

Figure S1. Positive correlation between plasma free immune complexes and anti-apoferritin IgG levels in CSS induced in CfH^{-/-} mice chimeric for the indicated bone marrow (**a**) and in those without BM transfers (**b**).

The lines fit by least squares analysis were:

(a) ICs = 17.3 + 0.743 x anti-apoferritin, $R = 0.71$, $P = 0.002$;

(b) ICs = 4.8 + 0.759 x anti-apoferritin, $R = 0.73$, $P = 0.001$.

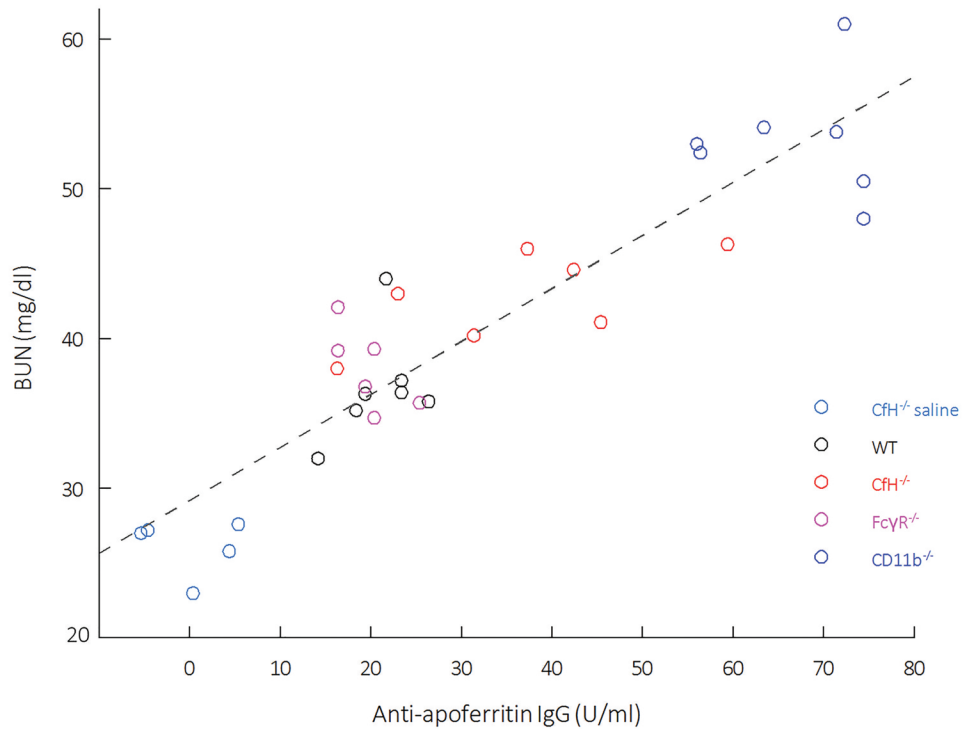


Figure S2. Positive correlation between measured BUN values and anti-apoferritin IgG levels. The line fit by least squares analysis was $BUN = 29.3 + (35.4 \times \text{anti-apoferritin})$. $R = 0.91$, $P < 0.001$.

The other significant correlations among disease measures were:
 $BUN = 20.9 + (0.47 \times \text{urinary albumin})$. $R = 0.97$, $P < 0.001$;
 $BUN = 26.8 + (8.16 \times \text{GN score})$. $R = 0.86$, $P = 0.002$; and,
 $\text{Urinary albumin} = 15.3 + (16.1 \times \text{GN score})$. $R = 0.81$, $P = 0.004$.

