

Supplementary Material

Figure S1. Malat1 ASO activity (ISH) and distribution (IHC) in the kidney. Mice were treated with PBS or 100 mg/kg Malat1 ASO for 4 wks and kidney Malat1 RNA expression was evaluated in (A) PBS-treated mice or (B) Malat1 ASO-treated mice by ISH. (C) The distribution of ASO accumulation by IHC using an anti-ASO Ab. Black arrows denote interstitial sites demonstrating Malat1 ASO activity (B) and accumulation (C). Scale bar = 50 μ m

Figure S2. Tolerability and activity evaluation of the lead CD40 ASO in inflammation-naïve mice. The lead CD40 ASO was evaluated in C57BL/6 mice after four weeks of SC administration of 50 and 100 mg/kg/wk. Basic measures of tolerability (body and spleen weights, plasma bilirubin, BUN and ALT) were assessed and CD40 RT-PCR was performed in the liver and kidney.

Figure S3. Dose responsive reduction of CD40-dependent inflammation with CD40 ASO treatments and leukocyte mRNA marker expression following CD40 mAb administration. (A) To evaluate dose-responsive CD40 ASO activity following CD40-dependent inflammation, C57BL/6 mice were given 40, 15 or 5 mg/kg ASO treatments on days 0 and 7, followed by 25 μ g of the activating CD40 mAb on day 10, kidneys were harvested on day 11 and CD40, CCL5 and IL12p40 mRNA levels were measured by RT-PCR. (B) mRNA markers of specific leukocyte populations (F4/80, CD19 and CD3d) were analyzed by RT-PCR in control mice and mice administered the CD40 mAb 24 hrs prior to sacrifice.

Figure S4. CD40 mRNA expression in blood PBMC cells 6 weeks post BMT. Verification of bone marrow reconstitution was performed by RT-PCR using two different CD40 primer probe sets that either amplify CD40 mRNA in WT and CD40 KO leukocytes (PPSet #3483) or only in WT

leukocytes (PPSet #1280). The six groups shown were treated with CD40 ASO and/or CD40 Ab as described in Figure 3f.

Table S1. CD40 ASO treatment effects on kidney T cell and macrophage markers in DOX nephropathy.

Table S2. Primer and probe sequences used for RT-PCR.

