### **Supplemental Information**

#### SUPPLEMENTARY FIGURE LEGENDS

Supplemental Figure 1. Characterization of Tgr5<sup>bm+/+</sup> and Tgr5<sup>bm-/-</sup> mice.

(A) Quantitative real time analysis of *Tgr5* on genomic DNA extracted from peripheral blood. (B) Daily food and (C) water intake measured in HFD-fed *Tgr5*<sup>bm+/+</sup> (white bar) and *Tgr5*<sup>bm-/-</sup> (black bar) mice. n=12/group. Results represent the mean±SEM. P values were calculated by using two tailed Student's t test and one-way ANOVA test followed by Bonferroni's post hoc analysis when appropriate.

Supplemental Figure 2. Comparable inflammation in gastrocnemius derived from  $Tgr5^{bm-/-}$  and  $Tgr5^{bm+/+}$  mice.

**(A)** mRNA levels of macrophage markers in gastrocnemius of HFD-fed  $Tgr5^{bm+/+}$  (white bar) and  $Tgr5^{bm-/-}$  (black bar) mice. n=12/group. **(B-C)** mRNA levels of **(B)** pro-inflammatory M1 and **(C)** anti-inflammatory M2 markers in gastrocnemius of HFD-fed  $Tgr5^{bm+/+}$  and  $Tgr5^{bm-/-}$  mice. n=12/group. Results represent the mean±SEM. P values were calculated by using two tailed Student's t test. \*statistically significant, p<0.05.

Supplemental Figure 3. *Glucose homeostasis and macrophage infiltration in eWAT of LysM-Cre Tgr5*<sup>fl/fl</sup> *mice.* 

(A) mRNA levels of Tgr5 in primary bone marrow-derived macrophages (BMDMs) isolated from *Tgr5*<sup>fl/fl</sup> or LysM-Cre *Tgr5*<sup>fl/fl</sup> mice. n=4/group. (B) Blood glucose levels during insulin tolerance test (ITT) (insulin; 0.50 U/kg) in chow diet-fed Tgr5<sup>fl/fl</sup> and LysM-Cre Tgr5<sup>fl/fl</sup> mice. n=12/group. (C) Plasma levels of insulin during an oral glucose tolerance test (OGTT) in chow diet-fed Tgr5<sup>fl/fl</sup> and LysM-Cre Tgr5<sup>fl/fl</sup> mice. n=12/group. (D) Blood glucose levels during OGTT in chow diet-fed Tgr5<sup>fl/fl</sup> and LysM-Cre Tgr5<sup>fl/fl</sup> mice. n=12/group. (E) mRNA levels of macrophage markers in eWAT of chow diet-fed Tgr5<sup>fl/fl</sup> and LysM-Cre Tgr5<sup>fl/fl</sup> mice. n=12/group. Results represent the mean±SEM. P values were calculated by using two tailed Student's t test and two-way ANOVA test followed by Bonferroni's post hoc analysis when appropriate. \*\*\*\*p<0.001.

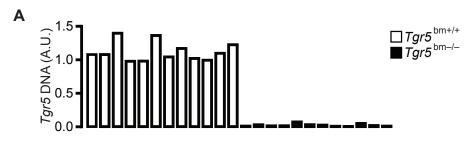
Supplemental Figure 4. Chemokine and chemokine receptor expression in Tgr5<sup>+/+</sup> and Tgr5<sup>-/-</sup> BMDMs.

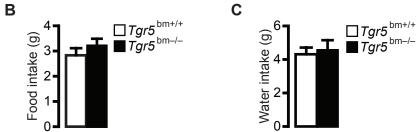
(A-E) mRNA levels of chemokines and their cognate receptors in primary bone marrow-derived macrophages (BMDMs) isolated from  $Tgr5^{+/+}$  or  $Tgr5^{-/-}$  mice and stimulated with LPS (10 ng/mL) for 24 hr, in presence or absence of the TGR5 agonist INT-777 (30 $\mu$ M). n=6/group. Results represent the mean±SEM. P values were calculated by using one-way ANOVA test followed by Bonferroni's post hoc analysis. \*p<0.05 \*\*p<0.01, \*\*\*p<0.001 versus genotype and #p<0.001 versus control.

Supplemental Figure 5. Expression of nuclear receptors and their target genes in Tgr5<sup>+/+</sup> and Tgr5<sup>-/-</sup> BMDMs.

