SUPPLEMENTAL MATERIAL



Figure S1. Systemic metabolic profile of MMP-9 deficient mice.

(A) Oxygen consumption. (B) Carbon dioxide production. (C) Heat / Energy expenditure (normalized to body weight). (D-F) Locomotor activity. (G) Body weight. (H) Total food consumption. The studies were conducted in metabolic cages. n=7 WT mice, n=5 Mmp9^{-/-} mice. *: $P \le 0.05$ vs. WT.



Figure S2. Supplemental quantitative analysis of lipids.

(A) Esterified lipids were elevated in livers from $Mmp9^{-/2}$ mice compared to livers from WT mice, particularly when the mice were fasted-and-refed a high carbohydrate diet. In contrast to Mmp9-/- mice, mice lacking MMP-7 ($Mmp7^{-/2}$) had a different hepatic lipid profile. n=4 mice *per* group, except n=3 for $Mmp7^{-/2}$ fasted-and-refed. $\dagger P \le 0.05 vs$. fasted for each genotype. $*:P \le 0.05 vs$. WT fasted-and-refed.

(B) Cholesterol excretion was unchanged in $Mmp9^{-/-}$ mice. n=4 mice per genotype.



Figure S3. Spleen is a likely source of plasma sPLA₂ in MMP-9-deficient mice. sPLA₂ activity was analyzed in duplicate using pools of the indicated tissues. n=3 WT and n=4 Mmp9^{-/-} mice. *: P<0.05 vs. WT.



Figure S4. SREBP-2 expression in response to dietary cholesterol and -/- **varespladib**. *Left*: Western blot showing amount of hepatic SREBP-2 protein in *Mmp9* mice in response to varespladib and dietary cholesterol supplementation. The experiment involved n=4 to 5 mice *per* group (or treatment). For analysis, livers were pooled, homogenized and the fraction containing nuclei was subjected to western blot analysis with SREBP-2 antibodies. *Right*: Quantitative analysis for two independent preparations and western blots. *: $P \le 0.05 vs$. untreated.



Figure S5. Proposed model. MMP-9 regulates cholesterol metabolism through PLA₂-dependent and PLA₂-independent mechanisms. Important elements are: peripheral organs (e.g., the spleen) acting as source of plasma sPLA₂ activity and MMP-9 (inhibitor of sPLA₂ release from peripheral organs). Once in the circulation, sPLA₂ acts on plasma lipoproteins or target organs (e.g., the liver) to release lipid mediators from phospholipids that ultimately influence cholesterol metabolism. Furthermore, the direct action of MMP-9 in the liver may influence hepatic cholesterol through as yet unclear PLA₂-independent pathways.



Figure S6. Supplement to Figure 7A containing the quantitative analysis of hepatic transcriptional responses to dietary cholesterol for the indicated genes and genotypes. n=6 WT mice, n=8 $Mmp2^{-7}$ mice, n=5 $Mmp7^{-7}$ mice, n=5 $Mmp9^{-7}$ mice and n=5 $Timp2^{-7}$ mice. *: $P \le 0.05$ vs. WT. †: $P \le 0.05$ vs. 0 days on cholesterol.



Figure S7. Extended quantitative analysis of the relative mRNA expression of cardiac and hepatic lipid metabolic and metalloproteinase genes and their response to dietary supplementation with 0.15% cholesterol for 2.5 days in mice deficient in one of several MMPs. n=6 WT mice, n=8 $Mmp2^{-7}$ mice, n=5 $Mmp7^{-7}$ mice, n=5 $Mmp9^{-7}$ mice. Slots without bars: Gene expression was not determined.