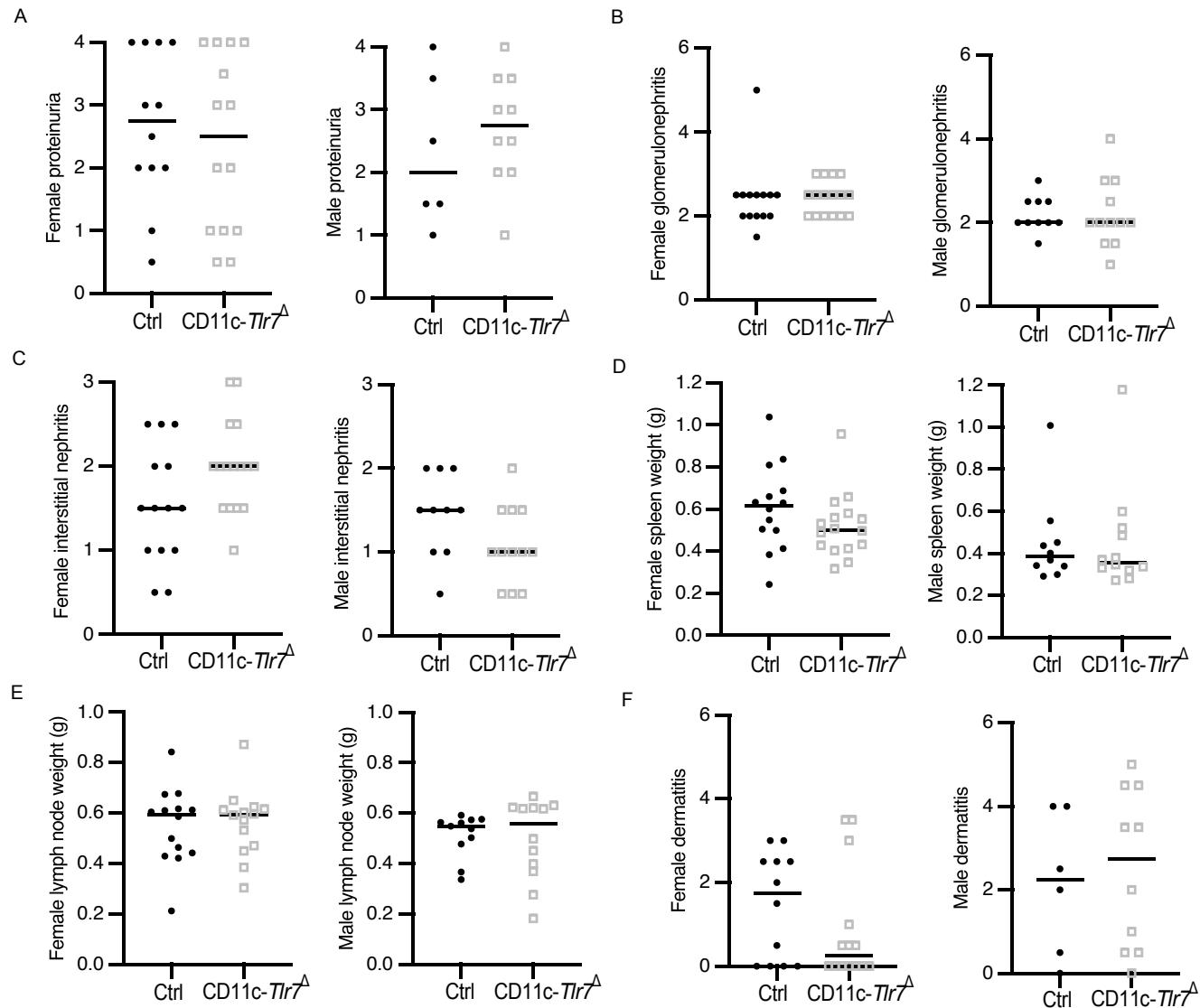
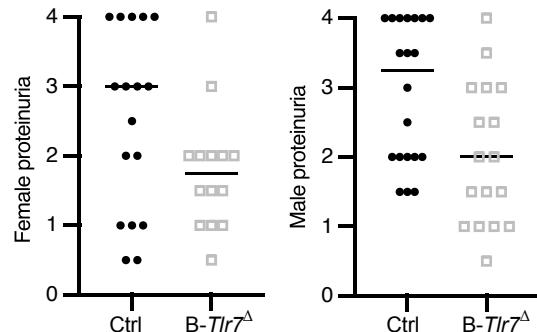


