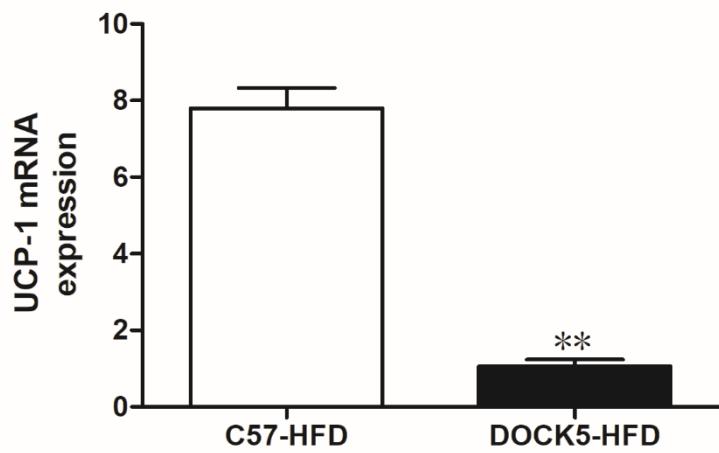


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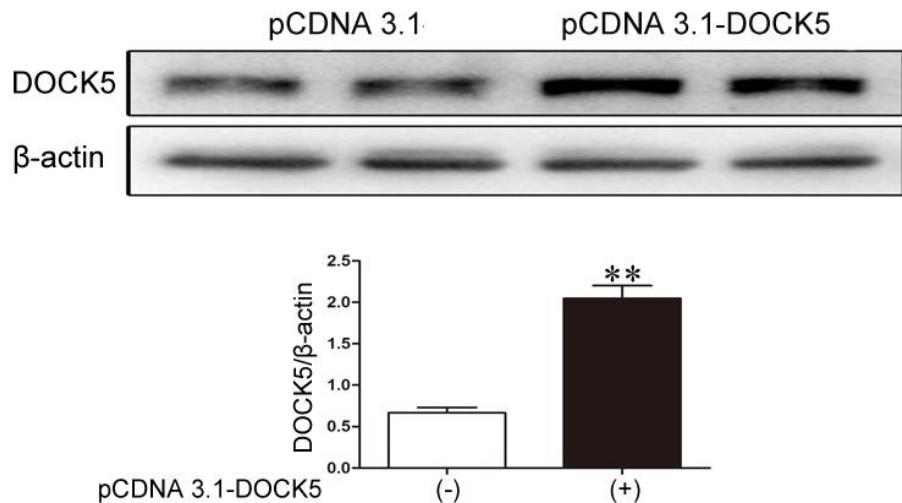
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Appendix Figure S1 UCP1 mRNA expression in the BAT of WT and DOCK5^{-/-} mice.

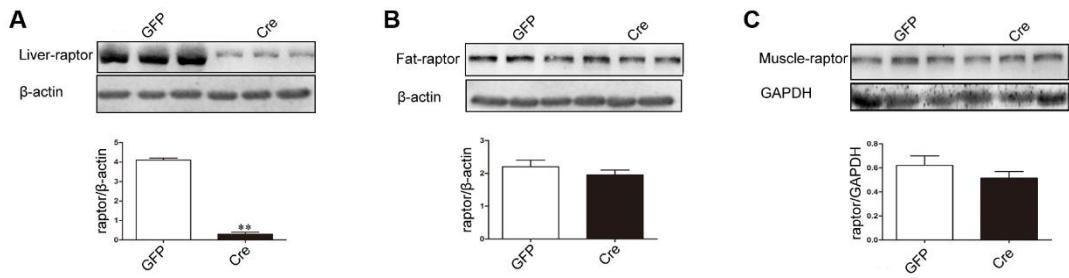
Data are expressed as the mean \pm SEM. P-values were determined with *t*-test, ** $p < 0.01$ vs.

WT.



Appendix Figure S2 DOCK5 protein expression in Hep1-6 cells.

Hep1-6 cells were infected with pCDNA3.1-DOCK5 (+) or pCDNA3.1 (-) for 48 h. Western blotting was conducted to examine DOCK5 protein expression. Data are presented as the mean \pm SEM. P-values were determined with *t*-test .** $p < 0.01$ vs. pCDNA3.1-DOCK5 (-).



Appendix Figure S3 Raptor protein expression in AAV8-Cre- or AVV8-GFP-treated Raptor^{flox/flox} mice.

The levels of Raptor protein expression in the

A liver

B adipose

C muscle. Data are presented as the mean ± SEM. P-values were determined with *t*-test. ** *p*

< 0.01 vs. GFP group.

Appendix Table S1 Biochemical parameters in WT and DOCK5^{-/-} mice

Parameters	3 months				16 months			
	SD -WT	SD-DOCK5 ^{-/-}	HFD -WT	HFD -DOCK5 ^{-/-}	SD -WT	SD -DOCK5 ^{-/-}	HFD -WT	HFD-DOCK5 ^{-/-}
FBG (mmol/L)	5.20 ± 0.33	5.05 ± 0.49	6.80 ± 0.52	7.68 ± 0.28**	6.33 ± 0.66	6.10 ± 0.32	11.6 ± 1.3	14.1 ± 0.9**
FINS (mIU/L)	11.2 ± 1.7	12.8 ± 3.3	31.9 ± 1.1	47.3 ± 5.1**	18.3 ± 2.5	16.5 ± 3.1	77.4 ± 6.30	88.5 ± 3.4**
TC (mmol/L)	2.18 ± 0.17	2.21 ± 0.14	2.99 ± 0.39	5.11 ± 0.57**	2.43 ± 0.24	2.39 ± 0.32	6.84 ± 0.73	8.20 ± 0.87*
TG (mmol/L)	0.49 ± 0.10	0.48 ± 0.10	0.75 ± 0.09	0.95 ± 0.11*	0.55 ± 0.07	0.53 ± 0.04	1.80 ± 0.31	2.46 ± 0.22**
FFA (mmol/L)	0.18 ± 0.03	0.22 ± 0.04	0.51 ± 0.14	0.82 ± 0.15*	0.23 ± 0.01	0.27 ± 0.03	1.51 ± 0.20	2.13 ± 0.32*

SD, standard diet; HFD, high fat diet; FBG, fasting blood glucose; FIns, fasting plasma insulin; TC, total cholesterol; TG, triglyceride; FFA, free fatty acids.

