

Supplementary

# Delayed Exercise Training Improves Obesity-Induced Chronic Kidney Disease by Activating AMPK Pathway in High-Fat Diet-Fed Mice

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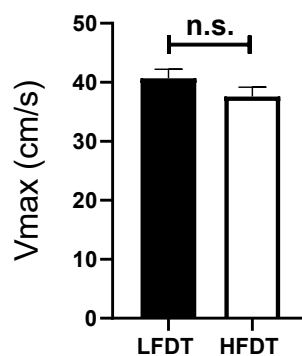
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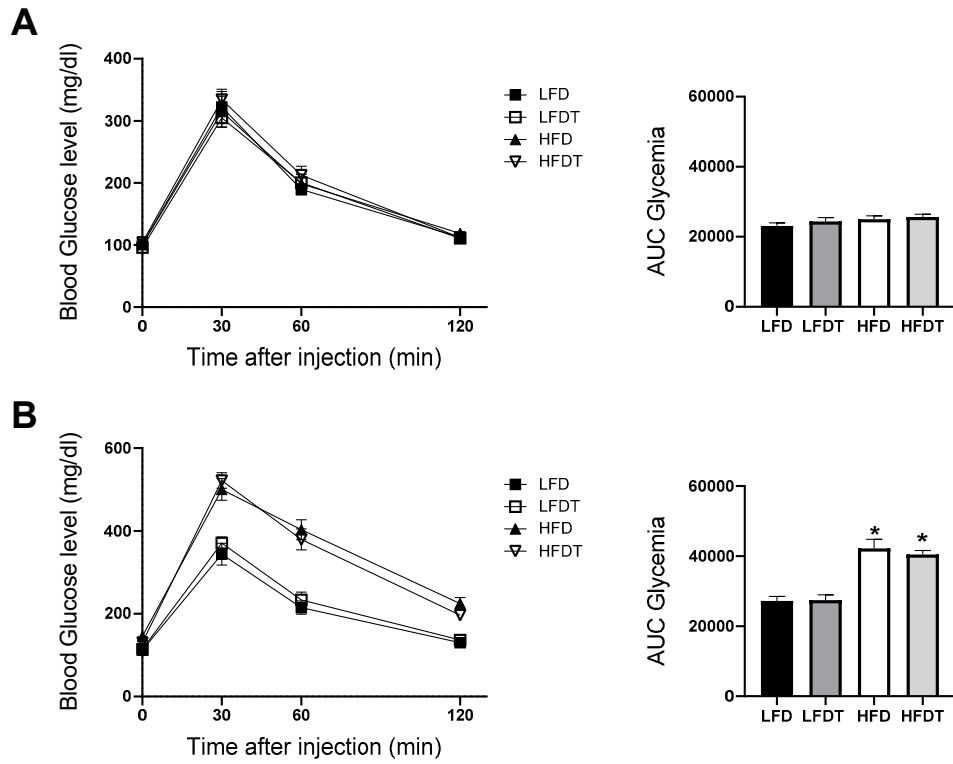
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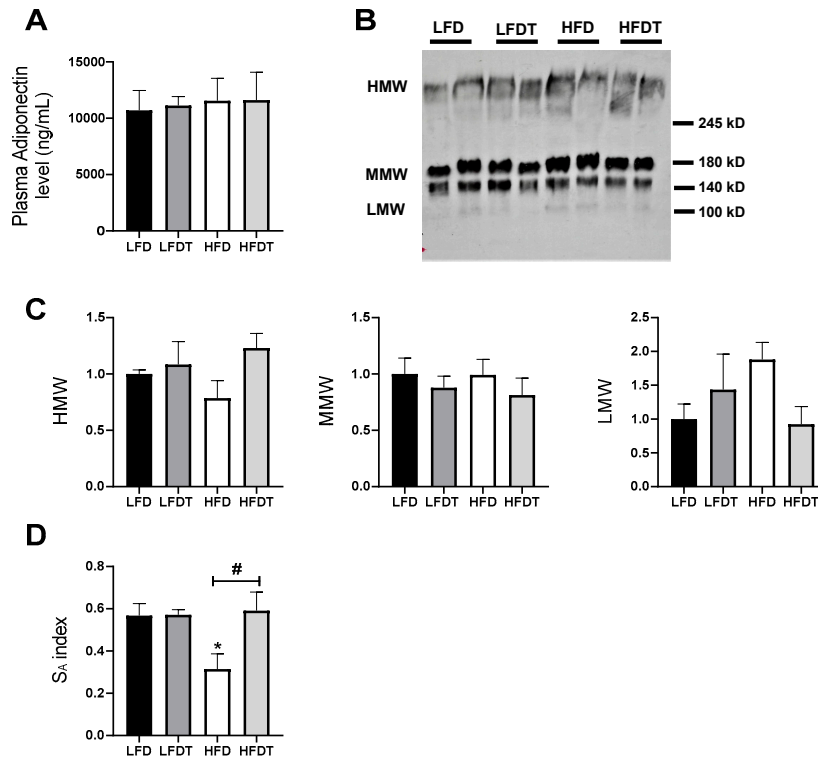
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**Figure S1.** Effects of endurance exercise training on running velocity. Representation of maximal running velocity performance after 14 weeks on diet. *t*-test. Data are presented as means  $\pm$  SEM. *ns* = non-significant. *n*=10 in each group.



