

## Supplementary material

### **Moderate L-lactate administration suppresses adipose tissue macrophage M1 polarization to alleviate obesity-associated insulin resistance**

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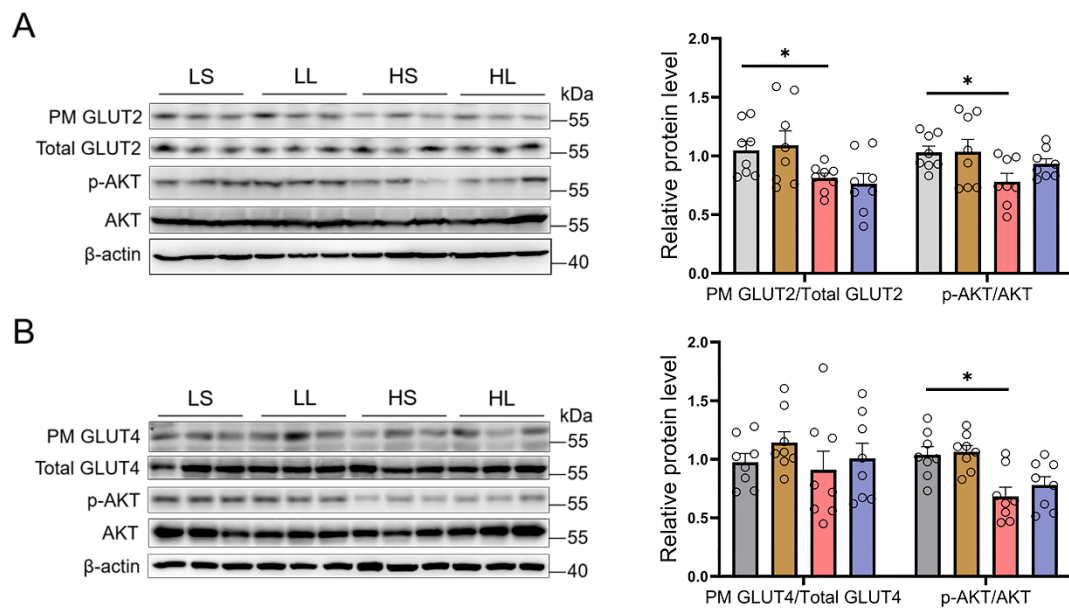
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The Supplemental material includes:

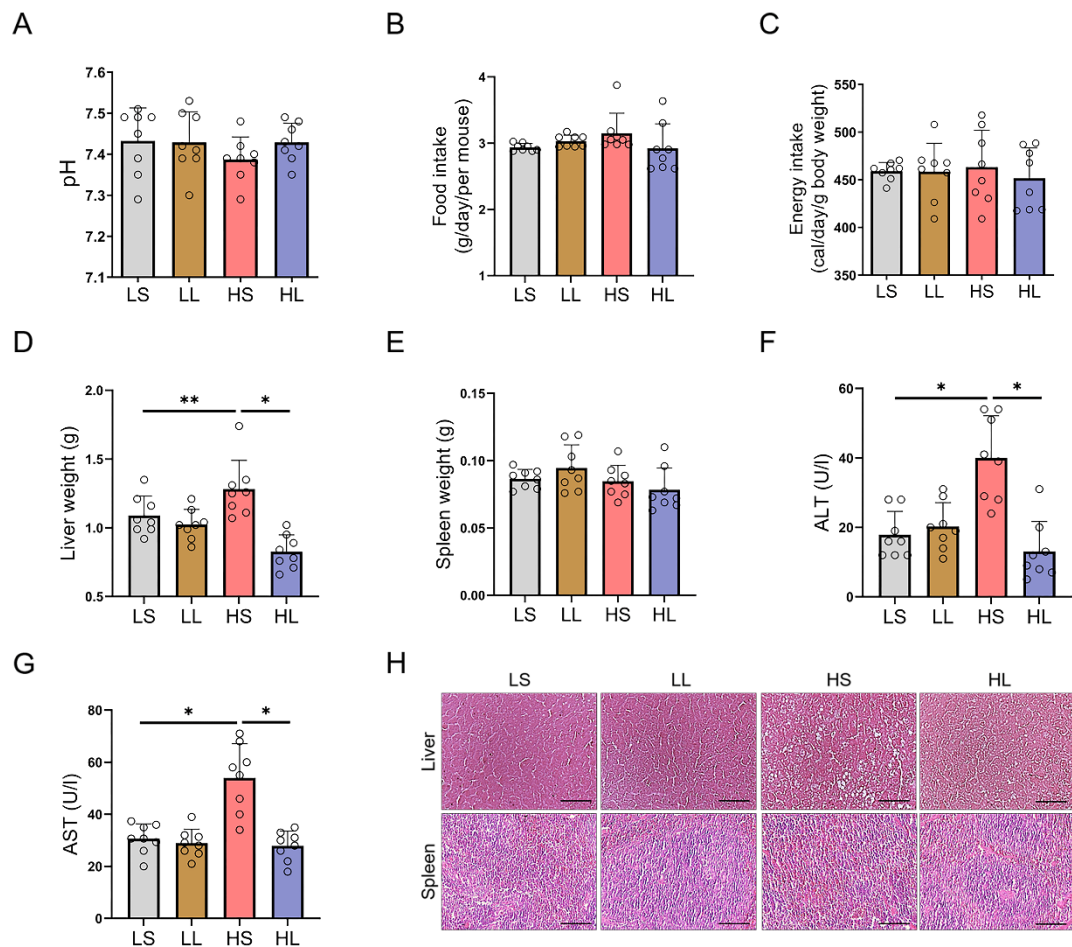
Fig. S1-10 (included in this file)

