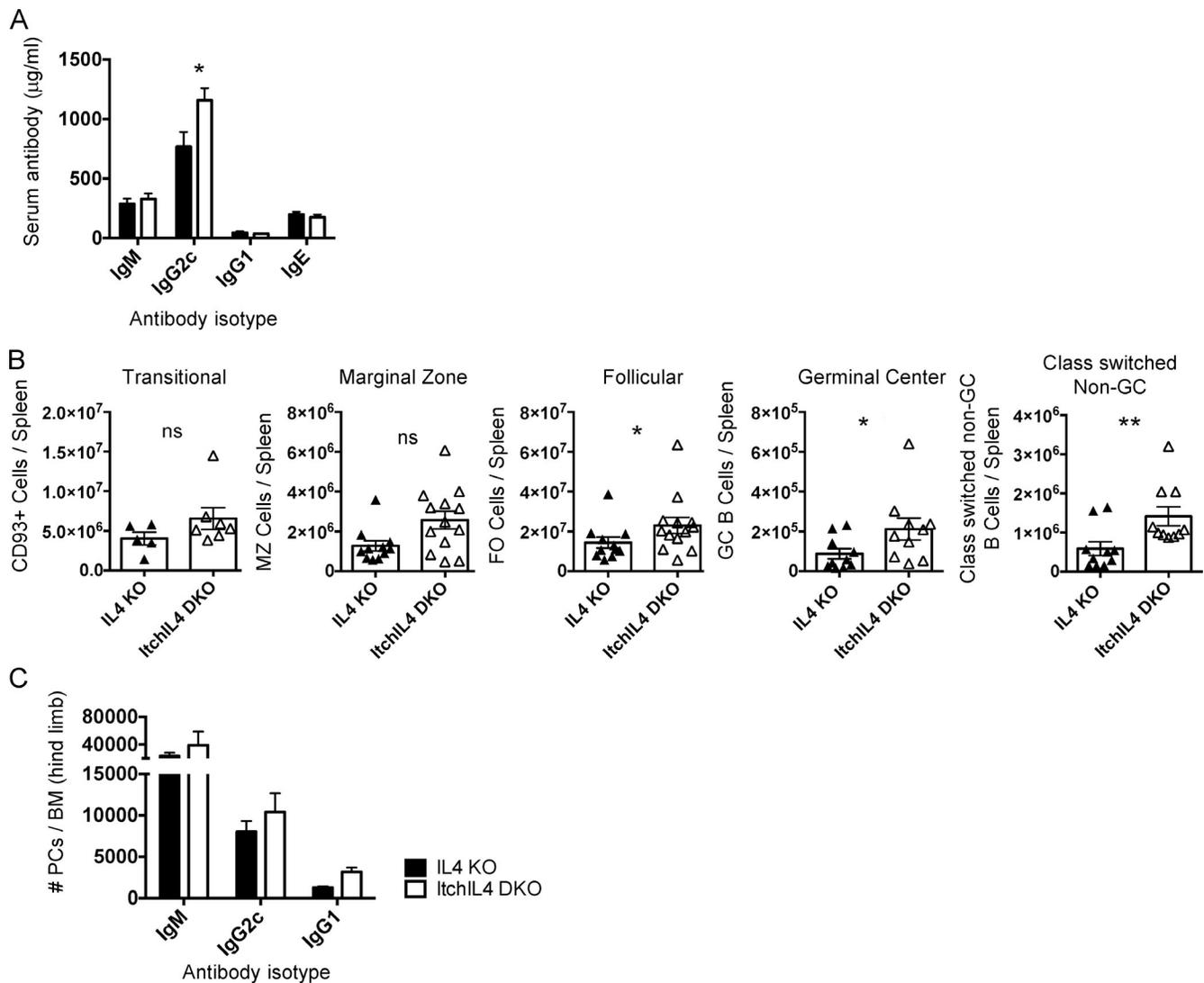


Figure S1. **Itch limits antibody and activated B cells in the absence of IL-4.** (A) Quantity of total antibody isotypes in serum of IL4 KO and Itch/IL4 DKO mice was determined by ELISA (mice between 8–12 wk old, for IgM $n = 6$ or 7 , for IgG1 $n = 12$ or 13 , for IgG1 $n = 4$, multiple t tests, Holm–Sidak correction). (B) Pre-immune and activated B cell subsets in spleen were quantified by flow cytometry. Cells were gated on live singlets, and then transitional B cells were CD19⁺CD93⁺, FO B cells were CD19⁺CD93⁺CD23⁺, and marginal zone B cells were CD19⁺CD23^{low}CD21⁺. For GC and class-switched non-GC B cells, cells were first gated on IgD⁺CD4⁺CD8⁺F480⁺GRI⁺CD19⁺IgM⁺, and then GC B cells were GL7⁺CD38⁺, and class-switched non-GC B cells were GL7⁺ ($n = 8–10$, compiled from three independent experiments, Mann–Whitney test). (C) BM IgM, IgG1, and IgG2c-secreting PCs were enumerated by ELISPOT ($n = 3$, two-way ANOVA with Sidak post-test, IL4 KO mice were on B6.SJL background [CD45.1]). Error bars indicate SEM. *, $P < 0.05$; **, $P < 0.01$; ns, not significant.

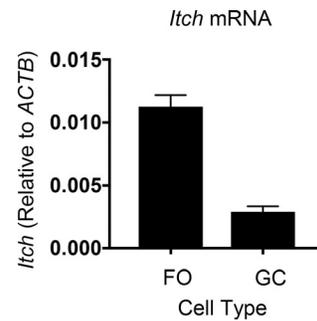


Figure S2. **Itch is expressed in naive and GC B cells.** RNA was extracted from purified splenic FO and GC B cells directly ex vivo. Reverse transcription and real-time PCR for *Itch* and β actin transcripts were performed ($n = 3$, two independent experiments; error bars indicate SEM; WT mice were C57Bl/6 [CD45.2]).