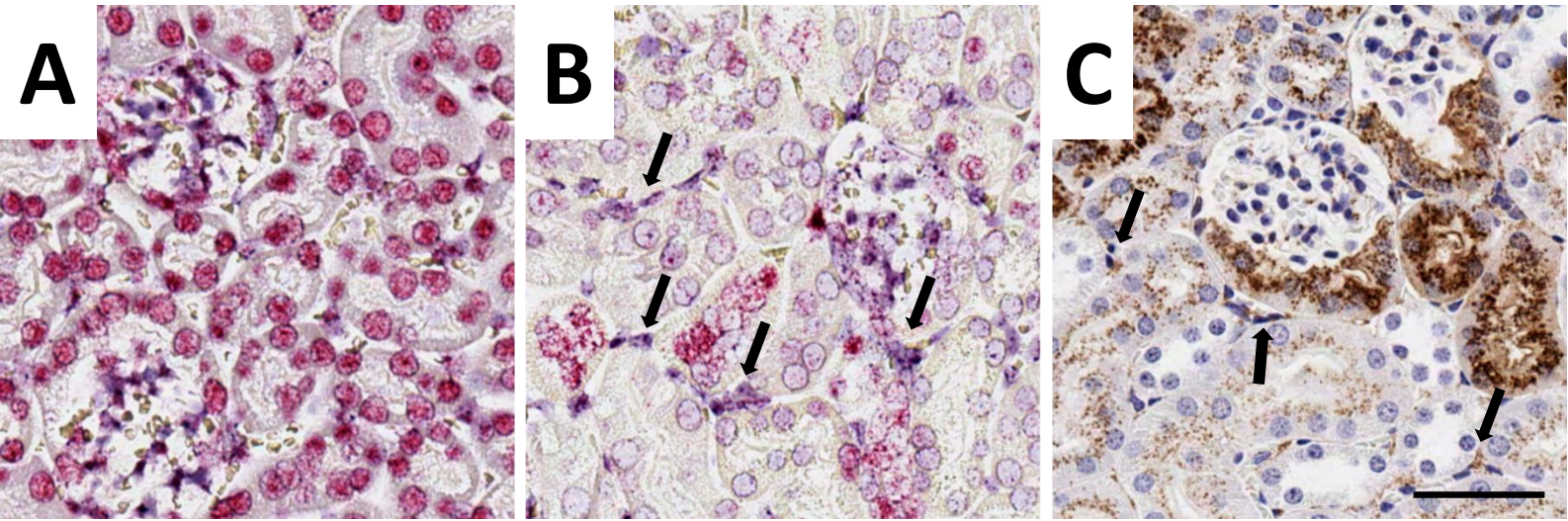


Figure S1

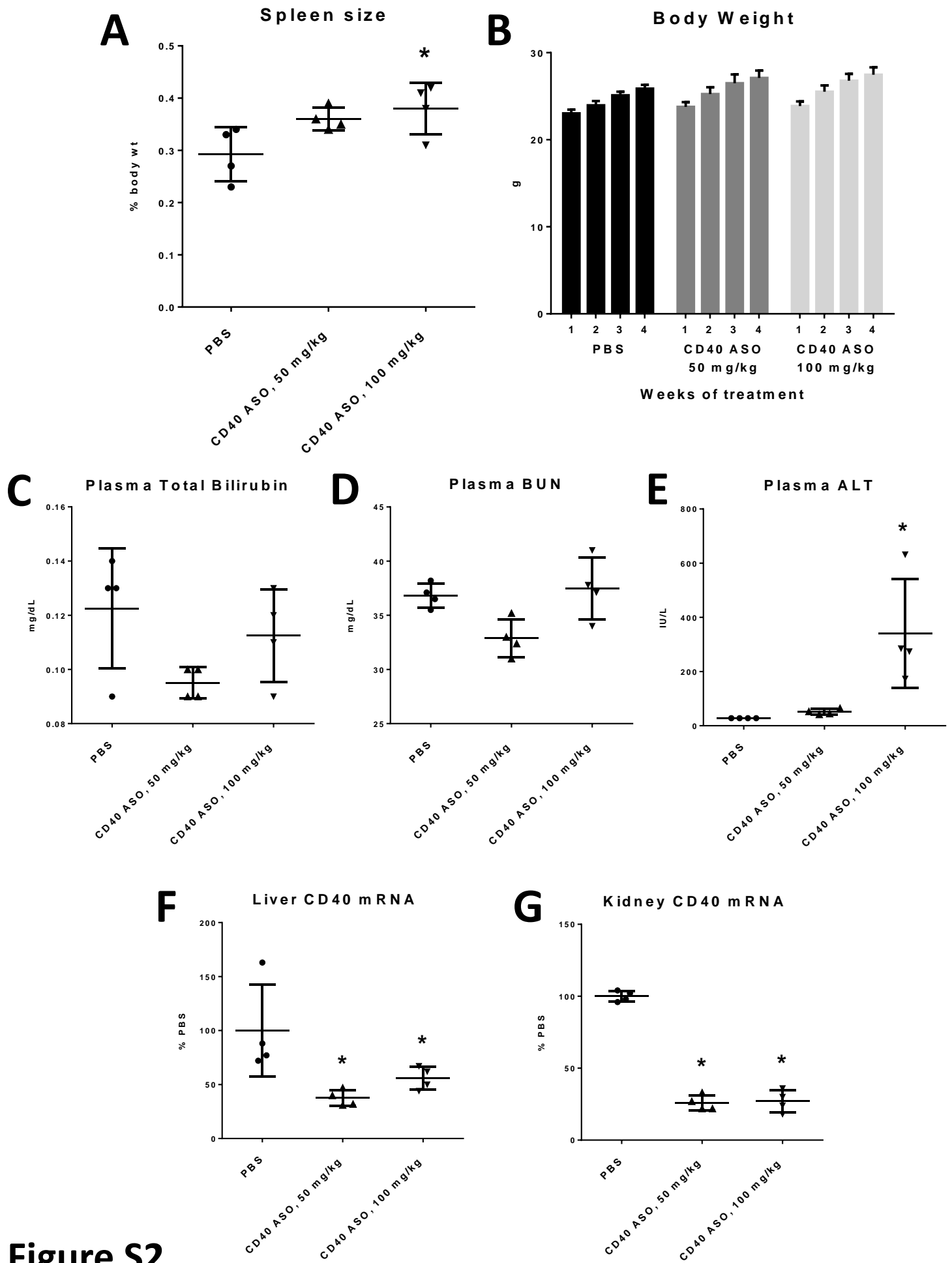
