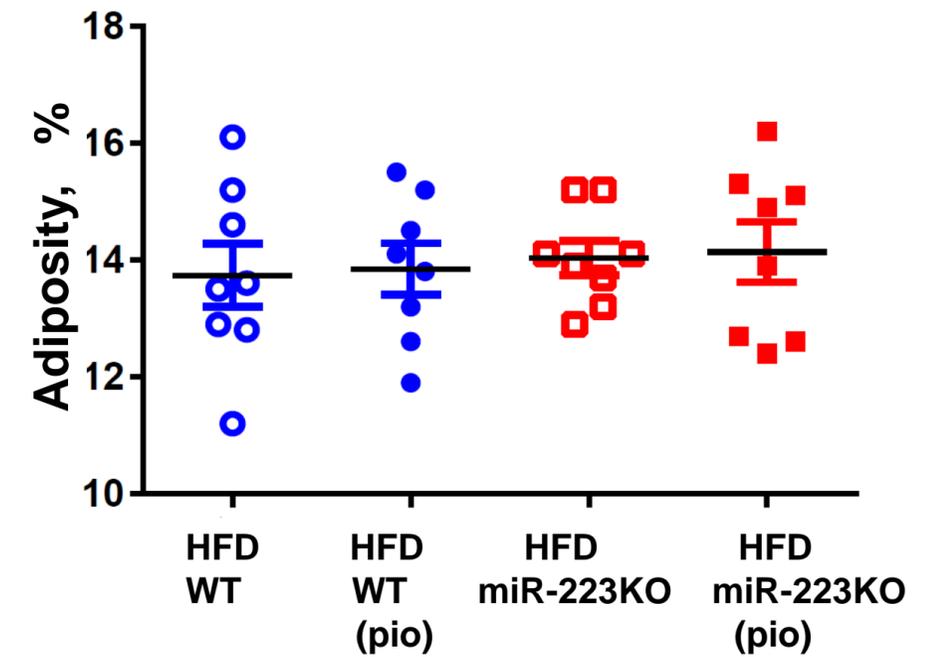
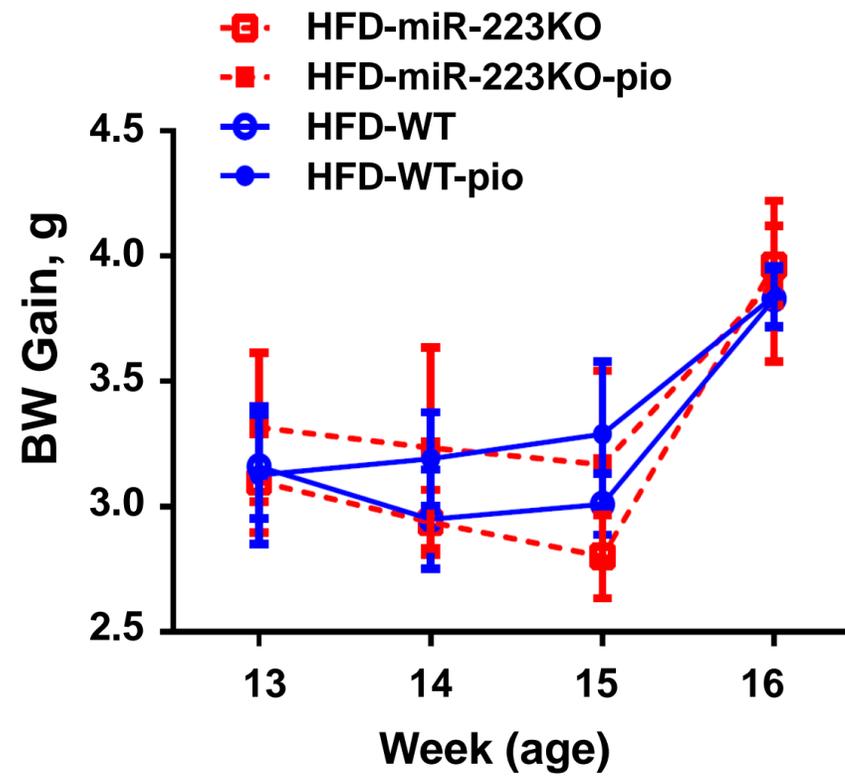
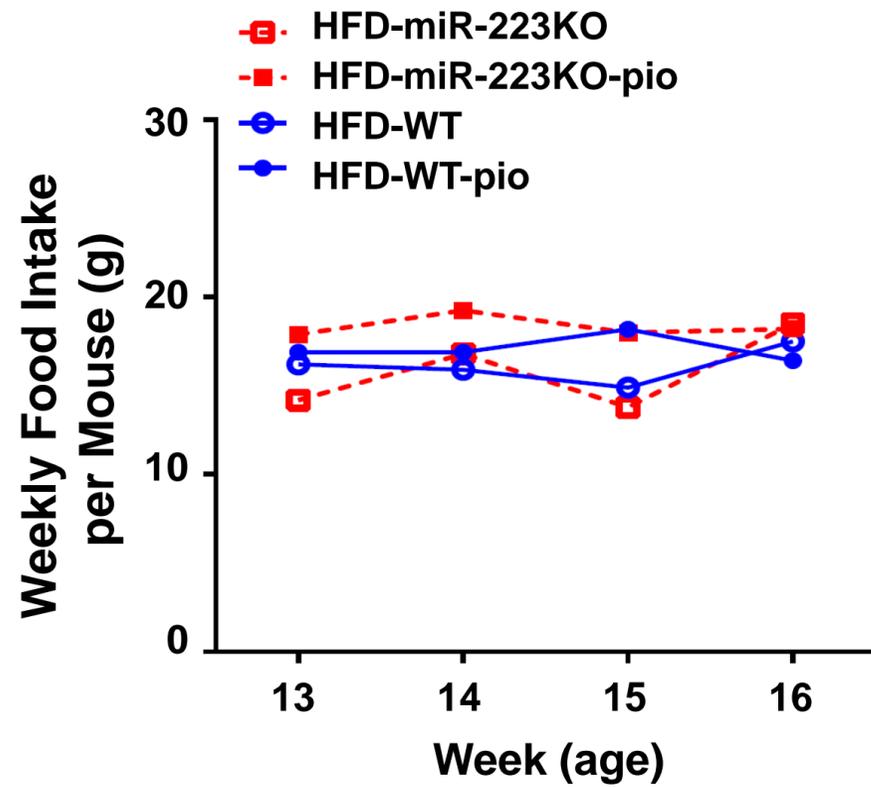
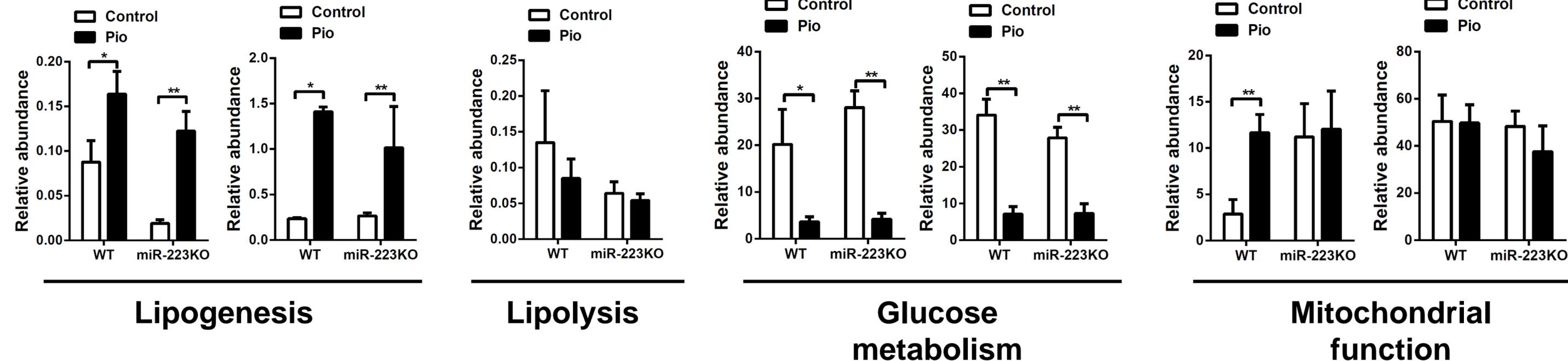


**Supplementary Figure S1. Restoring miR-223 rescues the M2 phenotypes of miR-223KO macrophages.**

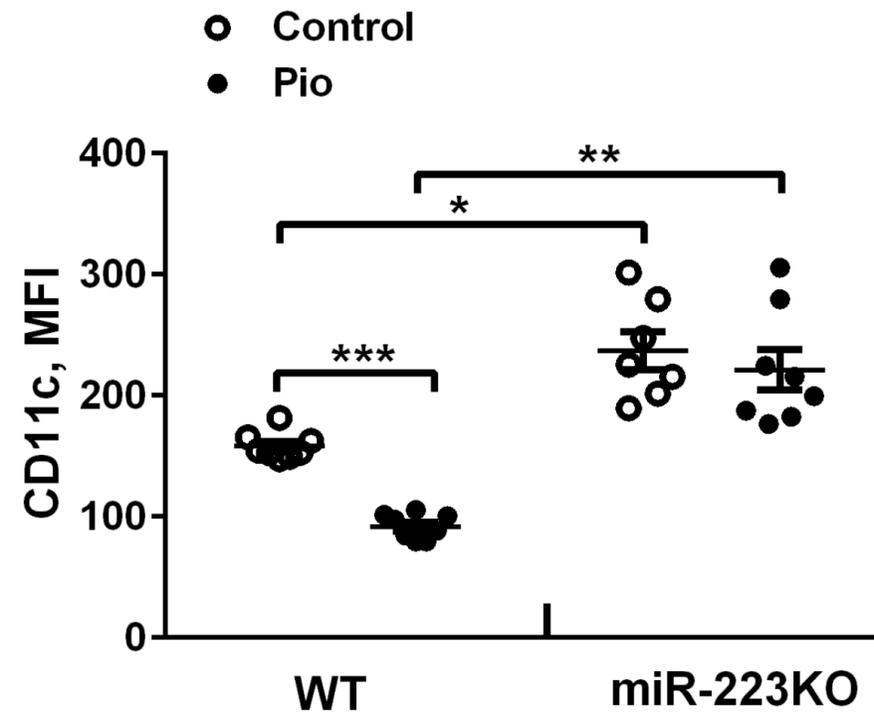
