

SUPPLEMENTAL MATERIAL

Deletion or inhibition of NOD1 favours plaque stability and attenuates atherothrombosis in advanced atherogenesis

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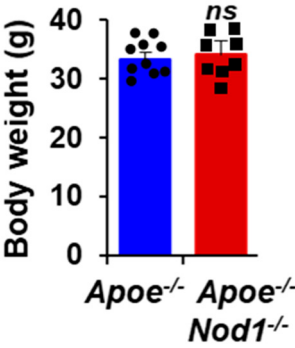
Running title: NOD1 deficiency promotes atheroma stability

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Supplementary Figures

a



b

mg/dL	Apoe ^{-/-}	Apoe ^{-/-} Nod1 ^{-/-}
Triglycerides	118.1 ± 2.3	112 ± 4.2
LDL	237.1 ± 4.6	244.6 ± 6.4
HDL	101.1 ± 4.5	103.6 ± 4.4
Total Cho.	372.1 ± 3.4	382.6 ± 8.1
Free Cho.	267.6 ± 6.7	279.9 ± 8.8

Figure S1. HFD-induced body weight increase and lipid profile changes in *Apoe*^{-/-} mice are not mediated by *Nod1*. (a) Body weight in *Apoe*^{-/-} (*n*=10) and *Apoe*^{-/-}*Nod1*^{-/-} (*n*=8) mice after 16 weeks on a HFD. (b) Serum concentration of triglyceride (TGA), high-density lipoproteins (HDL), low-density lipoproteins (LDL), total cholesterol (TCHO) and free cholesterol (FCHO) in the same cohort of mice. Data is represented as mean ± s.e.m. of the indicated number (*n*) of repeats. **ns**; Non-significant by Student's *t* test.

