Thrombin Inhibition with Dabigatran Protects against High-Fat Diet–Induced Fatty Liver Disease in Mice

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ABSTRACT

Nonalcoholic fatty liver disease (NAFLD) is the hepatic manifestation of obesity and metabolic syndrome. Robust coagulation cascade activation is common in obese patients with NAFLD. We identified a critical temporal relationship between thrombin generation and the manifestation of hepatic steatosis, inflammation, and injury in C57BL/6J mice fed a high-fat diet (HFD) for 1, 2, and 3 months. Mice fed a HFD exhibited dramatic increases in hepatocellular injury and inflammation over time. Hepatic fibrin deposition preceded an increase in serum alanine aminotransferase, and the most dramatic changes in liver histopathology occurred in conjunction with a detectable increase in plasma thrombin-antithrombin levels at 3 months. To directly determine whether thrombin activity promotes NAFLD pathogenesis, mice were fed a HFD and simultaneously treated with the direct thrombin inhibitor dabigatran etexilate for 3 months. Notably, dabigatran treatment significantly reduced hepatic fibrin deposition, hepatic inflammation, hepatocellular injury, and steatosis in mice fed a HFD. Of interest, dabigatran treatment also significantly attenuated HFD-induced body weight gain. Gene expression analysis suggested that thrombin potentially drives NAFLD pathogenesis by altering the expression of genes associated with lipid metabolism and bile acid synthesis. Collectively, the results suggest that thrombin activity is central to HFD-induced body weight gain, liver injury, and inflammation and provide the proof-of-principle evidence that pharmacological thrombin inhibition could be effective in limiting NAFLD and associated pathologies.

Introduction

Nonalcoholic fatty liver disease (NAFLD) is a common feature of obesity that increases the risk of developing cardiovascular disease and type II diabetes (Targher et al., 2010a). Atypical accumulation of triglycerides within hepatocytes occurs as a consequence of unbalanced lipid metabolism, which manifests histologically as micro- and macrovesicular hepatic steatosis. Hepatic inflammation accompanying the NAFLD spectrum varies in severity and can progress to severe steatohepatitis (NASH) and fibrosis, a leading cause of liver-related morbidity. Hepatic inflammation and injury dramatically increase the risk of cardiovascular disease, the leading cause of morbidity and mortality in patients with NAFLD/NASH (Targher et al., 2005, 2010a; Hotamisligil, 2006; Ndumele et al., 2011). Despite the identified relationship between liver inflammation and poor patient outcomes, the precise triggers of disease-amplifying processes, such as local inflammatory cell activation, are not completely understood.

Profound activation of the blood coagulation cascade is evident in patients with obesity and NAFLD, as indicated by increased plasma levels of thrombin, a serine protease (Ay et al., 2010; Beijers et al., 2010; Fritsch et al., 2010; Semeraro et al., 2012). Similarly, coagulation is evident in mouse models of NAFLD/NASH (Kassel et al., 2010, 2011; Luyendyk et al., 2010; Owens et al., 2012). Recent studies indicate that the coagulation cascade activation contributes to obesity-associated sequelae, including fatty liver disease (Kassel et al., 2011, 2012; Owens et al., 2012). Studies in mice suggest that procoagulant and signaling functions of tissue factor contribute to high-fat diet (HFD)–induced obesity and fatty liver disease (Kassel et al., 2011; Owens et al., 2012). Mice lacking the thrombin receptor, protease activated receptor (PAR-1) are protected from HFD-induced NAFLD, despite similar body weight gain (Kassel et al., 2011). However, the broader role of thrombin activity including the relationship between clotting function and HFD-induced NAFLD has not been completely addressed.

ABBRévIATIONS: A Fib, atrial fibrillation; ALT, alanine aminotransferase; CD, control diet; HFD, high-fat diet; NAFLD, nonalcoholic fatty liver disease; NASH, nonalcoholic steatohepatitis; PAR-1, protease activated receptor 1; TAT, thrombin-antithrombin.
In this study, we determined the temporal connection between coagulation cascade activation relative to the pathogenesis of systemic and hepatic features of HFD-induced NAFLD in mice. Moreover, we tested the hypothesis that prophylactic administration of the Food and Drug Administration–approved orally bioavailable thrombin inhibitor, dabigatran etexilate, would inhibit HFD-induced obesity and NAFLD.