Table. S1-2 (included in this file)

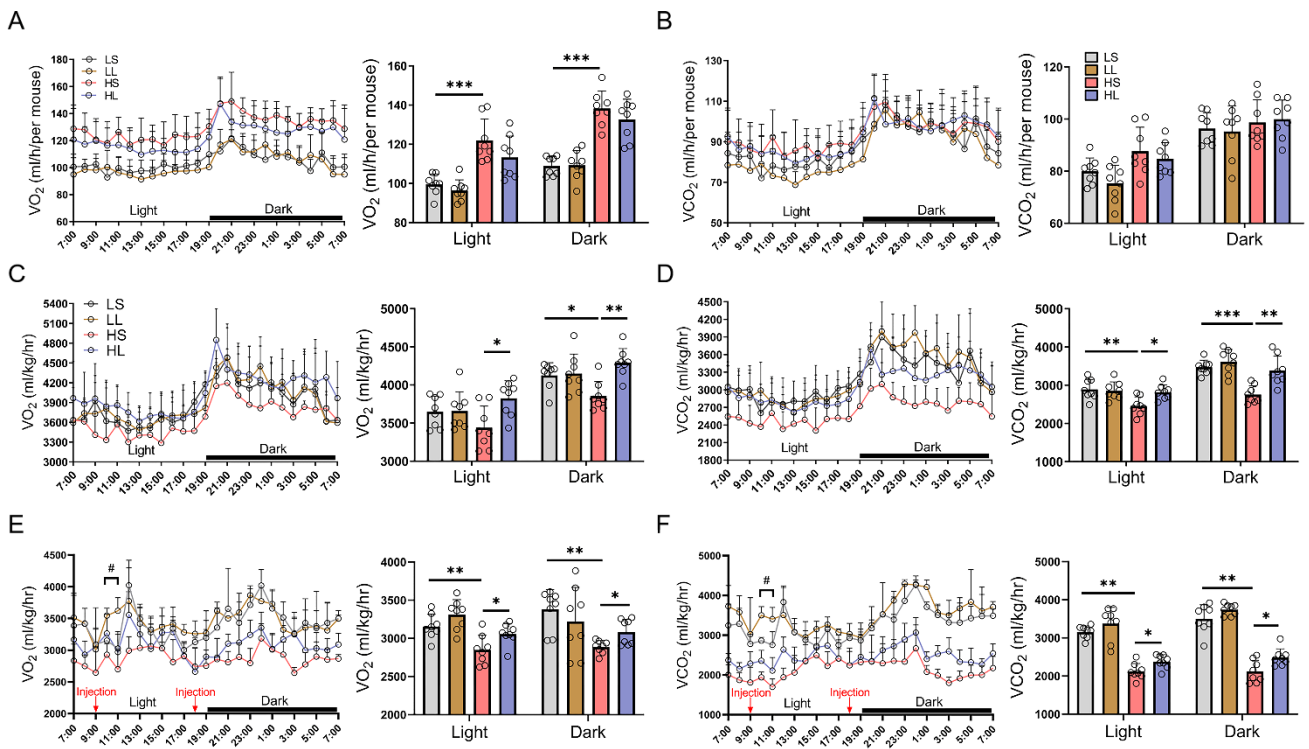
**Figure S1. The insulin signaling cascades in the liver and skeletal muscle.** The 6-week-old male mice were divided into four groups, including LS, LL, HS, and HL groups. **A** The GLUT2 translocation in the liver and quantification of plasma membrane GLUT2 to total GLUT2; Immunoblots for phosphorylation level of AKT in the liver. **B** The GLUT4 translocation in the skeletal muscle and quantification of plasma membrane GLUT4 to total GLUT4; Immunoblots for phosphorylation level of AKT in the skeletal muscle. Data are presented as means  $\pm$  SD of 8 mice per group, one-way ANOVA with Mann-Whitney test; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . LS, LFD-saline (i.p.); LL, LFD-lactate (i.p.); HS, HFD-saline (i.p.); HL, HFD-lactate (i.p.); PM, Plasma membrane.