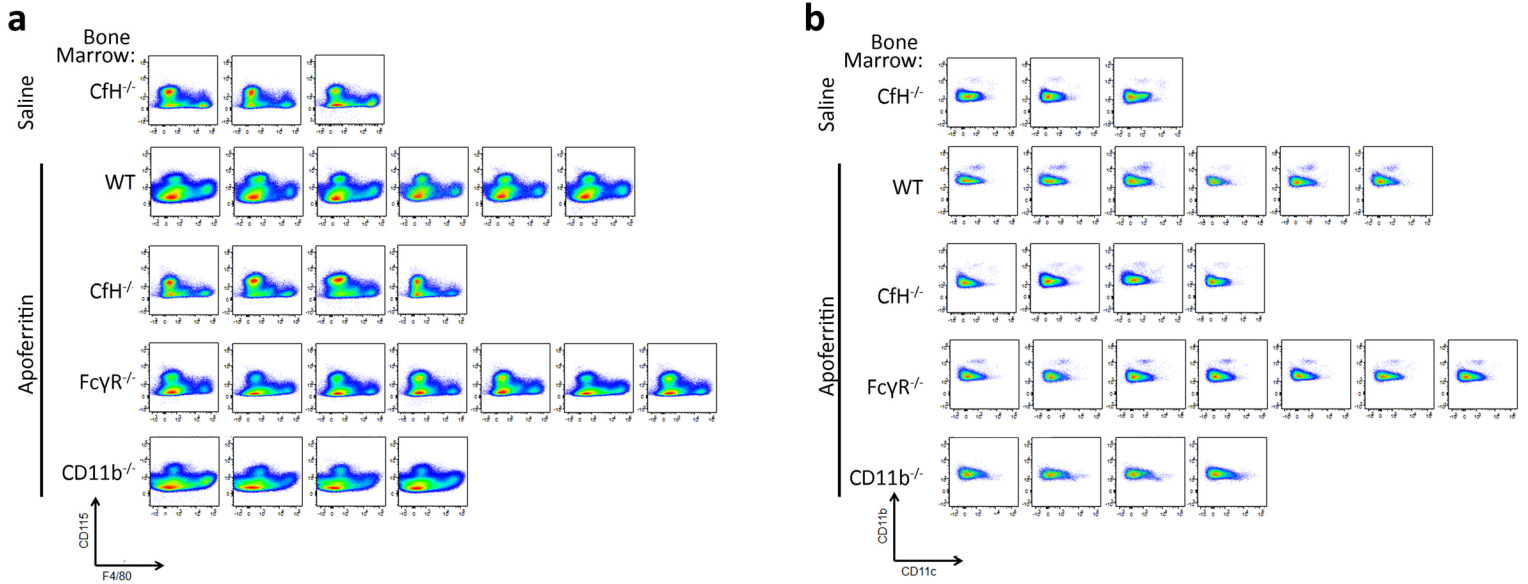


Figure S3. Analysis of CD3⁻CD19⁻ cells by flow cytometry. Cells were from kidneys of CfH^{-/-} mice chimeric for wildtype (WT), CfH^{-/-}, FcγR^{-/-}, and CD11b^{-/-} bone marrow immunized for 5 weeks with apoferritin, or saline as control. (a) Data for CD115 (y-axis) and F4/80 (x-axis). (b) CD115⁺ cells were analyzed for CD11b (y-axis) and CD11c (x-axis).

