Identification of a Novel Heart–Liver Axis: Matrix Metalloproteinase-2 Negatively Regulates Cardiac Secreted Phospholipase A2 to Modulate Lipid Metabolism and Inflammation in the Liver

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Background—Endocrine functions of the heart have been well established. We investigated the hypothesis that cardiac secretion of a unique phospholipase A2 recently identified by our laboratory (cardiac secreted phospholipase A2 [sPLA2]) establishes a heart–liver endocrine axis that is negatively regulated by matrix metalloproteinase-2 (MMP-2).

Methods and Results—In Mmp2–/– mice, cardiac (but not hepatic) sPLA2 was elevated, leading to hepatic inflammation, immune cell infiltration, dysregulation of the sterol regulatory element binding protein-2 and liver X receptor-α pathways, abnormal transcriptional responses to dietary cholesterol, and elevated triglycerides in very low-density lipoprotein and in the liver. Expression of monocyte chemoattractant protein-3, a known MMP-2 substrate, was elevated at both mRNA and protein levels in the heart. Functional studies including in vivo antibody neutralization identified cardiac monocyte chemoattractant protein 3 as a possible agonist of cardiac sPLA2 secretion. Conversely, systemic sPLA2 inhibition almost fully normalized the cardiohepatic phenotype without affecting monocyte chemoattractant protein-3. Finally, wild-type mice that received high-performance liquid chromatography–isolated cardiac sPLA2 from Mmp2–/– donors developed a cardiohepatic gene expression profile similar to that of Mmp2–/– mice.

Conclusions—These findings identified the novel MMP-2/cardiac sPLA2 pathway that endows the heart with important endocrine functions, including regulation of inflammation and lipid metabolism in the liver. Our findings could also help explain how MMP2 deficiency leads to cardiac problems, inflammation, and metabolic dysregulation in patients. (J Am Heart Assoc 2015;4:e002553 doi: 10.1161/JAHA.115.002553)

Key Words: heart • inflammation • liver • matrix metalloproteinase • metabolism • phospholipase A2

Matrix metalloproteinase-2 (MMP-2), sometimes referred to as gelatinase A or 72 kDa collagenase, is a member of a family comprising 25 different Zn-dependent endopeptidases. A spectrum of cellular processes covering cellular proliferation, angiogenesis, and inflammation is modulated through MMP-2–dependent cleavage and regulation of extracellular matrix components, cell membrane receptors, latent growth factors, and cytokines.1

Among the cytokines targeted by MMP-2 is monocyte chemoattractant protein-3 (MCP-3 encoded by Ccl7).2,3 MCP-3 is a CC-motif (N-terminus has 2 consecutive conserved cysteine residues) chemokine that is 76 amino acids long.

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Accompanying Figures S1 through S8 are available at http://jaha.ahajournals.org/content/4/11/e002553/suppl/DC1

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Binding of the hemopexin domain of MMP-2 to MCP-3 facilitates cleavage of MCP-3 at a glycine/isoleucine bond. The resultant MCP-3 peptide serves as a general antagonist of CC-chemokine receptors that inhibits inflammation signaling by intact MCP-3 and other chemokine receptor ligands.2

MCP-3 is also reported to be a physiological substrate of MMP-2 in arthritis.2 Interestingly, humans with genetic loss of MMP-2 activity suffer from crippling arthritis4–6 and a spectrum of cardiovascular problems including congenital cardiac malformations, transposition of the great arteries, mitral valve prolapse, bicuspid aortic valve, and atrial and ventricular septal defects.4,7 The molecular mechanisms underlying these pathologies are unknown, leaving clinicians without effective treatments.4 In addition to inflammatory diseases, obesity has been linked to single-nucleotide polymorphisms in the human MMP2 gene8 and MMP2 gene promoter.9 Genetic loss of MMP-2 (Mmp21/1) in mice results in an arthritis-like phenotype associated with skeletal, bone, and craniofacial defects.5 In addition, Mmp21/1 mice resist diet-induced obesity during development,10,11 exhibit cardiac inflammation at baseline,12 and are predisposed to hypertensive heart disease due to abnormal regulation of the sterol regulatory binding protein 2 (SREBP-2) pathway in the heart.13

The evident coexistence of inflammation and metabolic dysregulation with MMP-2 deficiency is of potential clinical significance but is very poorly understood. In this study, we report that the hepatic metabolic phenotype in Mmp21/1 mice can be largely explained by a novel heart–liver axis involving myocardial secretion of a unique phospholipase A2 (sPLA2), which we coined cardiac secreted PLA2 (sPLA2).12 Our findings identify a novel functional link between cardiac inflammation and hepatic metabolism.

**Materials and Methods**

Oil Red O stain, alkaline phosphatase, and cholesterol were from Sigma-Aldrich. Dulbecco minimum essential medium, antibodies against liver X receptor (LXR), TaqMan quantitative reverse transcription–polymerase chain reaction (qRT-PCR) primers, TRizol reagent, random primers, Superscript II, and penicillin/streptomycin were from Life Technologies. SREBP-2 primers, TRIzol reagent, random primers, Superscript II, and penicillin/streptomycin were from Life Technologies. Recombinant human pro–MMP-2 was from EMD Millipore. Collagen-coated cell dishes were from Greiner Bio-One. Varespladib was from Selleck Chemicals. PNGase F was from Promega. Enhanced chemiluminescence western blotting detection reagent was from GE Healthcare. Horseradish peroxidase–conjugated antirabbit antibodies and Bio-Rad Protein Assay were from Bio-Rad. The Pierce bicinechonic acid protein assay kit was from Thermo Scientific. MCP-3, neutralizing MCP-3 antibody, and control isotype-matched IgG1 were from R&D Systems, Inc. Densitometry was performed using ImageQuant 5.1 (Molecular Dynamics).

**Animals**

All protocols were approved by the University of Alberta animal care committee and conducted in accordance with institutional guidelines issued by the Canada Council on Animal Care. Except as otherwise stated, wild-type (WT) mice aged 10 to 15 weeks were purchased from Charles River Laboratories (Wilmington, DE) or Jackson Laboratory (Bar Harbor, ME) and compared with age- and sex-matched Mmp71/1− and Mmp91/1− mice purchased from Jackson Laboratory. Mmp21/1− and Mmp21/1− mice were bred at the University of Alberta. All mice were of a C57BL/6 background and were housed in the Health Sciences Laboratory Animal Services of the University of Alberta with a 12-hour light/dark cycle. Mice were fed a chow diet ad libitum (PicoLab Rodent Diet 20; Lab Diet), which was used in all experiments unless otherwise specified. At the end of the experiments, mice were euthanized with 65 mg/kg of sodium pentobarbital, the blood was collected with EDTA-coated syringes and tubes, and the organs were excised and snap-frozen in liquid nitrogen. Of note, we found that Mmp21/1− reproduced very slowly in our facility. Consequently, this study was conducted with limited numbers of mice available to us at any time. Typically, 4 to 5 mice were used per treatment group.

