

Table S1**Animals (*in vivo* studies)**

Species	Vendor or Source	Background Strain	Sex
Wild type (WT)	Clea Japan	C57BL/6j	M
<i>Dscr-1^{-/-}</i>	Ryeom and Folkman Lab	C57BL/6j	M
<i>ApoE^{-/-}</i>	Maeda Lab, Taconic, USA	C57BL/6j	M
<i>Dscr-1^{-/-} and ApoE^{-/-}</i>	Minami Lab	C57BL/6j	M
<i>DSCR-1-lacZ-hprt; ApoE^{-/-}</i>	Minami Lab	C57BL/6j	M

Genetically Modified Animals

	Species	Vendor or Source	Background Strain
Parent – Male and Female	<i>Dscr-1^{-/-}</i>	Breeding in our animal facility	C57BL/6j

	Species	Vendor or Source	Background Strain
Parent – Male and Female	<i>ApoE^{-/-}</i>	Breeding in our animal facility	C57BL/6j

	Species	Vendor or Source	Background Strain
Parent – Male and Female	<i>Dscr-1^{-/-} and ApoE^{-/-}</i>	Breeding in our animal facility	C57BL/6j

	Species	Vendor or Source	Background Strain
Parent – Male and Female	<i>DSCR-1-lacZ-hprt; ApoE^{-/-}</i>	Breeding in our animal facility	C57BL/6j

Antibodies

Target antigen	Vendor or Source	Catalog #	Working concentration	Persistent ID / URL
CD11b	BD Bioscience	550282	2.5 µg/ml	https://wwwbdbiosciences.com/ds/pm/tds/550282.pdf
Rat IgG Alexa Fluor 488	ThermoFisher	A21208	2 µg/ml	https://www.thermofisher.com/order/genome-database/dataSheetPdf?producttype=antibody&productssubtype=antibody_secondary&productId=A-21208&version=99
LDLR	Abcam	Ab52818	1.4 µg/ml	https://www.abcam.co.jp/ldl-receptor-antibody-ep1553y-ab52818.pdf
Actin	Sigma	A1978	1 µg/ml	https://www.sigmaaldrich.com/content/dam/sigma-aldrich/docs/Sigma/Datasheet/6/a1978dat.pdf
Rabbit IgG (H+L)-	ThremoFisher	A11008	2 µg/ml	https://www.thermofisher.com/order/genome-database/dataSheetPdf?producttype=antibody&

Alexa Fluor488				roductsubtype=antibody_secondary&productId=A-11008&version=99
Rat IgG Alexa Fluor 594	ThermoFisher	A21209	2µg/ml	https://www.thermofisher.com/order/genome-database/dataSheetPdf?producttype=antibody&productsubtype=antibody_secondary&productId=A-21209&version=99
Rabbit IgG-HRP	Sigma	A9169	1µg/ml	https://www.sigmaaldrich.com/content/dam/sigma-aldrich/docs/Sigma/Datasheet/7/a9169-074h4802dat.pdf
Mouse IgG-HRP	Sigma	A9044	1µg/ml	https://www.sigmaaldrich.com/content/dam/sigma-aldrich/docs/Sigma/Datasheet/6/a9044dat.pdf
CD31	BD Bioscience	550274	0.078µg/ml	https://wwwbdbiosciences.com/ds/pm/tds/550274.pdf
<i>lacZ</i>	MBL	PM049	5 µg/ml	https://ruo.mbl.co.jp/bio/dtl/dtlfiles/PM049-v3.pdf
DSCR-1	Institute of Immunology, Co.Ltd, Japan	N2838	5 µg/ml	Originally generated from Minami and Kodama Lab (<i>J.Biol.Chem.</i> 281: 20503-20 2006)
hSREBP2	Institute of Immunology, Co.Ltd, Japan		1 µg/ml	Originally generated from Sakai Lab. (jmsakai@med.tohoku.ac.jp)
mSREBP2	Abcam	ab30682	1 µg/ml	https://www.abcam.com/SREBP2-antibody-ab30682.html