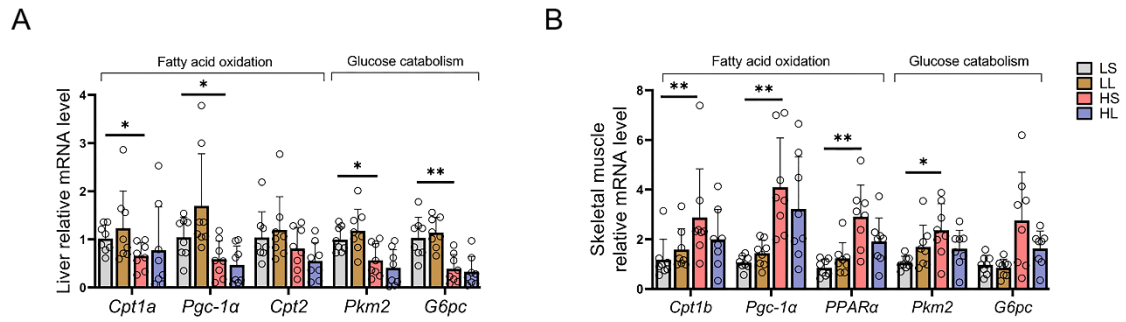
**Figure S2. Moderate L-lactate administration did not lead to the development of hyperlactatemia or lactic acidosis.** The 6-week-old male mice were divided into four groups, including LS, LL, HS, and HL groups. **A** The serum pH value. **B** Food intake. **C** Energy intake. **D** Liver weight. **E** Spleen weight. **F-G** The serum ALT (**F**) and AST (**G**). **H** The H&E staining of liver and spleen; scale bar, 100  $\mu$ m. Data are presented as means  $\pm$  SD of 8 mice per group, one-way ANOVA with Mann-Whitney test; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . LS, LFD-saline (i.p.); LL, LFD-lactate (i.p.); HS, HFD-saline (i.p.); HL, HFD-lactate (i.p.); ALT, Alanine transaminase; AST, Aspartate aminotransferase.



