## **Supplemental Data**

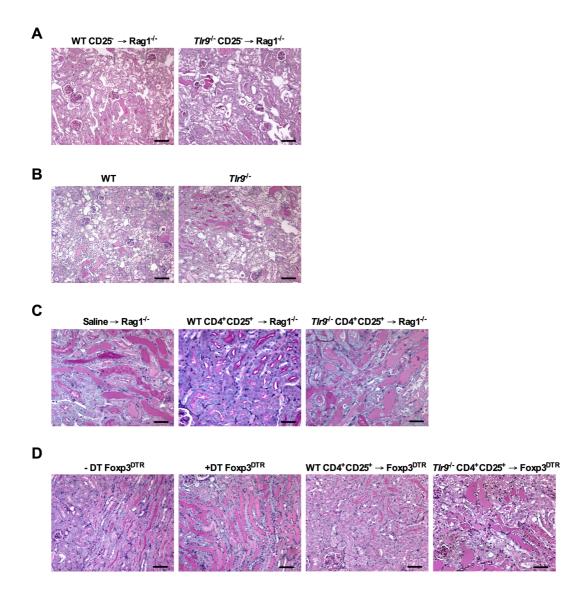
Supplemental Table 1. Gene and protein expression of chemokine receptors on CD4<sup>+</sup>CD25<sup>+</sup> cells from naïve WT and *Tlr9*<sup>-/-</sup> mice.

	Gene expression WT vs. <i>Tlr9</i> <sup>-/-</sup> CD4 <sup>+</sup> CD25 <sup>+</sup> cells (mRNA, RQ values)	Cell surface expression WT vs. <i>Tlr9</i> <sup>-/-</sup> (Flow cytometry, % of CD4 <sup>+</sup> CD25 <sup>+</sup> cells
CCR4	1.02±0.12 vs. 1.51±0.36, NS	10.1±1.17 vs. 9.51±0.93 <sup>a</sup> , NS
CCR6	1.00±0.05 vs. 1.38±0.07**	74.7±3.24 vs. 66.7±5.28 <sup>a</sup> , NS
CCR7	1.00±0.04 vs. 2.38±0.63, NS	98.9±0.18 vs. 99.1±0.18 <sup>b</sup> , NS
CXCR3	1.02±0.10 vs. 2.37±0.66, NS	34.9±1.92 vs. 37.9±1.12 <sup>b</sup> , NS

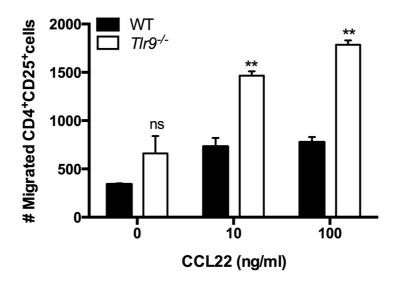
The left column shows mRNA gene expression data of chemokine receptors important in Treg recruitment to sites of inflammation on sorted  $CD4^+CD25^+$  cells from naïve WT and  $Tlr9^{-/-}$  spleens. n=4 in each group. Gene of interest is relative to 18S and presented as relative quantification (RQ) to WT  $CD4^+CD25^+$  cells. The right column shows flow cytometry data of cell surface expression of chemokine receptors on  $CD4^+CD25^+$  cells from the spleen or blood in naïve WT and  $Tlr9^{-/-}$  mice. n=5 in each group. Values represent mean  $\pm$  SEM (unpaired Student's t test). WT, wild-type, NS, non significant. \*\*, P < 0.01.

<sup>&</sup>lt;sup>a</sup> CD4<sup>+</sup>CD25<sup>+</sup> cells in the blood were investigated

<sup>&</sup>lt;sup>b</sup> CD4<sup>+</sup>CD25<sup>+</sup> cells in the spleen were investigated



**Supplemental Figure S1. Representative photomicrographs of PAS-stained kidney sections from reconsitution and depletion studies after cisplatin administration in Figure 2.** (A) Kidney sections after cisplatin administration showed equivalent tubular damage with tubular cast formation and matrix expansion in both reconstituted groups. Scale bar, 100 μm. (B) Kidney sections showed a severe but equivalent pattern and intensity of renal injury in CD25<sup>+</sup> cell-depleted WT and *Tlr9*<sup>-/-</sup> mice Scale bar, 100 μm. (C) Representative photomicrographs demonstrate more tubular injury in *Rag1*<sup>-/-</sup> mice reconstituted with *Tlr9*<sup>-/-</sup> CD4<sup>+</sup>CD25<sup>+</sup> regulatory cells and saline injection after cisplatin administration. Scale bar, 50 μm. (D) Kidney sections showing greater renal tubular damage in Foxp3<sup>DTR</sup> mice reconstituted with *Tlr9*<sup>-/-</sup> CD4<sup>+</sup>CD25<sup>+</sup> cells and control mice receiving diphtheria toxin (DT) injections compared to mice reconstituted WT CD4<sup>+</sup>CD25<sup>+</sup> cells and control mice not treated with DT. Scale bar, 100 μm.



Supplemental Figure S2.  $Tlr9^{-/-}CD4^+CD25^+$  T cells do not have reduced migration toward the inflammatory chemokine CCL22. The chemotactic response of WT and  $Tlr9^{-/-}CD4^+CD25^+$  T cells to increasing concentration of CCL22 was examined in an *in vitro* chemotaxis assay. The number of migrating cells was measured by flow cytometry acquisition of a fixed number of beads. Data represents mean  $\pm$  SEM (Two-way ANOVA). NS, not significant; \*\*, P < 0.01.