Materials and Methods

Mice and Diets. Wild-type male C57BL/6j mice were purchased from The Jackson Laboratory (Bar Harbor, ME). Mice were between 6 and 7 weeks of age at the start of each study. For the time course study mice were fed a control diet (AIN-93M (10% kcal from fat); Dyets Inc., Bethlehem, PA) or a HFD (Diet #100244 (40% kcal from milk fat); Dyets Inc.) for 1, 2, or 3 months. For the thrombin inhibitor study, mice were fed the same HFD formulated to contain the pulverized content of dabigatran etexilate capsules such that drug content was 10 g/kg for 3 months. An identical diet without drug material was fed for comparison. To avoid the challenge associated with bitter taste of the drug, both diets contained non-nutritive peanut flavoring and peanut butter (2 and 10 g/kg, respectively). After 1 week of acclimation, food intake for each cage and body weights of the mice were measured weekly. Animals were housed in Association for Assessment and Accreditation of Laboratory Animal Care–approved facilities at either the University of Kansas Medical Center or Michigan State University. Mice were housed at an ambient temperature of 22 ± 2°C with alternating 12-hour light/dark cycles and provided free access to purified drinking water. All animal procedures were approved by the University of Kansas Medical Center and/or Michigan State University Institutional Animal Care and Use Committees.

Sample Collection. Food was removed overnight before sample collection. Mice were anesthetized using isoflurane, and blood was collected from the caudal vena cava into a syringe containing sodium citrate (0.38% final) for collection of plasma and into an empty syringe for collection of serum. The liver was excised and washed in phosphate-buffered saline. A small section (~20 mg) of the right medial lobe was collected for RNA isolation and snap frozen in liquid nitrogen. The remainder of the right medial lobe was affixed to a cork using optimal cutting temperature compound (Fisher Scientific, Pittsburgh, PA) and immersed for approximately 3 minutes in liquid nitrogen–chilled 2-methylbutanate (Fisher Scientific). Multiple sections from the left lateral lobe were fixed in 10% neutral-buffered formalin for approximately 48 hours and then embedded in paraffin. The remaining liver was snap frozen in liquid nitrogen.

Serum Alanine Aminotransferase and Plasma Thrombin-Antithrombin Determination. Blood samples were spun at 4000g for 10 minutes or at 10,000g for 2 minutes for collection of plasma or serum, respectively. Serum alanine aminotransferase (ALT) activity was measured using a commercially available reagent (Infinity ALT/GPT; Thermo Fisher, Waltham, MA). Plasma thrombin-antithrombin (TAT) levels were determined using a commercial enzyme-linked immunosorbent assay kit (Enzygnost TAT micro; Siemens Healthcare Diagnostics, Deerfield, IL). For each assay, data were collected using an Infinite M200 plate reader (Tecan, Durham, NC).

Histopathology and Immunohistochemistry. Paraffin-embedded livers were sectioned at 5 μm and stained with H&E by the Michigan State University Investigative Histology Laboratory, a division of Human Pathology. At least 2 to 3 sections of liver from the left lateral lobe from each animal were evaluated by light microscopy to qualitatively determine the severity of micro- and macrovesicular steatosis. The average number of inflammatory foci (>5 clustered mixed inflammatory cells) in each 100× field was determined for all sections in each liver sample. Paraffin-embedded sections were stained for insoluble fibrin using a polyclonal rabbit anti-mouse fibrin antibody, as described previously (Flick et al., 2007), with alkaline phosphatase–conjugated anti-rabbit IgG (H+L) (Sigma-Aldrich, St. Louis, MO) used with Fast Red detection system (Sigma-Aldrich). Macrophage staining was performed by the Michigan State University Histopathology Laboratory. Briefly, paraffin-embedded livers were sectioned at 4 μm, deparaffinized, and subjected to enzymatic epitope retrieval using 0.3% Pronase E (Sigma-Aldrich). After blocking with rabbit serum, slides were incubated with rat anti-mouse F4/80 antibody (clone CI: A3-1; AbD Serotec, Raleigh, NC) at 1:100 dilution in normal antibody diluent (Scytek Laboratories, Logan, UT) for 60 minutes. Tissues were then incubated with biotinylated rabbit anti-rat IgG (H+L), mouse absorbed (Vector Laboratories, Burlingame, CA), antibody at 10 μg/ml in normal antibody diluent and incubated for 30 minutes, followed by R.T.U. Vector Elite Peroxidase Reagent (Vector Laboratories) for 30 minutes. Slides were then developed utilizing Vector NovaRED kit (Vector Laboratories) peroxidase chromogen incubation for 15 minutes followed by counterstain in Gill Hematoxylin (Thermo Fisher) for 30 seconds, differentiation, and dehydration, clearing, and mounting with Permount mounting media.