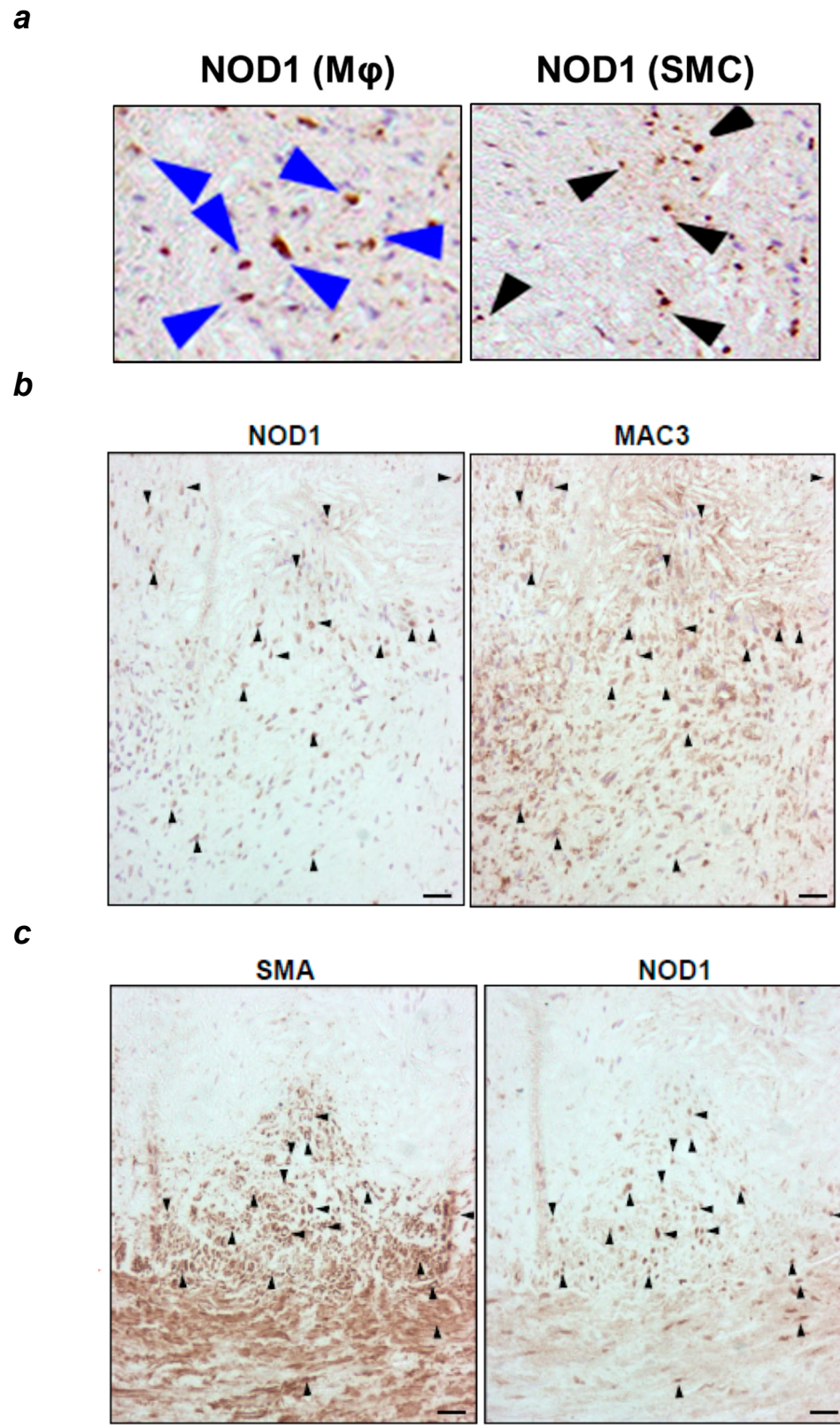


Figure S2. Inset of NOD1 from atherogenic human coronary arteries (from Fig. 1c) and NOD1 abundance in consecutive sections of human atherosclerotic coronary arteries presenting cholesterol crystals. (a) Amplification of atherogenic sections from Fig. 1c. **(b)** NOD1 co-localises with macrophage-rich areas (Mac3⁺-cells) in the media and intima of atherosclerotic human arteries. **(c)** Representative images showing the immunohistochemical analysis of SMA and NOD1. Blue arrows point to NOD1⁺ cells in macrophages; black arrows point out NOD1⁺ cells in the smooth muscle cells.