Supplemental Figure 1. Deletion analysis of TLR7 in aged experimental MRL/lpr mice. (A-C) Cre-lox mediated deletion in relevant cell subsets of experimental aged mice via representative histograms and tabulated data of anti-TLR7 PE MFI in the indicated subsets of cells: cDCs (CD19- CD11c⁺ MHCII⁺); pDCs (CD19- Siglec H⁺ CD317⁺); CD11c- B cells (TCRb⁻ CD19⁺ CD11c⁻); Marginal Zone B cells (TCRb⁻ CD19⁺ CD21/35⁺ CD23⁺); plasmablasts (TCRb⁻ CD44⁺ CD138⁺); and ABCs (CD19⁺ CD11c⁺ CD11b⁺). (A) Control (*Tlr7*^{+/+} and *Tlr7*^{+/+}) and CD11c-*Tlr7*^Δ (CD11c-Cre^{+/+} *Tlr7*^{+/+} and CD11c-Cre^{+/+} *Tlr7*^{+/+}) mice, (B) Control (*Tlr7*^{+/+} and *Tlr7*^{+/+}) and B-*Tlr7*^Δ (CD19-Cre^{+/+} *Tlr7*^{+/+} and CD19-Cre^{+/+} *Tlr7*^{+/+}) mice, and (C) Control (*Tlr7*^{+/+} *Tlr9*^{-/-} and *Tlr7*^{+/+} TLR9^{-/-}) and B-*Tlr7*^Δ *Tlr9*^{-/-} (CD19-Cre^{+/+} *Tlr7*^{+/+} *Tlr9*^{-/-} and CD19-Cre^{+/+} *Tlr7*^{+/+} *Tlr9*^{-/-}) mice. For all histograms the light line with open area is the control and the dark line with shaded area is the experimental (Cre x flox) group. Scatter plots display data from individual mice with black lines showing median values. *P<0.05, ** P<0.01, **** P<0.0001 by 2-tailed Mann-Whitney U test.

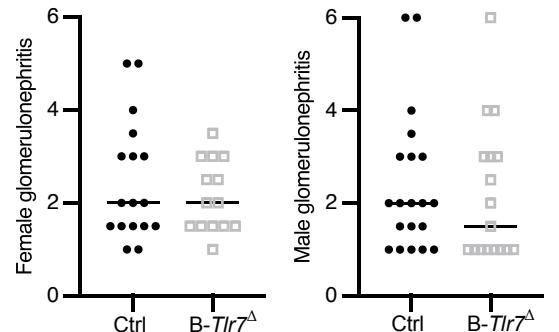


Supplemental Figure 2. Separate analysis of male and female clinical parameters in CD11c-Tlr7 Δ mice. Female control ($Tlr7^{IVL}$) and CD11c-Tlr7 Δ ($CD11c\text{-Cre}^{+/-} Tlr7^{IVL}$) mice were aged for 19 weeks. Male control ($Tlr7^{IVL}$) and CD11c-Tlr7 Δ ($CD11c\text{-Cre}^{+/-} Tlr7^{IVL}$) mice were aged for 22 weeks. **(A-F)** Evaluation of phenotypic disease markers in control and CD11c-Tlr7 Δ mice including: **(A)** proteinuria (females, n=12 and n=14; males, n=6 and n=10), **(B)** glomerulonephritis (females n=14 and n=16; males n=10 and n=12), **(C)** interstitial/perivascular infiltrates (females n=14 and n=16; males n=10 and n=12), **(D)** spleen weight (females n=14 and n=16; males n=10 and n=12), **(E)** lymph node weight (females n=14 and n=14; males n=10 and n=12), and **(F)** dermatitis (females n=14 and n=16; males n=6 and n=10). Scatter plots display data from individual mice with black lines showing median values. * P<0.05, ** P<0.01 by 2-tailed Mann-Whitney U test.