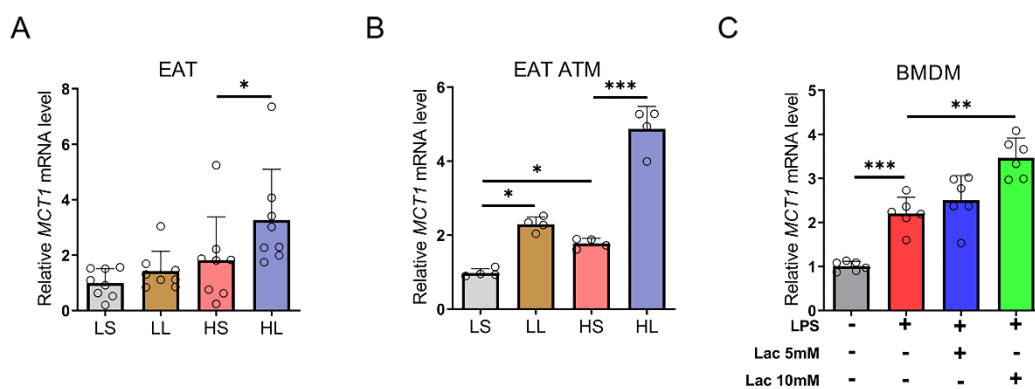
**Figure S3. The effects of moderate L-lactate administration on energy expenditure in LFD and HFD-fed mice.** At the age of 18 weeks, the metabolic capability in mice from four groups was measured. **A-B**  $VO_2$  (**A**) and  $VCO_2$  (**B**) per g body weight were detected without L-lactate injection during a 24-hour light-dark cycle. **C-D**  $VO_2$  (**C**) and  $VCO_2$  (**D**) per mouse were detected without L-lactate injection during a 24-h light-dark cycle. **E-F**  $VO_2$  (**E**) and  $VCO_2$  (**F**) per g body weight were detected during a 24-hour light-dark cycle and 400mg/kg L-lactate was injected at 9 a.m. and 6 p.m. Data are presented as means  $\pm$  SD of 8 mice per group, one-way ANOVA with Mann-Whitney test. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$  compared with HS group; # $p < 0.05$ , ## $p < 0.01$ , ### $p < 0.001$ , LL group was compared with LS group. LS, LFD-saline (i.p.); LL, LFD-lactate (i.p.); HS, HFD-saline (i.p.); HL, HFD-lactate (i.p.);  $VO_2$ , Oxygen consumption;  $VCO_2$ , Carbon dioxide production.



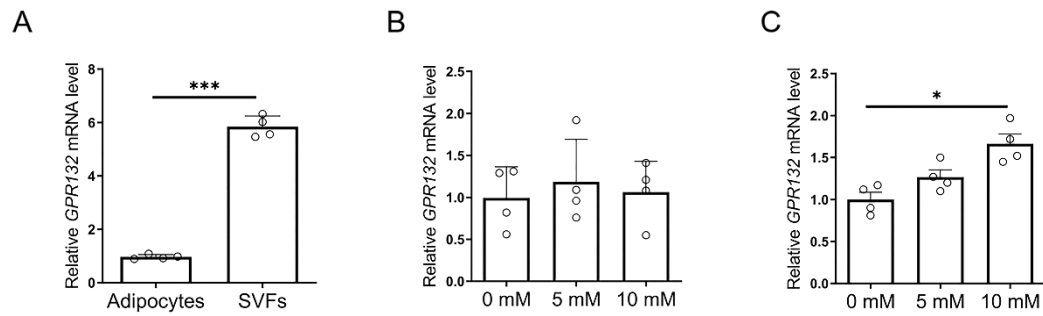
**Figure S4. The fatty acid oxidation and glucose catabolism in the liver and skeletal muscle.** The 6-week-old male mice were divided into four groups, including LS, LL, HS, and HL groups. **A-B** The relative genes of fatty acid oxidation and glucose catabolism in the liver (**A**) and skeletal muscle (**B**). Data are presented as means  $\pm$  SD of 8 mice per group, one-way ANOVA with Mann-Whitney test; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . LS, LFD-saline (i.p.); LL, LFD-lactate (i.p.); HS, HFD-saline (i.p.); HL, HFD-lactate (i.p.).