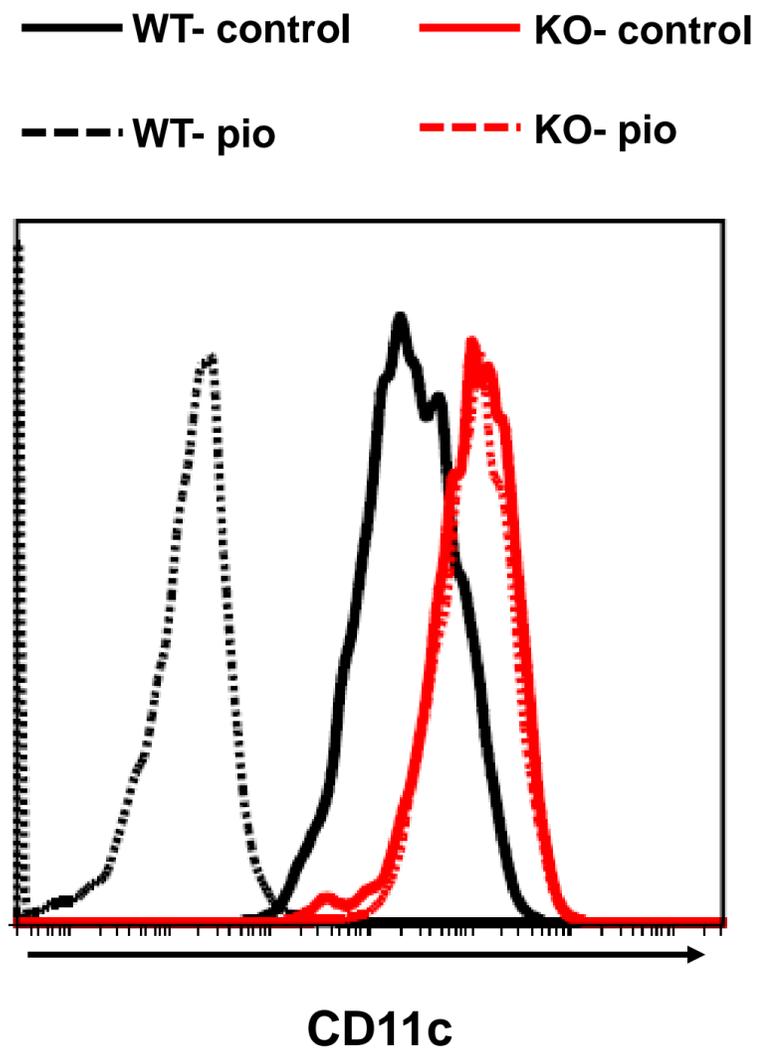
**A**, The expression of activation-related cell surface marker CD69 in WT or miR-223KO BMDMs with overexpression of miR-223 in the presence of pioglitazone IL4. ev, BMDMs transfected with empty vector; oe, BMDMs transfected with miR-223 overexpression construct. **B**, The expression of M2 related key genes *Arg1* and *Pgc1a* in miR-223KO BMDMs after introducing miR-223 overexpression construct for 48 hours. n=3. Data are presented as mean  $\pm$  SEM. \* $P$ <0.05, \*\* $P$ <0.01, \*\*\*\* $P$ <0.0001 One-way ANOVA with Bonferroni post-test (a), Student's *t* test (b).



