

Figure S1

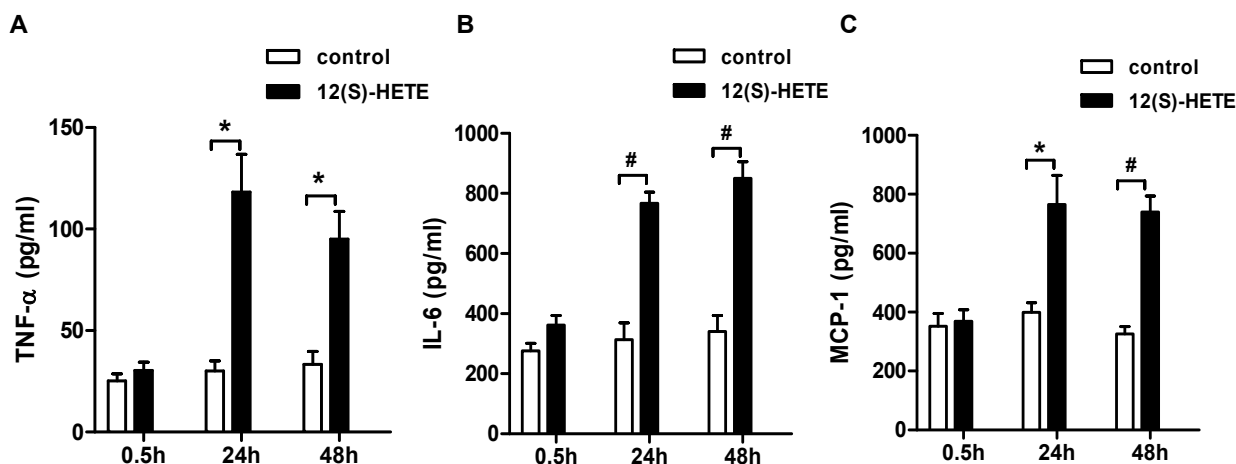


Figure S1. Effects of 12(S)-HETE on the production of adipose inflammatory factors. Equal numbers of subconfluent MCs were cultured in medium (RPMI1640+1%FBS) containing either buffer (control) or 12(S)-HETE ( $10^{-7}$ M) for 0.5–48 h. An aliquot supernatant was collected from each well and content of TNF- $\alpha$  (A), IL-6 (B), and MCP-1 (C) were measured by ELISA kits. Results are from three sets of experiments and shown as fold over control (mean $\pm$ SEM, n=3. \*p<0.05, #p<0.01 vs. control).

Figure S2

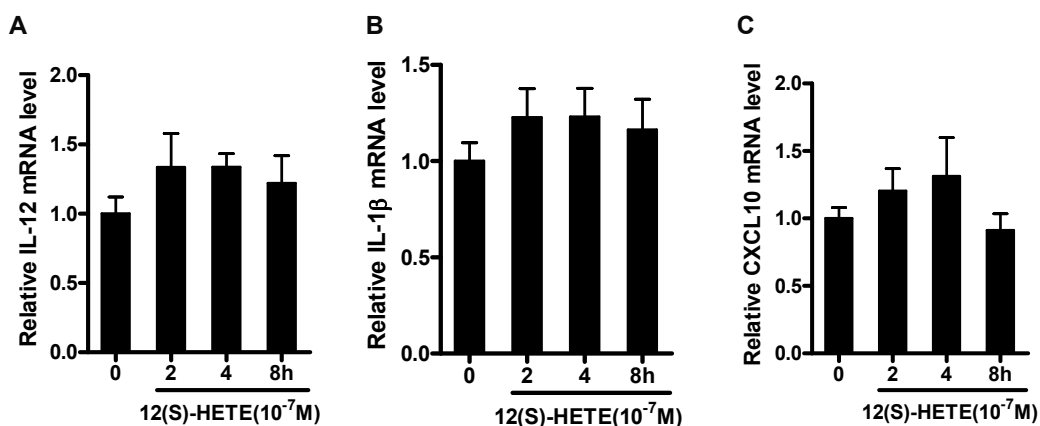


Figure S2. Effects of 12(S)-HETE on the expression of adipose inflammatory factors. MCs were stimulated with 12(S)-HETE ( $10^{-7}$ M) for 2–8 h and changes in intracellular mRNA levels of IL-12 (A), IL-1 $\beta$  (B), and CXCL-10 (C) were measured by RT-qPCR. Results are expressed as fold over none treatment of control (mean $\pm$ SEM, n=3).

