Hepatic Injury in Nonalcoholic Steatohepatitis Contributes to Altered Intestinal Permeability

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Hepatic Injury in Nonalcoholic Steatohepatitis Contributes to Altered Intestinal Permeability

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Abstract

BACKGROUND & AIMS—Emerging data suggest that changes in intestinal permeability and increased gut microbial translocation contribute to the inflammatory pathway involved in nonalcoholic steatohepatitis (NASH) development. Numerous studies have investigated the association between increased intestinal permeability and NASH. Our meta-analysis of this association investigates the underlying mechanism.

METHODS—A meta-analysis was performed to compare the rates of increased intestinal permeability in patients with NASH and healthy controls. To further address the underlying mechanism of action, we studied changes in intestinal permeability in a diet-induced (methionine- and choline-deficient; MCD) murine model of NASH. In vitro studies were also performed to investigate the effect of MCD culture medium at the cellular level on hepatocytes, Kupffer cells, and intestinal epithelial cells.

RESULTS—Nonalcoholic fatty liver disease (NAFLD) patients, and in particular those with NASH, are more likely to have increased intestinal permeability compared with healthy controls. We correlate this clinical observation with in vivo data showing mice fed an MCD diet develop intestinal permeability changes after an initial phase of liver injury and tumor necrosis factor-α (TNFα) induction. In vitro studies reveal that MCD medium induces hepatic injury and TNFα production yet has no direct effect on intestinal epithelial cells. Although these data suggest a role...
for hepatic TNFα in altering intestinal permeability, we found that mice genetically resistant to TNFα-myosin light chain kinase (MLCK)–induced intestinal permeability changes fed an MCD diet still develop increased permeability and liver injury.

**CONCLUSIONS**—Our clinical and experimental results strengthen the association between intestinal permeability increases and NASH and also suggest that an early phase of hepatic injury and inflammation contributes to altered intestinal permeability in a fashion independent of TNFα and MLCK.

**Keywords**

Meta-Analysis; Myosin Light Chain Kinase; Steatosis; Tight Junctions

Nonalcoholic fatty liver disease (NAFLD) has become an increasingly common clinical condition, with an estimated prevalence of 30% in the U.S. population. Despite this high prevalence, only a minority of NAFLD patients develop nonalcoholic steatohepatitis (NASH) and fibrosis, which account for significant morbidity and mortality. The clinical challenge remains identifying the patients who are more likely to develop NASH, as these patients are at greater risk for liver-related adverse events. Unfortunately, to date there is no reliable predictor of progression to NASH, nor are therapies approved by the U.S. Food and Drug Administration for this condition. Accordingly, a more fundamental understanding of the pathophysiology of NASH is critical to help identify high-risk NAFLD patients and therapeutic targets.

There are accumulating data that suggest a role for alterations in intestinal permeability in the pathogenesis of NASH. Specifically, it is hypothesized that an increase in intestinal epithelial cell permeability allows for translocation of microbial products into the portal vein, which propagate inflammation in a susceptible liver primed for injury. These data suggest that communication between the intestine and liver, the so-called gut-liver axis, plays a role in NASH development. Accordingly, there is building momentum to study the contribution of changes in intestinal homeostasis to liver injury and inflammation.

Although our understanding of the gut-liver axis is rapidly evolving, there remain many unanswered questions. Multiple studies have examined have examined the incidence of intestinal permeability changes in NASH patients; however, a comprehensive and systemic assessment of this relationship has yet to be performed. Furthermore, the inciting event responsible for intestinal permeability changes in patients with primary liver disease has not been identified. Obesity is common in patients with NASH and has been associated with intestinal inflammation and up-regulation of tumor necrosis factor-α (TNFα), both of which may contribute to intestinal leakage of microbial products. Additionally, differences in the gut microbiome in patients with NASH may alter intestinal permeability through inflammation-based and bacterial metabolite-driven pathways. The potential contribution of liver pathology to intestinal permeability in NASH, however, has yet to be investigated.

In an effort to comprehensively assess the rates of increased intestinal permeability in patients with and without NASH, we performed a meta-analysis of literature examining this question. We then sought to mechanistically explain the association between intestinal
permeability and NASH using an animal model for NASH as a means to eliminate possible confounders of the clinical data, such as antibiotic exposure and medical comorbidities. We focused our attention on the potential contribution of liver injury to intestinal permeability, as we hypothesized that an initial liver injury may lead to systemic disturbances, including an increase in intestinal permeability, which further propagates liver inflammation.