(A-C) mRNA levels of (A) PPARγ1 and its target genes (B) CD36 and (C) Lpl in primary bone marrow-derived macrophages (BMDMs) isolated from  $Tgr5^{+/+}$  or  $Tgr5^{-/-}$  mice and stimulated with LPS (10 ng/mL) for 24 hr, in presence or absence of the TGR5 agonist INT-777 (30μM). n=6/group. (D-I) mRNA levels of (D) LXRα, (E) LXRβ and their target genes (F) ApoE, (G) Abca1, (H) Abcg1 and (I) Srebp1c in primary bone marrow-derived macrophages (BMDMs) isolated from  $Tgr5^{+/+}$  or  $Tgr5^{-/-}$  mice and stimulated with LPS (10 ng/mL) for 24 hr, in presence or absence of the TGR5 agonist INT-777 (30μM). n=6/group. Results represent the mean±SEM. P values were calculated by using one-way ANOVA test followed by Bonferroni's post hoc analysis. \*p<0.05 \*\*p<0.01, \*\*\*p<0.001 versus genotype and #p<0.001 versus control.

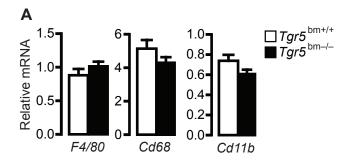
Supplemental Table 1. *Haematological parameters measured in Tgr5*<sup>bm+/+</sup> and *Tgr5*<sup>bm-/-</sup> mice after 18 week of HFD (n=6/group).

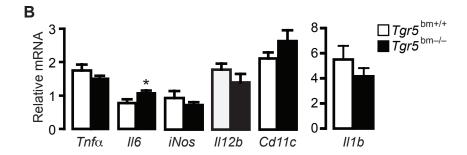


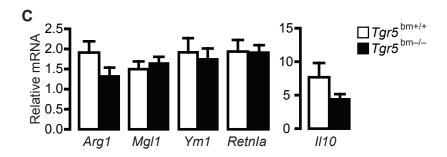


## **Supplemental Figure 1.**

Characterization of *Tgr5*<sup>bm+/+</sup> and *Tgr5*<sup>bm-/-</sup> mice. **(A)** Quantitative real time analysis of *Tgr5* on genomic DNA extracted from peripheral blood of *Tgr5*<sup>bm+/+</sup> and *Tgr5*<sup>bm-/-</sup> mice. n=12/group. **(B)** Daily food and **(C)** water intake measured in HFD-fed *Tgr5*<sup>bm+/+</sup> (white bar) and *Tgr5*<sup>bm-/-</sup> (black bar) mice. n=12/group. Results represent the mean±SEM. P values were calculated by using two tailed Student's t test and one-way ANOVA test followed by Bonferroni's post hoc analysis when appropriate.

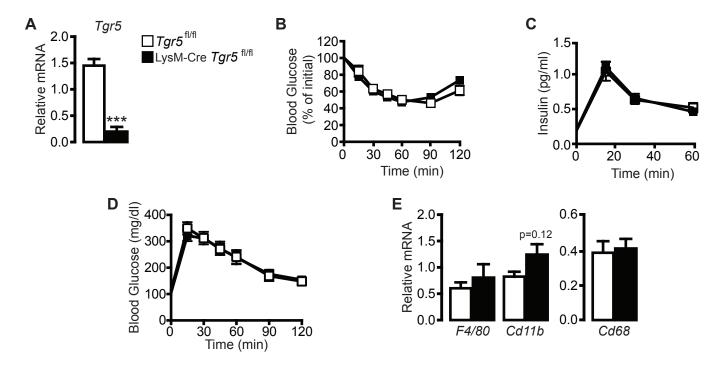






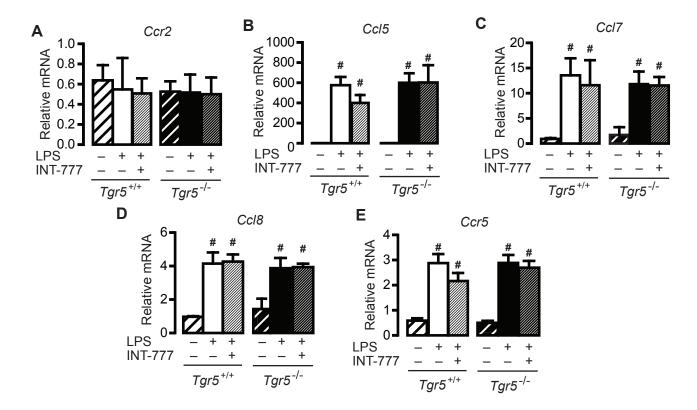
### Supplemental Figure 2.