**In Vivo Responses to Dietary Cholesterol, Fasting, and Fasting–Refeeding**

The dietary regimens in these studies followed previously described protocols.14 In the cholesterol supplementation studies, Mmp21/1− or WT mice (aged 12 to 14 weeks) were fed chow supplemented with 0% or 0.15% cholesterol for 2.5 or 6.5 days. In this study, the mice were not fasted before being euthanized. In the fasting and fasting–refeeding studies, mice (aged 10 to 22 weeks) were fasted for 16 hours or were fasted for 16 hours, fed a high-carbohydrate diet (TD.88122 mouse diet) for 4 hours, and then euthanized.

**In Vivo Pharmacological Studies**

To study the contribution of systemic sPLA2 to the inflammatory and lipid metabolic phenotype of MMP-2 deficiency, mice were gavaged with the sPLA2 inhibitor varespladib (10 mg/kg per day). The varespladib stock was prepared in dimethyl sulfoxide, as per the manufacturer’s instructions, and diluted to the working concentration before each experiment. Aqueous dimethyl sulfoxide solution of the same concentration as the varespladib working solution (equivalent to dimethyl sulfoxide 2.6 μL/kg per day) was used in...
control experiments (vehicle). Treatment was conducted for 2.5 or 5 days, followed by euthanasia. To study the contribution of MCP-3 to the phenotype of MMP-2 deficiency, Mmp2−/− mice were injected (intraperitoneally) with neutralizing MCP-3 antibody (0.6 mg/kg per day) for 2.5 days, and their responses were compared with those of WT mice that underwent exactly the same protocol. The dose regimen followed a previous report.3

Metabolic Studies

Metabolic caging studies were conducted at the Core Facility of the Cardiovascular Research Center, University of Alberta. Mice were individually housed in Oxymex/CLAMS metabolic chambers (Columbus Instruments) in which O2 consumption, CO2 production, food and water consumption, and movement were measured over 2 days and 2 nights.

Cell Culture Studies

Primary cardiomyocytes were isolated from WT or Mmp2−/− mouse hearts using the classical method of retrograde perfusion with collagenase through the coronary arteries to digest the extracellular matrix of the heart.15 This method resulted in the release of calcium-tolerant, rod-shaped mouse cardiomyocytes from adult hearts, as confirmed by microscopy. The isolated myocytes were seeded at a density of ≈10⁴ cells/cm² and cultured in serum-free Dulbecco minimum essential medium on collagen-coated dishes. The effects of extracellular MMP-2 or brefeldin A on sPLA2 activity were analyzed in primary cells cultured in Dulbecco minimum essential medium supplemented with 0.1% filter sterilized BSA in the absence or presence of human recombinant MMP-2 (40 nmol/L) or brefeldin A (1 μmol/L) for 16 hours.

qRT-PCR

Tissue was extracted by homogenizing 30 to 50 mg pieces of frozen tissue at 4°C in 1 mL of TRIzol reagent using the Bullet Blender (Next Advance). RNA was isolated from TRIzol according to the manufacturer’s instructions, and cDNA was generated from RNA using random primers and Superscript II reverse transcriptase. RNA isolated for each mouse’s tissue was quantitated in triplicate to obtain a value representative of the relevant tissue for each mouse. Expression analysis of the reported genes was performed by TaqMan qRT-PCR using an ABI 7900 HT sequence detection system (Applied Biosystems). For data normalization, both Gapdh and Actb (to confirm interpretation of data relative to Gapdh) were used as internal standards at steady state. The qRT-PCR data chosen for the figures are relative to Gapdh because we did not observe any significant quantitative differences in Gapdh versus Actb expression among WT, Mmp2−/−, Mmp7−/−, and Mmp9−/− mice fed chow. The genes chosen to characterize the cardiohepatic phenotype of Mmp2−/− mice and reported throughout the figures were found by experimentation to be differentially expressed across these genotypes of MMP deficiency and thus provide useful markers for studying the metabolic pathways modulated by these MMPs.

Protein Determinations

Colorimetric measurement of total protein was done using the Bio-Rad Protein Assay or Pierce bicinchoninic acid protein assay kit, according to the manufacturer’s instructions. Determination of hepatic liver LXR-α and SREBP-2 protein levels was conducted by western blotting. Briefly, 15- to 25-mg liver pieces were homogenized using the Bullet Blender at 4°C in a buffer of 5 mmol/L CaCl2, 150 mmol/L NaCl, 0.5 mmol/L NaN3, and 25 mmol/L Tris, pH 7.4, with complete protease inhibitor (Roche). The homogenate was incubated for 1 hour at 37°C with 50 units of alkaline phosphatase, then NP-40 was added to a concentration of 1%, and the samples were sonicated. The samples were incubated for 3 hours at 37°C with PNGase (10 units/μL). Homogenate was diluted at 1:5 (vol/vol) with SDS-PAGE loading buffer (15% SDS, 8 mol/L urea, 10% 2-mercaptoethanol, 25% glycerol, 0.2 mol/L Tris pH 6.8), heated at 37°C for 20 minutes, and subjected to 10% SDS-PAGE using the SE260 electrophoresis system (Hoefer). Following electrophoresis, proteins were transferred to a nitrocellulose membrane using the TE22 system (Hoefer). Membranes were visualized with Ponceau S acid stain; scanned to assess protein load; blocked in 5% BSA in 20 mmol/L Tris, 150 mmol/L NaCl, pH 7.4, containing 0.1% Tween-20; probed overnight with primary antibodies to LXR-α or SREBP-2; rinsed; probed for 30 minutes with secondary antibodies; and washed in 20 mmol/L Tris, 150 mmol/L NaCl, pH 7.4 containing 0.1% Tween-20 to remove excess antibody. Immunoreactivity was revealed using enhanced chemiluminescence detection reagent.

In Vitro Assays of PLA2 Enzymatic Activity and Inhibitor Profiles

PLA2 activity was measured by 2 different methods. Dr Fernandez-Patron’s laboratory used the commercial assay kit (Cayman) with diheptanoyl phosphatidyl choline as substrate, per the manufacturer’s instructions. Ex vivo tissue release of sPLA2 activity was equally measured using the Cayman kit, as described earlier.12 Confirmatory studies and the extended characterization of cardiac sPLA2 biochemical properties were conducted by Dr Lambeau’s laboratory using the highly sensitive [3H]-oleic acid–radiolabeled Escherichia coli membrane assay. Of note, E coli membranes are richer in...
Time-Resolved Fluoroimmunoassay for Detection of sPLA₂ Protein Levels

Time-resolved fluoroimmunoassay has equal or higher sensitivity than the [³H]-oleic acid–radiolabeled E coli membrane PLA₂ enzyme assay and was performed as described previously,¹⁷ using antibodies specific for the following sPLA₂: mGIB (PLA2G1B), mGIIA (PLA2G2A), mGIID (PLA2G2D), mGIE (PLA2G2E), mGIIF (PLA2G2F), mGV (PLA2G5), or mGX (PLA2G10)² with 0.5 μL of pooled plasma (20 μg total protein) or 5 μL of heart lysate (31 μg total protein). Samples from 2 WT and 2 Mmp2⁻/⁻ mice were used for time-resolved fluoroimmunoassay. Recombinant sPLA₂ proteins were prepared as described¹⁹ and served as positive controls.