Materials and Methods

Meta-analysis

We performed a meta-analysis to compare the rates of increased intestinal permeability in patients with and without NAFLD. For inclusion in the meta-analysis, a study had to meet the following criteria: 1) measurement in vivo of intestinal permeability with a validated test substance (monosaccharide, nonhydrolyzed or hydrolyzed disaccharide, $^{51}$Cr-EDTA, or $^{99m}$Tc-DTPA); 2) documentation of NAFLD with imaging or histology; 3) documentation of minimal to no alcohol use in both groups; and 4) lack of coexisting liver disease in both groups.

To find relevant articles, a systematic review of English and non-English articles was performed using PubMed (1946 to July 2014) and EMBASE (1988 to 2014 week 15). The PubMed search was performed by the authors (J.L., M.D., and S.J.P.), and an information library specialist at the Mayo clinic library performed the EMBASE search. To reduce reporting bias and error in data collection, two independent reviewers (J.L. and S.J.P.) extracted data from selected studies using standardized data extraction forms. We contacted the primary investigators of articles with questions that arose during data extraction. We identified additional studies by searching bibliographies of all the reviewed articles and abstracts presented at the Digestive Disease Week and the Liver Meeting from 2007 to 2014. We used crude odds ratio (OR) for increased intestinal permeability, comparing NAFLD patients to controls as our parameter of interest. We pooled estimates by random-effects meta-analysis according to the method of DerSimonian and Laird and fixed-effect meta-analysis by calculating the weighted average of study estimates with the inverse of estimates variance used as the study weight. Q statistic and $I^2$ values were estimated to evaluate the heterogeneity among the studies.

We used Comprehensive Meta-Analysis software for these analyses. The methodological quality of the studies was assessed by two investigators (J.L. and S.J.P.) independently using the Newcastle-Ottawa scale. For meta-analyses of a small number of studies (generally less than 10), the power to detect publication bias is poor and is not recommended.

In Vivo Induction of Nonalcoholic Steatohepatitis

Male C57/BL6 mice aged 8 to 10 weeks were obtained from Jackson Laboratory (Bar Harbor, ME). To study the role of TNF-α in NASH pathogenesis, male C57/BL6 mice genetically deficient in the long isoform of myosin light chain kinase (MLCK) were used. All animal experiments were reviewed and approved by the Massachusetts General Hospital and the University of Chicago subcommittees on research animal care. To induce NASH,
mice were fed a methionine-and-choline-deficient (MCD) diet (Research Diets, New Brunswick, NJ). Mice were euthanized at multiple time points up to 21 days.

**Biochemical Analysis of Liver Injury**

For the animal experiments, immediately after euthanasia, we collected systemic blood from the inferior vena cava. Serum was obtained by centrifugation of whole blood at 10,000 rpm for 10 minutes. For the in vitro experiments, cell-free culture supernatant was concentrated to 500 μL using Microcon centrifugal filter devices (Millipore, Billerica, MA). To determine the extent of hepatocyte injury, quantification of alanine aminotransferase (ALT) was performed from serum and concentrated culture supernatant using the Infinity ALT Liquid Stable Reagent (Thermo Scientific, Middletown, VA).

**Histologic Analysis of Liver Injury**

The intact liver was excised immediately after mouse sacrifice, fixed in formalin for 24 hours, and then embedded in paraffin. Histologic analysis was performed on liver sections stained with H&E.

The severity of liver injury was determined using the NAFLD activity score as previously described elsewhere with slight modifications. Briefly, using the most affected area of tissue, a steatosis score was determined by the extent of steatosis: 0 (<5%), 1 (5%–33%), 2 (34%–66%), and 3 (>66%). An inflammation score was determined based upon the number of inflammatory foci in a 200× field: 0 (0 foci), 1 (<2 foci), 2 (2–4 foci), and 3 (>4 foci). The degree of hepatocyte ballooning was not scored as we have not found this histologic feature to be a characteristic finding of the MCD model. The modified NAFLD activity score was calculated by summation of the steatosis and inflammation scores. The examinations were performed in a blinded fashion by two independent investigators (J.L. and V.D.).

**In Vivo Intestinal Permeability Assay**

To measure intestinal paracellular permeability, the serum level of fluorescein isothiocyanate-4 kD dextran was measured as previously described elsewhere. Briefly, mice were denied access to food and water for 8 hours before sacrifice. At 4 hours before sacrifice, fluorescein isothiocyanate-4 kD dextran (Sigma-Aldrich, St. Louis, MO) at a dose of 60 mg/100 g body weight was orally gavaged. Immediately after the mice were sacrificed, we collected serum and measured the fluorescence intensity in serum (excitation, 490 nm; emission, 525 nm) using the Synergy 2 plate reader (BioTek, Winooski, VT).