Comparable inflammation in gastrocnemius derived from  $Tgr5^{bm-/-}$  and  $Tgr5^{bm+/+}$  mice. **(A)** mRNA levels of macrophage markers in gastrocnemius of HFD-fed  $Tgr5^{bm+/+}$  (white bar) and  $Tgr5^{bm-/-}$  (black bar) mice. n=12/group. **(B-C)** mRNA levels of **(B)** pro-inflammatory M1 and **(C)** anti-inflammatory M2 markers in gastrocnemius of HFD-fed  $Tgr5^{bm+/+}$  and  $Tgr5^{bm-/-}$  mice. n=12/group. Results represent the mean±SEM. P values were calculated by using two tailed Student's t test. \*statistically significant, p<0.05.



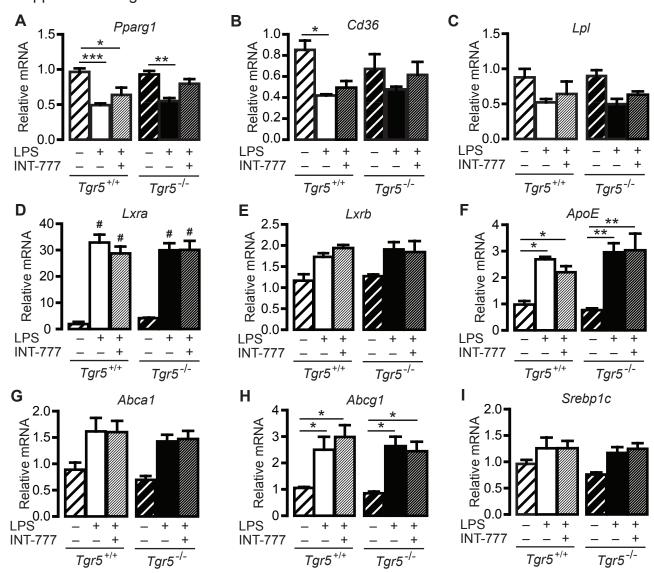
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# **Supplemental Figure 4.**

Chemokine and chemokine receptor expression in  $Tgr5^{+/+}$  and  $Tgr5^{-/-}$  BMDMs. (A-E) mRNA levels of chemokines and their cognate receptors in primary bone marrow-derived macrophages (BMDMs) isolated from  $Tgr5^{+/+}$  or  $Tgr5^{-/-}$  mice and stimulated with LPS (10 ng/mL) for 24 hr, in presence or absence of the TGR5 agonist INT-777 (30µM). n=6/group. Results represent the mean±SEM. P values were calculated by using one-way ANOVA test followed by Bonferroni's post hoc analysis. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001 versus genotype and #p<0.001 versus control.



## **Supplemental Figure 5.**

Expression of nuclear receptors and their target genes in  $Tgr5^{+/+}$  and  $Tgr5^{-/-}$  BMDMs. (A-C) mRNA levels of (A) PPAR $\gamma$ 1 and its target genes (B) CD36 and (C) Lpl in primary bone marrow-derived macrophages (BMDMs) isolated from  $Tgr5^{+/+}$  or  $Tgr5^{-/-}$  mice and stimulated with LPS (10 ng/mL) for 24 hr, in presence or absence of the TGR5 agonist INT-777 (30 $\mu$ M). n=6/group. (D-I) mRNA levels of (D) LXR $\alpha$ , (E) LXR $\beta$  and their target genes (F) ApoE, (G) Abca1, (H) Abcg1 and (I) Srebp1c in primary bone marrow-derived macrophages (BMDMs) isolated from  $Tgr5^{+/+}$  or  $Tgr5^{-/-}$  mice and stimulated with LPS (10 ng/mL) for 24 hr, in presence or absence of the TGR5 agonist INT-777 (30  $\mu$ M). n=6/group. Results represent the mean±SEM. P values were calculated by using one-way ANOVA test followed by Bonferroni's post hoc analysis. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001 versus genotype and #p<0.001 versus control.