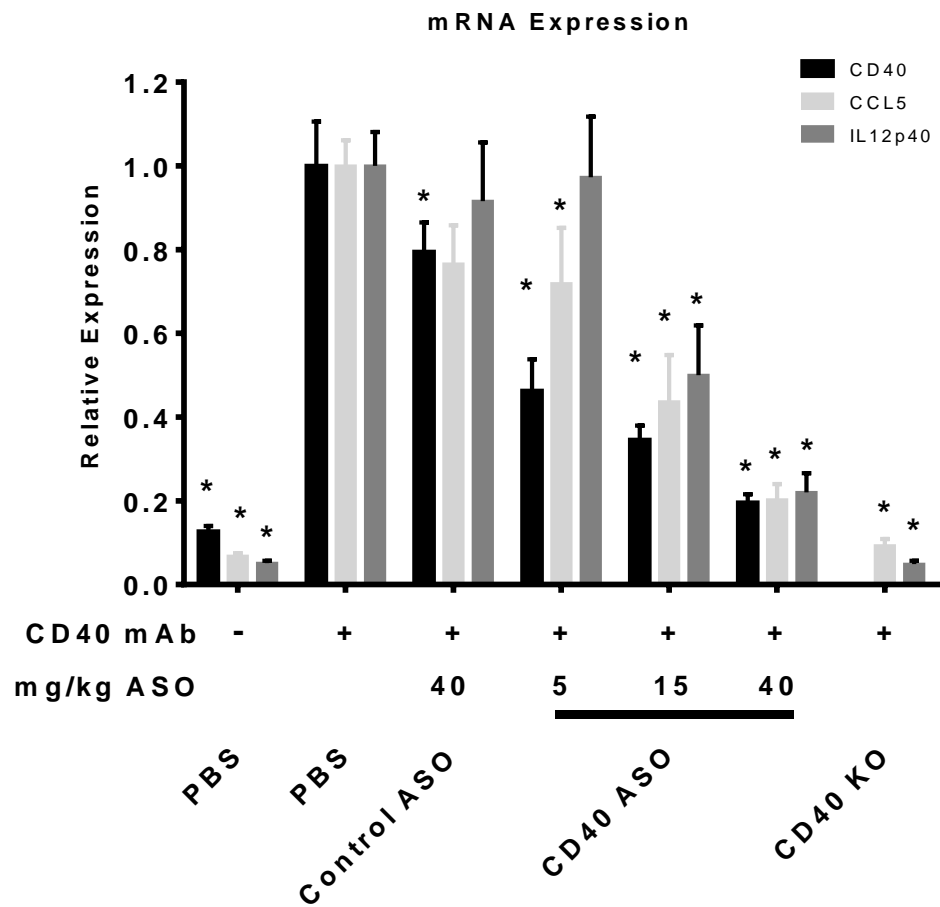
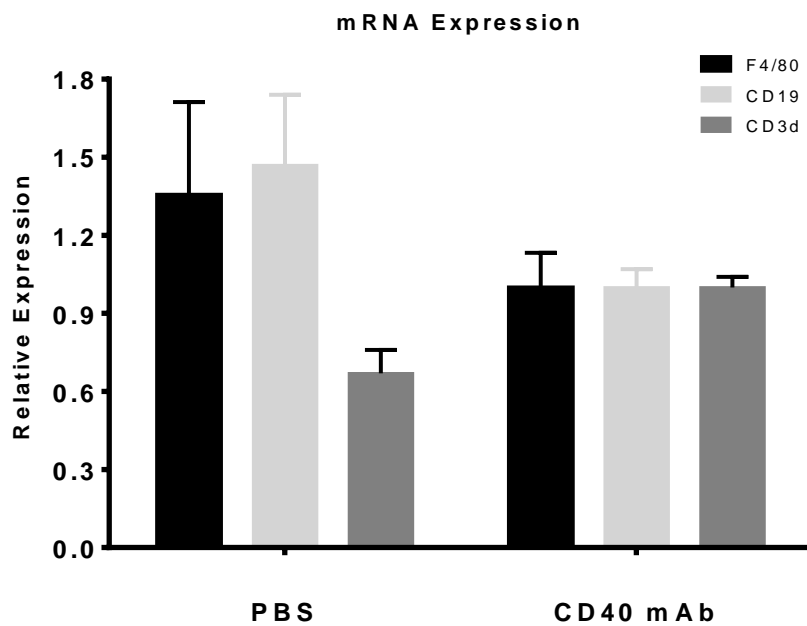


