

**Supplemental Table 1. Complete blood count of CT and EC-Myc KO mice.**

Test	CT	EC-Myc KO	p-value
<b>White Blood Cell Count (<math>10^3/\mu\text{L}</math>)</b>	$2.17 \pm 0.75$	$2.27 \pm 0.81$	0.828
<b>Red Blood Cell Count (<math>10^6/\mu\text{L}</math>)</b>	$8.23 \pm 0.60$	$8.32 \pm 0.71$	0.814
<b>Hemoglobin (g/dL)</b>	$11.02 \pm 0.72$	$11.08 \pm 0.83$	0.885
<b>Hematocrit (%)</b>	$39.33 \pm 3.67$	$40.00 \pm 2.37$	0.716
<b>MCV (fL)</b>	$48.00 \pm 1.10$	$48.00 \pm 1.41$	1.000
<b>MCH (pg)</b>	$13.50 \pm 0.55$	$13.33 \pm 0.52$	0.599
<b>MCHC (%)</b>	$28.17 \pm 1.17$	$27.83 \pm 0.75$	0.570
<b>Neutrophils Count (<math>10^3/\mu\text{L}</math>)</b>	$0.42 \pm 0.32$	$0.52 \pm 0.33$	0.618
<b>Neutrophils (%)</b>	$15.80 \pm 4.87$	$21.17 \pm 9.64$	0.290
<b>Lymphocytes Count (<math>10^3/\mu\text{L}</math>)</b>	$1.57 \pm 0.71$	$1.46 \pm 0.47$	0.766
<b>Lymphocytes (%)</b>	$78.40 \pm 6.07$	$66.17 \pm 10.76$	0.051
<b>Monocytes Count (<math>10^3/\mu\text{L}</math>)</b>	$0.17 \pm 0.08$	$0.26 \pm 0.14$	0.172
<b>Monocytes (%)</b>	$7.67 \pm 2.66$	$11.33 \pm 2.80$	<b>0.045*</b>
<b>Eosinophils Count (<math>10^3/\mu\text{L}</math>)</b>	$0.017 \pm 0.010$	$0.025 \pm 0.030$	0.536
<b>Eosinophils (%)</b>	$0.83 \pm 0.41$	$1.17 \pm 1.60$	0.632

Results are expressed as mean  $\pm$  standard deviation (n = 6). CT, control; EC-Myc KO, endothelial c-Myc knockout; MCV, mean corpuscular volume; MCH, mean cell hemoglobin; MCHC, mean corpuscular hemoglobin concentration. \*p<0.05.

**Supplemental Table 2. Flow cytometry analysis of bone marrow cells from CT and EC-Myc KO mice.**

Cell Type	Male		Female	
	CT	EC-Myc KO	CT	EC-Myc KO
<b>Long Term Hematopoietic Stem cell (LT-HSC)</b>	0.0076 ± 0.0016	0.0049 ± 0.0027	0.021 ± 0.026	0.022 ± 0.015
<b>Short Term Hematopoietic Stem cell (ST-HSC)</b>	0.0118 ± 0.0046	0.0136 ± 0.0061	0.074 ± 0.034*	0.069 ± 0.020§
<b>Multipotent Progenitors (MMP)</b>	0.146 ± 0.028	0.144 ± 0.057	0.274 ± 0.099*	0.266 ± 0.050§
<b>Lymphoid Progenitors (LP)</b>	0.060 ± 0.015	0.063 ± 0.036	0.166 ± 0.070*	0.146 ± 0.036§
<b>Common Myeloid Progenitors (CMP)</b>	0.112 ± 0.016	0.098 ± 0.013	0.074 ± 0.009*	0.057 ± 0.021§
<b>Granulocyte-Monocyte Progenitors (GMP)</b>	0.240 ± 0.023	0.204 ± 0.030	0.210 ± 0.055	0.204 ± 0.013
<b>Megakaryocyte-Erythroid Progenitors (MEP)</b>	0.310 ± 0.084	0.292 ± 0.073	0.176 ± 0.024*	0.224 ± 0.038
<b>Granulocytes</b>	52.7 ± 8.0	49.8 ± 7.4	48.1 ± 7.3	41.6 ± 3.5
<b>Erythrocytes</b>	10.1 ± 4.0	11.5 ± 4.0	5.9 ± 2.0	9.3 ± 2.5
<b>Monocytes/Macrophages</b>	5.30 ± 0.78	4.40 ± 0.62	3.72 ± 0.83*	3.41 ± 0.82
<b>B Lymphocytes</b>	4.94 ± 1.57	5.01 ± 1.99	6.07 ± 1.98	7.24 ± 2.34
<b>CD4+ T Lymphocytes</b>	0.38 ± 0.09	0.43 ± 0.26	0.20 ± 0.03	0.28 ± 0.20
<b>CD8+ T Lymphocytes</b>	0.42 ± 0.15	0.37 ± 0.19	0.11 ± 0.20	0.25 ± 0.51
<b>NK cells</b>	0.35 ± 0.12	0.39 ± 0.05	0.40 ± 0.13	0.45 ± 0.09

Values represent percentage of different cell types. Results are expressed as mean ± standard deviation (n = 5). CT, control; EC-Myc KO, endothelial c-Myc knockout. \*p<0.05 between genders within CT; §p<0.05 between genders within EC-Myc KO.