DNA/cDNA Clones

Clone Name	Sequence	Source / Repository
CA-mPCSK9	Primers for mutagenesis Pcsk9-D377Y-FW: TCCAGTTACTGCAGCACATGCTTCATG Pcsk9-D377Y-RV: GCTGCAGTAACTGGACGCTCCGATGAT	Minami lab

siRNA

Clone Name	Sequence	Source / Repository
Si-DSCR-1 (targeted both DSCR-1s and DSCR-1L)	5'-CAGAAACUCAGUCUUUAUGCAGCUGG-3' 5'-CCAGCUGCAUAAGACUAGUUUCUG-3'	Stealth siRNA, Specificity and efficacy was validated from <i>J.Biol.Chem.</i> 281: 20503-20 (2006)

Cultured Cells

Name	Vendor or Source	Persistent ID / URL
HEK293	ATCC	CRL-1573
HepG2	ATCC	CRL-10741

Data & Code Availability

Description	Source / Repository	Persistent ID / URL
Liver tissue microarray analysis	GSE172283	https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE172283
Aortic endothelial cells microarray analysis	GSE172283	https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE172283

Real-time PCR primers

Primer name	Sequence
<i>Dscr-1-Forward</i>	GACAAACAGTTCCTCATCTCCC
<i>Dscr-1-Reverse</i>	CATCTCTTCCTCTTCCTCCTCA
<i>Ceruloplasmin-Forward</i>	CAACAGTATGTGTATGTGCTGCAT
<i>Ceruloplasmin-Reverse</i>	AGATACCAGCTGAGATTTTCATCC
<i>Msr1-Forward</i>	CTGGACAAACTGGTCCACCT
<i>Msr1-Reverse</i>	TCCCCTTCTCTCCCTTTTGT
<i>Vcam-1-Forward</i>	GAACCCAAACAGAGGCAGAGTG
<i>Vcam-1-Reverse</i>	GGACTGCCCTCCTCTAGTATAGGA
<i>Icam-1-Forward</i>	GGTGACTGAGGAGTTCGACAGAA
<i>Icam-1-Reverse</i>	TCTGCGTCTCCAGGATCTGG
<i>Ppia-Forward</i>	ATCTTGTCCATGGCAAATGCTG
<i>Ppia-Reverse</i>	AAACGCTCCATGGCTTCCAC

Other

Description	Source / Repository	Persistent ID / URL
AAV-pro Helper Free system (AAV5)	Takara Bio	Cat#: 6650
pAd/CMV/V5-DEST Gateway Vector kit	ThermoFisher	Cat#: V49320