Determination of Triglycerides and Cholesterol in Serum and Liver and Plasma Glucose and Insulin Levels. Serum triglyceride and cholesterol levels were determined using Triglycerides GPO Liquid and Cholesterol Liquid reagent sets (Pointe Scientific Inc., Canton, MI), respectively, according to the manufacturer’s protocol. For hepatic sample analyses, lipids were extracted from 100 mg frozen livers as described previously (Luyendyk et al., 2010). Plasma insulin was determined using a commercial ultrasensitive insulin enzyme-linked immunosorbent assay (Alpco, Salem, NH), and plasma glucose was measured using commercial reagent (Pointe Scientific Inc.) according to the manufacturer’s protocols.

RNA Isolation, cDNA Synthesis, and Quantitative Real-Time Polymerase Chain Reaction. RNA isolation, cDNA synthesis, and quantitative real-time polymerase chain reaction were performed as described previously (Kassel et al., 2011, 2012). Gene expression was normalized to Hprt levels. A complete list of all gene names and primer sequences are provided in Supplemental Table 1.

Statistical Analyses. Comparison of two groups was performed using Student’s t test. Comparison of three or more groups was performed using two-way analysis of variance and Student-Newman-Keuls post hoc test. Data were considered significant when the P value was less than 0.05.

Results

HFD-Induced Systemic and Hepatic Coagulation Activity Track with the Severity of Hepatic Steatosis and Inflammation as well as Body Weight Gain, Liver Weight, and Lipid Profiles. To determine the temporal relationship between the development of fatty liver disease and evidence of local coagulation system activity, mice were fed either a control diet (CD) or HFD for 3 months, with subsets of animals taken at monthly intervals for analysis. Liver histology in mice fed a CD at each time was similar and unremarkable. Minimal microvesicular steatosis was evident in livers of mice fed a HFD for 1 month (Fig. 1A). Widespread microvesicular steatosis and infrequent macrovesicular steatosis was evident in livers of mice fed a HFD for 2 months (Fig. 1A), whereas marked pericentral mixed macro- and microvesicular steatosis was observed in livers of mice fed a HFD for 3 months (Fig. 1A). At the same time points and with samples from the same animals in which liver histology was evaluated, TAT levels, a stable marker of thrombin generation and robust indicator of coagulation activity, as well as hepatic fibrin deposition, were measured. Little to no hepatic fibrin deposition (red staining) was observed in mice fed a CD (Fig. 1B). In contrast, fibrin deposition was readily evident in mice fed a HFD for 2 and 3 months and was generally localized in the liver sinusoids (Fig. 1B). Compared with mice fed a CD, plasma TAT levels
increased significantly in mice only fed a HFD at 3 months (Fig. 1C). Together, these results suggest that local hepatic coagulation, as indicated by fibrin deposition, precedes a detectable increase in plasma TAT. Moreover, there is a temporal connection between HFD-induced fatty liver progression and activation of coagulation.

Hepatic inflammation, marked by an increase in foci of mixed-inflammatory cell infiltration (Fig. 2A) alongside marked elevations in serum ALT activity (Fig. 2B), was present in mice fed a HFD for 3 months. Immunohistochemical analysis confirmed distinct macrophage accumulation at 3 months (Fig. 2C), corresponding with increased gene expression of proinflammatory cytokines Ccl2 and Tnf at this time point (Fig. 2D). The increased hepatic inflammation and hepatic injury paralleled the robust increase in hepatic steatosis and coagulation activity at 3 months.

As expected, the development of hepatic steatosis, coagulation activity, and inflammation in mice fed a HFD coincided with significantly increased body weight, relative to mice fed a CD. The largest weight gain occurred between 2 and 3 months (Table 1). Whole liver weight and relative liver weight increased significantly at 2 and 3 months in mice fed a HFD (Table 1). Compared with mice fed a CD, serum triglyceride levels transiently decreased at 1 month in mice fed a HFD but were otherwise unaffected (Table 1). In contrast, liver triglyceride levels increased at both 2 and 3 months in HFD-fed mice (Table 1). A time-dependent increase in serum cholesterol occurred in HFD-fed mice (Table 1). In contrast, hepatic cholesterol levels were similarly elevated at each time point (Table 1). Plasma glucose and plasma insulin levels showed a time-dependent increase in animals fed a HFD, with significant increases at 2 and 3 months (Table 2).