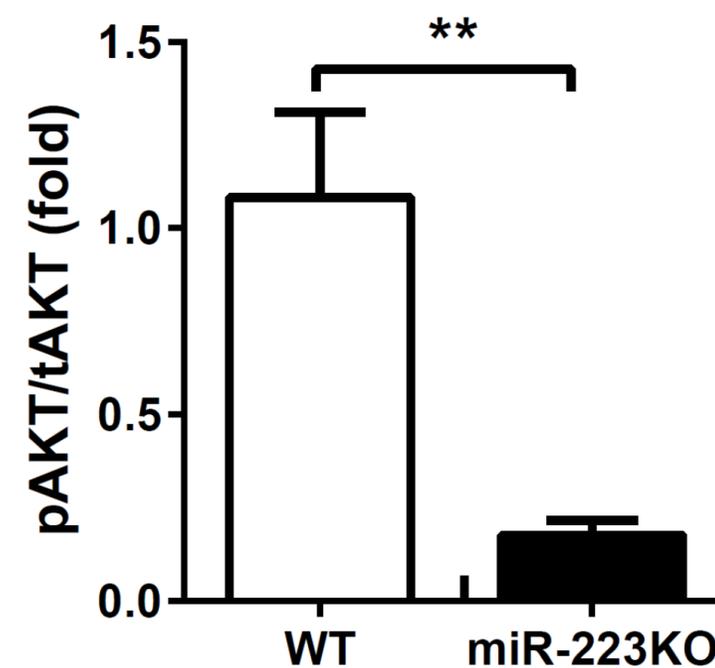
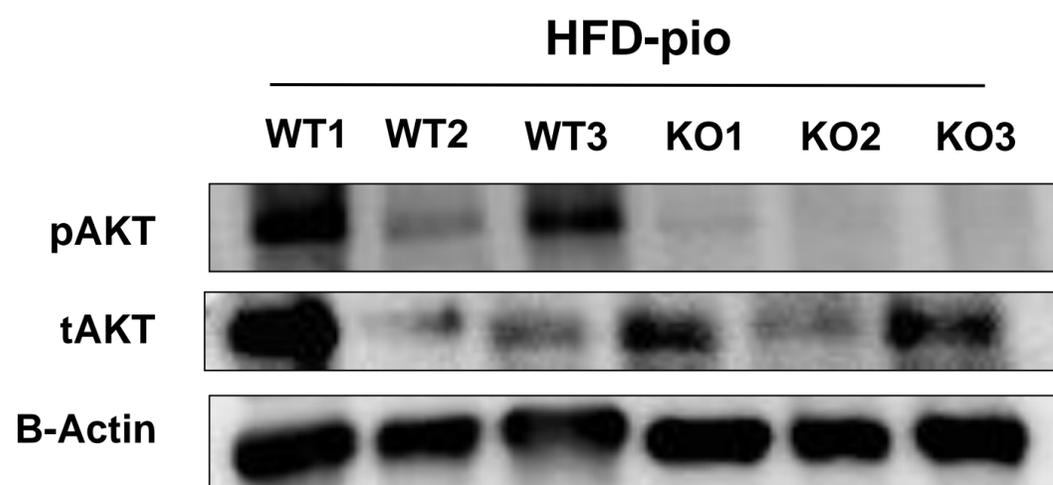
**Supplementary Figure S2. pioglitazone treatment does not affect food intake, body weight (BW) gain, and adiposity of HFD-fed mice.** Food intake, BW gain, and adiposity of HFD-fed WT or miR-223KO mice were measured during 4-week pioglitazone treatment (n=9-10). Data are presented as mean  $\pm$  SEM.