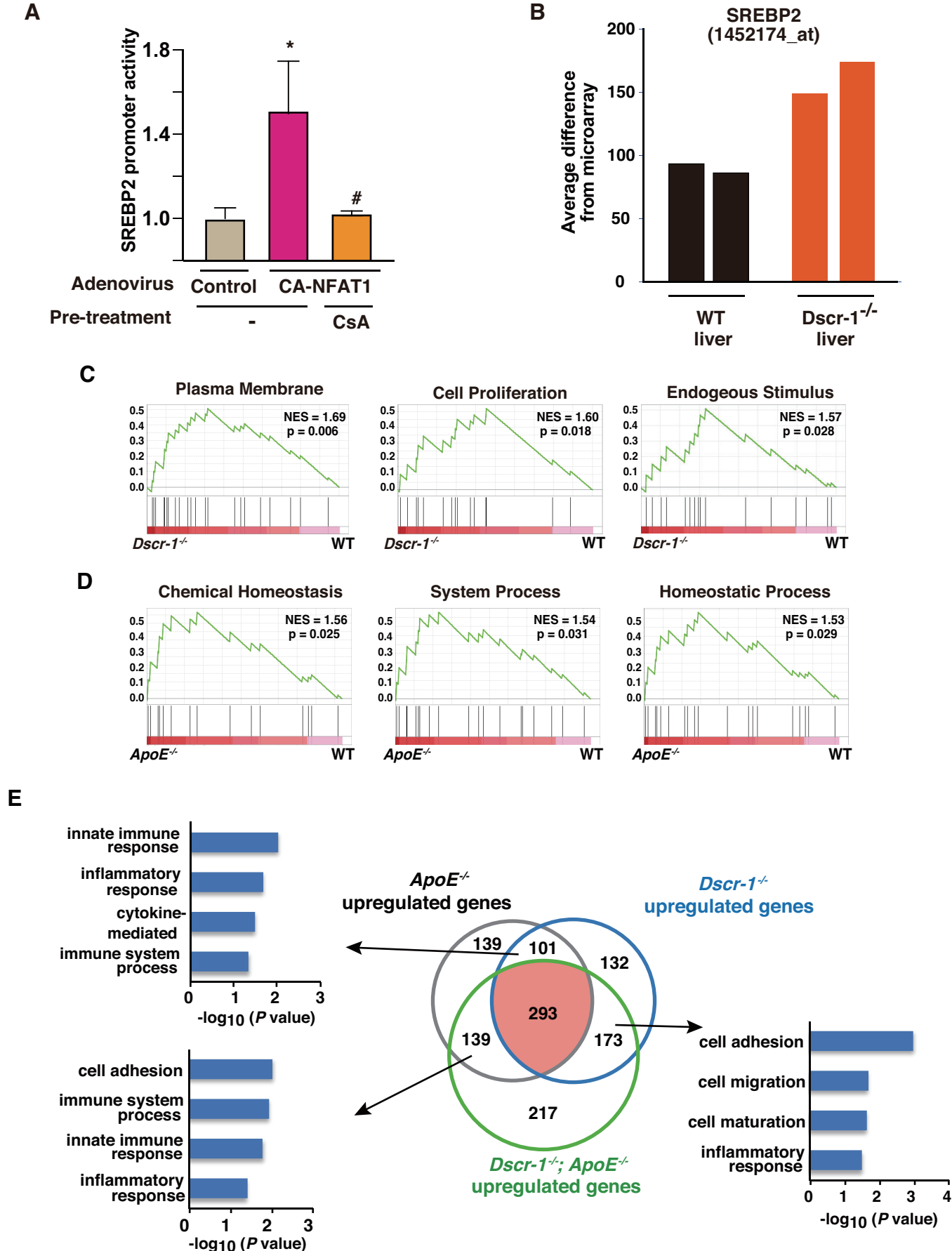


Figure S1 *Dscr-1* null mutation increases SREBP2 and changes cellular homeostasis gene set in liver.

(A) SREBP2-luc activities relative to Ad-control (n=3, mean± SD), **P*<0.05 compared with WT, #*P*<0.05 compared with CA-NFAT1 in the absence of CsA treatment. (B) Average differences of SREBP2 from two independent samples in liver. (C and D) GSEA pathway analysis of upregulated genes data set from *Dscr-1*^{-/-} (C) or *ApoE*^{-/-} (D) compared with WT. NES; ApoE, *Dscr-1* normalized enrichment score. (E) Venn diagrams of more than 2-fold upregulated gene volume in liver from null mutation of either and both of them. GO pathway analysis of commonly upregulated genes data set from *ApoE*^{-/-} and *Dscr-1*^{-/-} (upper-left), *ApoE*^{-/-} and *Dscr-1*^{-/-} plus *ApoE*^{-/-} double knockout (lower-left), and *Dscr-1*^{-/-} and *Dscr-1*^{-/-} plus *ApoE*^{-/-} knockout (lower-right).