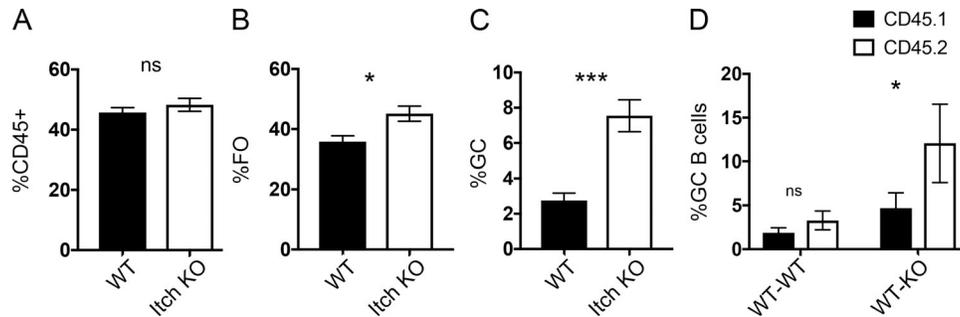


Figure S3. **Itch limits GC B cells in a B cell intrinsic manner.** (A–C) Mixed chimeras from Fig. 2 are analyzed in a different way. (A) Gated on live singlets, then the percent CD45.1⁺ or CD45.2⁺ is shown. (B) Gated on CD45.1⁺ or CD45.2⁺, then the percent IgD⁺ cells is shown ($n = 6$, two independent experiments, unpaired *t* test). (C) Cells were gated on CD45.1⁺IgD⁻ or CD45.2⁺IgD⁻, then the percent GC B cells is shown ($n = 6$, two independent experiments, unpaired *t* test). (D) Mixed chimeras were generated by injecting mixtures of equal numbers of T cell-depleted BM cells from WT (B6.SJL CD45.1) and WT (C57Bl/6 CD45.2) mice or WT (B6.SJL CD45.1) and Itch KO (CD45.2) mice into sublethally irradiated (400 rad) Rag KO mice. After 8 wk, spleens were harvested, and the cells were gated on CD45.1⁺CD19⁺ or CD45.2⁺CD19⁺, and the percent GC B cells is shown ($n = 4$ or 5, two independent experiments, two-way ANOVA with Sidak post-test). Error bars indicate SEM. *, $P < 0.05$; ***, $P < 0.001$; ns, not significant.

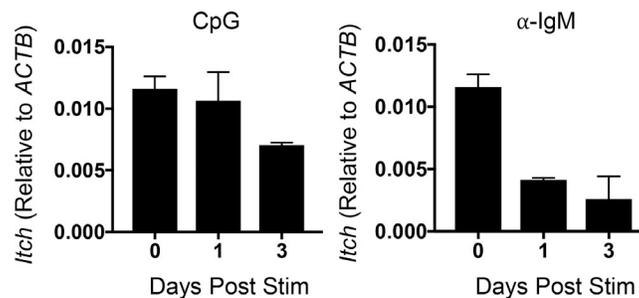


Figure S4. **Itch expression in B cells after in vitro stimulation.** FO B cells were purified from Itch-sufficient IL4 KO mice and cultured for 3 d with or without CpG or anti-IgM stimulation. RNA was extracted, cDNA was made by reverse transcription, and real-time PCR for *Itch* and *actb* transcripts was performed (for CpG, $n = 3$ or 4, two independent experiments, and for IgM, $n = 2$, one experiment. Error bars indicate SEM. IL4 KO mice were B6.SJL [CD45.1]). Post stim, post stimulation.

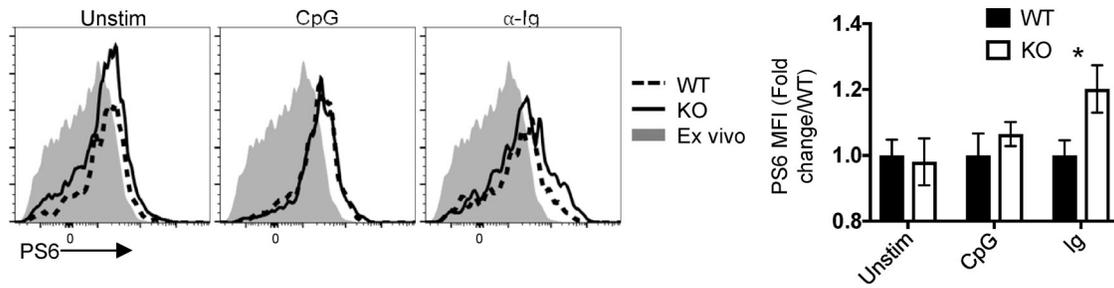


Figure S5. **Itch limits mTORC1 activity in spontaneous GC B cells.** Total B cells were isolated from WT or Itch KO spleens using negative selection. Cells were cultured in B cell media for 1 h, and were stimulated with 1 μ M CpG or 10 μ M anti-Ig. Cells were stained for GC markers and P-S6, then analyzed by flow cytometry. GC B cells were gated on CD19⁺IgD⁻IgM⁻GL7⁺CD38⁻, and MFI of P-S6 was determined. Fold change indicates MFI/average MFI of WT for each independent experiment ($n = 6$, three independent experiments, multiple t tests; error bars indicate SEM). *, $P < 0.05$. Unstim, unstimulated.