Measurements of Eicosanoids

Prostaglandin E₂ in the heart and liver was measured by the Prostaglandin E₂ Express EIA Kit (Cayman).

Oil Red O Stain

Oil Red O stain was used to image droplets containing neutral lipids (triglycerides, diglycerides, and cholesteryl esters) in 10-μm-thick sections of frozen liver. The sections were cut with a cryostat from Optimal Cutting Temperature-embedded liver pieces. The sections were dehydrated and mounted with a coverslip for microscopic inspection.

Extraction of Lipids From Tissues

Lipids were extracted from tissues, as described previously.²⁰ Tissue was homogenized in 8 to 30 μL of water/mg of tissue using the Bullet Blender. A total of 500 to 1000 μL of homogenate or lysate was mixed at a ratio of 3:4:8 aqueous homogenate/lysate:methanol:chloroform and vortexed for 1 minute and then centrifuged for 15 minutes to separate phases. The bottom phase containing lipids was transferred into a new tube, dried down under argon, and resuspended in 100 μL chloroform to be used for downstream lipid analysis.

Chromatographic Determinations of Lipid Content

High-performance liquid chromatography (HPLC) and fast-performance liquid chromatography analyses were performed at the Lipid and Lipid Metabolite Analysis Core Facility, part of the Women and Children’s Health Research Institute and Faculty of Medicine and Dentistry at the University of Alberta. Hepatic lipids were extracted in the presence of 50 μg dipalmitoyl-phosphatidyl dimethylethanolamine as an internal standard and separated by HPLC, as described previously.²¹ Lipid species were separated in an Onyx monolithic silica normal-phase column (Phenomenex) using a 3-solvent system in an 1100 series HPLC system (Agilent Technologies) and quantified using in-line detection with an Alltech ELSD2000 evaporative light-scattering detector (W. R. Grace). Plasma lipoprotein particles were resolved in a Superose 6 10/300 gel-filtration fast-performance liquid chromatography column isocratically with 50 mmol/L NaCl buffer using a 1200 series HPLC system (Agilent Technologies). Cholesterol or triglycerides in lipoproteins were detected enzymatically by in-line reaction at 37°C.
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Hepatic Cell Isolation and Flow Cytometric Analysis

Hepatic cells were isolated according to a previously described method\(^{22}\) with minor modifications. To enumerate absolute number of immune cell subsets, mice were fully perfused with PBS to minimize contamination of peripheral blood, and livers were cut into same-sized (same-volume) segments. To enumerate absolute number of immune cell subsets, mouse liver was cut into same-sized segments. Each segment was incubated with enzymatic mixture containing 675 U/mL collagenase I, 18.75 U/mL collagenase XI, and 9 U/mL hyaluronidase (Sigma-Aldrich) in calcium- and magnesium-containing Hanks’ balanced salt solution for 40 minutes at 37°C with gentle shaking. After digestion with enzyme mix, cells were washed and further processed by percoll (Roche) gradient to enrich CD45\(^{+}\) immune cells. Single-cell suspensions were incubated in culture supernatant from the 2.42G2 hybridoma (Fc receptor block, ATCC HB-197) prior to staining with the indicated antibodies (BioLegend). Stained cells were acquired using a BD LSQRFortessa flow cytometer (BD Biosciences) and analyzed with FlowJo (Tree Star). Side-scattered light (SSC\text{low}) and forward scattering light (FSC\text{low}) dead cells and doublet cells were gated out and analyzed with FlowJo.

Isolation of Cardiac sPLA\(_2\) by a Multidimensional Orthogonal Strategy

Cardiac tissue was washed 2 times with sterile PBS, finely minced, and homogenized in sterile PBS (1 mL per 100 mg of cardiac tissue) using the Bullet Blender. The homogenate was next subjected to differential centrifugal fractionation. First, we produced a 300 g pellet by centrifuging at 4°C for 10 minutes. The resultant supernatant was centrifuged at 10 000g at 4°C for 30 minutes. The resultant pellet was resuspended in 250 µL of the Cayman sPLA\(_2\) activity kit buffer (25 mmol/L Tris-HCl, pH 7.5, 10 mmol/L CaCl\(_2\), 100 mmol/L KCl, and 0.3 mol/L Triton X-100). An sPLA\(_2\)–rich supernatant was obtained after centrifugation at 100 000g at 4°C for 60 minutes.

Analysis of sPLA\(_2\) activity distribution reproducibly showed that the 100 000g supernatant contained most of the cardiac sPLA\(_2\), regardless of whether frozen or fresh cardiac tissue was subjected to the centrifugal fractionation scheme. We concentrated cardiac sPLA\(_2\) at −20°C by subjecting the 100 000g supernatant to precipitation with acetone. Briefly, supernatant was mixed 1:4 (vol/vol) with acetone, and protein was left to precipitate at −20°C for 2 hours. Following centrifugation at 28 893 g, the organic phase was removed, and precipitated protein was dried at 37°C for 10 minutes, sonicated, and resuspended in 250 µL of Cayman sPLA\(_2\) activity kit buffer. Aliquots were kept at −80°C for later use.

In studies to establish cardiac sPLA\(_2\) as a biological mediator and determinant of the cardiohepatic phenotype of MMP-2–deficient mice, we subjected hearts from 5 different Mmp2\(^{−/−}\) mice to the earlier fractionation scheme. Next, 60 µL of the acetone concentrate was resolved using an HPLC system equipped with a size exclusion column (25 cm × 4.6 mm) packed with Superdex 75 prep grade (GE Healthcare Life Sciences). The separation was carried out under isocratic conditions using sterile PBS supplemented with 10 mmol/L CaCl\(_2\) as the mobile phase and a flow rate of 200 µL/min. Fractions were collected every 10 minutes. The described combination of differential centrifugation, acetone precipitation, and size exclusion chromatography reproducibly yielded a distinct low-molecular-weight (<20 kDa) fraction with high cardiac sPLA\(_2\) activity and little protein content, confirmed by both SDS-PAGE and Bradford assays. The sPLA\(_2\) activity detected when WT hearts were subjected to the above fractionation protocol was negligible. Size exclusion fractions with the most sPLA\(_2\) activity typically corresponded to fractions F6 to F8. In experiments described in this paper, fractions F6 and F7 were combined and used to intraperitoneally inject WT mice (denoted recipient mice) with or without the pan-sPLA\(_2\) inhibitor varespladib (10 mg/kg per day) for 5 consecutive days.

Statistical Analysis

Unless otherwise indicated, the results are reported as mean±SEM. Results were analyzed with the SigmaPlot 11 software (Systat Software) using 1-way ANOVA or Student \(t\) test, as appropriate. ANOVA with repeated-measures analysis was conducted to determine differences among groups in time-course experiments. Whenever the normality and equal variance assumptions were not met, statistical significance (\(P<0.05\)) was determined by applying nonparametric Wilcoxon signed rank sum tests and Kruskal–Wallis tests.