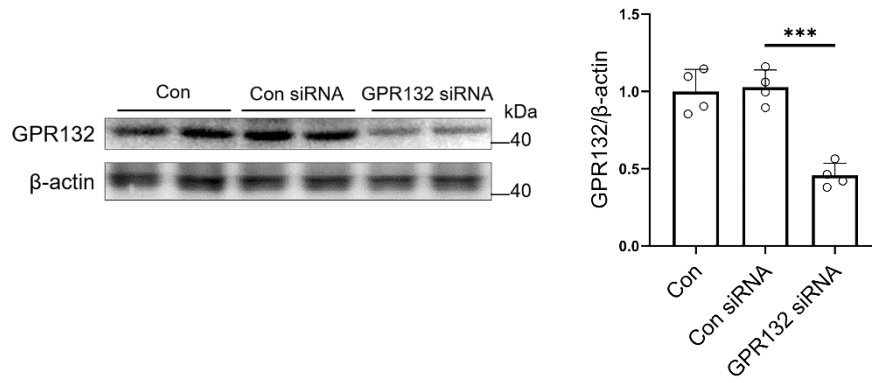
**Figure S5. The expression of MCT1 in ATMs and BMDMs after inflammatory stimulation.** The 6-week-old male mice were divided into four groups, including LS, LL, HS, and HL groups. **A** The mRNA level of *MCT1* in EATs. **B** The mRNA level of *MCT1* in ATMs from EATs. BMDMs were treated with vehicle or L-lactate (5 mM or 10 mM), and then LPS was added. **C** The mRNA level of *MCT1* in BMDMs. Data are presented as means  $\pm$  SD of 8 mice per group in vivo, 4 parallel cell samples per group in ATMs and 6 parallel cell samples per group in BMDMs, one-way ANOVA with Mann-Whitney test for mice and two-tailed Student's t-test for cell samples; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . LS, LFD-saline (i.p.); LL, LFD-lactate (i.p.); HS, HFD-saline (i.p.); HL, HFD-lactate (i.p.); Lac, Lactate; EAT, Epididymal adipose tissue; ATM, Adipose tissue macrophage.



**Figure S6. The expression of GPR132 in adipocytes and SVFs.** **A** The mRNA level of *GPR132* in adipocytes and SVFs from EATs in HFD-fed mice. **B** The mRNA level of *GPR132* in primary adipocytes after L-lactate treatment. **C** The mRNA level of *GPR132* in SVFs after L-lactate treatment. Data are presented as means  $\pm$  SD of 4 parallel samples per group, one-way ANOVA with Mann-Whitney test for mice and two-tailed Student's t-test for cell samples; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