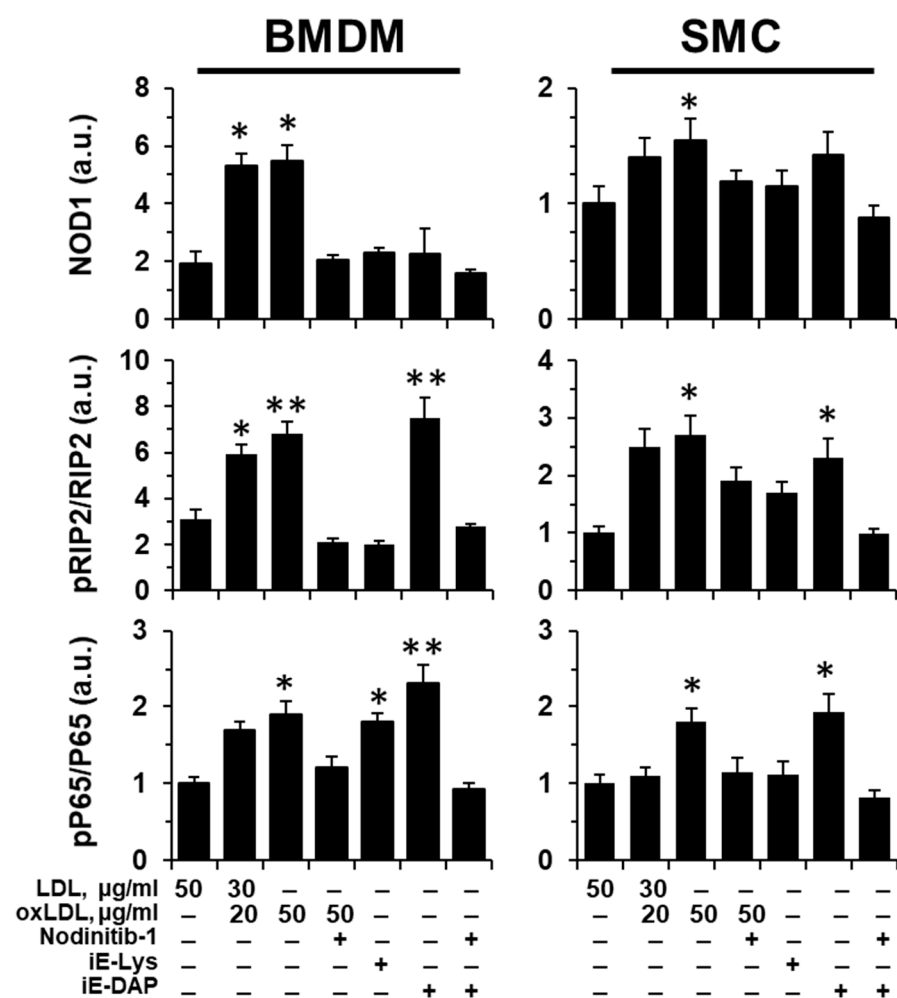


Figure S3. Quantification of the immunoblots shown in Fig. 1d. Analysis of NOD1, pRIP2/RIP2, pP65/P65 in *Wt* BMDM ($n=6$) and SMC ($n=4$) pre-treated with the NOD1 inhibitor Nodinitib-1 and/or stimulated with native LDL (as control for lipid load in the medium), oxLDL, c12-iE-DAP (an agonist for NOD1) and iE-Lys (an iE-DAP analogue that fails to activate NOD1) and c12-iE-DAP (an agonist for NOD1) for 24h or 48h, respectively. Protein levels were normalized to tubulin. Data are represented as mean $\pm$ s.e.m. of the indicated number (n) of independent experiments. * $P < 0.05$, ** $P < 0.01$ vs. the LDL condition by Student's t test. a.u., arbitrary units.

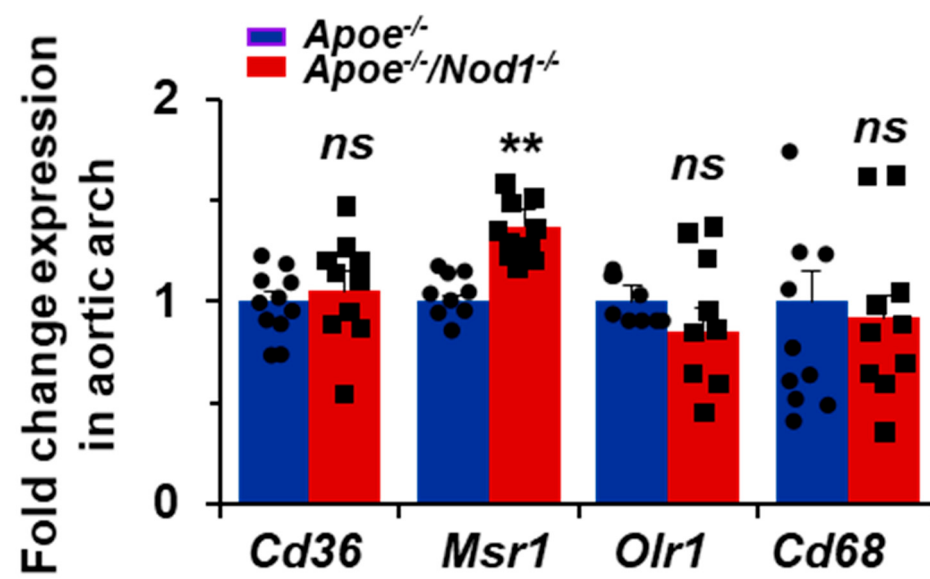


Figure S4. mRNA levels of scavenger receptors in athero-prone *Apoe*^{-/-} mice in the presence and absence of *Nod1*. Quantitative PCR analysis of the scavenger receptors *Cd36*, mouse scavenger receptor 1 (*Msr1*), oxidised low-density receptor 1 (*Olr1*) and the low-density lipoprotein receptor *Cd68* expression in the aortic arch of *Apoe*^{-/-} (*n*=10) and *Apoe*^{-/-}/*Nod1*^{-/-} (*n*=10) mice after 16 weeks on HFD. mRNA levels were normalised to *mRplp0* expression. Data are represented as mean ± s.e.m. of the indicated number (*n*) of repeats. ***P* < 0.01 vs. *Apoe*^{-/-} by Student's *t* test. **ns**; Non-significant by Student's *t* test.

