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Supplemental Information

Aryl Hydrocarbon Receptor Interacting

Protein Maintains Germinal Center

B Cells through Suppression of BCL6 Degradation

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Figure S1. RAG1 Cre deletion of Aip and phenotype of GC B cells; related to Figure 1

(A) qPCR and (B) Western blot analysis of Aip from GC B cells from $Aip^{fl/fl}$ Cre⁺ and $Aip^{fl/fl}$ Cre⁻ mice. Data are plotted as mean \pm SEM. (C) gating strategy to identify and analyze GC B cells. (D) size (FSC), (E) BCL2 expression, (F) proliferation of GC B cells from Cre⁻ and Cre⁺ mice.



Figure S2. Increased B cells in the periphery of *Aip^{fl/fl}*Cre⁺ mice; related to Figure 2

(A) Percentage of B cells in spleen, lymph nodes and circulating blood of B220⁺ IgD⁺ B cells from $Aip^{fl/fl}$ Cre⁺ mice compared to WT mice. Data shown as percentage \pm SEM. Total number of B220⁺ IgD⁺ B cells (**B**) and total number of splenocytes (**C**). Total number of CD3⁺ T cells (**D**). Results are from 4-5 independent experiments, 2-3 mice per experimental group.



Figure S3. No difference in Marginal Zone or Follicular B cells or B cell development in *Aip^{fl/fl}* Cre⁺ mice; related to Figure 2

(A) Marginal zone (MZ) (CD21⁺ CD23⁻) and Follicular (FO) CD21⁻ CD23⁺) B cells were analysed by flow cytometry. (B) Expression of IgM or IgD on B cells from WT and *Aip^{fl/fl}* Cre⁺ mice. (C) Bone marrow from the femurs of Aip^{fl/fl} Cre⁺ and Cre⁻ mice and were analyzed for different stages of B cell development by flow cytometry. B220⁺ cells were stained with CD43 and CD24. The CD24⁺ cells were divided into different subpopulations depending upon their expression of IgM and IgD. (D) Bone marrow from the femurs of *Aip^{fl/fl}* Cre⁺ and Cre⁻ mice and were analyzed for B220^{hi} IgM^{+/-} B cells by flow cytometry. There was a significant reduction in the percentage of B220^{hi} IgM⁺ B cells from *Aip^{fl/fl}* Cre⁺ mice compared to Cre⁻ mice. (E) serum from Cre- and Cre+ mice were examined by ELISA for immunoglobulin isotypes. B cells from Cre- and Cre+ were isolated and stimulated in vitro in the presence of anti-IgM, anti-CD40 and IL-4 and the percentage of proliferating cells (blasts) (F), IgG1 expression (G) and CD138 (H) examined at 48, 72 and 96 hours post stimulation by flow cytometry. 2-3 independent experiments, Results are from 2-3 mice per experimental group.



Figure S4. *AHR*^{∆TB} mice have normal GC B cells and immune responses towards SRBCs; related to Figure 2

(A) $AHR^{\Delta TB}$ and Cre⁻ mice were immunized with SRBCs and the percentage of GC B cells (B220⁺ IgD⁻ GL-7⁺) examined. (B) BCL6 expression determined from $AHR^{-/-}$ and $AHR^{+/+}$ IgD⁻ B cells. (C) DZ (CXCR4⁺) and LZ (CD86⁺) GC B cells from $AHR^{\Delta TB}$ and Cre⁻ B cells and the DZ/LZ ratio. (D) Serum from $AHR^{\Delta TB}$ and Cre⁻ mice immunized with SRBCs were serially diluted and incubated with SRBCs and the amount of IgG bound determined by flow cytometry. Results are from two independent experiments, 2-3 mice per experimental group. Data are plotted as mean \pm SEM.



Figure S5. *Aip^{fl/fl}* Cre⁺ mice decreased extra-follicular immune responses; related to Figure 1

 $Aip^{f/f!}$ Cre⁺ and Cre⁻ mice were immunized with NP-Ficoll and analyzed 14 days later NP-specific serum (A) IgM, (B) IgG and (C) IgG3 analyzed by ELISA. Data are plotted as mean \pm SEM. (D-E) Percentage of NP-specific plasmablasts (B220^{+/-} CD138⁺) analyzed by flow cytometry. Representative FACS plots from at least 2 independent experiments, 2-3 mice per experimental group.



Figure S6. *Aip^{fl/fl}* Cre⁺ mice have no differences in ERK or SYK phosphorylation following BCR stimulation; related to Figure 3

IgD⁻ B cells from *Aip^{fl/fl}* Cre⁺ and Cre⁻ mice were stimulated with anti-IgM (10µg/ml) and examined for the expression of phosphorylated (**A**) ERK and (**B**) SYK from 5 to 60 minutes post-stimulation. Grey histograms show expression at time 0 (T0). Data expressed as percentage from time 0. Results are from two independent experiments with 1-2 mice per experimental group.