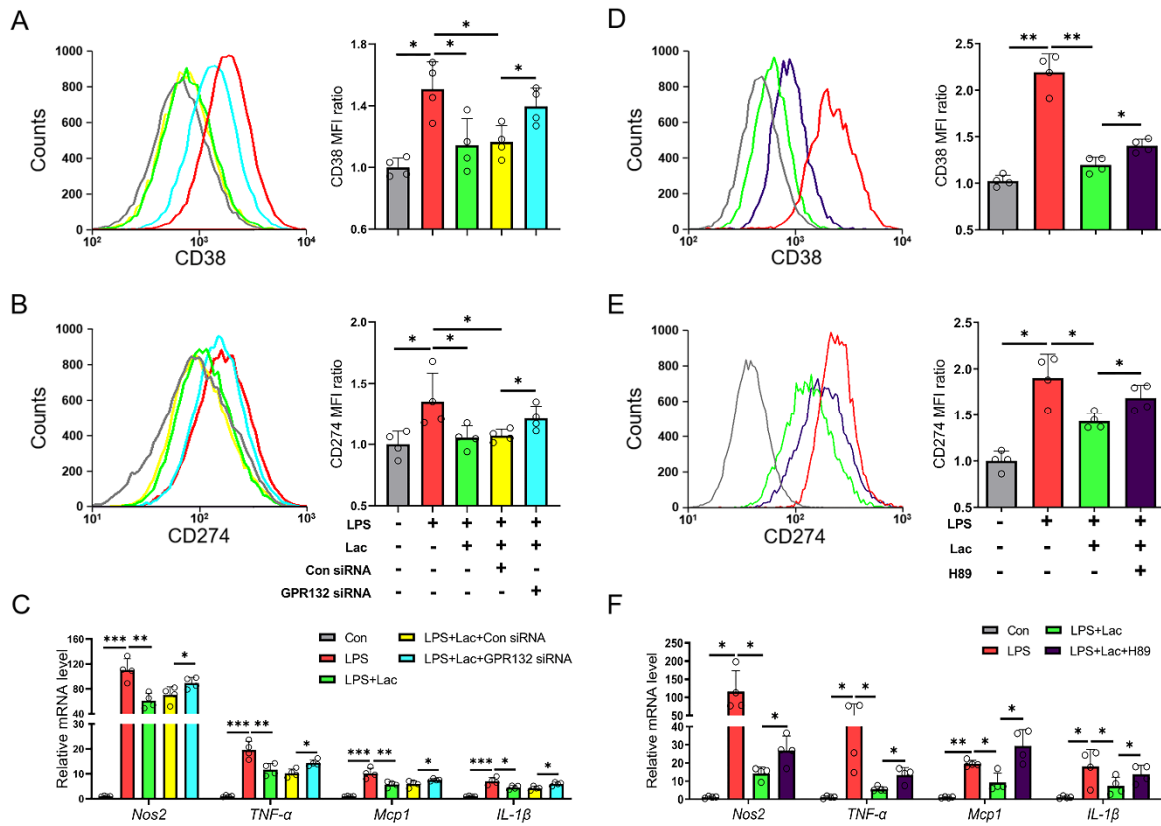


**Figure S7. siRNA-mediated knockdown of GPR132 in BMDMs.** BMDMs were transfected with a negative control siRNA or a siRNA targeting GPR132 for 48 h and lysed for immunoblots. Immunoblots for GPR132 in BMDMs and quantification of GPR132 to  $\beta$ -actin. Data are presented as means  $\pm$  SD of 4 parallel samples per group, two-tailed Student's t-test; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