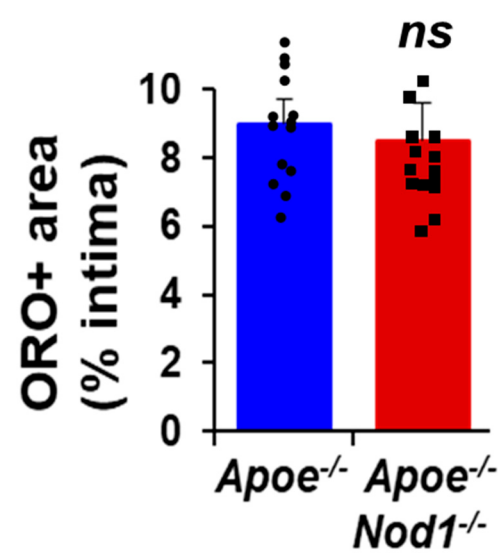


Figure S5. *Nod1* does not affect lipid atherosclerotic area in *Apoe*^{-/-} mice. Quantification of positive Oil Red O (ORO) lesion area in the semilunar valve cups of *Apoe*^{-/-} (*n*=13) and *Apoe*^{-/-}/*Nod1*^{-/-} (*n*=13) mice fed HFD for 16 weeks. Data are represented as mean ± s.e.m. of the indicated number (*n*) of repeats. Non-significant by Mann-Whitney U test. **ns**; Non-significant by Student's *t* test.

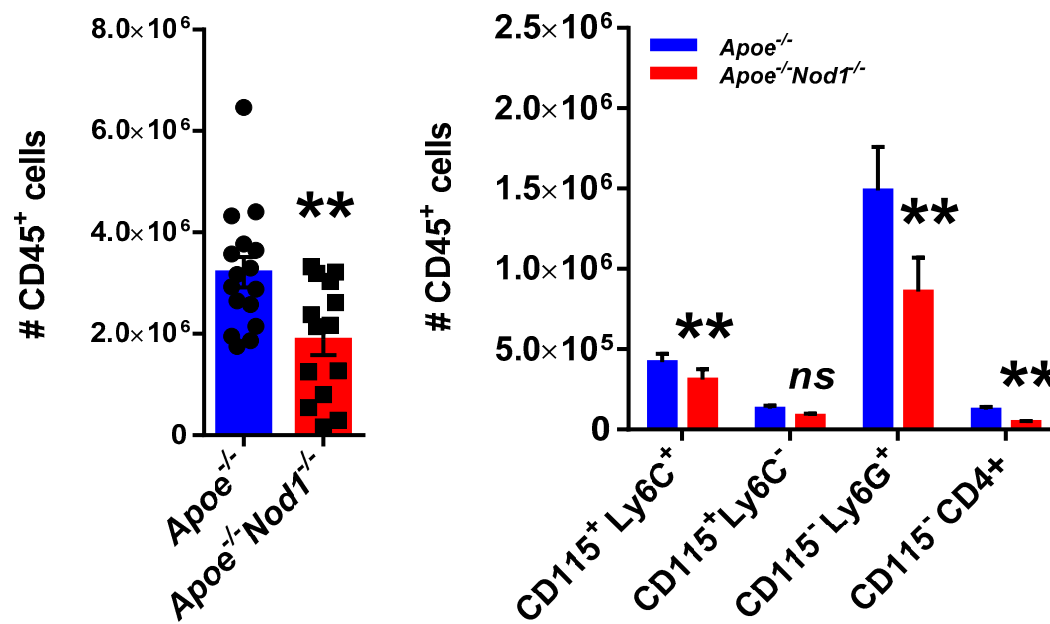


Figure S6. *Apoe*^{-/-}*Nod1*^{-/-} mice present less leukocyte cell content in blood comparing to their *Apoe*^{-/-} counterparts. Quantification of leukocyte, monocyte/macrophage, neutrophil and CD4⁺ lymphocyte cells in blood collected from *Apoe*^{-/-} (*n*=16) and *Apoe*^{-/-}*Nod1*^{-/-} (*n*=16) mice (16 weeks HFD-fed), performed by flow cytometry. Data are represented as mean ± s.e.m. of the indicated number (*n*) of repeats. **P* < 0.05, ***P* < 0.01 vs. *Apoe*^{-/-} by Student's *t* test. *ns*; Non-significant by Student's *t* test.