**Immunofluorescence Microscopy**

The small intestine was excised and immediately placed in optimal cutting temperature compound (Tissue Tek; Sakura, Torrance, CA) and frozen in liquid nitrogen. We collected 5-μm frozen sections on coated slides and fixed them with 4% paraformaldehyde. Immunostaining was performed as previously described elsewhere. Briefly, the fixed sections were permeabilized with 0.1% Triton X-100 in phosphate-buffered saline (PBS) for 20 minutes and washed five times with 1% bovine serum albumin in PBS. Zona occludens-1 (ZO-1) was localized by incubating with rabbit anti-ZO-1 rabbit polyclonal antisera (Upstate...
Biotechnology, Waltham, MA) at a 1:200 dilution for 1 hour at room temperature. Subsequently, tissue sections were incubated with fluorescein isothiocyanate (FITC)-conjugated anti-rabbit IgG antibody (Sigma-Aldrich) at a 1:200 dilution for 1 hour. Standard epifluorescence microscopy was performed using a fluorescence microscope (EVOS FL; Life Technologies, Grand Island, NY).

**Quantitative Real-Time Polymerase Chain Reaction**

Mouse liver tissues were mechanically homogenized using the PowerGen 125 Homogenizer (Fisher Scientific, Fair Lawn, NJ). Total RNA was extracted using the RNeasy Mini Kit (Qiagen, Valencia, CA). Total RNA (500 ng) was converted into cDNA using the RT² First Strand Kit (SA Biosciences, Valencia, California). Quantitative real-time polymerase chain reaction (qRT-PCR) was performed using the RT² Master Mix Kit (SA Biosciences, Valencia, CA) and the iQ 5 system (Bio-Rad Laboratories, Hercules, CA). Quantitative RT-PCR was performed for mRNA expression of β-actin and TNF-α, interleukin-1β (IL1β), and IL6 using primers designed by SA Biosciences (Qiagen). Expression of β-actin was used to standardize the samples, and the results are expressed as a ratio relative to control.

**Detection of Cytokines in the Systemic Circulation**

The levels of multiple cytokines (TNF-α, IL1β, and IL6) was assessed from serum using magnetic bead-based multiplex assays (Millipore) coupled with the Luminex-200 System Analyzer (Luminex, Austin, TX), as recommended by the manufacturer’s overnight protocol. The mean fluorescence intensity was expressed as the ratio relative to the average mean fluorescence intensity value obtained at day 0.

**Cell Lines and Primary Cell Isolation**

We maintained the H35 hepatocyte-derived cells as previously described elsewhere.14 The Caco2 cells were maintained in Dulbecco’s modified Eagle’s medium supplemented with 10% fetal bovine serum and antibiotics (100 U/mL penicillin G and 100 U/mL streptomycin) (Life Technologies, Grand Island, NY).

Rat Kupffer cells were separated from the non-parenchymal cell (NPC) fraction using centrifugal elutriation, as previously described elsewhere.15 Briefly, the NPC fraction was centrifuged at 300g for 15 minutes. The supernatant was discarded, and the pellet was resuspended in ice-cold PBS and passed through a cell strainer (40-μm pore) to remove any large debris. The elutriator was sterilized by circulating water (45 mL/min, 0g, 5 minutes), 6% H₂O₂ (10 mL/min, 50g, 5 minutes), 15 mg/mL catalase solution (10 mL/min, 0g, 5 minutes), sterile water (10 mL/min, 40g, 5 minutes), and Hank’s balanced salt solution (10 mL/min, 40g, 5 minutes). The elutriator was then ramped to 600g, and a flow of 10 mL/min was maintained. We introduced 10 mL of the NPC fraction into the elutriator and washed for 10 minutes to remove any cell debris while we maintained the rotor at 600g. Kupffer cells were eluted at 45 mL/min, 600g, and 100 mL of the cell suspension was collected. The cells were pelleted at 500g for 7 minutes (no brake) and used as an enriched fraction in experiments.
In Vitro Induction of Steatohepatitis

Media identical to standard Dulbecco’s modified Eagle’s medium culture yet deficient in L-methionine and choline chloride (Life Technologies) (MCD medium) was used to induce steatohepatitis in vitro as previously described elsewhere. Briefly, H35 cells were cocultured with rat primary Kupffer cells at a 2:1 ratio in 12-well plates in standard medium for 24 hours, after which the standard medium was replaced by MCD medium for 24 hours. Cell-free culture supernatant was harvested for the ALT measurement. Additionally, the protein level of TNF-α in the culture supernatant was measured using a TNF-α Platinum ELISA kit (eBioscience, San Diego, CA).