Pharmacologic Thrombin Inhibition Protects against HFD-Induced NAFLD and Weight Gain in Mice. Our time course study indicated that increased cellular inflammation, severe steatosis, and elevated serum ALT levels were closely preceded by hepatic fibrin deposition and tightly linked to significantly increased plasma TAT levels. This suggested a direct connection between thrombin activity and fatty liver pathogenesis in HFD-fed mice. Accordingly, we hypothesized that administration of a direct thrombin inhibitor would reduce primary features of HFD-induced fatty liver disease in mice. To test this hypothesis we used dabigatran etexilate, a novel, orally bioavailable direct thrombin inhibitor, used clinically to prevent thrombosis in nonvalvular atrial fibrillation (A-Fib) (Hankey and Eikelboom, 2011). After treatment with dabigatran, liver histopathology revealed a near complete prevention of steatosis relative to mice fed only a HFD (Fig. 3A). Thrombin inhibition with dabigatran significantly reduced serum ALT levels (Fig. 3B), focal hepatic mixed inflammatory cell infiltration (Fig. 3C), macrophage accumulation (Fig. 3D), and hepatic mRNA expression of proinflammatory cytokines Ccl2 and Tnf (Fig. 3E) in mice fed a HFD for 3 months. Moreover, compared with mice fed a HFD for 3 months, plasma thrombin time was markedly prolonged by dabigatran treatment (not shown). Finally, dabigatran also dramatically reduced hepatic fibrin deposition in mice fed through aortic coarctation 3 months (Fig. 3F).
a HFD (Fig. 3F), consistent with previous studies demonstrating a dose of 10 g/kg chow ad libitum is anticoagulant in mice (Bogatkevich et al., 2011; Antoniak et al., 2013). Surprisingly, dabigatran significantly reduced body weight gain in mice fed a HFD (Fig. 4, A and B). This could not be attributed to a difference in food consumption (not shown). Dabigatran significantly decreased liver weight and relative liver weight, indicating a body weight-independent effect of dabigatran on liver size (Fig. 4, C and D). Moreover, thrombin inhibition significantly reduced serum and hepatic triglyceride levels (Fig. 4, E and F), as well as serum and hepatic cholesterol in mice fed a HFD (Fig. 4, G and H). Finally, dabigatran did not have any effect on plasma insulin or plasma glucose in mice fed a HFD (Fig. 4, I and J).

Thrombin Inhibition Suppresses Genes Controlling Lipid Metabolism and Induces Genes Implicated in Bile Acid Synthesis in Mice Fed a HFD. Thrombin inhibition with dabigatran etexilate significantly reduced body and liver weights and hepatic steatosis in mice fed a HFD for 3 months. To anchor these phenotypic differences to underlying gene expression changes, we used quantitative real-time polymerase chain reaction to assess hepatic expression of putative genes involved in lipid metabolism and bile acid synthesis.