**Supplemental Table 3. Multiplex analysis of serum samples from CT and EC-Myc KO mice.**

Target	CT	EC-Myc KO	p-value
<b>CCL2</b>	n.d.	n.d.	
<b>CCL3</b>	n.d.	n.d.	
<b>CCL4</b>	n.d.	n.d.	
<b>CCL5</b>	n.d.	n.d.	
<b>CCL7</b>	37 ± 19	74 ± 40	<b>0.012*</b>
<b>CCL11</b>	761 ± 315	743 ± 492	0.920
<b>CCL12</b>	85 ± 67	75 ± 28	0.661
<b>CCL21</b>	2291 ± 700	2875 ± 2494	0.463
<b>TNFSF11</b>	121 ± 21	90 ± 24	<b>0.004**</b>
<b>TWEAK</b>	n.d.	n.d.	
<b>IL1β</b>	n.d.	n.d.	
<b>IL6</b>	n.d.	n.d.	
<b>IL-5</b>	n.d.	n.d.	
<b>IL-16</b>	519 ± 255	563 ± 206	0.662
<b>Osteopontin</b>	139986 ± 17031	197414 ± 65967	<b>0.011*</b>

Values represent concentration in pg/ml calculated using standard curves for each marker. Results are expressed as mean ± standard deviation (n = 11). CT, control; EC-Myc KO, endothelial c-Myc knockout; n.d., not detected. \*p<0.05, \*\*p<0.01.

**Supplemental Table 4. Phenotypic analysis of CT and EC-Myc KO mice after long-term exposure to HFD.**

Male	CT-CTD	EC-Myc KO-CTD	CT-HFD	EC-Myc KO-HFD
<b>Body Weight (g)</b>	32.8 ± 3.7	37.6 ± 4.5*	27.2 ± 3.2†	27.9 ± 4.2‡
<b>Liver Weight (g)</b>	1.26 ± 0.20	1.49 ± 0.22	1.91 ± 0.25†	2.00 ± 0.30‡
<b>Liver/Body Weight Ratio</b>	0.038 ± 0.004	0.039 ± 0.002	0.071 ± 0.011†	0.073 ± 0.013‡
<b>Plasma Triglycerides (mg/dL)</b>	56 ± 15	65 ± 20	55 ± 13	58 ± 14
<b>Plasma Cholesterol (mg/dL)</b>	104 ± 15	119 ± 17	150 ± 29†	<b>177 ± 38\$§</b>
<b>Liver Triglycerides (mg/g tissue)</b>	17.2 ± 14.1	30.1 ± 13.2	19.1 ± 5.5	21.3 ± 8.8
<b>Liver Cholesterol (mg/g tissue)</b>	3.03 ± 0.67	3.34 ± 0.61	22.67 ± 7.07†	21.54 ± 6.80‡
Female	CT-CTD	EC-Myc KO-CTD	CT-HFD	EC-Myc KO-HFD
<b>Body Weight (g)</b>	22.1 ± 1.4	23.2 ± 3.1	21.7 ± 1.5	22.2 ± 2.0
<b>Liver Weight (g)</b>	0.87 ± 0.09	0.92 ± 0.15	1.86 ± 0.24†	1.93 ± 0.21‡
<b>Liver/Body Weight Ratio</b>	0.040 ± 0.004	0.039 ± 0.003	0.086 ± 0.006†	0.087 ± 0.006‡
<b>Plasma Triglycerides (mg/dL)</b>	67 ± 21	67 ± 25	75 ± 29	55 ± 24
<b>Plasma Cholesterol (mg/dL)</b>	78 ± 12	81 ± 12	160 ± 41†	<b>191 ± 27\$§</b>
<b>Liver Triglycerides (mg/g tissue)</b>	26.2 ± 5.0	24.7 ± 8.3	20.3 ± 2.1	21.6 ± 5.5
<b>Liver Cholesterol (mg/g tissue)</b>	4.35 ± 0.50	3.55 ± 0.52	30.75 ± 2.30†	31.20 ± 4.54‡

Results were collected after long-term exposure to experimental diet and are expressed as mean ± standard deviation (n = 8-13). CT, control; EC-Myc KO, endothelial c-Myc knockout; CTD, low-fat control diet; HFD, high-fat diet. \*p<0.05 between genotypes within CTD; †p<0.05 between diets within CT; ‡p<0.05 between diets within EC-Myc KO; §p<0.05 between genotypes within HFD.