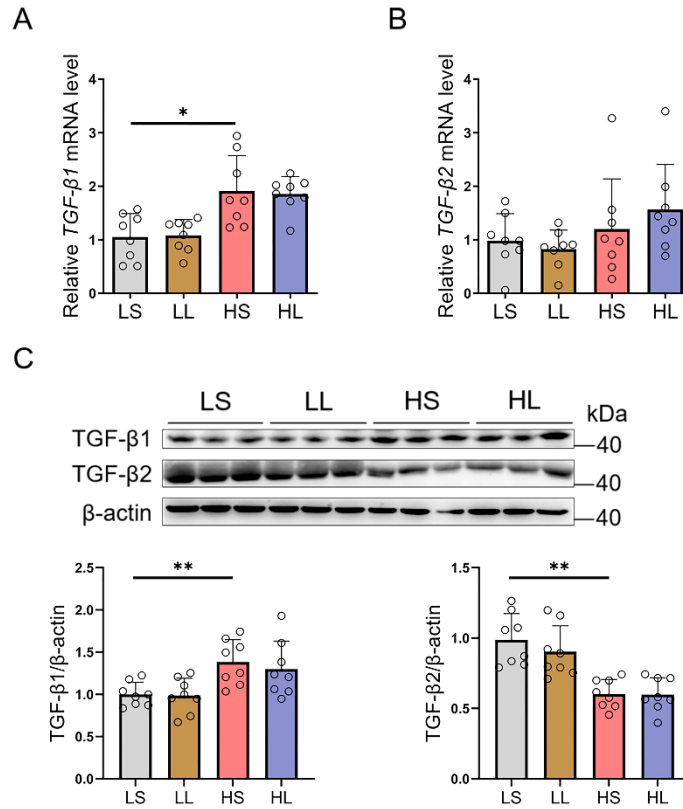




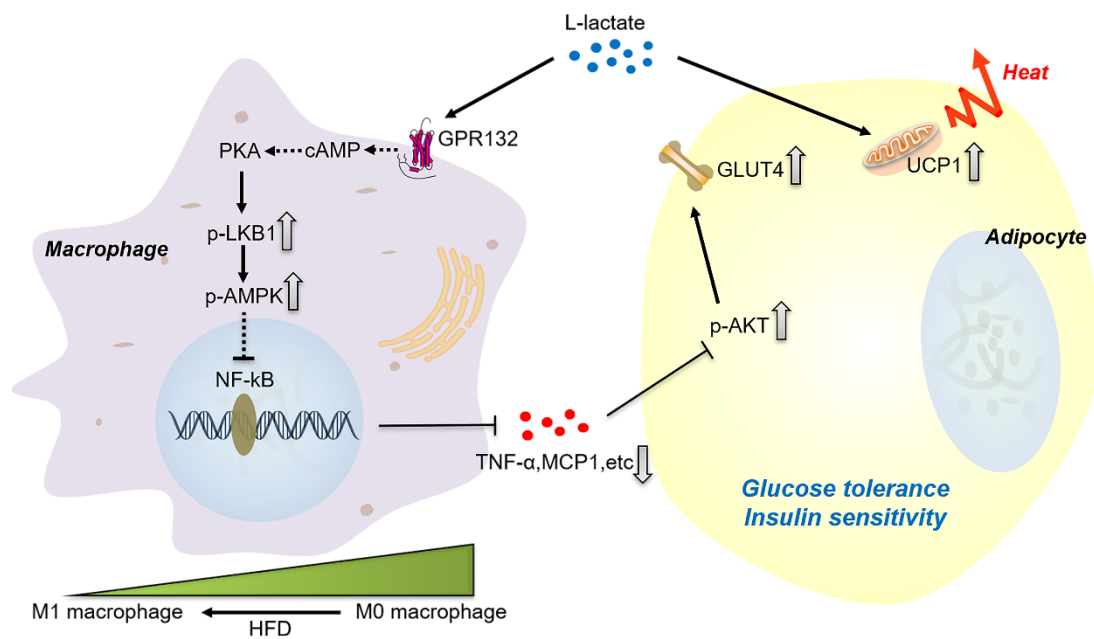
**Figure S8. GPR132-PKA participated in the inhibition of L-lactate on macrophage M1 polarization.** BMDMs were treated with vehicle, L-lactate, Con siRNA, or GPR132 siRNA, and then LPS was added. **A-B** Flow cytometry analyses of M1 surface marker CD38 (**A**) and CD274 (**B**). **C** The mRNA levels of pro-inflammatory genes. In another independent study, BMDMs were treated with vehicle or L-lactate, and then LPS was added. H89 was used as a PKA inhibitor. **D-E** Flow cytometry analyses of M1 surface marker CD38 (**D**) and CD274 (**E**). **F** The mRNA levels of pro-inflammatory genes. Data are presented as means  $\pm$  SD of 4 parallel samples per group, two-tailed Student's t-test; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . Lac, L-lactate.



**Figure S9. The expression of TGF- $\beta$ 1 and TGF- $\beta$ 2 in EATs.** The 6-week-old male mice were divided into four groups, including LS, LL, HS, and HL groups. **A-B** The mRNA levels of *TGF- $\beta$ 1* (**A**) and *TGF- $\beta$ 2* (**B**). **C** Immunoblots for TGF- $\beta$ 1 and TGF- $\beta$ 2 in EATs and quantification of TGF- $\beta$ 1 and TGF- $\beta$ 2 to  $\beta$ -actin. Data are presented as means  $\pm$  SD of 8 mice per group, one-way ANOVA with Mann-Whitney test; \* $p$  < 0.05, \*\* $p$  < 0.01, \*\*\* $p$  < 0.001. LS, LFD-saline (i.p.); LL, LFD-lactate (i.p.); HS, HFD-saline (i.p.); HL, HFD-lactate (i.p.).



**Figure S10. The proposed mechanism for moderate L-lactate administration improves adipose tissue insulin resistance.** On the one hand, moderate L-lactate administration elevates adipose tissue mitochondrial thermogenic protein UCP1 expression. On the other hand, L-lactate could bind to the GPR132 on the membrane of macrophages and activates the downstream PKA-LKB1-AMPK $\alpha$ 1 signal, which subsequently inhibits the NF- $\kappa$ B signal and the secretion of inflammatory cytokines. The suppression of moderate L-lactate administration on macrophage pro-inflammatory M1 polarization further promotes AKT phosphorylation and GLUT4 translocation in adipocytes. Collectively, moderate L-lactate administration activates adipose tissue macrophage GPR132-PKA-AMPK $\alpha$ 1 pathway to alleviate obesity-associated insulin resistance in mice.