Thrombin Inhibition In Diet-Induced Obesity and Fatty Liver

**TABLE 1**

<table>
<thead>
<tr>
<th>Time Point</th>
<th>Diet</th>
<th>Body Weight</th>
<th>Body Weight Gain</th>
<th>Liver Weight</th>
<th>Relative Liver Weight</th>
<th>Serum Triglycerides</th>
<th>Hepatic Triglycerides</th>
<th>Serum Cholesterol</th>
<th>Hepatic Cholesterol</th>
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<tr>
<td></td>
<td>g</td>
<td>%</td>
<td>g</td>
<td>g</td>
<td>%</td>
<td>mg/dl</td>
<td>mg/g liver</td>
<td>mg/dl</td>
<td>mg/g liver</td>
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<tr>
<td>1 month</td>
<td>CD</td>
<td>25.7 ± 1.6</td>
<td>22.3 ± 10.7</td>
<td>0.86 ± 0.08</td>
<td>3.3 ± 0.2</td>
<td>115.5 ± 13.4</td>
<td>20.3 ± 5.5</td>
<td>133.4 ± 8.9</td>
<td>5.5 ± 0.4</td>
</tr>
<tr>
<td></td>
<td>HFD</td>
<td>27.3 ± 1.6</td>
<td>37.6 ± 11.0</td>
<td>1.00 ± 0.14</td>
<td>3.7 ± 0.4</td>
<td>91.5 ± 7.4*</td>
<td>15.3 ± 5.5</td>
<td>126.3 ± 9.0</td>
<td>7.8 ± 1.2*</td>
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<tr>
<td>2 months</td>
<td>CD</td>
<td>28.5 ± 1.5</td>
<td>46.3 ± 11.2</td>
<td>0.90 ± 0.07</td>
<td>3.2 ± 0.1</td>
<td>110.8 ± 11.1</td>
<td>19.7 ± 4.1</td>
<td>132.1 ± 11.4</td>
<td>5.0 ± 0.4</td>
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<tr>
<td></td>
<td>HFD</td>
<td>32.5 ± 3.4*</td>
<td>73.0 ± 18.8*</td>
<td>1.44 ± 0.45*</td>
<td>4.4 ± 0.9*</td>
<td>103.4 ± 12.8</td>
<td>26.3 ± 7.6*</td>
<td>156.1 ± 26.4*</td>
<td>7.3 ± 1.4*</td>
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<tr>
<td>3 months</td>
<td>CD</td>
<td>29.4 ± 2.5</td>
<td>57.6 ± 5.7</td>
<td>0.91 ± 0.11</td>
<td>3.1 ± 0.1</td>
<td>111.7 ± 17.5</td>
<td>23.1 ± 2.7</td>
<td>131.4 ± 8.9</td>
<td>5.3 ± 0.3</td>
</tr>
<tr>
<td></td>
<td>HFD</td>
<td>43.1 ± 3.2*</td>
<td>120.6 ± 23.2*</td>
<td>2.58 ± 0.66*</td>
<td>5.9 ± 1.2*</td>
<td>108.0 ± 5.1</td>
<td>31.5 ± 4.8*</td>
<td>199.8 ± 22.2*</td>
<td>7.6 ± 0.7*</td>
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</tbody>
</table>

*P < 0.05 compared with time-matched CD measured by analysis of variance followed by Student-Newman-Keuls post hoc test.
TABLE 2

<table>
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<tr>
<th>Time Point</th>
<th>Diet</th>
<th>Plasma Insulin (pmol/l)</th>
<th>Plasma Glucose (mg/dl)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1 month</td>
<td>CD</td>
<td>17.8 ± 4.9</td>
<td>150.4 ± 17.8</td>
</tr>
<tr>
<td></td>
<td>HFD</td>
<td>19.2 ± 4.0</td>
<td>164.4 ± 34.3</td>
</tr>
<tr>
<td>2 months</td>
<td>CD</td>
<td>18.0 ± 7.0</td>
<td>133.8 ± 31.1</td>
</tr>
<tr>
<td></td>
<td>HFD</td>
<td>37.0 ± 21.2*</td>
<td>196.8 ± 60.2*</td>
</tr>
<tr>
<td>3 months</td>
<td>CD</td>
<td>20.7 ± 9.1</td>
<td>112.3 ± 17.2</td>
</tr>
<tr>
<td></td>
<td>HFD</td>
<td>51.9 ± 17.0*</td>
<td>187.6 ± 13.4*</td>
</tr>
</tbody>
</table>

*P < 0.05 compared with time-matched CD measured by analysis of variance followed by Student-Newman-Keuls post hoc test.

Temporal effects of HFD and CD on plasma insulin and plasma glucose levels in C57BL/6J mice

Data are mean ± S.D. for 7–9 mice per group.

gene products known to control diet-induced obesity and fatty liver, including the fatty acid desaturase Scd1 (Fig. 5A), fatty acid transporter Cd36 (Fig. 5B), fatty acid “sensor” Cidea (Fig. 5C), and nuclear receptor Pparg (Fig. 5D) (Scaglia et al., 2005; Zhou et al., 2008, 2012; Moran-Salvador et al., 2011). Analysis revealed a temporal connection between the onset of hepatic fibrin deposition and upregulation of each gene (Fig. 5, A–D) in HFD-fed mice, with the largest magnitude of gene induction occurring at 3 months, a time coinciding with a significant increase in plasma TAT levels. Importantly, thrombin inhibition with dabigatran significantly suppressed induction of each gene in HFD-fed mice (Fig. 5, A–D). Notably, dabigatran treatment did not universally reverse changes in hepatic lipogenic gene expression, because expression of several genes coordinating lipid metabolism, including fatty acid synthesis and fatty acid beta oxidation (e.g., Fasn, Acoxl, and Acaca) were not significantly impacted by dabigatran treatment (not shown).

