

Table S1: Primers used for quantitative real-time PCR

Gene	Forward primer	Reverse primer
<i>Mmp12</i>	CTGCTCCCATGAATGACAGTG	AGTTGCTTCTAGCCCAAAGAAC
<i>Rplp0</i>	AGATTCGGGATATGCTGTTGGC	TCGGGTCCTAGACCAGTG TTC
<i>Tnfa</i>	CCACCACGCTCTTCTGTCTAC	AGGGTCTGGGCCATAGA ACT
<i>Ccl2</i>	TTAAAAACCTGGATCGGAACCAA	GCATTAGCTTCAGATTTACGGGT
<i>Arg</i>	TGGCTTGCGAGACGTAGAC	GCTCAGGTGAATCGGCCTTTT
<i>Gro1</i>	CTGGGATTACCTCAAGAACATC	CAGGGTCAAGGCAAGCCTC
<i>Cox2</i>	TGAGCAACTATTCCAAACCAGC	GCACGTAGTCTTCGATCACTATC
<i>Emr1</i>	CTTTGGCTATGGGCTTCCAGTC	GCAAGGAGGACAGAGTTTATCGTG
<i>Cd68</i>	AACAGGACCTACATCAGAGC	TCAAGGTGAACAGCTGGAGA
<i>Adipoq</i>	GGAGATGCAGGTCTTCTTGG	CGAATGGGTACATTGGGAAC
<i>Mcp5</i>	ATTTCCACACTTCTATGCCTCCT	ATCCAGTATGGTCCTGAAGATCA
<i>Il6</i>	CTGCAAGAGACTTCCATCCAG	AGTGGTATAGACAGGTCTGTTGG
<i>Il1b</i>	GAAATGCCACCTTTTGACAGTG	TGGATGCTCTCATCAGGACAG

Table S2: Fold change in Log2 and p-value of selected downregulated proteins in eWAT of DKO mice

Protein symbol	Protein name	Log ₂ FC	p-value
ELAVL1	ELAV like RNA binding protein 1	-6.29	3.496E-06
PADI2	Peptidyl arginine deiminase, type II	-4.89	2.223E-02
BAX	BCL2-associated X protein	-4.88	4.363E-03
PSAP	Prosaposin	-4.48	1.972E-02
COTL1	Coactosin like F-actin binding protein 1	-4.27	1.805E-06
NCK1	NCK adaptor protein 1	-3.93	2.176E-02
NPM1	Nucleophosmin 1	-3.18	2.204E-02
USP14	Ubiquitin specific peptidase 14	-1.92	1.788E-02
ACSL1	Acyl-CoA synthetase long-chain family member 1	-1.38	4.900E-04

Table S3: Fold change in Log2 and p-value of selected upregulated proteins in eWAT of DKO mice

Protein symbol	Protein name	Log2FC	p-value
MFN2	Mitofusin 2	6.64	6.145E-03
LRP6	Low density lipoprotein receptor-related protein 6	5.58	2.522E-02
SOD3	Superoxide dismutase 3, extracellular	3.66	1.858E-02
FGFR1	Fibroblast growth factor receptor 1	3.53	3.089E-02
CREG1	Cellular repressor of E1A-stimulated genes 1	3.41	1.195E-02
PPM1A	Protein phosphatase 1A, magnesium dependent, alpha isoform	3.15	4.063E-02
AOC3	Amine oxidase, copper containing 3	3.10	2.046E-02
CAVIN1	Caveolae associated 1	2.74	9.590E-04
PLIN1	Perilipin 1	2.46	2.484E-03
MYDGF	Myeloid derived growth factor	2.43	1.335E-03
CES1D	Carboxylesterase 1D	1.79	2.145E-02
AFDN	Afadin, adherens junction formation factor	1.70	1.475E-02
PCLO	Piccolo (presynaptic cytomatrix protein)	1.66	2.535E-02
HPX	Hemopexin	1.56	1.478E-03

Table S4: Fold change in Log2 and p-value of selected downregulated proteins in aortas of DKO mice

Protein symbol	Protein name	Log ₂ FC	p-value
COL11A1	Collagen, type XI, alpha 1	-4.78	5.833E-03
C2	Complement C2	-3.68	3.152E-02
ACTR10	ARP10 actin-related protein 10	-3.49	1.320E-02
COL1A1	Collagen, type I, alpha 1	-2.82	4.062E-02
TFRC	Transferrin receptor	-2.49	1.300E-02
XDH	Xanthine dehydrogenase	-2.40	2.121E-02
CNN1	Calponin 1	-1.76	2.121E-02
DIAPH1	Diaphanous related formin 1	-1.69	4.725E-02
IDH1	Isocitrate dehydrogenase 1 (NADP+), soluble	-1.45	7.696E-04
CYB5R3	Cytochrome b5 reductase 3	-0.86	4.954E-02
LMOD1	Leiomodin 1 (smooth muscle)	-0.71	4.171E-02

Table S5: Fold change in Log2 and p-value of selected upregulated proteins in aortas of DKO mice

Protein symbol	Protein name	Log ₂ FC	p-value
DBN1	Drebrin 1	4.10	2.126E-02
LONP1	Ion peptidase 1, mitochondrial	3.27	1.555E-02
SWAP70	SWA-70 protein	3.26	1.838E-02
MBL2	Mannose-binding lectin (protein C) 2	3.14	9.937E-04
CSAD	Cysteine sulfinic acid decarboxylase	3.08	3.989E-02
LRP8	LDR receptor-related protein 8, apolipoprotein e receptor	2.80	4.032E-02
CTSB	Cathepsin B	1.80	3.688E-02
CYB5A	Cytochrome b5 type A (microsomal)	1.78	2.085E-04
DDX3X	DEAD box helicase 3, X-linked	1.59	1.500E-02
PARK7	Parkinson disease (autosomal recessive, early onset) 7	1.39	2.189E-02
ITIH4	Inter alpha-trypsin inhibitor, heavy chain 4	1.22	4.129E-02
CALR	Calreticulin	1.15	8.883E-03
ARPC1B	Actin related protein 2/3 complex, subunit 1B	0.99	2.245E-02
FN1	Fibronectin 1	0.90	1.728E-02
ANXA11	Annexin A11	0.64	6.333E-03

Table S6: Average Ct values of pro-inflammatory markers in aortas after 20-week feeding of HFSC diet

Gene	LdlrKO	DKO
<i>Ccl2</i>	31.87	31.96
<i>Il1b</i>	31.12	30.50
<i>Cox2</i>	28.85	28.98
<i>Mrc5</i>	31.55	32.31
<i>Tnf</i>	>40	>40
<i>Il6</i>	>40	>40