Results

Loss of MMP-2 Affects Systemic Metabolism

Compared with age-matched WT mice, male Mmp2\(^{−/−}\) mice aged 10 to 14 weeks exhibited normal food intake and reduced locomotor activity that were associated with decreased body weight and increased energy expenditure (Figure 1).

Loss of MMP-2 was associated with changes in hepatic and plasma lipids: Fasted Mmp2\(^{−/−}\) mice had elevated triglycerides in the liver (Figure 2A and 2B) and very low-density lipoprotein (Figure 2C, top) and decreased cholesterol in high-density lipoprotein (Figure 2C, bottom). Nonfasted Mmp2\(^{−/−}\) mice also had elevated hepatic triglycerides (Figure 2D).
The hepatic gene expression phenotype in \textit{Mmp2}^{-/-} mice was complex, with some LXR-\(\alpha\) target genes being significantly upregulated, for example, cholesterol 7\(\alpha\)-hydroxylase (the rate-limiting enzyme in the classical bile acid biosynthesis pathway, encoded by \textit{Cyp7a1}) and ATP-binding cassette G5 and G8 (2 cholesterol transporters, encoded by \textit{Abcg5} and \textit{Abcg8}) (Figure 3C).

Moreover, we detected increased hepatic expression of the transcription factor SREBP-2 (encoded by \textit{Srebf2}) and many of its target genes including \textit{Hmgcr}, \textit{Ldlr}, and \textit{Pcsk9} in \textit{Mmp2}^{-/-} mice (Figure 3C). \textit{Hmgcr} encodes 3-hydroxy-3-methylglutaryl-coenzyme A reductase, the rate-limiting enzyme in the cholesterol and isoprenoid synthesis pathways. \textit{Ldlr} encodes the low-density lipoprotein receptor, which is involved in clearance of low-density lipoprotein from circulation. \textit{Pcsk9} encodes proprotein convertase subtilisin/kexin type 9, a protein that binds and negatively regulates hepatic low-density lipoprotein receptor protein levels.\textsuperscript{26} The upregulation of SREBP-2 protein and its proteolytically processed transcriptionally active form (nuclear SREBP-2) was confirmed by western blotting (Figure 3D) and was restricted to cardiac and hepatic tissues (Figure S1). LXR-\(\alpha\) and SREBP-2 expression profiles were similar in \textit{Mmp2}^{-/-} and haploinsufficient (\textit{Mmp2}^{+//-}) mice (Figure 3E, left panel). Furthermore, male and female \textit{Mmp2}^{-/-} mice exhibited similar gene expression profiles, demonstrating lack of sexual dimorphism in the regulation of LXR-\(\alpha\) or SREBP-2 by MMP-2 (Figure 3E, right panel).

Loss of MMP-2 (or MMP-7 or MMP-9, as determined in \textit{Mmp2}^{-/-} mice) (Figure S2) had only a limited impact on carbohydrate and nitrogen metabolism and bile acid production. Of note, insulin resistance was evident in the stress phase of the insulin-tolerance test, which is driven by adrenocorticotropic hormone–mediated signals such as cortisol release from the adrenals.\textsuperscript{27} These data revealed MMP-2 as a modulator of the hepatic LXR and SREBP pathways.

**MMP-2 Modulates Transcriptional Responses to Dietary Cholesterol**

To investigate the contribution of MMP-2 to hepatic transcriptional responses to metabolic cues, we subjected mice to dietary cholesterol supplementation. Dietary cholesterol inhibits the SREBP-2 pathway to decrease hepatic synthesis and uptake of cholesterol; at the same time, oxysterols activate LXR-\(\alpha\) signaling to increase the clearance of hepatic cholesterol.\textsuperscript{28,29} Consistently, WT mice fed chow supplemented with 0.15% cholesterol exhibited decreased hepatic expression of \textit{Srebf2} and \textit{Hmgcr} along with a powerful induction of \textit{Cyp7a1} and \textit{Cyp27a1}, 2 rate-limiting enzymes in the bile acid synthesis pathways. In contrast, in \textit{Mmp2}^{-/-} mice, the same genes (ie, \textit{Srebf2}, \textit{Hmgcr}, \textit{Cyp7a1}, and

The qRT-PCR analysis revealed \textit{Nr1h3} (encoding LXR-\(\alpha\)) to be the most downregulated gene in the liver of \textit{Mmp2}^{-/-} mice (Figure 3A). Reduced LXR-\(\alpha\) protein was confirmed by western blotting (Figure 3B). Consistent with LXR-\(\alpha\) promotion of the transcription of genes involved in fatty acid synthesis,\textsuperscript{23} mRNA expression of hepatic \textit{Fasn} (encoding fatty acid synthase) was repressed in \textit{Mmp2}^{-/-} mice (Figure 3A). Likewise, expression of another LXR-\(\alpha\) target, lysophosphatidylcholine acyltransferase 3 (LPCAT3, encoded by \textit{Lpcat3}), was downregulated (Figure 3A). LPCAT3 catalyzes the attachment of polyunsaturated fatty acids to membrane phospholipids.\textsuperscript{24,25} Downregulation of LXR-\(\alpha\) and \textit{Lpcat3} (one of its target genes) suggested a potential link between the metabolic phenotype of MMP-2 deficiency and hepatic membrane phospholipid remodeling, a process potentially associated with inflammation signaling in MMP-2-deficient mice\textsuperscript{12} and that was investigated in the experiments described below.

**Figure 1.** Systemic metabolic abnormalities of matrix metalloproteinase 2–deficient mice. A, Heat/energy expenditure (normalized to body weight). B and C, Locomotor activity. D, Body weight. E, Total food consumption. *P\(\leq\)0.05 vs WT. The studies were conducted in metabolic cages. WT indicates wild type.