In Vitro Intestinal Permeability Assay

To assess the effect of MCD medium on intestinal permeability in vitro, Caco2 cells were added to the apical side of collagen-coated Transwells (Corning, Lowell, MA). Caco2 cells were grown for 21 days to allow for monolayer and tight junction formation. We then added the MCD medium to the culture system. After 24 hours, the intestinal permeability was assessed by measuring the transepithelial electrical resistance (TEER) using the Millicell-ERS electrical resistance system (Millipore). The resistance obtained from each experimental well was subtracted from a blank value obtained by inserting the electrodes in a transwell harboring a cell-free medium. This value was multiplied by the area of the membrane to obtain TEER (Ω × cm²).

Statistical Analyses for In Vivo and In Vitro Experiments

Data are expressed as mean ± standard error and analyzed by unpaired Student t tests. Two-tailed P values were calculated, and P < .05 was considered statistically significant.

Results

Nonalcoholic Steatohepatitis Patients Are More Likely to Have Increased Intestinal Permeability Compared With Healthy Controls

The initial search strategy for the meta-analysis yielded 288 potential articles for inclusion (Supplementary Table 1). Although there were multiple reasons for exclusion, the most common included studies involved animals and studies that did not directly measure intestinal permeability but rather used a surrogate measure such as serum lipopolysaccharide. After analysis of the selected articles, nine were reviewed in further detail. Subsequently, five studies (N = 128 patients) met our inclusion criteria (Supplementary Figure 1). The characteristics of the included studies are summarized in Table 1. Normal values for tests measuring intestinal permeability were determined by the included study or from previously defined values.

We found that 39.1% of NAFLD patients in our analysis had evidence for increased intestinal permeability, compared to 6.8% of healthy controls. The OR of NAFLD patients having increased intestinal permeability compared with controls was 5.08 (95% confidence interval [CI], 1.98–13.05) (Figure 1A). There was minimal heterogeneity in the included studies (I² = 9.8%, Q statistic 4.43, P = .35), which was further strengthened by an exclusion sensitivity analysis (Figure 1B). The assessment of study quality is outlined in Table 2.
We next performed a subgroup analysis comparing the frequency of increased intestinal permeability in patients with NASH with that of the healthy controls. Four of the five studies (n = 83 patients) allowed for this analysis. We found that patients with NASH are also more likely to have increased intestinal permeability compared to healthy controls (OR 7.21; 95% CI, 2.35–22.13). The incidence of permeability changes in this subgroup (49.2%) was higher as compared to NAFLD patients as a whole, which includes patients with bland steatosis (Figure 2A). Although no statistically significant heterogeneity was detected (P = .09), there was moderate heterogeneity (I² = 54.5%) across the studies. Interestingly, exclusion sensitivity analysis revealed the study by Wigg et al17 markedly reduced the observed OR (OR when excluded: 34.3; 95% CI, 28.3–161.65) (Figure 2B). Overall, these data show that NAFLD patients, and in particular those with NASH, are more likely to exhibit increased intestinal permeability compared with healthy controls.

A Methionine-Choline-Deficient Diet Induces Intestinal Permeability Changes After Liver Injury

To further investigate the clinical association between intestinal permeability and NASH, we examined this relationship in a dietary animal model (the MCD model) for NASH. The MCD model was chosen for its well-established ability to induce steatosis and inflammation and do so reproducibly in a relatively short period of time. However, it must be noted that the MCD model does not induce the weight gain, lipid abnormalities, or insulin resistance that are common in NASH patients.

We found that mice fed an MCD diet developed liver injury rapidly, as early as 6 days into the diet, at which time serum levels of ALT were elevated and histologic evidence for hepatic steatosis and inflammation was also observed (Figure 3A–C). This early phase of liver injury and inflammation was followed by a secondary phase of injury, with peak ALT and histologically based modified NAFLD activity scores occurring at day 21. In parallel, we found that both hepatic and systemic levels of TNFα rose at an early time point with a subsequent secondary rise at day 21 (Figure 3D and E). Interestingly, we were unable to detect a difference in hepatic or systemic levels of IL6 or IL1β over the 21-day course, suggesting that these cytokines, which have been implicated in NASH pathogenesis, may contribute to a later phase of injury and inflammation.23,24