**A*****Acc******Fas*****B*****Cpt1*****C*****G6pase******Pepck*****D*****Scd1******Pgc1b***

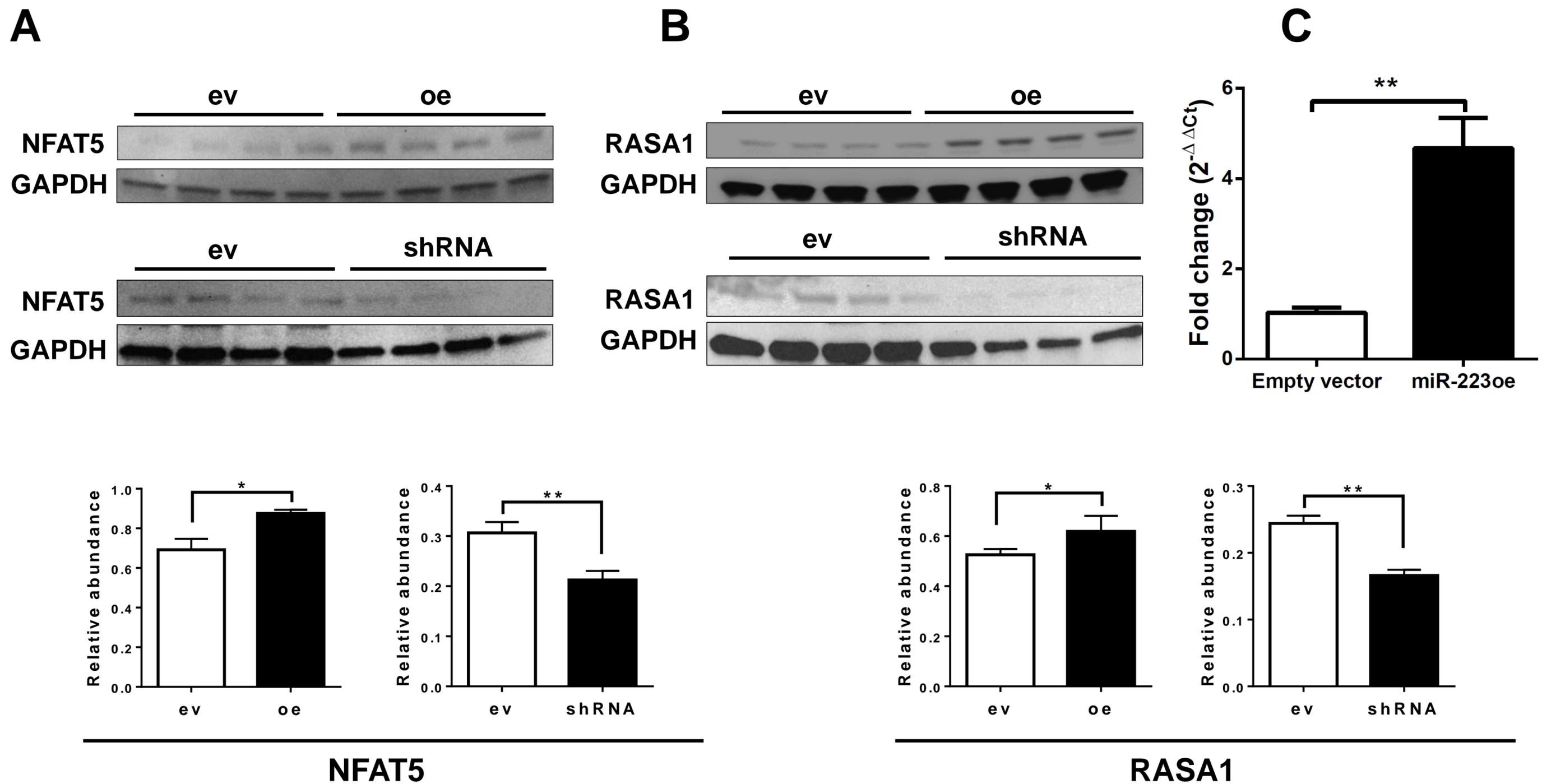
**Supplementary Figure S3. The effects of pioglitazone on key genes related to liver metabolic function of HFD-fed WT or miR-223KO mice.** The expression of key regulators for lipogenesis (A), lipolysis (B), glucose metabolism (C), mitochondrial function (D) was measured in the liver collected from HFD-fed WT or miR-223KO mice without (Control) or with pioglitazone (Pio) administration (normalized to  $\beta$ -actin; n=3). *Acc*, acetyl-CoA carboxylase; *Fas*, fatty acid synthetase; *Scd1*, stearoyl-CoA desaturase-1; *pgc1b*, peroxisome proliferator-activated receptor  $\gamma$ , coactivator 1 $\beta$ ; *Cpt1*, carnitine palmitoyltransferase 1; *G6pase*, glucose 6-phosphatase; *Pepck*, phosphoenolpyruvate carboxykinase. Data are means  $\pm$  SEM. \* $P$ <0.05, \*\* $P$ <0.01 Student's  $t$  test.