**Table S1.** The primer sequences used for qRT-PCR.

Gene	Forward primer	Reverse primer
<i>β-actin</i>	CATCCGTAAAGACCTCTATGCCAAC	ATGGAGCCACCGATCCACA
<i>TNF-α</i>	ACGGCATGGATCTCAAAGAC	AGATAGCAAATCGGCTGACG
<i>IL-1β</i>	CTTCCCAGGGCATGTTAAG	ACCCTGAGCGACCTGTCTTG
<i>IFN-γ</i>	ATGAACGCTACACACTGCATC	CCATCCTTTTGCCAGTTCCTC
<i>MCP1</i>	CCCCAAGAAGGAATGGGTCC	GGTTGTGGAAAAGGTAGTGG
<i>F4/80</i>	TGACTCACCTTGTGGTCCTAA	CTTCCCAGAATCCAGTCTTTCC
<i>Nos2</i>	CCAAGCCCTCACCTACTTCC	CTCTGAGGGCTGACACAAGG
<i>Arg1</i>	CTCCAAGCCAAAGTCCTTAGAG	AGGAGCTGTCATTAGGGACATC
<i>Ucp1</i>	CACTCAGGATTGGCCTCTACG	GGGGTTTGATCCCATGCAGA
<i>Prdm16</i>	CCACCAGACTTCGAGCTACG	ACACCTCTGTATCCGTCAGCA
<i>Pgc-1α</i>	CCCTGCCATTGTTAAGACC	TGCTGCTGTTCTGTTTTTC
<i>Cidea</i>	TGACATTCATGGGATTGCAGAC	GGCCAGTTGTGATGACTAAGAC
<i>GPR132</i>	GTGCCATTGTGGATCATCTACA	CTCTCCAGTGCATAGACCACG
<i>MCT1</i>	TGTTAGTCGGAGCCTTCATTTC	CACTGGTCGTTGCACTGAATA
<i>TGF-β1</i>	CTCCCGTGGCTTCTAGTGC	GCCTTAGTTTGGACAGGATCTG
<i>TGF-β2</i>	CTTCGACGTGACAGACGCT	GCAGGGGCAGTGTAAACTTATT
<i>Cpt1a</i>	CTCCGCCTGAGCCATGAAG	CACCAGTGATGATGCCATTCT
<i>Cpt2</i>	CAGCACAGCATCGTACCCA	TCCCAATGCCGTTCTCAAAT
<i>Pkm2</i>	GCCGCCTGGACATTGACTC	CCATGAGAGAAATTCAGCCGAG
<i>G6pc</i>	CGACTCGCTATCTCCAAGTGA	GTTGAACCAGTCTCCGACCA
<i>CPT1b</i>	GCACACCAGGCAGTAGCTTT	CAGGAGTTGATTCCAGACAGGTA
<i>PPARα</i>	AGAGCCCCATCTGTCTCTC	ACTGGTAGTCTGCAAAACCAA

**Table S2.** The antibodies used in this study.

	<b>Antibody</b>	<b>Source</b>	<b>Catalog number</b>	<b>Dilution</b>
<b>Western blot</b>	$\beta$ -actin	Cell Signaling Technology	4967	5000
	AKT	Cell Signaling Technology	4691S	5000
	Phosphor-AKT	Cell Signaling Technology	9271S	1000
	AMPK $\alpha$ 1	Cell Signaling Technology	2532S	1000
	Phosphor-AMPK $\alpha$ 1	Cell Signaling Technology	2531S	1000
	LKB1	Cell Signaling Technology	3047	1000
	Phosphor-LKB1	Cell Signaling Technology	3482	1000
	UCP1	Abcam	ab209483	5000
	TGF- $\beta$ 1	Abcam	ab215715	1000
	TGF- $\beta$ 2	Abcam	ab36495	1000
	GPR132	Santa	sc-137112	200
	GLUT2	Proteintech	66889-1-Ig	5000
	GLUT4	Millipore	07-1404	1000
<b>Flow cytometry</b>	APC anti-mouse F4/80	BioLegend	123116	200
	FITC anti-mouse CD206	BioLegend	141704	300
	PE anti-mouse CD11c	eBioscience	12-0114	200
	APC anti-mouse CD274	BioLegend	124312	200
	PE anti-mouse CD38	BioLegend	102708	200