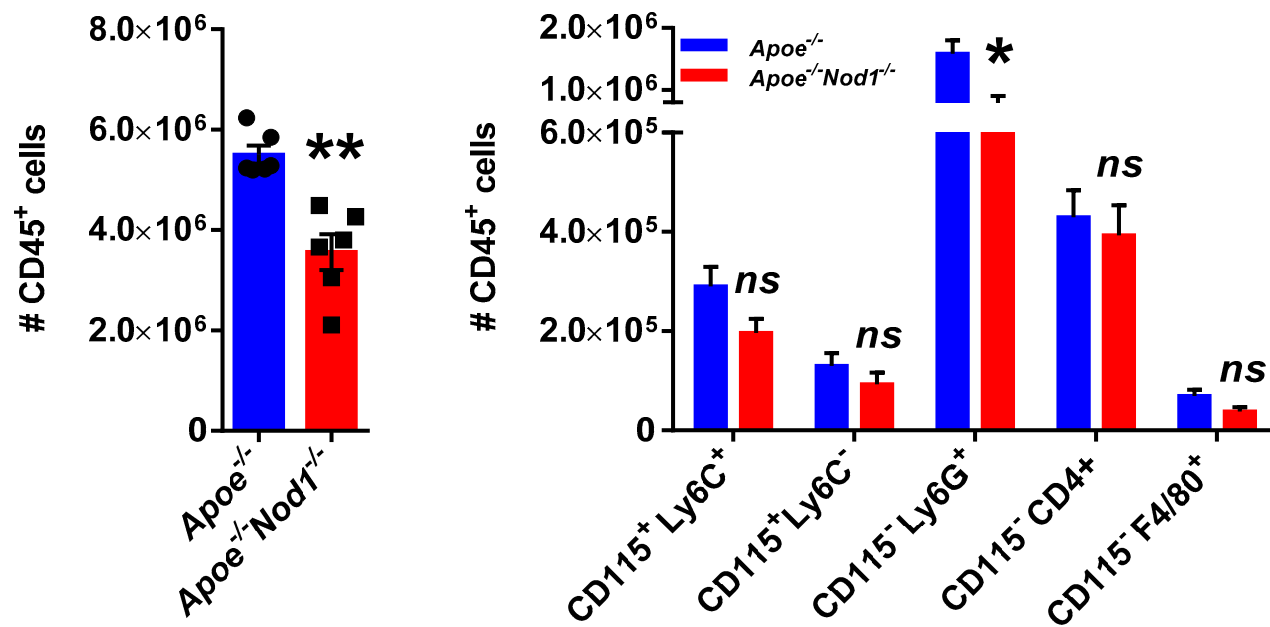


Figure S7. *Nod1* inactivation in mice results in a lower number of total CD45⁺ cells and neutrophils in spleen, while there are no significant changes in other white cells subsets. Leukocyte, monocyte/macrophage, neutrophil and CD4⁺ lymphocyte number of cells was assessed in the spleen obtained from *Apoe*^{-/-} (*n*=6) and *Apoe*^{-/-}*Nod1*^{-/-} (*n*=6) mice (16 weeks HFD-fed), using flow cytometry techniques. Data are represented as mean ± s.e.m. of the indicated number (*n*) of repeats. **P* < 0.05, ***P* < 0.01 vs. *Apoe*^{-/-} by Student's *t* test. *ns*; Non-significant by Student's *t* test.