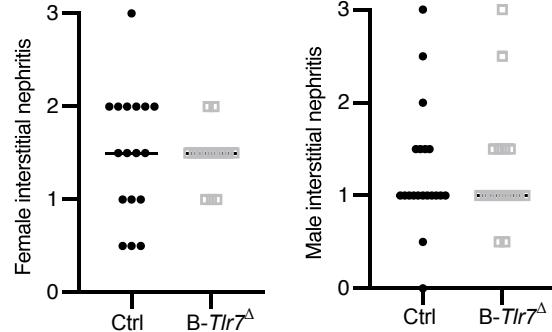
A



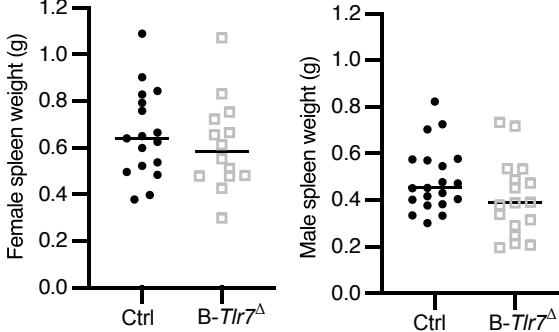
B



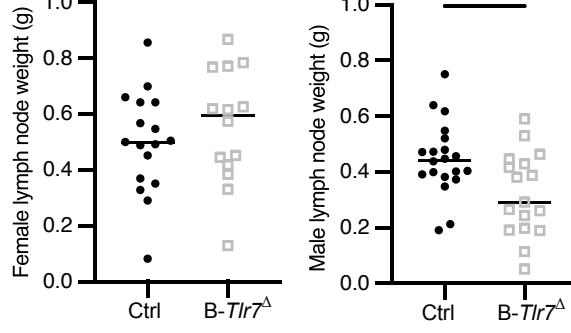
C



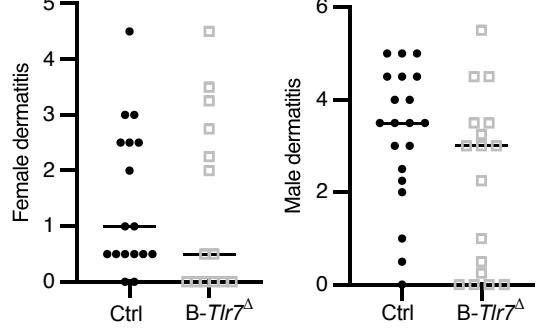
D



E

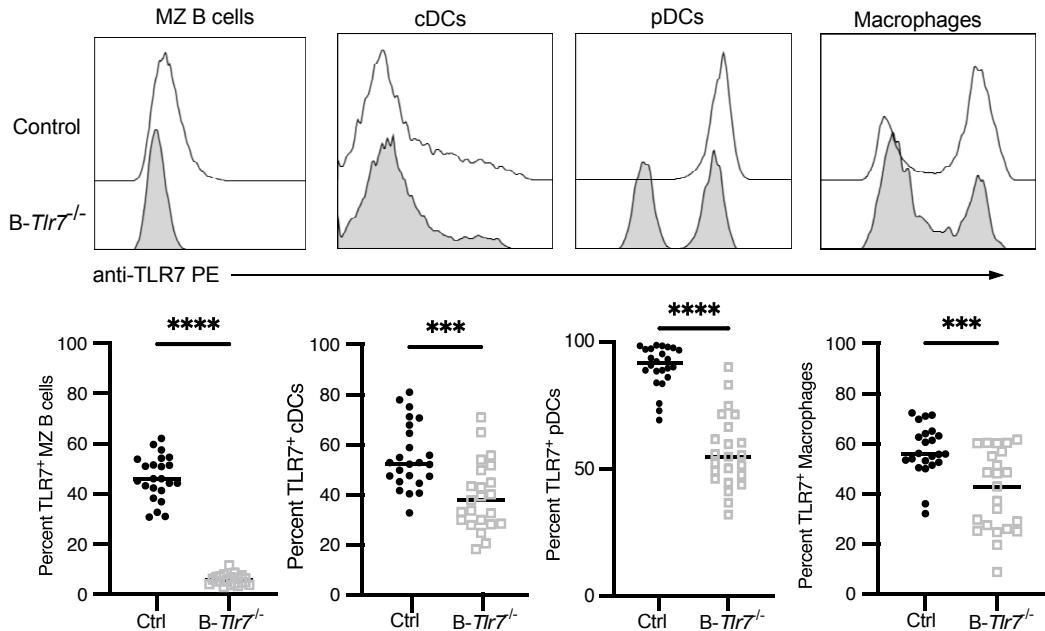


F

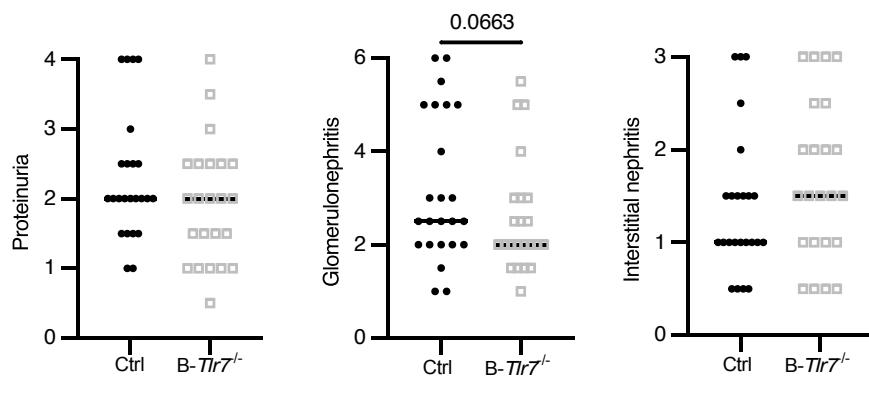


Supplemental Figure 3. Separate analysis of male and female clinical parameters in B-Tlr7 $^{\Delta}$ mice. Female control ($Tlr7^{IVL}$) and B-Tlr7 $^{\Delta}$ (CD19-Cre $^{+/-}$ $Tlr7^{IVL}$) were aged for 19 weeks. Male control ($Tlr7^{IVL}$) and B-Tlr7 $^{\Delta}$ (CD19-Cre $^{+/-}$ $Tlr7^{IVL}$) were aged for 22 weeks. (A-F) Evaluation of phenotypic disease markers in control and B-Tlr7 $^{\Delta}$ mice including: (A) proteinuria (females, n=17 and n=14; males, n=20 and n=17), (B) glomerulonephritis (females n=17 and n=14; males n=20 and n=17), (C) interstitial/perivascular infiltrates (females n=17 and n=14; males n=20 and n=17), (D) spleen weight (females n=17 and n=14; males n=20 and n=17), (E) lymph node weight (females n=17 and n=14; males n=20 and n=17), and (F) dermatitis (females n=17 and n=14; males n=20 and n=17). Scatter plots display data from individual mice with black lines showing median values. * $P < 0.05$, ** $P < 0.01$ by 2-tailed Mann-Whitney U test.