Figure S2

A**B****Figure S3**

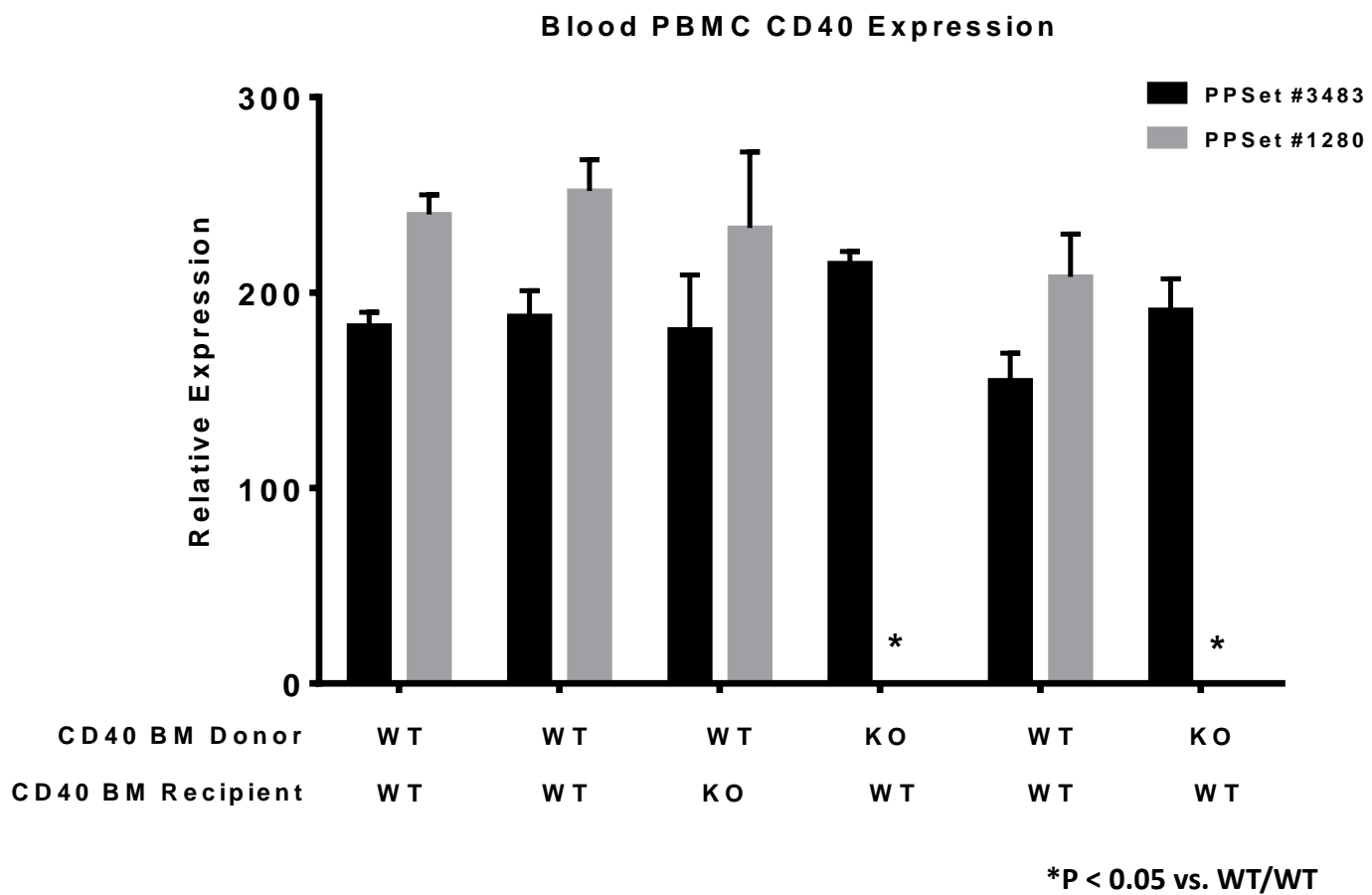


Figure S4

Fold-change mRNA levels \pm SEM (§ except CD3d IHC), relative to PBS Dox animals

	Dox				No Dox
	PBS	Con ASO-25	CD40 ASO-12.5	CD40 ASO-25	PBS
CD3d	1.00 \pm 0.18	0.86 \pm 0.11	0.87 \pm 0.11	0.61 \pm 0.04*	0.56 \pm 0.04*
CD4	1.00 \pm 0.12	0.86 \pm 0.14	0.90 \pm 0.14	0.57 \pm 0.08*	0.36 \pm 0.03*
CD8b	1.00 \pm 0.13	0.90 \pm 0.15	0.90 \pm 0.15	0.56 \pm 0.08*	0.25 \pm 0.01*
IL-2	1.00 \pm 0.12	0.58 \pm 0.11*	0.60 \pm 0.11*	0.34 \pm 0.04*	0.24 \pm 0.05*
CD3d IHC§	40 \pm 23	37 \pm 23	35 \pm 24	24 \pm 12	24 \pm 8
CD68	1.00 \pm 0.18	0.99 \pm 0.13	1.26 \pm 0.17	1.10 \pm 0.10	0.34 \pm 0.02*
IL-12	1.00 \pm 0.12	0.75 \pm 0.14	0.91 \pm 0.08	0.91 \pm 0.11	0.38 \pm 0.03*
TNF- α	1.00 \pm 0.17	0.69 \pm 0.10	0.93 \pm 0.08	0.80 \pm 0.07	0.24 \pm 0.01*
TGF- β	1.00 \pm 0.08	0.94 \pm 0.08	1.22 \pm 0.06*	1.00 \pm 0.05	0.67 \pm 0.06*
CTGF	1.00 \pm 0.06	0.97 \pm 0.11	0.69 \pm 0.05*	0.73 \pm 0.05*	0.46 \pm 0.04*
CD206	1.00 \pm 0.16	0.95 \pm 0.11	0.88 \pm 0.08	0.86 \pm 0.15	0.45 \pm 0.04*

*P < 0.05 vs. PBS (DOX)

Table S1

Sequence information for the probe and primers used to generate the RT-PCR data

*** Sequence information not available from manufacturer (Life Technologies). Unique identifier listed.**

Target mRNA	Forward Primer	Reverse Primer	Probe