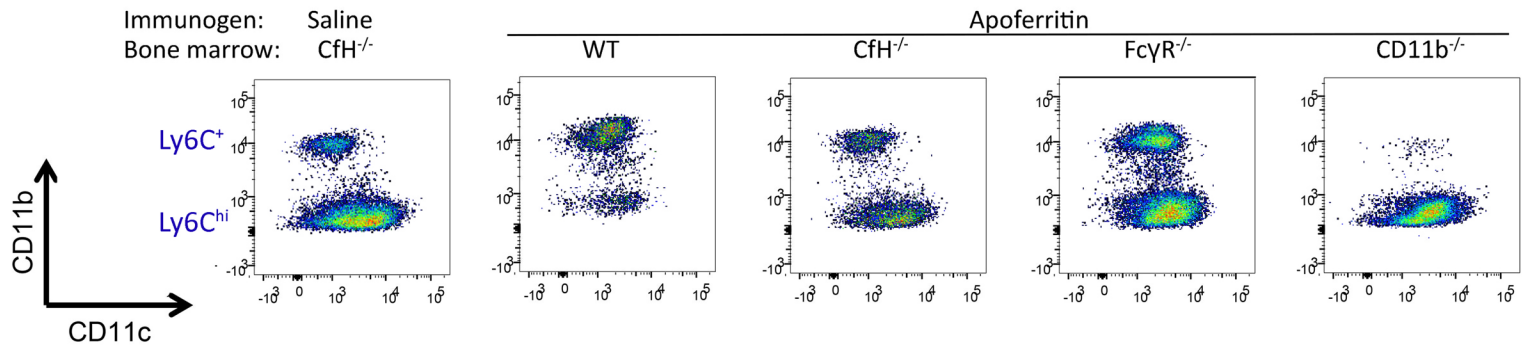


Figure S4. Analysis of F4/80⁺Ly6C⁺CCR2⁻ cells by flow cytometry. Cells were from kidneys of CfH^{-/-} mice chimeric for wildtype (WT), CfH^{-/-}, FcγR^{-/-}, and CD11b^{-/-} bone marrow immunized for 5 weeks with apoferritin, or saline as control. Representative data for CD11b (y-axis) and CD11c (x-axis) are shown. Ly6C^{hi} cells were CD11b⁻, while Ly6C⁺ cells were CD11b⁺, except in the CD11b^{-/-} chimeras.

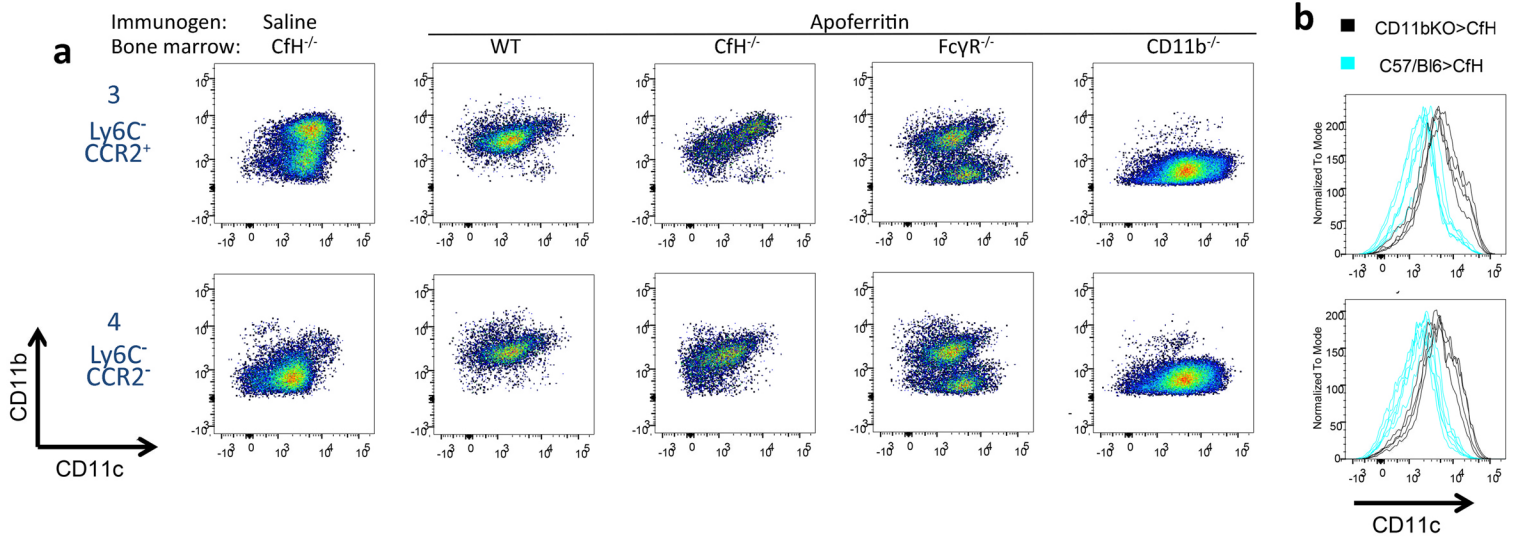


Figure S5. Analysis of F4/80⁺Ly6C⁻ cells for CD11b and CD11c by flow cytometry. Cells were from kidneys of Cfh^{-/-} mice chimeric for wildtype (WT), Cfh^{-/-}, FcγR^{-/-}, and CD11b^{-/-} bone marrow immunized for 5 weeks with apoferritin, or saline as control. CCR2⁺ and CCR2⁻ cells were analyzed separately; these are from quadrants 3 and 4, respectively, from Figure 4. (a) Representative data for CD11b (y-axis) and CD11c (x-axis) are shown. (b) CD11c expression is shown in histograms for all wildtype and CD11b^{-/-} chimeras.