Previous studies found that transgenic overexpression of Cyp7a1, the rate-limiting enzyme in hepatic bile acid metabolism, inhibited HFD-induced obesity coincident with a reduction in hepatic triglycerides and hepatic cholesterol (Li et al., 2010). In agreement, we identified that pharmacologic thrombin inhibition increased the expression of genes related to bile acid synthesis (Fig. 5, E–H). This included induction of the “classic pathway” genes with the rate-limiting enzyme Cyp7a1 (Fig. 5E) and increase in Cyp8b1 (Fig. 5F) expression, as well as induction of the “alternative pathway” genes, including Cyp7b1 and Cyp39a1 (Fig. 5, G and H). In contrast, expression of these genes was suppressed in mice fed a HFD at 3 months relative to CD-fed animals (Fig. 5, E–H). This finding suggests one important mechanism whereby thrombin inhibition with dabigatran protects against diet-induced obesity and NAFLD by increasing bile acid synthesis.

Discussion

Previous studies have linked a procoagulant state to metabolic syndrome and NAFLD development in patients (Ay et al., 2010; Beijers et al., 2010; Fritsch et al., 2010; Targher et al., 2010b; Semeraro et al., 2012). Additionally, perturbations in the expression and activity of coagulation factors has been observed in mice fed a HFD (Kassel et al., 2011; Owens et al., 2012). Here, we defined the temporal relationship between the onset of coagulation cascade activation and HFD-induced progression of fatty liver. Mild changes in serum lipids and hepatic steatosis were evident before activation of the coagulation cascade in mice fed a HFD. Marked increases in hepatic inflammation and injury coincided with coagulation cascade activation, suggesting a temporal link between coagulation cascade activation and exacerbation of HFD-induced NAFLD in mice.

Our studies with the thrombin inhibitor dabigatran, demonstrate that coagulation system activity is not simply a biomarker of fatty liver disease severity, but also that thrombin-mediated proteolysis is mechanistically linked to the development and progression of NAFLD. This likely relates in part to the exciting observation that HFD-induced body weight gain is reduced in mice treated with dabigatran. The protection from HFD-induced body weight gain and hepatic steatosis mediated by dabigatran did not appear to be linked to altered glucose handling, at least given the statistical power of the current study. However, more definitive studies of putative effects of dabigatran on glucose metabolism are ongoing. Intriguingly, the reduction in liver weight in dabigatran-treated mice could not be fully explained by body weight changes, raising the possibility that thrombin-mediated induction of genes controlling de novo lipogenesis contributes directly to hepatic steatosis. Indeed, HFD-induced hepatic expression of several genes (e.g., Cidea, Pparg, Scd1, Cd36) is attenuated in PAR-1–null mice, which are protected from fatty liver, despite equivocal weight gain when fed a HFD (Kassel et al., 2011). Collectively, the results support the hypothesis that coagulation is central to the pathogenesis of HFD-induced NAFLD, driven through both local changes in gene expression and systemic effects including body weight gain.

Previous studies found that reducing hematopoietic cell tissue factor expression decreased thrombin generation, body weight gain, and NAFLD in HFD-fed Ldlr−/− mice (Kassel et al., 2011; Owens et al., 2012). Our results with prophylactic administration of a clinically used thrombin inhibitor are consistent with a reduction in tissue factor-driven thrombin generation conferring protection against HFD-induced obesity and NAFLD. Thus, it is surprising that short-term (4 weeks) thrombin inhibition with argatroban did not significantly impact body weight gain in obese hypercholesterolemic Ldlr−/− mice (Kassel et al., 2012). However, argatroban is a thrombin inhibitor of limited utility and potency due to poor pharmacokinetics (e.g., t1/2 of only 50 minutes) (Kondo et al., 2001), a requirement of parenteral delivery and a complex drug formulation. It is conceivable that in the previous study the degree or duration of thrombin inhibition with argatroban was insufficient to impact body weight gain, despite the observed reduction in hepatic inflammation and correction of serum cholesterol in argatroban-treated mice (Kassel et al., 2012). Nevertheless, the current and previous studies highlight the critical role of thrombin signaling in the regulation of HFD-induced fatty liver and associated pathologies.