**Figure S2.** Effects of LFD and HFD on glucose tolerance in mice. Glucose tolerance test at week 0 (**A**) and week 12 (**B**). Fasted mice were submitted to an intraperitoneal injection of glucose (2 g/ kg b.w.). Glycemia was measured before (0) and 30, 60 and 120 min after injection. Histogram represents the area under the curve (AUC) of glycemia from 0 to 120 min



**Figure S3.** Total plasma adiponectin and adiponectin multimers distribution analysis at week 20. **A.** Adiponectin plasma level. The total plasma Adiponectin concentration was measured using indirect ELISA. **B.** Representative immunoblots of Adiponectin multimers expression in plasma sample. **C.** Relative densitometry of the immunoblots representing respectively HMW (High Molecular Weight), MMW (Medium Molecular Weight) and LMW (Low Molecular Weight) multimers of Adiponectin normalized with total Adiponectin protein level. **D.** SA index was calculated as the ratio  $HMW/(HMW + LMW)$ . Data are presented as means  $\pm$  SEM. \*  $P \leq 0.05$  versus LFD #  $P \leq 0.05$  versus HFD.  $n=6-8$  in each group.

**Table S1.** Primer sequences for RT-qPCR analysis of mRNA expression

<b>Gene</b>		<b>Primer Sequences (5'-3')</b>
<i>COL1</i>	Fw	CTTGCCCCATTCATTTGTCT
	Rv	GCAGGTTACCTACTCTGTTCT
<i>COL3A3</i>	Fw	TGAGTCGAATTGGGGAGAAT
	Rv	TCCCCTGGAATCTGTGAATC
<i>TGFβ</i>	Fw	TGGAGCAACATGTGGAACTC
	Rv	GTCAGCAGCCGTTACCA
<i>MCP-1</i>	Fw	CTTCTGGGCCTGCTGTTCA
	Rv	CCAGCCTACTCATTGGGATCA
<i>IL1β</i>	Fw	AGTTGACGGACCCCAAAAG
	Rv	AGCTGGATGCTCTCATCAGG
<i>TNFα</i>	Fw	TACTGAACTTCGGGGTGATTGGTCC
	Rv	CAGCCTTGTCCTTGAAGAGAACC
<i>IL6</i>	Fw	GCTACCAAACCTGGATATAATCAGGA
	Rv	CCAGGTAGCTATGGTACTCCAGAA
<i>ACC (Acaca)</i>	Fw	ATGGGCGGAATGGTCTCTTTC
	Rv	TGGGGACCTTGTCTTCATCAT
<i>FAS (FASN)</i>	Fw	GGAGGTGGTGATAGCCGGTAT
	Rv	TGGGTAATCCATAGAGCCCAG
<i>CPT-1 (CPT1a)</i>	Fw	CTCCGCCTGAGCCATGAAG
	Rv	CACCAGTGATGATGCCATTCT
<i>18S</i>	Fw	CGCCGCTAGAGGTGAAATTCT
	Rv	CGAACCTCCGACTTTCGTTCT

**Table S2.** Effects of delayed EET on systolic, diastolic and mean blood pressure in mice fed a LFD, LFDT or HFD and HFDT.

	<b>LFD</b>	<b>LFDT</b>	<b>HFD</b>	<b>HFDT</b>
<b><i>Systolic blood pressure (mmHg)</i></b>	128,6 ± 10,99	138,0 ± 4,988	119,5 ± 10,86	119,3 ± 5,850
<b><i>Diastolic blood pressure (mmHg)</i></b>	106,8 ± 10,60	117,6 ± 2,584	96,33 ± 8,750	101,7 ± 6,035
<b><i>Mean blood pressure (mmHg)</i></b>	117,7 ± 10,78	127,8 ± 3,763	107,9 ± 9,768	110,5 ± 5,570

Measurement were performed during the last week of the experimental protocol (week 20). Systemic, diastolic and mean blood pressures were measured using a non-invasive CODA tail-cuff blood pressure occlusion system (Kent Scientific, Torrington, USA). During the week 20, measurements were taken for each animal that were acclimatized for a 1-hour period before experiments into restraining chambers. The animals were placed onto a preheated pad maintained at 30°C in a designed quiet area and blood pressure measurements were initiated when tail temperature reached 30°C (measured using an infrared sensor) and recorded at least 5 times. Mice were acclimated for at least 3 consecutive days before baseline blood pressure measurements. No statistical difference was found by One-way ANOVA analysis.  $n=5$  in each group.

**Table S3.** Effects of delayed EET on renal gene expression in mice fed a LFD, a LFDT, a HFD and a HFDT.

	<b>LFD</b>	<b>LFDT</b>	<b>HFD</b>	<b>HFDT</b>
<b><i>Lipid metabolism markers</i></b>				
<i>ACC</i>	1,000 ± 0,0918	0,9636 ± 0,1356	1,067 ± 0,1404	1,028 ± 0,0796
<i>FAS</i>	1,000 ± 0,0265	1,023 ± 0,1480	1,218 ± 0,2496	1,112 ± 0,1884
<i>CPT-1</i>	1,000 ± 0,0326	0,9017 ± 0,1308	0,8483 ± 0,1201	1,222 ± 0,2337

Real-time quantitative qPCR for Acetyl-CoA carboxylase (*ACC*), Fatty acid synthase (*FAS*) and Carnitine palmitoyltransferase I (*CPT1*). mRNA expressions were performed on kidney tissue from LFD, LFDT, HFD and HFDT mice normalized against 18S. Statistical analyses were performed by one-way ANOVA followed by Newman-Keuls post hoc test. Data are presented as means ± SEM. \*  $P \leq 0.05$  versus LFD #  $P \leq 0.05$  versus HFD.  $n=6$  in each group.