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Figure 2. Matrix metalloproteinase 2 deficiency is associated with abnormalities in lipid storage and transport. A, Oil Red O stain of neutral lipid species from frozen liver sections of fasted mice. Scale bar=30 μm. B, Fasting levels of hepatic lipids in Mmp2−/− vs WT mice as assessed by high-performance liquid chromatography with in-line evaporative light-scattering detection. n=3 per genotype. *P<0.05 vs WT. C, Fasted TG and cholesterol levels in lipoparticle fractions of plasma separated by fast-performance liquid chromatography (left). Traces correspond to pools of plasma from Mmp2−/− and WT. D, Hepatic TGs were elevated in nonfasting mice. Gavage treatment with vehicle or pan-secreted phospholipase A2 inhibitor varespladib; normalized hepatic TG levels in nonfasting Mmp2−/− mice. n=4 individual mice per genotype per group (or treatment). *P≤0.05 vs WT. †P≤0.05 vs Mmp2−/− administered vehicle. CE indicates cholesteryl esters; FA, free fatty acids; FC, free cholesterol; HDL, high-density lipoprotein; mAU, milliabsorbance units; LDL, low-density lipoprotein; PC, phosphatidyl choline; PE, phosphatidyl ethanolamine; PI, phosphatidyl inositol; PS, phosphatidyl serine; SM, sphingomyelin; TG, triglyceride; VLDL, very low-density lipoprotein; WT, wild type.
Figure 3. Lipid metabolic gene expression profile analysis of the hepatic phenotype of MMP-2–deficient mice. A, qRT-PCR analysis identified several downregulated metabolic genes in Mmp2–/– mice. These results are representative of 4 mice per genotype. The data for the LXR-α (encoded by Nr1h3) pathway genes were confirmed in multiple studies and different batches of WT and Mmp2–/– mice. B, Western blot confirming reduced hepatic LXR-α protein in Mmp2–/– mice. Pool of 4 per genotype. C, qRT-PCR analysis identified several upregulated metabolic genes in Mmp2–/– mice. These results are representative of 4 mice per genotype. The data for the SREBP-2 (encoded by Sreb2) pathway genes were confirmed in multiple studies and different batches of WT and Mmp2–/– mice. D, Western blot confirming elevated hepatic SREBP-2 protein in Mmp2–/– mice. Pool of 4 per genotype. E, The major lipid metabolic gene expression abnormalities of MMP-2 deficiency are present in both sexes and in haploinsufficient mice. (Left) Male mice with both copies (Mmp2+/+), lacking 1 copy (Mmp2+/–), or lacking both copies (Mmp2–/–) of Mmp2, n=4 for Mmp2+/+, and n=4 for Mmp2+/–, n=3 for Mmp2–/–. *P<0.05 vs WT. †P<0.05 vs Mmp2+/+. (Right) Female mice. n=5. *P<0.05 vs WT. LXR-α indicates liver X receptor α; MMP-2, matrix metalloproteinase 2; ND, not detected; qRT-PCR, quantitative reverse transcription–polymerase chain reaction; SREBP-2, sterol regulatory binding protein 2; WT, wild type.

Cyp27a) showed blunted transcriptional responses reminiscent of a hepatic insensitivity to dietary cholesterol, whereas other genes such as Sreb1 and Fasn, were clearly uncoupled in their responses (Figure 4). These data identified MMP-2 as a modulator of hepatic transcriptional responses to dietary cholesterol.
A Heart–Liver Axis Mediated by Cardiac sPLA2 and Negatively Regulated by MMP-2

Cardiac sPLA2 localizes to cardiomyocytes

Recently, we reported an MMP-2/cardiac sPLA2 mechanism that modulates blood pressure homeostasis, cardiac inflammation, and lipopolysaccharide-induced fever.12 We confirmed the heart as a major source of systemic sPLA2 activity in Mmp2−/− mice (Figure 5A). We also detected elevated sPLA2 activity in the aorta but not in skeletal muscle or the liver (Figure 5A). Ex vivo release of sPLA2 activity was evident from the heart but not from the liver (Figure 5A and inset). Cardiac sPLA2 activity was significantly elevated in Mmp2−/− mice (versus WT) whether the substrate was diheptanoyl thio-phosphatidyl choline or [3H]-oleic acid–radiolabeled E. coli membranes. Analysis of plasma sPLA2 activity with the highly sensitive E. coli membrane assay indicated a 1000-fold elevation in Mmp2−/− mice (Figure 5A inset, Figure S3). Cardiac sPLA2 activity was evenly distributed across left and right cardiac atria and ventricles (Figure 5B). Cardiomyocytes isolated from Mmp2−/− whole hearts displayed significantly elevated intracellular sPLA2 activity (Figure 5C). Culture of Mmp2−/− cardiomyocytes with recombinant human MMP-2 at 40 nmol/L, a concentration that approximates that of plasma MMP-2, reduced sPLA2 activity (Figure 5C, left panel). Brefeldin A (1 μmol/L), an inhibitor of the classical secretory pathway at the level of the endoplasmic reticulum to Golgi transition, blunted sPLA2 activity in Mmp2−/− cardiomyocytes (Figure 5C, right panel).

Cardiac sPLA2, an elusive unique enzyme, circulates in plasma

We reported previously that cardiac and plasma sPLA2 had similar molecular weights (≈18–20 kDa), had similar apparent Michaelis–Menten constants for diheptanoyl thio-phosphatidyl choline of ≈235 μmol/L, required calcium for activity, and were not cleaved or inactivated by incubation with MMP-2.12 Cardiac and plasma sPLA2 had strikingly similar activity profiles against a panel of inhibitors indicative that the cardiac enzyme is present in the circulation (Figure 5D, top panel). The pan-sPLA2 blocker varespladib30 decreased cardiac sPLA2 activity with a half maximal inhibitory concentration of 10 μmol/L, similar to what we reported for indoxam (≈2 μmol/L)12 and different from PLA2G5 (Figure 5D, bottom panel). Using a time-resolved fluorescence immunoassay with highly specific antibodies against various mouse sPLA2 isoforms, we excluded PLA2G1B, PLA2G2A/2D/2E/2F, PLA2G5, and PLA2G10 as major components of either cardiac or plasma sPLA2 in Mmp2−/− mice (Figure S4).

Systemic sPLA2 activity contributes to the hepatic inflammatory and lipid metabolic phenotype of MMP-2 deficiency

Treatment of Mmp2−/− mice with the pan-sPLA2 inhibitor varespladib (10 mg/kg per day) reduced sPLA2 activity time...
Identification of a Novel Heart–Liver Axis

Hernandez-Anzaldo et al

ORIGINAL RESEARCH

Ex vivo release of sPLA2 activity

WT
Mmp2–/–

Ventricles
Aorta
Pancreas
Small intestine
Liver
Kidney
Skeletal muscle
Adipose

Ex vivo release of sPLA2 activity

sPLA2 activity (µmol/min/mg of tissue)

Liver
Heart

Ex vivo release of sPLA2 activity (% of WT)

WT
Mmp2–/–

Cardiomyocytes

sPLA2 activity (% of WT)

C

Legend:
0: No inhibitor
1: EDTA, non-specific inhibitor of Ca²⁺-dependent PLA₂
2: DTT, SH redox agent
3: MJ33, PLA₂ inhibitor
4: KH064, sPLA₂ inhibitor
5: YM 26734, sPLA₂ (PLAG2A, PLA2G5) inhibitor
6: AACOCF3, cPLA₂ and iPLA₂ inhibitor
7: ACA, PLA₂ inhibitor
8: BEL, iPLA₂ inhibitor
9: Heparin, Inhibits some sPLA₂s

sPLA2 activity (% of vehicle)

[D] [C] [B] [A]

Heart
Plasma

sPLA2 activity (µmol/min/ml)

Heart
Plasma

Remaining PLA2 activity (%)

0 1 2 3 4 5 6 7 8 9

Remaining PLA2 activity (µmol/min/ml)

sPLA2 activity (% of vehicle)

[Varespladib] (µmol/l)