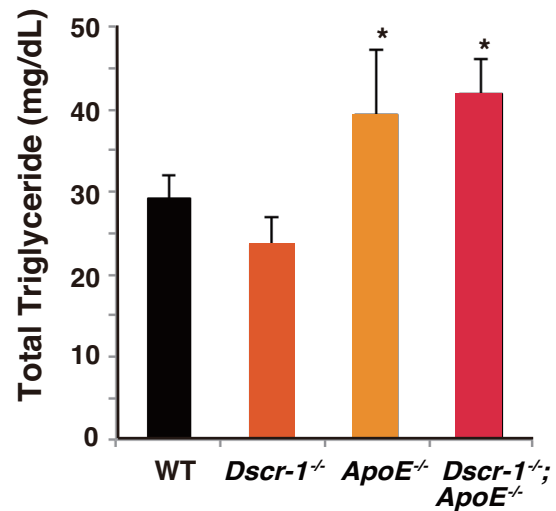
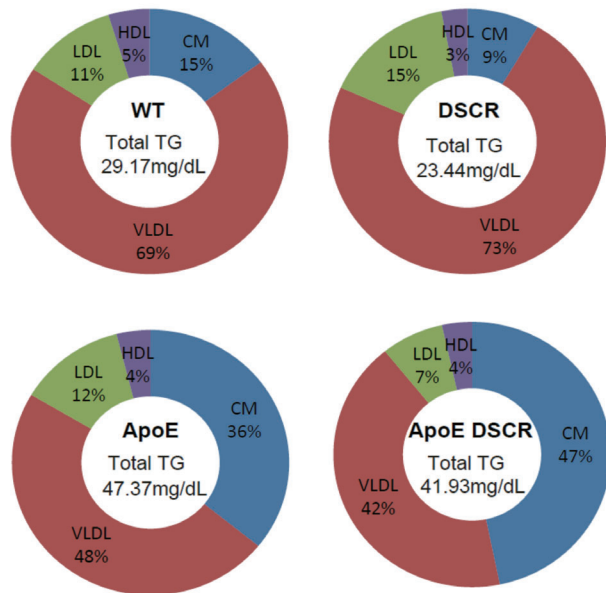
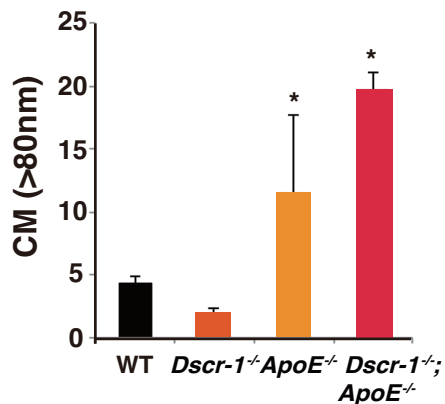
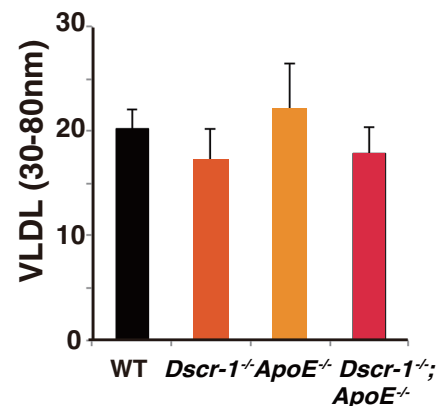
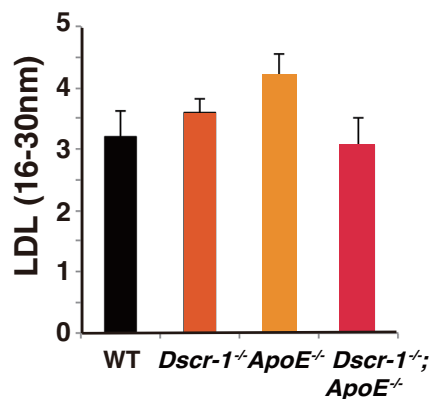
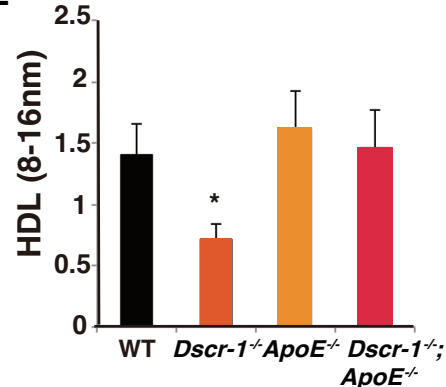
A**B****C****D****E**

Figure S2 Lipid profiles in *Dscr-1*, *ApoE*, and the double null mutation mice.

(A) Pie chart of total triglycerides (TG) from either or combined null mutation of *Dscr-1* and *ApoE*, and littermate control (WT) after a high fat diet fed for 8 weeks. (B-E). Total TG volume and contents (B: chylomicron (CM), C: VLDL, D: LDL, E: HDL) were shown in pie chart. Graph value means total or indicated each content volume (mg/dL) (n=12, mean ± SD) *, $P < 0.05$ compared with WT.

