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Supplemental Information

The persistence of interleukin-6

is regulated by a blood buffer system

derived from dendritic cells

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4 h post injection

Follicle

Follicle

Figure S1. Zbtb46-DTR Chimeras lack cDC but take up antigen from the blood and show cDCdependent TCR-independent control over the antibody responses to the protein antigens gp120 and ovalbumin, related to Figure 1. (A) Representative flow plots in untreated and DT-treated chimeras. The cDCs (B220⁻/CD3⁻/CD11c^{hi}/MHCII⁺) were reduced upon addition of DT (presented are mean \pm SD, n= 3 animals per group; **P<0.002, Student's T test). Naïve B cells (CD3⁻/B220⁺/IgM⁺/IgD⁺) are also presented in a side-by-side comparison. (B) Capture and uptake of HA-APC following intravenous injection. Spleen follicles were demarcated by CD169 metallophilic macrophages (bar = 50 μ m). The spleen sections were imaged at one and four hours post-injection and the amount of antigen taken into the follicles (intensity per spleen follicle at four hours) within DT treated versus untreated Zbtb46-DTR chimeras was quantified by imaging software (10 replicates within each of n=3 DT-treated or n=3 untreated chimeras). (C) Serum IgM versus IgG responses against HIV gp120 in *Tcra*^{-/-} versus WT mice, and Zbtb46-DTR chimeras receiving or not receiving DT. (D) Serum IgM versus IgG responses against ovalbumin in *Tcra*^{-/-} versus WT, and Zbtb46-DTR chimeras receiving or not receiving DT. (E) Serum IgM versus IgG responses to HA in WT versus *Ifnar1*^{-/-} mice. The dilution curves (presented as mean \pm SD, n=5 animals per group), were quantified by endpoint dilution (*P<0.02, **P<0.03, ***P<0.0001, Student's T test).





Figure S2. B cell, T cell, DC, and macrophage phenotypes from spleen are comparable between WT, cDC-*Irf5*^{-/-}, and cDC-*Il6ra*^{-/-} animals, related to Figure 2. Their representative flow plots and gating schemes are presented along with a corresponding quantification of proportions of these immune cell types (mean \pm SD, n= 5 animals per genotype).



Figure S3. IL6R measured in bone marrow chimeras and in Batf3-/- mice, related to Figure 2. (A) Concentration of circulating sIL6R in 100% Zbtb46-DTR chimeras or 50% Zbtb46-DTR +50% MHCII deficient (homozygous $H2^{\overline{dlAbl-Ea}}$), each \pm DT. Presented is the mean \pm SD where ***P<0.001, Student's T-test. (B) Surface expression of IL6R in cDC1 and cDC2 within WT, Batf3-/- (in which cDC1 is largely absent), and cDC-*Il6ra*^{-/-} (*P<0.05, ***P<0.001, ANOVA with Tukey's Test). Total cDC is made up by 27.7 \pm 2.54% cDC1 and 72.0 \pm 2.50% cDC2 (see also Figure S2). (C) Serum sIL6R was measured in WT, $Batf3^{-/-}$, and cDC-*Il6ra*^{-/-} genotypes (mean \pm SD where *P<0.05, Student's T test). (D) The modest reduction in serum sIL6R seen in the Batf3^{-/-} genotype does not impact IgM or IgG responses to HA, as measured in WT vs Batf3^{-/-} mice.



Figure S4. Addition of sIL6R can rescue dysregulated IL-6 capture in vivo, related to Figure 4. (Ai, Bi, Ci) sIL6R concentration in cDC-*Il6ra*^{-/-} versus mice of the same genotype that received AAV-sIL6R (three weeks post delivery). (Aii) Animals were then injected with intravenously with bacterial lipid A to promote IL-6 release. The circulating concentration of IL-6 was measured at days 3, 7 and 14 (***P<0.001, ANOVA, with Tukey's Test). (Bii) Animals were infected with NC99 influenza virus to promote IL-6 release. The circulating concentration of IL-6 was measured at days 3, 7 and 14 (***P<0.001, ANOVA, with Tukey's Test). NC99 infects and propagates within mice but does not cause disease. (Cii) Animals were challenged with a lethal dose of PR8 influenza virus and morbidly was indexed by measuring weight loss over the next four days. The rate of decline (slope) was compared using F Tests. If body weights were 80% or less (horizontal bar) the mice were euthanized.