Mmp2–/– hearts (isolation 1)
Mmp2–/– hearts (isolation 2)
mPLA2G5
Figure 5. Tissue profiling and intracardiac localization of sPLA2. A, The heart is the predominant source of sPLA2 activity in Mmp2−/− mice. The sPLA2 activity (pools of 4 mice per genotype) was analyzed in duplicate. *P<0.05 vs WT (statistical unit was the duplicate). (Inset) Ex vivo release of sPLA2 activity was elevated for specimens of left ventricle, but not liver, from Mmp2−/− mice. Ex vivo release results are representative of 3 determinations conducted on different days with different mice.*. B, Intracardiac distribution of sPLA2 activity in Mmp2−/+ mice. Frozen heart sections (≈1 mm thick) were divided into pieces with approximately equal surface area (≈1 mm²), homogenized, and individually assessed for sPLA2 activity. Diagrams present mean values of sPLA2 activity normalized to protein content in each piece of tissue. n=3 mice per genotype. C, Cardiac sPLA2 localizes to cardiomyocytes. Isolated left) and in plasma (Figure 6A, right). Furthermore, varespladib lowered prostaglandin E2 levels in the heart and liver (Figure S5) and fasting nor fasting and refeeding with a high-carbohydrate diet (Figure S7) or dietary supplementation with polyunsaturated fatty acids (data not shown) or cholesterol affected cardiac sPLA2 activity. We observed small increases in the mRNA levels for conventional sPLA2 isoforms in Mmp2−/+ mice (Figure 7). Similarly, varespladib lowered hepatic triglycerides in Mmp2−/+ mice (Figure 2D).

Adaptive transfer of cardiac sPLA2 evokes the cardiohepatic phenotype of MMP-2−/− deficient mice

When HPLC-isolated cardiac sPLA2 from Mmp2−/− donors was injected into WT mice (fractions F6 and F7), the recipient WT mice displayed highly elevated plasma and cardiac sPLA2 activity (Figure 8). Moreover, the cardiohepatic gene expression profile of WT mice that received HPLC-isolated cardiac sPLA2 from Mmp2−/− donors was strikingly similar to that of Mmp2−/− mice. The phenotypic changes evoked by cardiac sPLA2 from Mmp2−/− donors were blunted in mice that also received the pan-sPLA2 inhibitor varespladib (Figure 9).

Evidence in support of MCP-3 as an agonist of cardiac sPLA2

The data to date indicate that cardiac sPLA2 is a unique enzyme produced and released by cardiomyocytes into plasma, where it circulates and affects liver function. Furthermore, cardiac sPLA2 intracellular activation is inhibited by extracellular MMP-2, likely by blocking an agonist.

To identify possible triggers of cardiac sPLA2 activity, we first examined the impact of feeding and fasting. Neither fasting nor feeding and refeeding with a high-carbohydrate diet (Figure S7) or dietary supplementation with polyunsaturated fatty acids (data not shown) or cholesterol affected cardiac sPLA2 activity.

Second, we examined the influence of circadian rhythm on cardiac sPLA2 activity. We observed small increases in the mRNA levels for conventional sPLA2 isoforms in Mmp2−/+ mice euthanized at night but no differences in cardiac sPLA2 activity at night (9:30 PM±0.5 hour) versus day (11:30 AM±0.5 hour) (Figure S8).

Third, we examined whether MCP-3 (a proinflammatory CC-chemokine bound, cleaved, and inactivated by MMP-2) had the potential to induce cardiac sPLA2 activation or secretion.

Although Ccl7 mRNA (encodes MCP-3) was elevated at baseline in both Mmp2−/− liver and heart (Figure 6B and 6C), MCP-3 protein levels were significantly elevated only in the heart, not in the liver or plasma, of Mmp2−/− mice (Figure 10). The reason for the cardiac-specific overexpression of MCP-3 was not investigated further.

To assess the in vivo significance of MCP-3 for cardiac sPLA2 activity, we injected mice with either neutralizing MCP-3 monoclonal antibody or isotype-matched IgG1 at a dose (0.6 mg/kg per day, intraperitoneally) validated in a previous study. Neutralizing MCP-3 antibody (but not control IgG1) treatment normalized cardiac MCP-3 protein levels in Mmp2−/− mice (Figure 10). Ex vivo incubation during 3 hours
Figure 6. High systemic sPLA2 activity sustains the inflammatory and lipid metabolic gene expression phenotype of matrix metalloproteinase 2–deficient mice. A, Cardiac and plasma sPLA2 activity in Mmp2−/− mice administered the pan-sPLA2 inhibitor varespladib (10 mg/kg per day). B, Hepatic expression of inflammatory and lipid metabolic genes in mice administered varespladib (10 mg/kg per day). C, Cardiac expression of inflammatory and lipid metabolic genes in mice administered varespladib (10 mg/kg per day). Analysis of 12 mice per genotype (n=4 per time point). *P≤0.05 vs WT (time 0 days). †P≤0.05 vs Mmp2−/− (time 0 days). sPLA2 indicates secreted phospholipase A2; WT, wild type.
with MCP-3 (200 nmol/L) enhanced the release of sPLA2 activity by 87% from WT heart and 39% from Mmp2/C0/C0 heart; however, sPLA2 release from liver was unchanged regardless of genotype (Figure 11). Most important, neutralizing MCP-3 antibody (but not IgG1) fully normalized cardiac and plasma sPLA2 activity in Mmp2/C0/C0 mice (Figure 12A). Neutralizing MCP-3 antibody partially restored gene expression of Mmp2/C0/C0 mice including LXR-a, SREBP-2, and inflammatory markers in the heart (Figure 12B) and liver (Figure 12C). In contrast to neutralizing MCP-3 antibody, systemic sPLA2 inhibition with varespladib did not normalize MCP-3 protein expression in the heart of Mmp2/C0/C0 mice (Figure 10). These results identified cardiac MCP-3 as a possible agonist acting upstream of cardiac sPLA2; however, the possibility that other cytokines or mechanisms contribute to cardiac sPLA2 activation or release warrants further research.