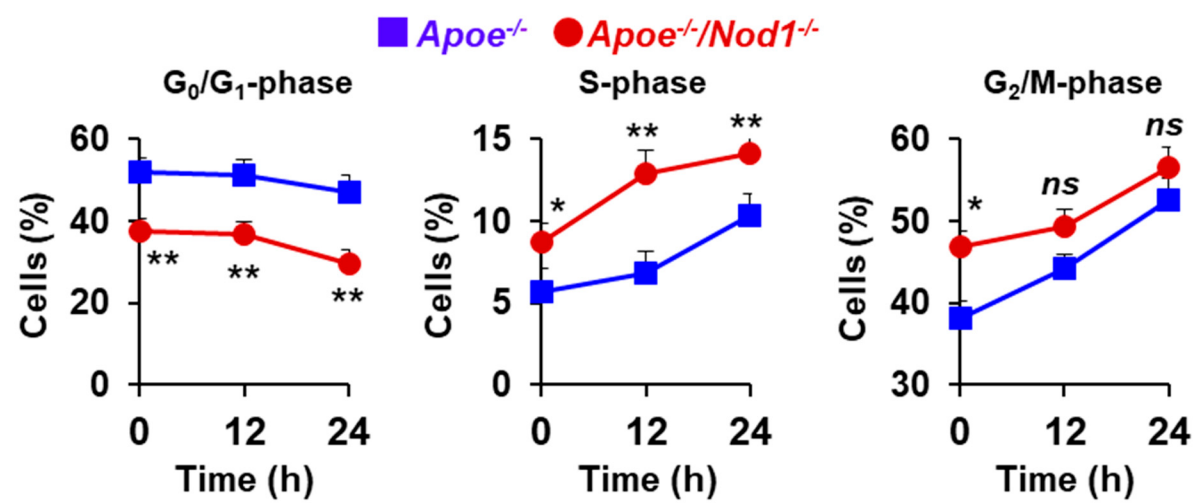


Figure S8. *Nod1* inactivation in cultured SMCs promotes faster cell cycle entry into S-phase. Cell cycle profile of primary vascular SMCs from *Apoe*^{-/-} (*n*=10) and *Apoe*^{-/-}/*Nod1*^{-/-} (*n*=10) mice was assessed in a flow cytometer by propidium iodide staining. Cells were synchronised in G₀/G₁ phase by 72-h serum deprivation and restimulated for the indicated times. Results are average representative of three independent experiments. Data are represented as mean ± s.e.m. of the indicated number (*n*) of repeats. **P* < 0.05, ***P* < 0.01 vs. *Apoe*^{-/-} by Student's *t* test. *ns*; Non-significant by Student's *t* test.

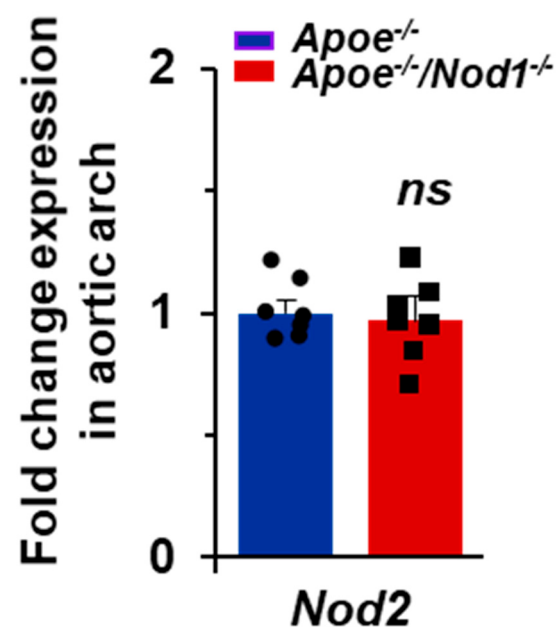


Figure S9. mRNA levels of *Nod2* in the aortic arch of athero-prone *Apoe*^{-/-} mice in the presence and absence of *Nod1*. Quantitative PCR analysis of *Nod2* expression in the aortic arch of *Apoe*^{-/-} (*n*=7) and *Apoe*^{-/-}/*Nod1*^{-/-} (*n*=7) mice after 16 weeks on HFD. mRNA levels were normalised to *mRplp0* expression. Data are represented as mean ± s.e.m. of the indicated number (*n*) of repeats. *ns*; Non-significant by Student's *t* test.