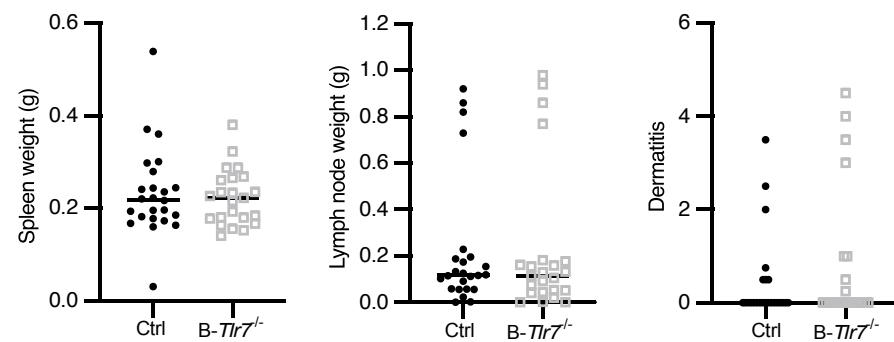
A



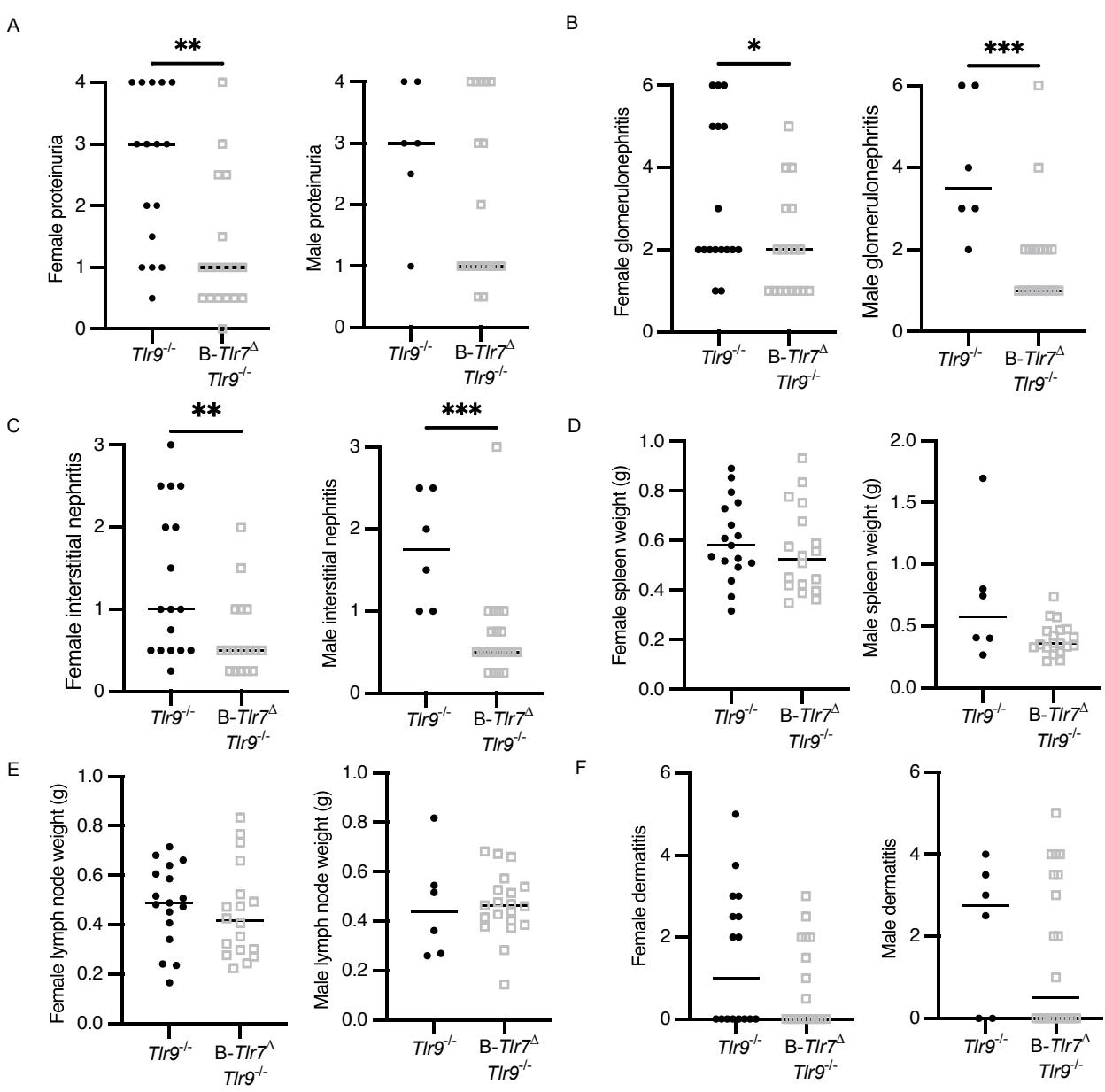
B



C



Supplemental Figure 4. TLR7 expression in B-Tlr7^{-/-} bone marrow chimeras and combined male and female clinical parameters in the chimeras. (A) Representative histograms showing TLR7 expression by flow cytometry and tabulated data of percent TLR7⁺ cells in control (no fill) and B-Tlr7^{-/-} (gray fill) mice in relevant cell subsets: Marginal Zone B cells (TCRb- CD19⁺ CD21/35⁺ CD23⁻); cDCs (CD19⁻ CD11c⁺ MHCII⁺); pDCs (CD19⁻ Siglec H⁺ CD317⁺); and macrophages (CD19⁻ CD11b⁺ LyG/C-F4/80⁺). (B) Combined male and female clinical SLE parameters in control vs B-Tlr7^{-/-} mice, n=24 and n=23 respectively. Scatter plots display data from individual mice with black lines showing median values. *** P < 0.001, **** P < 0.0001 by 1-tailed Mann-Whitney U test.