We next examined the temporal pattern of intestinal permeability changes during the course of the MCD diet. Although permeability steadily increased as the experiment progressed, a significant increase in intestinal permeability, as measured by both the serum signal intensity of FITC-dextran, was not observed until day 10 (Figure 3F), well after the initial phase of liver injury, inflammation, and TNFα production. Furthermore, we first noted evidence for disruption of ZO-1 localization in the small intestine at day 10, suggesting injury to the epithelial tight junction complex. These findings are consistent with the clinical association between increased intestinal permeability and NASH, but suggest that increased intestinal permeability is an effect rather than cause of NASH. One potential explanation could be that an initial phase of hepatic injury and inflammation releases intermediates, such as cytokines, that increase intestinal permeability, and that the latter further contribute to the progression of liver injury.
Methionine and Choline Deficient Culture Medium Directly Induces Hepatocyte Injury and Inflammation but Does Not Affect Intestinal Epithelial Paracellular Permeability

To address our in vivo observation of early liver injury and TNF$\alpha$ production subsequently influencing intestinal permeability, we studied the effect of MCD medium on hepatocytes, Kupffer cells, and intestinal epithelial cells in vitro. We cocultured rat hepatocytes (H35 cells) and primary rat Kupffer cells together in an attempt to more appropriately model the physiologic inflammatory dynamic within the liver. We found that exposing hepatocytes and Kupffer cells to MCD medium induced significant hepatocyte injury and TNF$\alpha$ production as compared with cells grown in standard medium (Figure 4A and B). These data reinforced our in vivo observation that MCD directly induces liver injury and inflammation.

In contrast, we did not observe any deleterious effect of exposing intestinal epithelial cells to MCD medium. Specifically, intestinal epithelial cells exposed to the MCD medium showed no reduction in TEER, suggesting that the MCD medium does not directly disrupt intestinal tight junction function. Taken together, these in vitro data show that MCD medium induces hepatocyte injury and TNF$\alpha$ production yet does not independently disrupt intestinal epithelial tight junction function, suggesting that our in vivo observation of increased intestinal permeability in the MCD model may rely on MCD-induced hepatic injury and TNF$\alpha$ production.

Mice Resistant to TNF$\alpha$-Induced Intestinal Permeability Changes Develop Significant Intestinal Permeability and Liver Injury When Fed a NASH-Inducing Diet

Our data thus far suggest that MCD-induced hepatic TNF$\alpha$ production may play a role in altering intestinal permeability and contribute to ongoing liver injury and inflammation. To test this hypothesis, we exposed mice deficient in the long isoform of myosin light chain kinase (MLCK) to the MCD diet. These mice are resistant to acute-TNF$\alpha$-induced intestinal permeability increases,$^{13,25}$ and also are protected from barrier loss in experimental inflammatory bowel disease.$^{26}$ We hypothesized that long MLCK-deficient mice would exhibit early hepatic injury induced by the MCD diet but be protected against intestinal permeability changes and ongoing hepatic injury and inflammation.

However, we found that MCD-fed MLCK-deficient mice developed equally significant liver injury compared with wild-type mice fed an MCD diet for 21 days (Figure 5A and B). Furthermore, long MLCK-deficient mice exhibited increased intestinal permeability compared with the MLCK-deficient mice fed a standard chow (Figure 5C). Taken together, these data strongly suggest that TNF$\alpha$ is not predominantly responsible for the observed changes in intestinal permeability seen during the MCD diet.

Discussion

In this study we demonstrate that NAFLD patients, and in particular patients with NASH, are more likely to exhibit increased intestinal permeability compared with healthy controls. We correlate this clinical observation with in vivo data showing that MCD-fed mice develop intestinal permeability changes after an initial phase of liver injury and TNF$\alpha$ induction.
In vitro studies reveal that MCD medium induces hepatic injury and TNFα production yet has no direct effect on intestinal epithelial cells. Although these data suggest a role for hepatic TNFα in altering intestinal permeability, we found that MCD-fed mice genetically resistant to TNFα-induced intestinal permeability changes were not protected against liver injury. Taken together, our clinical and experimental results strengthen the association between intestinal permeability changes and NASH as well as suggest that an early phase of hepatic injury and inflammation contributes to altered intestinal permeability in a TNFα- and MLCK-independent fashion.