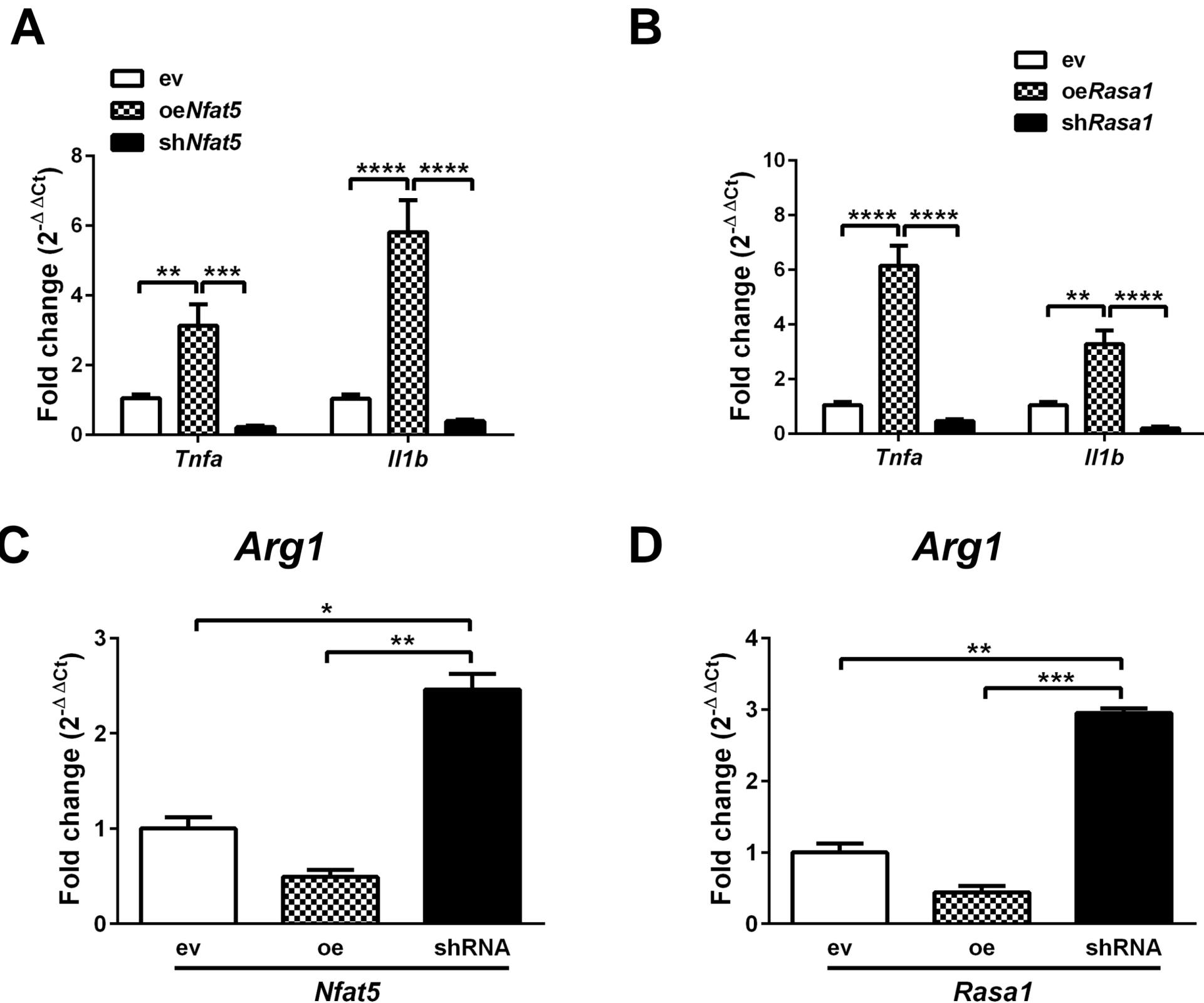
**Supplementary Figure S4. The effect of pioglitazone on expression of inflammatory marker CD11c in ATMs.** The level of CD11c on the cell surface of F4/80+CD11b+ ATMs in HFD-fed mice was measured by flow cytometry analysis with antibody against CD11c. n=7-9. Data are means  $\pm$  SEM. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$  One Way ANOVA with Bonferroni post-test .