Supplemental Figure 5. Separate analysis of male and female clinical parameters in B-Tlr7^Δ Tlr9^{-/-} mice. Female control (*Tlr7^{fl/fl}/Tlr9^{-/-}*) and B-Tlr7^Δ (*CD19-Cre^{+/+} Tlr7^{fl/fl}/Tlr9^{-/-}*) were aged for 16 weeks. Male control (*Tlr7^{fl/fl}/Tlr9^{-/-}*) and B-Tlr7^Δ (*CD19-Cre^{+/+} Tlr7^{fl/fl}/Tlr9^{-/-}*) were aged for 19 weeks. **(A-F)** Evaluation of phenotypic disease markers in control and B-Tlr7^Δ mice including: **(A)** proteinuria (females, n=17 and n=18; males, n=6 and n=19), **(B)** glomerulonephritis (females n=17 and n=18; males n=6 and n=19), **(C)** interstitial/perivascular infiltrates (females n=17 and n=18; males n=6 and n=19), **(D)** spleen weight (females n=17 and n=18; males n=6 and n=19) **(E)** lymph node weight (females n=17 and n=18; males n=6 and n=19), and **(F)** dermatitis (females n=16 and n=18; males n=6 and n=19). Scatter plots display data from individual mice with black lines showing median values. * P<0.05, ** P<0.01 by 2-tailed Mann-Whitney U test.

Autoantibody	Ctrl	CD11c- <i>Tlr7</i> ^Δ	p-value	Ctrl	B- <i>Tlr7</i> ^Δ	p-value	<i>Tlr9</i> ^{-/-}	B- <i>Tlr7</i> ^Δ <i>Tlr9</i> ^{-/-}	p-value
Anti-Sm	1367.00 ± 452.90	536.90 ± 357.40	0.0037	1382.00 ± 513.60	0.10 ± 0.00	<0.0001	1995.00 ± 860.80	7.27 ± 7.17	<0.0001
Anti-RNA	87.74 ± 18.68	39.80 ± 12.14	0.0140	22.80 ± 7.53	3.80 ± 1.34	<0.0001	125.40 ± 24.34	35.81 ± 13.77	0.0002
Anti-nucleosome	2510.00 ± 926.0	3485.00 ± 728.50	0.5796	2288.00 ± 511.20	1997.00 ± 462.40	0.8416	5.07 ± 1.70	1.62 ± 0.64	0.0213

Supplemental Table 1. Autoantibody titers of MRL/lpr serum from indicated strains. Data represent mean ± SEM. Gray boxes indicate statistically significant differences by 2-tailed (CD11c-*Tlr7*^Δ and B-*Tlr7*^Δ cohorts) or 1-tailed (B-*Tlr7*^Δ *Tlr9*^{-/-}) Mann-Whitney *U* test.