Although the pathogenesis of NASH is likely multifactorial, there are emerging data that suggest alterations in intestinal permeability can contribute to liver injury. The emergence of the "gut-liver axis" implicates changes in intestinal physiology in the pathogenesis of liver injury. For example, Gabele et al demonstrated that induction of colitis in mice fed a high-fat diet caused more severe steatohepatitis as compared with mice fed a high-fat diet without colitis. They further showed that increased intestinal translocation of microbial products into the portal vein of colitis-affected mice occurred, which suggests that disruption of the intestinal barrier allowed proinflammatory substrates access to the liver. Similarly, Henao-Mejia et al associated changes in gut physiology and microbiota with elevated levels of Toll-like receptor agonists in the portal vein and more severe steatohepatitis in an MCD model.

In parallel with these animal data, clinical experience also supports the notion of a gut-liver axis. Pathological intestinal conditions with documented changes in intestinal permeability have been associated with NASH, including celiac disease and inflammatory bowel disease. Furthermore, probiotic treatment has been shown to improve liver aminotransferases in patients with NAFLD, suggesting a role for intestinal microbes in NASH pathogenesis. On the other hand, there is less evidence suggesting that end-stage liver disease or cirrhosis can affect intestinal physiology, and the mechanism for this is unclear.

Our results further strengthen the potential role of intestinal permeability in the pathogenesis of NASH, but uniquely suggest that early changes in liver physiology may also affect intestinal homeostasis. We show that liver injury is induced early in the course of the MCD diet, before any change in intestinal permeability, which suggests that the initial liver injury phase may be contributing to the observed intestinal permeability changes in a TNFα-independent manner.

The mechanism by which hepatic injury may affect intestinal permeability, however, remains elusive. We focused our study on TNFα as higher levels of this cytokine have been associated with NASH in clinical studies. Additionally, its role in modulation of intestinal permeability has been well-established through its activation of long MLCK and the subsequent effect on tight junction permeability. However, despite the initially encouraging data implicating hepatic TNFα, ultimately it does not appear to be responsible for altering intestinal permeability through an MLCK-dependent mechanism.
Alternatively, another liver-produced mediator may be responsible for the observed intestinal changes. Besides TNFα, several other cytokines have been implicated in altering intestinal permeability. We did not find any differences in hepatic and systemic levels of IL1β, which can alter intestinal permeability through an MLCK-dependent mechanism.\(^{38}\) Furthermore, there were no differences in IL6, which has been associated with liver-related mortality.\(^{39}\) and can alter intestinal permeability through claudin-2 up-regulation, although there is no expected increase in FITC-dextran permeability.\(^{40}\) Our findings should not underscore the potential importance of these cytokines to NASH pathogenesis, as elevated serum and hepatic levels of IL1β have been reported at different time points in the MCD model.\(^{41}\) Several other mediators, such as transforming growth factor β and plasminogen activator inhibitor-1, deserve study. Further investigation into identifying a novel hepatic-produced modulator of intestinal permeability is desirable.

Although the contribution of TNFα to NASH pathogenesis is debated,\(^{42}\) there are data highlighting its key role. Tomita et al\(^{43}\) showed mice deficient in both TNF receptors 1 and 2 developed attenuated liver steatosis and fibrosis when fed the MCD diet, implicating the TNFα/TNF-receptor-mediated pathway in NASH pathogenesis. Koca et al\(^{44}\) demonstrated that administration of an antibody to TNFα lessened the severity of steatohepatitis induced by the MCD diet. Thalidomide, an immunosuppressant with anti-TNFα properties, has also been shown to reduce the inflammatory profile induced in a murine NAFLD model.\(^{45}\) Furthermore, NAFLD patients receiving etanercept, an antibody to TNF, have significant reductions in the AST/ALT ratio and serum C-reactive protein levels.\(^{46}\)

Our data demonstrate that hepatic TNFα is not responsible for altering intestinal permeability through an MLCK-dependent mechanism. It is possible, however, that TNFα modulates intestinal permeability independent of MLCK and tight junctions by accelerating intestinal epithelial cell turnover.\(^{26}\) Additionally, it is also possible that TNFα alters permeability through activation of TNF-receptor 1, which has been recently been shown as important to the pathogenesis of alcoholic liver disease.\(^{47}\) Therefore, the potential role for TNFα in NASH pathogenesis and intestinal barrier loss should not be fully discounted.