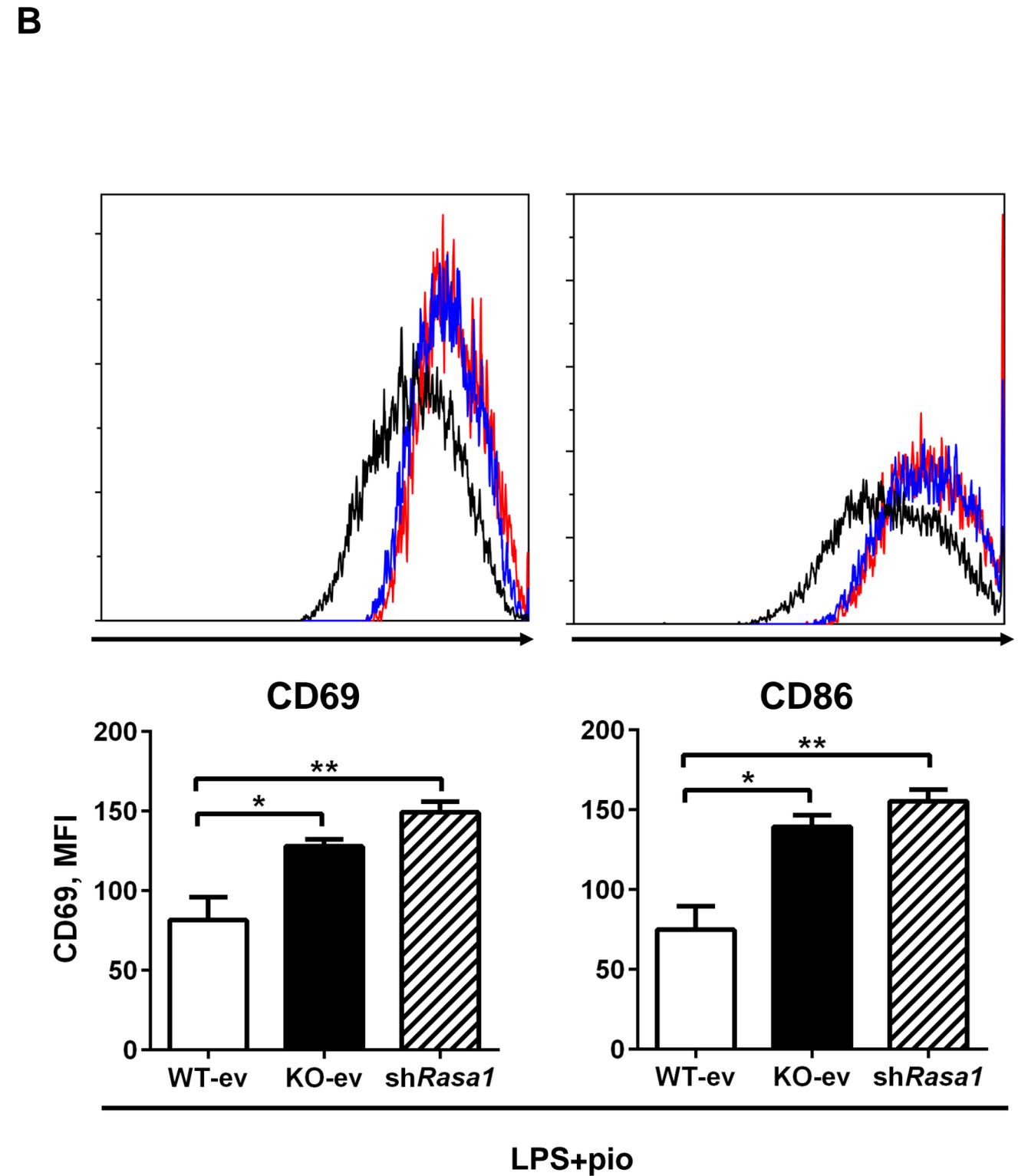
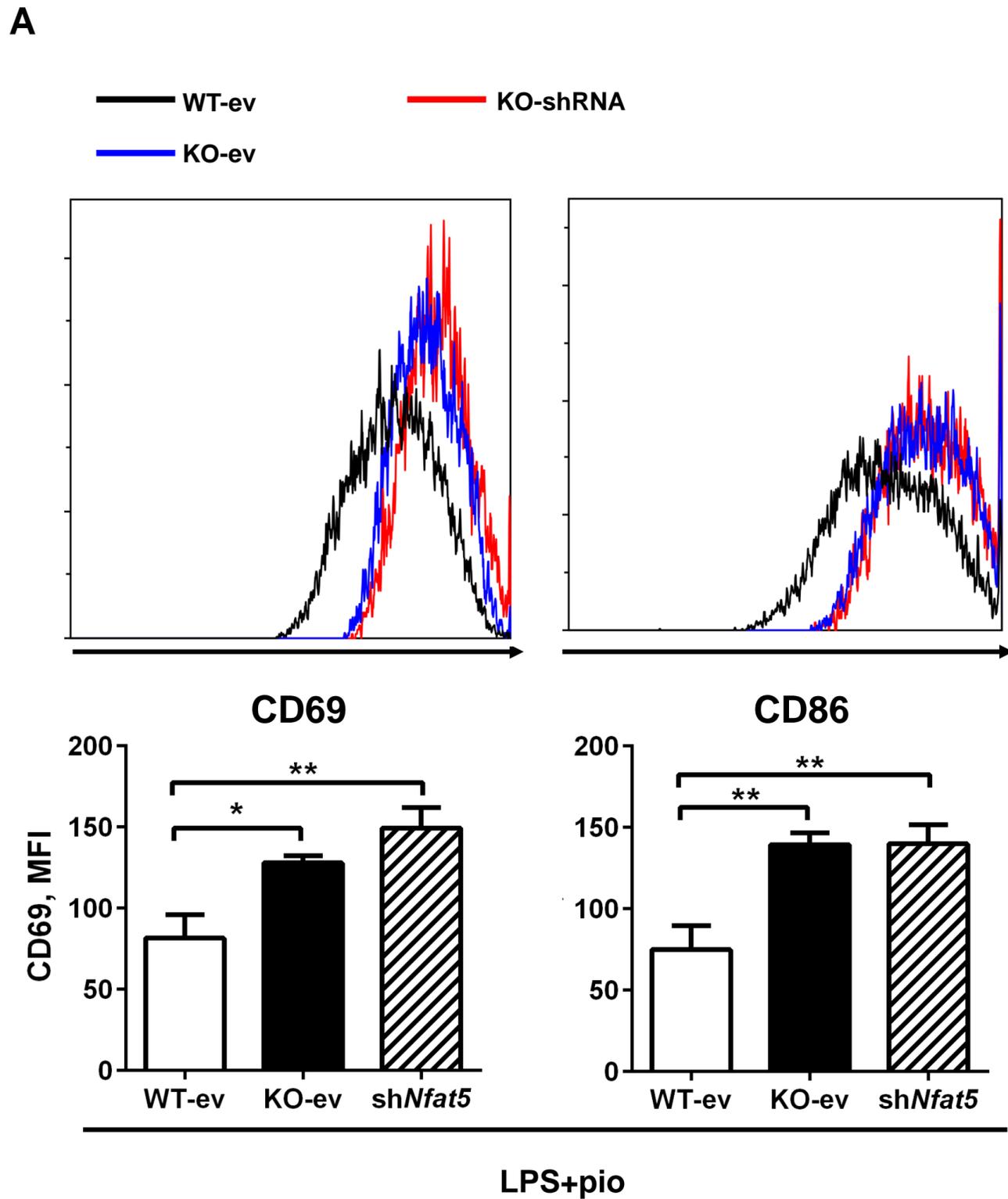
**Supplementary Figure S5. The effect of pioglitazone on activation of AKT signaling in visceral adipose tissues (VAT) of HFD-fed WT and miR-223KO mice.** The level of AKT activation in VAT of HFD-Fed mice with pioglitazone treatment was measured by western blots with antibodies against phosphorylated AKT (pAKT) and total AKT (tAKT).  $n=3$ . Data are means  $\pm$  SEM. \*\* $P<0.01$ , Student's  $t$  test.



**Supplementary Figure S6. The expression level of NFAT5, RASA1, or miR-223 after knockdown or overexpression assay.** Overexpression (oe) or knockdown of miR-223 target genes *Nfat5* (A) and *Rasa1* (B) expression in miR-223KO BMDMs by short hairpin RNA (shRNA) assay were validated by western blots. Cells transfected with empty vectors (ev) were used as control. C, The expression of miR-223 was validated by RT-PCR after ectopic expression assay in BMDMs. n=4. Data are means  $\pm$  SEM. \*\* $P < 0.001$ , Student's *t* test.



**Supplementary Figure S7. The effects of *Nfat5* and *Rasa1* on macrophage activation. A and B,** The expression of M1 activation-related key genes after knockdown of *Rasa1* or *Nfat5* in miR-223KO BMDMs. n=3. **C and D,** The expression of M2 activation-related gene *Arg1* in miR-223KO BMDMs with knockdown of miR-223 target genes. n=3. Data are means ± SEM. \**P*<0.05, \*\**P*<0.01, \*\*\**P*<0.001, \*\*\*\**P*<0.0001 One-way ANOVA with Bonferroni post-test .



**Supplementary Figure S8. The effect of pioglitazone on M1 activation of macrophages with knockdown of *Nfat5* or *Rasa1*.** **A and B,** The expression of M1 activation-related cell surface marker CD69 and CD86 in BMDMs after knockdown of *Nfat5* or *Rasa1* in the presence of pioglitazone (pio) and LPS. WT-ev, wild type BMDMs transfected with empty vector; KO-ev, miR-223KO BMDMs transfected with empty vector. n=4. Data are presented as mean  $\pm$  SEM. \* $P$ <0.05, \*\* $P$ <0.01 One-way ANOVA with Bonferroni post-test .