Cell Type	Ctrl	CD11c- <i>Tlr7</i> ^Δ	Ctrl	B- <i>Tlr7</i> ^Δ	B- <i>Tlr7</i> ^{+/+}	B- <i>Tlr7</i> ^{-/-}	<i>TLR9</i> ^{-/-}	B- <i>Tlr7</i> ^Δ <i>Tlr9</i> ^{-/-}
B cells	16.36 ± 1.87	17.35 ± 1.57	11.77 ± 1.21	10.47 ± 0.98	8.42 ± 1.29	10.78 ± 1.47	10.54 ± 1.80	16.93 ± 1.72
Marginal Zone (% of B cells)	32.48 ± 2.25	33.15 ± 2.73	40.96 ± 3.11	46.23 ± 2.50	10.20 ± 1.76	14.47 ± 1.54	21.92 ± 2.23	32.43 ± 1.84
Follicular (% of B cells)	35.11 ± 3.95	33.02 ± 2.57	25.85 ± 2.04	27.68 ± 2.22	6.18 ± 0.83	10.20 ± 1.76	38.86 ± 2.43	38.60 ± 1.77
CD11b ⁺ CD11c ⁺ B cells (% of B cells)	2.21 ± 0.23	2.56 ± 0.28	6.28 ± 0.69	6.25 ± 0.72	3.33 ± 0.40	4.46 ± 0.56	3.05 ± 0.32	3.07 ± 0.29
Plasmablasts (% of TCRβ ⁺)	2.18 ± 0.34	1.51 ± 0.17	1.85 ± 0.44	1.79 ± 0.47	2.26 ± 0.29	2.56 ± 0.33	2.16 ± 0.39	2.05 ± 0.30
cDCs (% of CD19 ⁺)	0.54 ± 0.06	0.41 ± 0.04	1.44 ± 0.25	1.19 ± 0.18	1.59 ± 0.15	1.07 ± 0.10	0.50 ± 0.07	0.68 ± 0.05
pDCs (% of CD19 ⁺)	0.25 ± 0.03	0.22 ± 0.02	0.15 ± 0.02	0.16 ± 0.02	0.36 ± 0.03	0.36 ± 0.04	0.20 ± 0.03	0.28 ± 0.02
T cells	49.54 ± 2.76	41.45 ± 3.16	46.16 ± 3.23	45.68 ± 3.14	37.15 ± 1.66	35.85 ± 1.40	49.64 ± 3.86	45.81 ± 2.19
CD4 T cells (% of T cells)	24.64 ± 1.40	30.52 ± 1.88	30.28 ± 2.03	32.04 ± 2.04	57.74 ± 0.90	56.31 ± 1.71	25.98 ± 3.01	28.91 ± 1.44
CD4 naïve T cells (% of CD4 T cells)	4.49 ± 0.47	4.24 ± 0.44	5.64 ± 0.93	6.43 ± 1.00	3.29 ± 0.37	3.94 ± 0.45	2.82 ± 0.29	4.11 ± 0.32
CD8 T cells (% of T cells)	22.31 ± 1.21	20.58 ± 1.03	19.94 ± 1.16	21.70 ± 1.23	21.47 ± 0.87	20.66 ± 0.71	14.81 ± 1.78	23.16 ± 1.43
CD8 naïve T cells (% of CD8 T cells)	11.76 ± 0.79	12.20 ± 0.92	10.05 ± 0.62	10.65 ± 0.63	11.46 ± 1.48	12.57 ± 1.41	6.30 ± 0.55	8.56 ± 0.48
Double negative T cells (% of T cells)	50.48 ± 2.08	45.57 ± 1.94	46.69 ± 2.25	43.56 ± 2.00	16.01 ± 0.78	18.55 ± 1.56	52.66 ± 4.55	43.86 ± 2.57
Macrophages (% of CD11b ⁺ CD19 ⁺)	8.91 ± 0.78	8.05 ± 1.06	7.82 ± 1.01	9.57 ± 1.26	5.80 ± 0.93	4.75 ± 0.90	9.74 ± 1.31	9.82 ± 0.75
Monocytes (% of CD11b ⁺ CD19 ⁺)	12.12 ± 1.22	14.14 ± 1.66	7.47 ± 0.91	7.41 ± 0.88	16.12 ± 0.92	15.69 ± 0.91	8.99 ± 1.05	12.08 ± 0.97
Neutrophils (% of CD11b ⁺ CD19 ⁺)	22.33 ± 3.50	34.53 ± 4.24	33.62 ± 3.36	31.03 ± 3.17	51.77 ± 2.82	44.60 ± 4.60	37.76 ± 3.06	39.52 ± 2.26

Supplemental Table 2. Percent cell compositions of MRL/lpr spleens from indicated strains. Data represent mean ± SEM. Gray boxes indicate statistically significant differences by 2-tailed (CD11c-*Tlr7*^Δ and B-*Tlr7*^Δ cohorts) or 1-tailed (B-*Tlr7*^{-/-} and B-*Tlr7*^Δ *Tlr9*^{-/-} cohorts) Mann-Whitney *U* test.