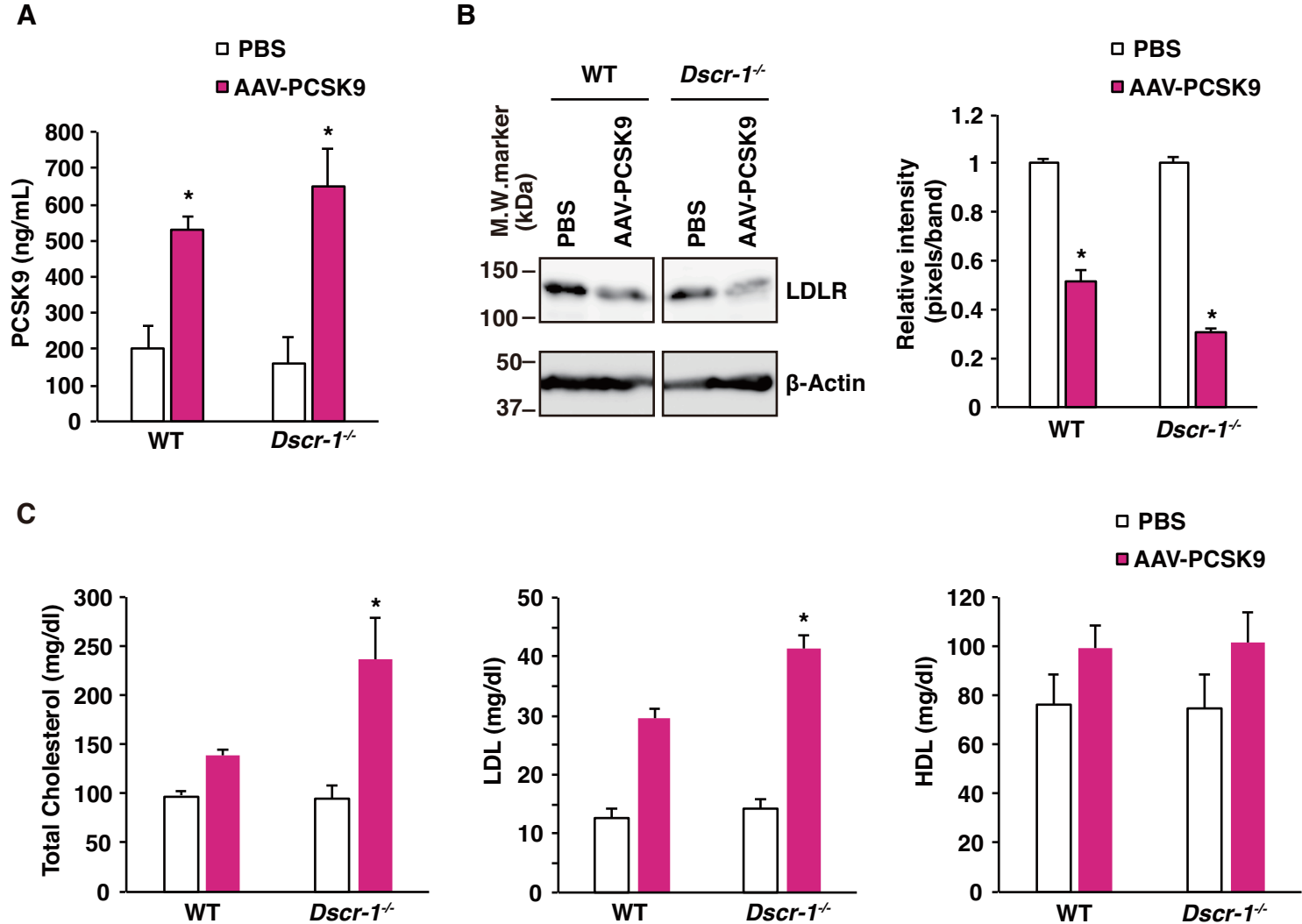


Figure S3 Loss of *Dscr-1* exaggerates hypercholesterolemia with adeno-associated virus (AAV)-mediated PCSK9 overexpression.

(A) Serum PCSK9 levels at 7 days after the AAV administration. (B) Western blot of anti-LDLR antibody from total liver extracts. β -Actin indicates the loading control. *Right* bar graph; LDLR quantification from the western blots (*left*) ($n=3$, mean \pm SD). *, $P < 0.05$ compared in the mock (PBS) treatment. (C) Quantification of total cholesterol (*left*), LDL (*middle*), and HDL (*right*) from indicated conditions ($n=3$, mean \pm SD). *, $P < 0.05$ compared with the AAV treatment in WT mice.