Data are presented as means ± SD, P-values were determined with *t*-test .**p* < 0.05, ***p* < 0.01 vs. HFD-WT.

Appendix Table S2 Biochemical parameters in WT and DOCK5^{-/-} mice during the HEC

Index	SD-WT (N = 6)	SD-DOCK5 ^{-/-} (N = 6)	HFD-WT (N = 6)	HFD-DOCK5 ^{-/-} (N = 6)
Basal-state				
BG (mmol/L)	5.43 ± 0.15	5.91 ± 0.19	7.02 ± 0.16	7.92 ± 0.31**
Ins (mIU/L)	11.0 ± 1.7	11.2 ± 2.0	30.9 ± 2.0	48.8 ± 3.1**
TC (mmol/L)	2.21 ± 0.14	2.23 ± 0.09	2.87 ± 0.21	5.06 ± 0.51**
TG (mmol/L)	0.47 ± 0.07	0.49 ± 0.09	0.73 ± 0.06	0.97 ± 0.10**
FFA (mmol/L)	0.21 ± 0.04	0.23 ± 0.05	0.48 ± 0.07	0.79 ± 0.10**
Steady-state				
BG (mmol/L)	5.91 ± 0.19	6.02 ± 0.20	5.81 ± 0.36	5.98 ± 0.19
Ins (mIU/L)	80.5 ± 10.7	108.6 ± 19.3	240.6 ± 26.9	356.3 ± 32.1**
TC (mmol/L)	1.05 ± 0.10	1.11 ± 0.08	1.39 ± 0.12	2.41 ± 0.06**
TG (mmol/L)	0.16 ± 0.07	0.18 ± 0.04	0.37 ± 0.09	0.52 ± 0.08*
FFA (mmol/L)	0.11 ± 0.06	0.13 ± 0.04	0.24 ± 0.06	0.39 ± 0.07**

FBG, fasting blood glucose; FIns, fasting plasma insulin; TC, total cholesterol; TG, triglyceride; FFA, free fatty acids; Data are presented as means ± SD, P-values were determined with *t*-test .***P* < 0.01 vs. HFD-WT.

Appendix Table S3 Biochemical parameters in HFD-fed Raptor^{f/f} mice during the HEC

Index	GFP (n=6)	<i>shDOCK5</i> (n=6)	<i>shDOCK5+Cre</i> (n=6)
Basal-state			
FBG (mmol/L)	7.40 ± 0.31	8.78 ± 0.20**	7.90 ± 0.30▲▲
FIns (mIU/L)	33.3 ± 2.51	44.3 ± 4.10**	38.2 ± 2.84▲▲
TC (mmol/L)	2.83 ± 0.22	5.01 ± 0.31**	3.85 ± 0.42▲▲
TG (mmol/L)	0.71 ± 0.08	0.86 ± 0.06**	0.76 ± 0.07▲
FFA (mmol/L)	0.56 ± 0.04	0.81 ± 0.05**	0.68 ± 0.06▲▲
Steady-state			
FBG (mmol/L)	5.91 ± 0.32	5.94 ± 0.16	5.85 ± 0.17
Ins (mIU/L)	256.3 ± 27.3	348.4 ± 35.2**	297.2 ± 24.3▲▲
TC (mmol/L)	1.23 ± 0.14	2.35 ± 0.08**	1.89 ± 0.09▲▲
TG (mmol/L)	0.31 ± 0.07	0.48 ± 0.06**	0.40 ± 0.05▲
FFA (mmol/L)	0.16 ± 0.03	0.41 ± 0.05**	0.30 ± 0.06▲▲

GFP, AAV8-GFP; *shDOCK5*, AAV8-*shDOCK5*; Cre, AAV8-Cre; FBG, fasting blood glucose; FIns, fasting plasma insulin; TC, total cholesterol; TG, triglyceride; FFA, free fatty acids; Data are presented as means ± SD, P-values were determined with one way ANOVA ** $P < 0.01$ vs. GFP group; ▲ $P < 0.05$, ▲▲ $P < 0.01$ vs. *shDOCK5* group.

Appendix Table S4 Specific primers used for RT-PCR analyses

Gene	Forward (5'-3')	Reverse (5'-3')
DOCK5	GGGGTTGCCATCTACAACTAC	AGCCCTCATACATCTCCAGGA
G6Pase	GTATTGTCTCATCACCATCTTC	CGCTCACACCATCTCTTAG
PEPCK	TAGGAGCAGCCATGAGAT	CGAAGTTGTAGCCGAAGA
UCP1	CTTCCCCTGGACACT	TTTCGGCAATCCTTCT
GAPDH	TGTGTCCGTCGTGGATCTGA	CCTGCTTCACCACCTTCTTGAT
β actin	GCTGTCCCTGTATGCCTCT	GATGTCACGCACGATTCC

DOCK5, dedicator of cytokinesis 5; G6Pase, glucose-6-phosphatase; PEPCK, phosphoenolpyruvate carboxykinase; UCP1, uncoupling protein 1; GAPDH, Glyceraldehyde 3-phosphate dehydrogenase; β actin, beta Actin

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