Our study does have limitations. First, defining normal values for intestinal permeability is difficult and was not uniform throughout the clinical studies included in our meta-analysis. For example, the normal value chosen for 51Cr-EDTA testing in the study by Miele et al\(^{19}\) of 4.88%, which was based on the median value obtained from the NAFLD group in the study, is higher than normal values previously reported.\(^{48}\) As another example, the normal value for lactulose-rhamnose intestinal permeability testing of 0.18 used to analyze data from the study by Wigg et al\(^{17}\) has been shown to have sensitivity and specificity values of 78%.\(^{22}\) Along these lines, intestinal permeability is difficult to assess in vivo. Although multiple methodologies have been used to assess permeability, measurement of serum FITC dextran is commonly employed. These analyses, however, may not sensitively detect small changes in intestinal permeability to large probes, such as FITC-dextran, and are unable to measure paracellular permeability increases limited to small solutes such as ions. Last, although the MCD diet as a model for NASH is effective in recapitulating the histologic changes seen in NASH, it does not induce weight gain, lipid abnormalities, or insulin
resistance, which are common in NASH patients. Therefore, it is difficult to comment on the potential impact of these clinical variables on our results.

In summary, our clinical and experimental results strengthen the association between intestinal permeability increases and NASH and also suggest that an early phase of hepatic injury and inflammation contributes to altered intestinal permeability in a TNF-α- and MLCK-independent fashion. As evidence continues to mount supporting the notion of gut-liver cross-talk, further research is needed to decipher the mechanism by which this cross-talk occurs. It is likely that the findings of such research will have significant clinical implications for NASH patients.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Abbreviations used in this paper

ALT alanine aminotransferase
CI confidence interval
FITC fluorescein isothiocyanate
IL interleukin
MCD methionine and choline deficient
MLCK myosin light chain kinase
NAFLD nonalcoholic fatty liver disease
NASH nonalcoholic steatohepatitis
NPC nonparenchymal cells
OR odds ratio
PBS phosphate-buffered saline
qRT-PCR quantitative real-time polymerase chain reaction
TEER transepithelial electrical resistance
TNF-α tumor necrosis factor-α
ZO-1 zona occludens-1
References


SUMMARY

This study comprehensively defines the clinical association between intestinal permeability increases and nonalcoholic steatohepatitis. The results suggest that early-phase hepatic injury and inflammation contribute to altered intestinal permeability in a fashion independent of tumor necrosis factor-α and myosin light chain kinase.
Figure 1. Meta-analysis of increased intestinal permeability rates in nonalcoholic fatty liver disease (NAFLD) patients versus healthy controls
(A) Forest plot of increased intestinal permeability in patients with NAFLD as compared to healthy controls using a fixed-effects model. (B) Exclusion sensitivity plot of increased intestinal permeability in NAFLD patients versus healthy controls. CI, confidence interval; OR, odds ratio.
Figure 2. Meta-analysis of increased intestinal permeability rates in nonalcoholic steatohepatitis (NASH) patients versus healthy controls

(A) Forest plot of increased intestinal permeability in patients with NASH as compared with healthy controls using a fixed-effects model. 

(B) Exclusion sensitivity plot of increased intestinal permeability in NASH patients versus healthy controls. CI, confidence interval; OR, odds ratio.
Figure 3. Temporal characterization of liver injury and intestinal permeability changes in a murine dietary nonalcoholic steatohepatitis (NASH) model

C57BL/6 mice (N = 5 mice/group) were fed a diet deficient in methionine and choline (MCD) and were sacrificed at multiple time points up to 21 days. We found evidence for significant MCD-induced liver injury as early as day 6 based on both (A) serum alanine aminotransferase (ALT) and (B, C) H&E liver histology (original magnification: 20×) that progressed to a peak value at day 21. Specifically, liver histologic examination revealed a progressively increasing number of inflammatory foci and steatosis throughout the experiment. In parallel, hepatic mRNA expression and systemic levels of tumor necrosis

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factor-α (TNF-α) were elevated at an early phase of the diet (D, E). Temporal evaluation of intestinal permeability changes, based on (F) fluorescein isothiocyanate (FITC) dextran serum measurements, and tight junction architecture based on (G) immunofluorescence staining for zona-occludens-1 (ZO-1), revealed evidence for a significant increase in intestinal permeability and disruption of normal tight junction architecture (loss of chicken-wire appearance of ZO-1: arrows) at day 10. DAPI, 4′,6-diamidino-2-phenylindole.
Rat hepatocytes (H35) were grown in coculture with primary rat Kupffer cells (KC) at a ratio of 2:1. Cells were exposed to standard medium (SM) or methionine-and-choline-deficient (MCD) medium for 24 hours, after which the supernatant was harvested for further analysis. (A) Coculturing of hepatocytes and KCs in the presence of MCD caused the most significant elevation in hepatocyte injury, based on supernatant levels of alanine aminotransferase (ALT). (B) Further, MCD-exposed hepatocytes and KCs produced significantly more tumor necrosis factor-α (TNF-α) compared to cells grown in standard medium. (C) Intestinal epithelial cells (Caco2 cells) were grown to confluence and allowed to form strong tight junctions, after which they were exposed to either MCD medium or SM. We found no difference in tight junction function between cells grown in MCD medium versus SM, suggesting MCD medium is not directly toxic to these cells. TEER, transepithelial electrical resistance.
Figure 5. In vivo assessment of tumor necrosis factor-α (TNF-α) on intestinal permeability in nonalcoholic steatohepatitis (NASH) pathogenesis