Clinical Parameter	Sex	Ctrl	CD11c- <i>Tlr7^Δ</i>	p-value	Ctrl	B- <i>Tlr7^Δ</i>	p-value	Ctrl	B- <i>Tlr7^{-/-}</i>	p-value	B- <i>Tlr9^{-/-}</i>	<i>Tlr7^Δ</i> <i>Tlr9^{-/-}</i>	p-value
Proteinuria	All	2.56 ± 0.28	2.52 ± 0.25	0.9232	2.73 ± 0.19	1.92 ± 0.17	.0034	2.29 ± 0.19	1.94 ± 0.18	0.1143	2.63 ± 0.25	1.64 ± 0.21	0.0014
	Male	2.33 ± 0.49	2.70 ± 0.28	0.5366	2.93 ± 0.23	2.03 ± 0.25	.0014	1.96 ± 0.14	2.10 ± 0.19	0.2632	2.92 ± 0.45	2.00 ± 0.32	0.0781
	Female	2.67 ± 0.35	2.40 ± 0.38	0.6358	2.50 ± 0.32	1.79 ± 0.24	.0643	2.58 ± 0.30	1.81 ± 0.29	0.0316	2.56 ± 0.32	1.25 ± 0.25	0.0011
Glomerulonephritis	All	2.33 ± 0.14	2.34 ± 0.11	0.7041	2.43 ± 0.23	2.16 ± 0.21	.2137	3.19 ± 0.32	2.57 ± 0.25	0.0663	3.39 ± 0.38	1.92 ± 0.21	0.0001
	Male	2.20 ± 0.13	2.21 ± 0.23	0.7545	2.40 ± 0.34	2.18 ± 0.36	.4891	2.69 ± 0.38	2.35 ± 0.39	0.2449	4.00 ± 0.68	1.79 ± 0.29	0.0009
	Female	2.43 ± 0.22	2.44 ± 0.10	0.4412	2.47 ± 0.31	2.14 ± 0.21	.6821	3.62 ± 0.48	2.55 ± 0.30	0.0594	3.18 ± 0.45	2.06 ± 0.30	0.0204
Interstitial Nephritis	All	1.50 ± 0.12	1.61 ± 0.13	0.6271	1.37 ± 0.11	1.36 ± 0.09	.4650	1.40 ± 0.16	1.67 ± 0.18	0.1175	1.44 ± 0.18	0.69 ± 0.09	0.0001
	Male	1.45 ± 0.16	1.08 ± 0.14	0.1136	1.25 ± 0.15	1.27 ± 0.16	.9698	1.00 ± 0.12	1.40 ± 0.23	0.0972	1.75 ± 0.28	0.70 ± 0.13	0.0003
	Female	1.54 ± 0.19	2.00 ± 0.14	0.0693	1.50 ± 0.17	1.46 ± 0.08	.7661	1.73 ± 0.24	1.89 ± 0.25	0.3562	1.32 ± 0.22	0.65 ± 0.11	0.0052
Spleen Weight	All	0.54 ± 0.04	0.49 ± 0.04	0.3382	0.57 ± 0.03	0.50 ± 0.04	.1532	0.23 ± 0.02	0.22 ± 0.01	0.3541	0.63 ± 0.06	0.47 ± 0.03	0.0056
	Male	0.45 ± 0.07	0.45 ± 0.07	0.7713	0.49 ± 0.03	0.41 ± 0.04	.1035	0.19 ± 0.01	0.19 ± 0.02	0.3523	0.72 ± 0.21	0.40 ± 0.03	0.0674
	Female	0.61 ± 0.05	0.52 ± 0.04	0.1536	0.66 ± 0.71	0.61 ± 0.05	.4043	0.27 ± 0.03	0.25 ± 0.02	0.2577	0.60 ± 0.04	0.55 ± 0.04	0.3558
LN Weight	All	0.53 ± 0.03	0.53 ± 0.03	0.5601	0.47 ± 0.03	0.43 ± 0.04	.2426	0.23 ± 0.06	0.23 ± 0.07	0.3194	0.48 ± 0.04	0.46 ± 0.03	0.2742
	Male	0.51 ± 0.03	0.50 ± 0.05	0.5824	0.45 ± 0.03	0.32 ± 0.04	.0157	0.024 ± ±0.10	0.22 ± 0.11	0.1618	0.46 ± 0.09	0.46 ± 0.03	0.3722
	Female	0.55 ± 0.04	0.56 ± 0.04	0.9554	0.50 ± 0.04	0.56 ± 0.06	.4743	0.22 ± 0.07	0.24 ± 0.08	0.4259	0.48 ± 0.04	0.45 ± 0.04	0.2338
Dermatitis	All	1.69 ± 0.33	1.56 ± 0.36	0.8025	2.43 ± 0.27	1.84 ± 0.32	.0961	0.43 ± 0.19	0.77 ± 0.30	0.2741	1.67 ± 0.36	1.26 ± 0.26	0.2013
	Male	2.17 ± 0.69	2.50 ± 0.61	0.6924	3.24 ± 0.33	2.22 ± 0.45	.0740	0.27 ± 0.23	0.50 ± 0.40	0.7243	2.17 ± 0.71	1.60 ± 0.41	0.2887
	Female	1.46 ± 0.37	0.89 ± 0.36	0.3898	1.47 ± 0.32	1.38 ± 0.43	.4537	0.56 ± 0.29	0.98 ± 0.44	0.6205	1.48 ± 0.42	0.81 ± 0.25	0.1423

Supplemental Table 3. Clinical data of MRL/lpr mice from indicated strains. Data represent mean ± SEM. Gray boxes indicate statistically significant differences and bold text indicates data trending towards significance by 2-tailed (CD11c-*Tlr7^Δ* and B-*Tlr7^Δ* cohorts) or 1-tailed (B-*Tlr7^{-/-}* and B-*Tlr7^Δ* *Tlr9^{-/-}*) Mann-Whitney U test.