Inflammatory cells and the mediators they produce have emerged as critical regulators of obesity and metabolic disease (Hotamisligil et al., 1993; De Taeye et al., 2007; Huang et al., 2010; Stienstra et al., 2010; Henao-Mejia et al., 2012). Obesity is associated with a pathologic state, termed metabolic inflammation, in which control of energy usage and hepatic and adipose metabolite fate are directed by accumulated immune cells and mediators they produce (e.g., cytokines) (Hotamisligil, 2006; Gregor and Hotamisligil, 2011; Cao, 2014). In liver, accumulation and activation of inflammatory cells, particularly activation of Kupffer cells and recruitment of monocytes/macrophages, has been shown to promote hepatic lipid dysmetabolism and tissue injury in NAFLD/NASH (Baffy, 2009; Huang et al., 2010;
Thus, one possible mechanism whereby thrombin activity could promote fatty liver disease and obesity is by exacerbating inflammatory cell activity and associated metabolic inflammation. Indeed, coagulation and inflammatory pathways are deeply integrated, and thrombin is well appreciated to increase macrophage proinflammatory function (Esmon, 2003, 2004; Opal, 2003). Thrombin signaling through PAR-1 expressed by hepatic nonparenchymal cells, such as macrophages and stellate cells, has the potential to trigger additional inflammatory cell recruitment and activation via upregulation of cytokine expression (Gaca et al., 2002; Copple et al., 2003; Fiorucci et al., 2004; Chen and Dorling, 2009). We found that thrombin inhibition with dabigatran markedly reduced hepatic inflammation, alongside decreased mRNA expression of macrophage chemotactic factors (e.g., MCP-1/CCL2) and macrophage-derived proinflammatory cytokines (e.g., TNF). PAR-1−/− mice were also protected from HFD-induced NAFLD and displayed a similar dramatic reduction in hepatic TNF.
and MCP-1/CCL2 expression and macrophage accumulation (Luyendyk et al., 2010). Collectively, these results suggest that potentiation of hepatic inflammation by thrombin–PAR-1 signaling is one mechanism whereby thrombin activity alters hepatic lipid metabolism and contributes to NAFLD/NASH pathology.

In addition to thrombin signaling, fibrin deposition is associated with obesity and multiple components of the metabolic syndrome (Rugman et al., 1994; Nienaber et al., 2008). Notably, a key functional link to metabolic disease sequelae is the concept that fibrin can exert powerful pro-inflammatory activities. Specifically, cross-linked hepatic
Fibrin deposition may direct macrophage localization and activation through direct engagement of the αMβ2 integrin expressed by macrophages in the liver (Ugarova et al., 1998). Thrombin activation of factor XIII and thrombin-activatable fibrinolysis inhibitor can support inflammation via stabilization of fibrin (Myles et al., 2003). Moreover, outside of its traditional targets in hemostasis, thrombin increases the proinflammatory potential of osteopontin, a molecule linked to obesity and NAFLD (Sahai et al., 2004). Taken together, these studies suggest multiple pathways whereby thrombin targets drive inflammation that promotes the pathogenesis of NAFLD.

The finding that dabigatran treatment markedly reduces hepatic inflammation sets the stage for additional mechanistic studies determining the contribution of multiple thrombin targets to NAFLD pathology. Specifically, our proof-of-principle findings with dabigatran reinforce the possibility that pharmacologic inhibition of thrombin or factor Xa, or their targets, may form a novel treatment strategy for obesity and NAFLD/NASH. Clinical investigation of this connection is highly feasible, insofar as patients with NAFLD/NASH and/or obesity often have a concurrent indication for anticoagulation, such as nonvalvular A-Fib. For example, existing clinical evidence suggests that compared with warfarin, patients taking dabigatran

**Fig. 5.** Thrombin inhibition with dabigatran etexilate (DE) alters genes associated with hepatic metabolism and bile acid synthesis in C57BL/6J mice fed a HFD. Gene expression analysis by quantitative real-time polymerase chain reaction was performed on the HFD time course and thrombin inhibition study hepatic samples and identified changes in Scd1 (A), Cd36 (B), Cidea (C), and Pparg (D) levels. Moreover, HFD suppressed bile acid metabolism genes, including Cyp7a1 (E), Cyp8b1 (F), Cyp7b1 (G), and Cyp39a1 (H). Data are expressed as mean ± S.E.M., n = 4 or 7–9 mice per group. Time course: fold change is expressed relative to CD fold change at 1 month. *P < 0.05 compared with time-matched CD measured by one-way analysis of variance followed by Student-Newman-Keuls post hoc test. Thrombin inhibition study: fold change in HFD + DE is expressed relative to the HFD fold change at 3 months, which is set to 1. *P < 0.05 compared with HFD measured by Student’s t-test.
have significantly lower plasma apolipoprotein B levels (Joseph et al., 2013). These findings demonstrate yet another metabolic benefit of direct thrombin inhibition in managing stroke and potentially decreasing atherosclerotic complications in patients with A-Fib.