CCL5	GCAGTCGTGTTTGTCACTCGAA	GATGTATTCTTGAACCCACTTCTCTC	AACCGCCAAGTGTGTGCCAACCC
CD206	GAATACACAGCACTAGCGTCTTAACAC	GGGCTTACGTGGTTGTTTCTAGA	TAGGTCTCATGTCAACCCTGCAGATTCAAG
CD3d	GAGGTTTCTCAAAGTGCCATAGC	CGGAAGGAAACGTCTTTATTGG	ACGGCGCCTTCCCCTGTGATC
CD4	Mm00442754*		
CD40 (#3483)	CCATCTAGGGCAGTGTGTTACG	CCTGGCTGGCACAATCAC	ACAAACAGTACCTCCACGATGGCCAGTG
CD40 (#1280, to genotype the CD40 KO)	AGCCAGGAAGCCGACTGA	CCCTTGATTGGGTTACAGTGT	ACTCAGGCGAATTC
CD40L	Mm00441911*		
CD68	TTGGGAACACTACAGTGGGC	CGGATTGAATTGGGCTTG	AACGGCTCCAGCCTTGTGTTGAG
CD8b	Mm00438116*		
COL1A1	TGGATTCCCGTTCGAGTACG	TCAGCTGGATAGCGACATC	AAGCGAGGGCTCCGACCCGA
E Selectin	Mm01310197*		
F4/80	GGCATTGCCAGATTTTC	CGGTTGAGCAGACAGTGAATGA	CCAGATTGGCCCCTTGCAAGC
GAPDH	GGCAAATCAACGGCACAGT	GGGTCTCGCTCCTGGAAGAT	AAGGCCGAGAATGGGAAGCTTGTGATC
ICAM1	CCGCAGGTCCAATTCACACT	CAGAGCGGCAGAGCAAAG	CAGCTCGGAGGATCACAACGAAGCT
IL-12p40 (IL12/IL23)	GCCAGTACACCTGCCACAAA	GACCAAATTCATTTTCTTCTTG	AGGCGAGACTCTGAGCCACTCACATCTG
IL-2	Mm00434256*		
MCP1	AGTTGACCCGTAATCTGAAGCTAA	CACACTGGTCACTCCTACAGAAGTG	CATCCACTACCTTTCCACAACCACCTCA
Nephrin	Mm01176615*		
NGAL	GGCCTCAAGGACGACAACA	ACCACCCATTGAGTTGTCAATG	CATCTTCTGTCCCCACCGACCAA
TGF-β	AAACGGAAGCGCATCGAA	GGGACTGGCGAGCCTTAGTT	CCATCCGTGGCCAGATCCTGTCC
CTGF	Mm01192933_g1		
TNF-α	CAGGTTCTGTCCCTTCACTCACT	CTGTGCTCATGGTGTCTTTTCTG	CCCAAGGCGCCACATCTCCCT
VCAM1	GGTGCTGTGACAATGACCTGTT	ATTATCTAACTTCTGCCCCAGAAAAT	CAGCGAGGGTCTACCAGCTCCTGAGA

Table S2