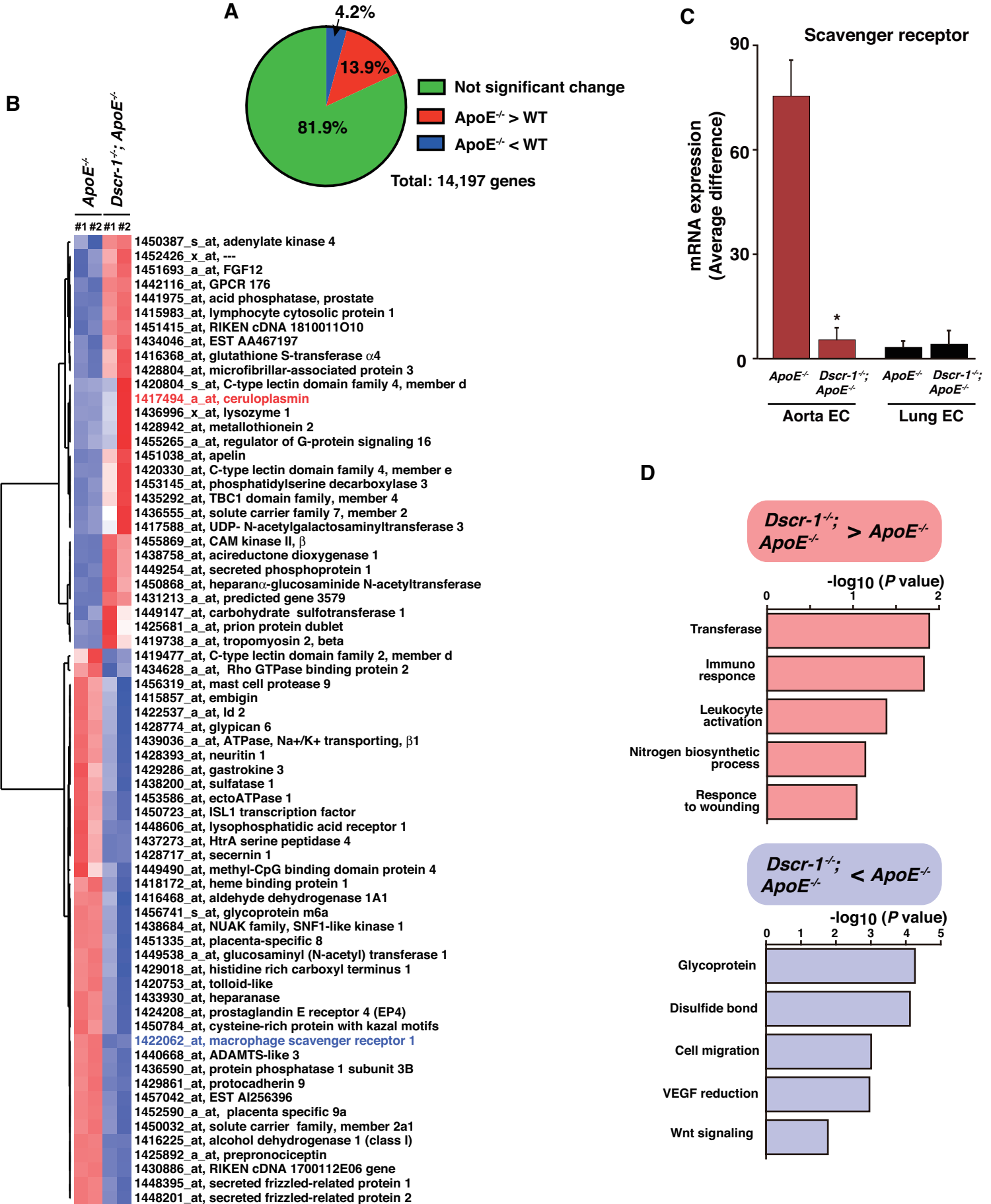


Figure S4 Loss of Dscr-1 in atherosclerotic aorta changes the unique gene expression pattern.

(A) Pie chart illustration of changed gene set of aortic endothelial cell (EC)s from ApoE^{-/-} and WT. More than two-fold reduced (blue), induced (red), and not changed (green) populations were shown in ApoE^{-/-} compared to WT. (B) Complete heatmap illustration of more than 2-fold up-or downregulated genes derived from two independent microarrays. (C) Average difference of scavenger receptor mRNA expression among ApoE^{-/-} and Dscr-1^{-/-} plus ApoE^{-/-} knockout mice (n=3, mean \pm SD) *, P<0.01 compared with ApoE^{-/-}. (D) GO analysis of commonly upregulated (upper graph) or downregulated (lower graph) gene set in Dscr-1 plus ApoE combined mutation, compared to ApoE single mutation.