Discussion

Our study identified a novel endocrine function of the heart: the production and secretion of a unique PLA2 by cardiomyocytes. Secretion of cardiac sPLA2 into the plasma enables a previously unknown heart–liver axis that is activated, at least in part, by the proinflammatory CC-chemokine MCP-3 and negatively regulated by extracellular MMP-2, which cleaves MCP-3.2 This heart–liver endocrine axis profoundly influences the inflammatory and lipid metabolic gene expression characteristics of the liver, effectively increasing the levels of hepatic and very low-density lipoprotein triglycerides. Because the heart is a major user of very low-density lipoprotein triglycerides, the MMP-2/cardiac sPLA2 system may enable the heart to signal the liver to satisfy the energy needs (Figure 13). Our findings identify cardiac sPLA2 as a
Figure 8. HPLC isolation of biologically active cardiac sPLA₂. A, Strategy for isolation of biologically active cardiac sPLA₂ for in vivo studies. Hearts from 5 Mmp2⁻/⁻ or WT mice were pooled to obtain sufficient cardiac sPLA₂ for in vivo characterization. B, Representative centrifugal fractionation. Note that the 100 000g supernatant fraction is rich in active cardiac sPLA₂, and WT hearts express negligible cardiac sPLA₂ activity, as opposed to Mmp2⁻/⁻ hearts. C, Representative HPLC isolation of active cardiac sPLA₂ and Coomassie blue–stained 16% T, 3% C Tricine–SDS-PAGE gels of 100 000g supernatant before and after size-exclusion fractionation. D, Phenotypes induced by HPLC fractions F6 and F7 from Mmp2⁻/⁻ donors vs WT donors in recipient WT mice. Injection of HPLC fractions F6 and F7 from Mmp2⁻/⁻ donors (but not from WT donors) evoked high sPLA₂ activity in both plasma and the heart of WT recipient mice after 5 consecutive injection-days. For comparison, we show the levels of activity in intact Mmp2⁻/⁻ mice. WT plus vehicle: Mice received 100 μL of HPLC mobile phase (vehicle) consistent with sterile PBS, 10 mmol/L CaCl₂. WT plus fractions F6 and F7: Mice received HPLC fractions F6 and F7 from WT donors and from Mmp2⁻/⁻ donors. *P<0.05 vs WT. n=4 mice per treatment. AU indicates arbitrary units; HPLC, high-performance liquid chromatography; sPLA₂, secreted phospholipase A₂; WT, wild type.
key biological mediator of the cardiohepatic phenotype associated with MMP-2 deficiency in mice. These findings could also help explain how MMP2 deficiency leads to cardiac problems and delayed growth in patients.4–6,32

The endocrine function of the heart was first suggested in the days of the discovery of the circulatory system by William Harvey in the 17th century.33 The concept has since evolved through the work of many. In 1921, Loewi discovered the
release of acetylcholine from cardiac sympathetic nerves. Investigations conducted between 1971 and 1983 by de Bold’s team revealed the secretion of cardiac natriuretic peptides from granules present in the atria (but absent in ventricles). In addition to natriuresis, cardiac natriuretic peptides mediate vasodilation in the failing heart, increase gluconeogenesis in the liver, regulate lipolysis in adipocytes, and promote thermogenesis and energy expenditure. The heart is also source, albeit not exclusively, of other polypeptide hormones, including adrenomedullin and endothelins, which—acting in autocrine, paracrine, and endocrine fashion—provide additional cardiovascular and metabolic regulatory mechanisms.

Likewise, the heart may regulate white adipose tissue and liver function through as-yet-unidentified factors produced and released downstream of mediator complex subunit 13, a protein with transcription that is negatively regulated by the heart-specific microRNA-208a. This latter mechanism may promote resistance to obesity induced by a high-fat diet and improve insulin sensitivity. Bidirectional interactions between the heart and the liver have been demonstrated in mice bearing a hypertrophic cardiomyopathy-causing mutation in myosin (R403Q). This mutation is associated with decreased cardiac lipid uptake and increased plasma and hepatic lipid content together with increased gluconeogenesis and blood glucose that may ultimately exacerbate cardiac disease.

Our previous studies revealed autocrine/paracrine actions of cardiac sPLA2 involving the modulation of inflammatory and lipid metabolic gene expression in the heart. Furthermore, we documented the systemic effects of cardiac sPLA2 on systolic blood pressure and fever. Extending these observations, our current research identifies the liver as the first target organ of cardiac sPLA2.

Pioneering work on MMP-2–deficient mice revealed lipodystrophy as an important trait. In this study, we identified new metabolic traits of MMP-2 deficiency. These include (1) hepatic lipid deposition; (2) altered expression of transcription factors and enzymes involved in sensing, biosynthesis, and catabolism of lipids in the liver; (3) altered transcriptional responses to dietary cholesterol; and (4) abnormal oxygen consumption and locomotor activity patterns.

**Figure 10.** Cardiac MCP-3 expression is upregulated in Mmp2−/− mice and unrelated to systemic secreted phospholipase A2 activity. A, MCP-3 protein levels in the heart, plasma, and liver of mice administered nMCP-3 Ab or isotype-matched IgG1 (0.6 mg/kg per day). Results are representative of 19 to 20 mice per genotype (n=3 to 4 mice per time point). *P<0.05 vs WT at day 0. †P<0.05 vs Mmp2−/− plus IgG1, at day 5. Note that the data presented for day 2.5 refers to pools of 3 to 4 mice for which MCP-3 levels were measured in duplicate; for simplicity, no statistical analysis is indicated at day 2.5. B, MCP-3 protein levels in the heart, plasma, and liver of mice administered varespladib (10 mg/kg per day). Results are representative of 12 to 13 mice per genotype (3 to 5 mice per time point were pooled and MCP-3 levels were measured in duplicate). *P<0.05 vs WT at day 0. †P<0.05 vs Mmp2−/− at day 5. MCP-3 indicates monocyte chemoattractant protein 3; nMCP-3 Ab, neutralizing MCP-3 monoclonal antibody; WT, wild type.

**Figure 11.** MCP-3 enhances sPLA2 activity release from myocardium. MCP-3 enhanced the ex vivo release of sPLA2 activity from heart but not liver. Freshly dissected specimens of the indicated organs (≈2 mm3) were subjected to the ex vivo release bioassay described previously in the presence or absence of MCP-3 (200 nmol/L), n=3 mice. Arrows indicate when MCP-3 or PBS (vehicle) was added. *P<0.05 vs time 0 hours. †P<0.05 vs vehicle (passive ex vivo release). MCP-3 indicates monocyte chemoattractant protein 3; sPLA2, secreted phospholipase A2; WT, wild type.
Our study provides several lines of evidence indicating that this complex phenotype is caused, at least in part, by activation of a heart–liver axis mediated by cardiac sPLA₂. This axis is broken down as follows. First, systemic absence of MMP-2 increases the bioavailability of MCP-3; this was shown by the cardiac-specific excess of MCP-3 protein observed in this study.