**Supplemental Table 5. Multiplex analysis of liver lysates from CT and EC-Myc KO mice after short-term exposure to HFD.**

Target	Male		Female	
	CT	EC-Myc KO	CT	EC-Myc KO
<b>Eotaxin/CCL11</b>	11.9 ± 1.9	13.3 ± 3.3	13.6 ± 3.6	<b>18.4 ± 5.2**‡</b>
<b>G-CSF</b>	0.81 ± 0.92	0.73 ± 0.49	0.85 ± 0.34	0.85 ± 0.45
<b>GM-CSF</b>	n.d.	n.d.	n.d.	n.d.
<b>IFNγ</b>	7.56 ± 1.78	8.79 ± 2.60	9.65 ± 0.89†	10.51 ± 1.73
<b>IL-1α</b>	158.7 ± 17.8	168.1 ± 18.0	177.8 ± 19.3†	191.0 ± 9.9‡
<b>IL-1β</b>	6.19 ± 2.17	5.95 ± 1.74	6.95 ± 1.62	7.83 ± 0.88‡
<b>IL-2</b>	32.7 ± 5.0	34.6 ± 7.3	36.5 ± 5.4	<b>43.4 ± 2.9**‡</b>
<b>IL-3</b>	0.35 ± 0.22	0.34 ± 0.15	0.35 ± 0.12	0.48 ± 0.20
<b>IL-4</b>	0.32 ± 0.12	0.32 ± 0.12	0.43 ± 0.13†	0.47 ± 0.05‡
<b>IL-5</b>	n.d.	n.d.	n.d.	n.d.
<b>IL-6</b>	1.55 ± 0.37	1.85 ± 0.84	1.94 ± 0.48	2.06 ± 0.67
<b>IL-7</b>	5.31 ± 1.53	5.69 ± 0.97	6.94 ± 0.65†	<b>8.26 ± 0.91**‡</b>
<b>IL-9</b>	43.0 ± 9.2	42.6 ± 9.8	50.8 ± 7.4	57.2 ± 7.6‡
<b>IL-10</b>	30.7 ± 10.8	35.3 ± 7.1	39.3 ± 8.4†	43.2 ± 4.4‡
<b>IL-12p40</b>	42.8 ± 18.7	51.5 ± 21.7	63.3 ± 18.9†	72.5 ± 13.6‡
<b>IL-12p70</b>	4.02 ± 2.97	3.89 ± 1.08	3.16 ± 1.37	4.01 ± 1.90
<b>IL-13</b>	1.81 ± 1.09	2.06 ± 0.71	2.70 ± 0.55†	3.35 ± 0.78‡
<b>IL-15</b>	23.2 ± 8.3	25.6 ± 6.4	31.8 ± 5.2†	<b>39.3 ± 6.3**‡</b>
<b>IL-17</b>	0.39 ± 0.04	0.42 ± 0.09	0.52 ± 0.04†	<b>0.60 ± 0.08**‡</b>
<b>KC/CXCL1</b>	53.0 ± 14.3	54.8 ± 5.8	58.7 ± 13.6	<b>70.4 ± 7.1**‡</b>
<b>LIX/CXCL5</b>	146.8 ± 67.5	141.8 ± 47.3	164.8 ± 56.7	201.0 ± 34.7‡
<b>MCP-1/CCL2</b>	16.9 ± 6.4	14.8 ± 7.0	17.1 ± 12.9	12.8 ± 5.5
<b>M-CSF</b>	2.00 ± 0.38	1.61 ± 0.38	1.92 ± 0.40	2.02 ± 0.41

<b>MIP-1<math>\alpha</math>/CCL3</b>	25.4 $\pm$ 2.7	28.0 $\pm$ 3.0	27.8 $\pm$ 3.4	30.1 $\pm$ 2.5
<b>MIP-1<math>\beta</math>/CCL4</b>	n.d.	n.d.	n.d.	n.d.
<b>MIP-2/CXCL2</b>	195.2 $\pm$ 37.3	197.4 $\pm$ 42.9	213.1 $\pm$ 32.2	243.9 $\pm$ 27.1 $^{\ddagger}$
<b>RANTES/CCL5</b>	3.20 $\pm$ 0.63	3.64 $\pm$ 1.15	4.14 $\pm$ 1.36	3.81 $\pm$ 0.75
<b>TNF<math>\alpha</math></b>	1.75 $\pm$ 0.34	1.43 $\pm$ 0.27	2.11 $\pm$ 0.64	2.37 $\pm$ 0.84 $^{\ddagger}$
<b>VEGF</b>	14.5 $\pm$ 2.0	14.0 $\pm$ 1.3	14.8 $\pm$ 2.1	16.7 $\pm$ 3.8

Values represent concentration in pg/ug total protein calculated using standard curves for each marker. Results are expressed as mean  $\pm$  standard deviation (n = 9-10). CT, control; EC-Myc KO, endothelial c-Myc knockout; n.d., not detected. \*p<0.05 between genotypes within female; †p<0.05 between genders within CT;  $^{\ddagger}$ p<0.05 between genders within EC-Myc KO.