TNF-α-induced increases in intestinal permeability are mediated through myosin light chain kinase (MLCK); therefore, genetic deletion of MLCK renders mice impervious to intestinal permeability changes caused by TNF-α. We tested whether mice deficient in the long isoform of MLCK would be protected against MCD-induced liver injury and intestinal permeability changes. MLCK-knockout (KO) and wild-type (WT) mice were fed the MCD diet for 24 days and then euthanized (N = 5 mice per group). There were no differences in (A) serum levels of ALT or (B) in H&E liver histology (original magnification: 20×) between MLCK-KO and WT mice. (C) Furthermore, we were unable to detect a difference in intestinal permeability changes between the MCD-fed MLCK-KO and WT mice.
Table 1
Characteristics of Included Nonalcoholic Fatty Liver Disease (NAFLD) Studies

<table>
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<tr>
<th>Author</th>
<th>Year</th>
<th>Location</th>
<th>NAFLD Diagnosis</th>
<th>Intestinal Permeability Assay</th>
<th>Single versus Multicenter</th>
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<th>Control (n)</th>
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<td>Wigg et al</td>
<td>2001</td>
<td>Australia</td>
<td>Histology</td>
<td>Urinary lactulose/rhamnose at 5 hours</td>
<td>Single</td>
<td>18</td>
<td>16</td>
</tr>
<tr>
<td>Farhadi et al</td>
<td>2008</td>
<td>United States</td>
<td>Histology</td>
<td>Urinary lactulose/mannitol at 5 hours</td>
<td>Single</td>
<td>16</td>
<td>12</td>
</tr>
<tr>
<td>Miele et al</td>
<td>2009</td>
<td>Italy</td>
<td>Histology</td>
<td>Urinary chromium-51 ethylene diamine excretion over 24 hours</td>
<td>Single</td>
<td>35</td>
<td>24</td>
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<tr>
<td>Volynets et al</td>
<td>2012</td>
<td>Germany</td>
<td>Ultrasound and serology</td>
<td>Urinary lactulose/mannitol at 6 hours</td>
<td>Single</td>
<td>20</td>
<td>10</td>
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<tr>
<td>Giorgio et al</td>
<td>2014</td>
<td>Italy</td>
<td>Histology</td>
<td>Urinary lactulose/mannitol at 6 hours</td>
<td>Single</td>
<td>39</td>
<td>21</td>
</tr>
</tbody>
</table>
Table 2

Newcastle-Ottawa Scale for Assessment of Quality of Included Studies

<table>
<thead>
<tr>
<th>Quality Assessment Criteria</th>
<th>Wigg et al 17</th>
<th>Farhadi et al 18</th>
<th>Miele et al 19</th>
<th>Volynets et al 20</th>
<th>Giorgio et al 21</th>
</tr>
</thead>
<tbody>
<tr>
<td>Is the case definition adequate? (Yes, with independent validation)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Representatives of cases? (Consecutive or obviously representative series of cases)</td>
<td>+</td>
<td>–</td>
<td>+</td>
<td>–</td>
<td>+</td>
</tr>
<tr>
<td>Selection of controls? (Community controls)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>–</td>
<td>+</td>
</tr>
<tr>
<td>Definition of controls? (No history of studied end point)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>–</td>
<td>+</td>
</tr>
<tr>
<td>Study controls for age/sex? (Matching or multivariable analysis)</td>
<td>+</td>
<td>–</td>
<td>+</td>
<td>–</td>
<td>+</td>
</tr>
<tr>
<td>Study controls for at least three additional factors?</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>Ascertainment of exposure? (Secure record, structured interview by health-care practitioner, blind to case/control status)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Same method of ascertainment of cases/controls? (Yes)</td>
<td>–</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Nonresponse rate? (Same for both groups)</td>
<td>–</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Total overall score (Maximum = 9)</td>
<td>6</td>
<td>6</td>
<td>8</td>
<td>5</td>
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</table>