Figure S3

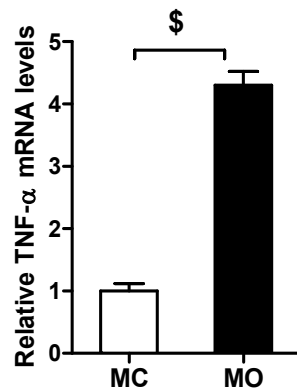


Figure S3. Quiescent MCs and MOs were collected for experiment. Intracellular mRNA expression of TNF- $\alpha$  was measured by RT-qPCR,  $\beta$ -Actin was used as internal control. Results are expressed as fold over MCs (mean $\pm$ SEM, n=3. \$p<0.001 vs. MC).

Figure S4

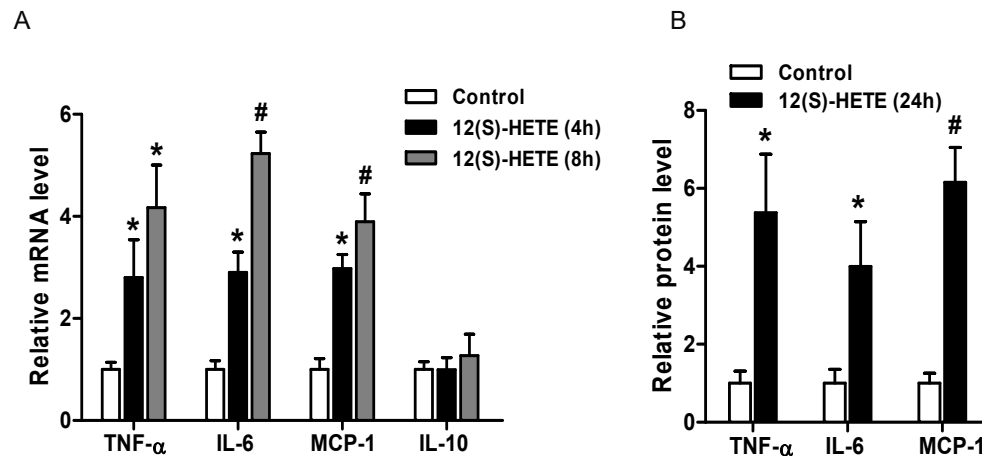


Figure S4. Effects of 12(S)-HETE on the expression of adipose inflammatory factors in MOs. (A) Quiescent MOs (RAW264.7) were stimulated with 12(S)-HETE ( $10^{-7}$ M) for 4–8 h and changes in intracellular mRNA levels of TNF- $\alpha$ , IL-6, MCP-1 and IL-10 were measured by RT-qPCR. Results are expressed as fold over none treatment of Control (mean $\pm$ SEM, n=3). (B) Equal numbers of subconfluent MOs were cultured in medium (DMEM+1%FBS) containing either buffer (Control) or 12(S)-HETE ( $10^{-7}$ M) for 24 h. An aliquot supernatant was collected from each well and content of TNF- $\alpha$ , IL-6, and MCP-1 were measured by ELISA kits. Results are from three sets of experiments and shown as fold over Control (mean $\pm$ SEM, n=3. \*p<0.05, #p<0.01 vs. Control).

**Table S1. Primer sequences used in RT-qPCR analysis**

Primers (5'→3')		
Gene (species)	Sense	Anti-sense
IL-1 $\beta$ (M)	TTGACGGACC CAAAAAG ATG	AGAAGGTGCTCATG TCCTCA
IL-12 (M)	TCTGCAGAGAAGGTCACACT	ATGAAGAAGCTGGTGCTGTA
CXCL10 (M)	CTCATCCTGCTGGGTCTGAG	CCTATGGCCCTCATTCTCAC
M, Mouse		