Thrombin inhibition led to induction of bile acid synthesis genes that catalyze steps in the classic and alternative pathways of bile acid synthesis from cholesterol. Others have shown that bile acid activation of the nuclear receptor farnesoid X receptor inhibits progression of NAFLD/NASH, and activation of the G protein–coupled receptor TGR5 promotes thermogenesis in brown adipose (Kong et al., 2009; Zhang et al., 2009; Pols et al., 2011). In agreement with these functions of bile acids, transgenic mice over-expressing hepatic Cyp7a1 are resistant to HFD-induced obesity (Li et al., 2010). Thus, the protective effects of dabigatran on body weight gain could relate to enhanced expression of bile acid synthesizing enzymes. Indeed, PAR-1 deficiency was not associated with upregulation of Cyp7a1 (not shown), providing potential explanation as to why deficiency in this thrombin target failed to impact body weight gain (Kassel et al., 2011). Moreover, induction of bile acid synthesis could be the mechanism by which thrombin inhibition reduces serum cholesterol. We demonstrated herein, and previously in hypercholesterolemic mice (Kassel et al., 2012), that inhibition of thrombin reduces serum cholesterol levels. Additional studies are required to both fully characterize the impact of thrombin activity on the size and composition of the bile acid pool and evaluate whether the thrombin contribution to bile acid dynamics is direct or secondary to other thrombin-mediated changes in the microenvironment (e.g., enhancing local hepatic inflammation).

The coagulation system has emerged as an important contributor to multiple facets of metabolic disease (Ay et al., 2010; Beijers et al., 2010; Fritsch et al., 2010; Targher et al., 2010b; Semeraro et al., 2012), with mouse models suggesting a connection to HFD-induced obesity, insulin resistance, and fatty liver disease (Kassel et al., 2011; Antoniak et al., 2013). Identifying the mechanisms whereby thrombin and its targets promote NAFLD/NASH could identify novel strategies to inhibit this disease process without affecting hemostasis. Our data also suggest another potential therapeutic benefit of direct thrombin inhibitors in patients with an existing indication, such as A-Fib. Not surprisingly, NAFLD/NASH is tightly linked to obesity (Fabbriini et al., 2010), the latter being an important risk factor for A-Fib (Badheka et al., 2010). Indeed, elevated liver enzymes (e.g., ALT) were recently found to be associated with the risk of A-Fib in the Framingham Heart Study (Targher et al., 2013). Viewed in the context of our findings, the use of direct thrombin inhibitors to reduce stroke in obese patients with A-Fib may also correct NAFLD/NASH and potentially have beneficial effects on obesity. Furthermore, a reduction in liver enzymes (and potentially hepatic inflammation) could in turn reduce the risk of A-Fib. However, the comparative benefit of anticoagulant therapy with either novel oral direct inhibitors of thrombin or factor Xa or with conventional agents (e.g., warfarin, heparin) in models of obesity or obese patients has not yet been evaluated. Additional mechanistic studies elucidating the effects of direct thrombin inhibitors on limiting obesity and associated sequelae including fatty liver, hepatic inflammation, and injury in mouse models have the potential to guide future clinical investigation related to anticoagulant indications and drug selection.

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Author Contributions

Participated in research design: Kopec, Joshi, Kassel, Sullivan, Luyendyk.

Conducted experiments: Kopec, Joshi, Towery, Kassel, Sullivan, Luyendyk.

Performed data analysis: Kopec, Joshi, Towery, Kassel, Sullivan, Luyendyk.

Wrote or contributed to the writing of the manuscript: Kopec, Joshi, Luyendyk.

References


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Thrombin inhibition with dabigatran protects against high fat diet-induced fatty liver disease in mice

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**Supplementary Table 1.** Gene names and primer sequences (5'-> 3') for transcripts verified by qPCR.

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<th>Gene Name</th>
<th>Gene Symbol</th>
<th>Gene ID</th>
<th>Forward primer 1</th>
<th>Forward primer 2</th>
<th>Reverse primer 1</th>
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