Figure 12. MCP-3 affects the cardiohepatic phenotype of Mmp2<sup>−/−</sup> mice mimicking effects of sPLA₂. A, Cardiac and plasma sPLA₂ activity levels in mice administered nMCP-3 Ab or isotype-matched IgG₁ (0.6 mg/kg per day). B, Cardiac gene expression in mice administered nMCP-3 Ab or IgG₁ (0.6 mg/kg per day). C, Hepatic gene expression in mice administered nMCP-3 Ab or IgG₁ (0.6 mg/kg per day). Analysis of 9 to 12 mice per genotype (n=3 to 4 per time point). *P≤0.05 vs WT (time 0 days). †P≤0.05 vs Mmp2<sup>−/−</sup> (time 0 days). AU indicates arbitrary units; MCP-3, monocyte chemoattractant protein 3; nMCP-3 Ab, neutralizing MCP-3 monoclonal antibody; sPLA₂, secreted phospholipase A₂; WT, wild type.
Second, MCP-3 enhances cardiac sPLA$_2$ release. Once secreted, cardiac sPLA$_2$ reaches distant target organs (eg, liver) where hydrolysis of the outer leaflet of membrane phospholipids releases sn-2 esterified fatty acids. Such fatty acids include precursors of eicosanoids and ligands of nuclear metabolic transcription factors such as LXR-$\alpha$. Downstream changes in the transcription of multiple inflammatory and lipid metabolic genes would profoundly affect cardiac, hepatic, and systemic metabolism. Indeed, we detected abnormally high prostaglandin $E_2$ levels and significant dysregulation of the LXR-$\alpha$ and SREBP-2 pathways in the liver of MMP-2–deficient mice. Moreover, these mice exhibited hepatic immune cell infiltration, specifically, monocytes and macrophages, which are key cells for inflammation. We also detected elevated numbers of dendritic cells, T cells, B cells, and natural killer cells; however, the contribution of these cell types to the observed phenotype requires further investigations.

Importantly, treatment of mice with the pan-sPLA$_2$ inhibitor varespladib almost fully normalized the altered expression of inflammatory and lipid metabolic genes would profoundly affect cardiac, hepatic, and systemic metabolism.

MMP-2 serves as a metabolism modulatory signal and may exert multipronged anti-inflammatory actions. Consequenlty, one may expect that pharmacological MMP-2 inhibi-

Figure 13. Proposed model for a novel heart–liver axis mediated by cardiac sPLA$_2$. Key elements of the proposed model are the heart (source and autocrine/paracrine target of cardiac sPLA$_2$), MCP-3 (agonist of cardiac sPLA$_2$), MMP-2 (endogenous inhibitor), cardiac sPLA$_2$ (signaling molecule), and liver (a noncardiac target of cardiac sPLA$_2$). We propose that, under physiological conditions, MMP-2 activity maintains low levels of certain chemokines, such as MCP-3, by proteolysis. MMP-2 deficiency (due to functional blockade or genetic deletion) and MCP-3 may trigger cardiac sPLA$_2$ release from the heart. Cardiac sPLA$_2$ circulates in the plasma, reaching distant target organs (eg, liver) and affecting the inflammatory and lipid metabolic phenotype of these organs. The MMP-2/cardiac sPLA$_2$ system may serve multiple purposes including signaling to the liver to provide the heart with TGs and maintaining systemic blood pressure homeostasis. LXR-$\alpha$ indicates liver X receptor-$\alpha$; MCP-3 indicates monocyte chemoattractant protein 3; MMP-2, matrix metalloproteinase 2; sPLA$_2$, secreted phospholipase A$_2$; SREBP-2, sterol regulatory binding protein 2; TG, triglyceride; VLDL, very low-density lipoprotein.

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mation in the liver. The key component of this heart–liver axis is secretion of cardiac sPLA2, a unique enzyme the release of which from cardiomyocytes is enhanced by chemokines that are substrates of MMP-2–mediated proteolysis, such as MCP-3. We propose that the MMP-2/cardiac sPLA2 system enables the heart to signal the liver to satisfy the energy needs. Our findings may support understanding of the development of metabolic, inflammatory, and cardiac disturbances associated with MMP-2 deficiency in patients, and may provide a rationale for the development of future therapies.

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Author Contributions: Hernandez-Anzaldo: Conducted vast majority of experiments, conducted statistical analysis. Berry: Conducted phospholipase A2 (PLA2) and hepatic gene determinations and contributed to data analysis. Brigelj, Lambeau: Characterized cardiac and plasma PLA2 inhibition profiles, time-resolved fluoroimmunoassay measurements and contributed to data analysis. Leung: Measured cardiac secreted PLA2 distribution in ventricles and atria. Yun, Lee, Cheong: Determined and analyzed immune cell infiltration. Kassiri: Provided the MMP-2 deficient mice and contributed to data analysis. Filep, Cheong, Kassiri, Lambeau and Lehner: Provided critical conceptual input and proof-read the manuscript. Fernandez-Patron: Directed the research, provided most of the funding and wrote the manuscript.

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Disclosures

None.

References


Supplemental Figure S1. qRT-PCR analysis of *Srebf2* and *Hmgcr* in various tissues from WT and *Mmp2*−/− mice. *n=4 per genotype. *: *P* < 0.05 vs. WT.
Supplemental Figure S2.

(A) Quantitative analysis of mRNA levels for genes involved in glucose metabolism (related to regular Figure 2A, C). *: P<0.05 vs. WT. 

(B) Glucose tolerance test. n=4 individual mice per genotype. No difference between genotypes.

(C) Insulin tolerance test. *: P<0.05 vs. WT.

(D) Quantitative analysis of metabolites. Pools of n=4 per genotype. No difference was observed between genotypes.
Supplemental Figure S3.
PLA$_2$ activity of individual WT and $Mmp2^{-/-}$ hearts and pooled plasma samples determined using the *in vitro* $[^3]$H-oleic acid radiolabelled *E.coli* membrane assay. $n=4$ (WT) and $n=5$ ($Mmp2^{-/-}$).
Supplemental Figure S4. Time-resolved fluorescence immunoassay of mouse sPLA$_2$ isoforms.

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Supplemental Figure S5. The inhibition of systemic sPLA$_2$ by varespladib lowered cardiac as well as hepatic PGE$_2$ levels. Data are related to our earlier study\textsuperscript{1}. The PGE$_2$ levels of pools of $n=4$ mice per genotype were analyzed in duplicate. $*: P \leq 0.05$ vs. WT (-). †: $P \leq 0.05$ vs. $Mmp2^{-/-}$ (-).
Supplemental Figure S6. Phenotype induced by HPLC-fractions F6/F7 from WT donors in recipient WT mice.

(A) Injection of the HPLC-fractions F6/F7 had no impact on plasma or cardiac sPLA₂ of WT recipient mice.

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Supplemental Figure S7. Effect of feeding on cardiac sPLA2 activity.
Neither overnight fasting alone nor overnight fasting followed by re-feeding a high carbohydrate diet for 5 hours did alter systemic sPLA2 activity in WT or Mmp2-/- mice. Pools of n=4 per genotype. *:P<0.05 vs. WT.
Supplemental Figure S8. Effect of circadian rhythm on cardiac sPLA₂ activity.

(A) Comparison of diurnal and nocturnal sPLA₂ activity for WT and Mmp2⁻/⁻ hearts. n=4 mice per genotype. *: P≤0.05 vs. WT (Day).

(B) qRT-PCR analysis for a panel of individual mouse sPLA₂ isoforms at day (11:30 AM ± 0.5 hr) and night (9:30 PM ± 0.5 hr). n=3 for WT and n=4 for Mmp2⁻/⁻. *: P≤0.05 vs. Day.
REFERENCE

Identification of a Novel Heart–Liver Axis: Matrix Metalloproteinase–2 Negatively Regulates Cardiac Secreted Phospholipase A2 to Modulate Lipid Metabolism and Inflammation in the Liver

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