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C/EBP homologous protein modulates liraglutide-mediated attenuation of non-alcoholic steatohepatitis

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Abstract

The CCAAT/enhancer-binding protein (C/EBP) homologous protein (CHOP), a major transcriptional regulator of endoplasmic reticulum (ER) stress-mediated apoptosis, is implicated in lipotoxicity-induced ER stress and hepatocyte apoptosis in non-alcoholic fatty liver disease (NAFLD). We have previously demonstrated that the glucagon like peptide 1 (GLP-1) agonist, liraglutide, protects steatotic hepatocytes from lipotoxicity-induced apoptosis by improved handling of free fatty acid (FFA)-induced ER stress. In the present study, we investigated whether CHOP is critical for GLP-1 mediated restoration of ER homeostasis and mitigation of hepatocyte apoptosis in a murine model of NASH (non-alcoholic steatohepatitis). Our data show that despite similar caloric intake, CHOP KO (CHOP-/-) mice fed a diet high in fat, fructose, and cholesterol (HFCD) for sixteen weeks developed more severe histological features of NASH compared with wild type (WT) controls. Severity of NASH in HFCD-fed CHOP-/- mice correlated with significant decrease in peroxisomal β -oxidation, and increased *de novo* lipogenesis and ER stressmediated hepatocyte apoptosis. Four weeks of liraglutide treatment markedly attenuated steatohepatitis in HFCD-fed WT mice by improving insulin sensitivity, and suppressing de novo lipogenesis and ER stress-mediated hepatocyte apoptosis. However, in the absence of CHOP, liraglutide did not improve insulin sensitivity, nor suppress peroxisomal β -oxidation or ER stressmediated hepatocyte apoptosis. Taken together, these data indicate that CHOP protects hepatocytes from HFCD-induced ER stress, and plays a significant role in the mechanism of liraglutidemediated protection against NASH pathogenesis.

Keywords

non-alcoholic fatty liver disease; non-alcoholic steatohepatitis; C/EBP homologous protein; apoptosis

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Non-alcoholic fatty liver disease (NAFLD) represents a broad spectrum of disorders including benign non-alcoholic fatty liver-or bland steatosis (NAFL), to the more severe non-alcoholic steatohepatitis (NASH) and steatofibrosis^{1, 2}. According to recent reports, the prevalence of NAFL ranges from 20-30% in the general population and as high as 75-100% in obese individuals^{3, 4}. Although most NAFL patients remain asymptomatic, one fifth of NAFL patients progress to chronic hepatitis (NASH), which can further progress to cirrhosis, portal hypertension, and promote development of hepatocellular carcinoma (HCC)^{4, 5}. Despite its high prevalence, and established morbidity and mortality associated with NAFLD, mechanisms leading to NAFLD progression remain poorly understood.

Systemic insulin resistance, lipotoxicity, and oxidative stress are central players in the pathogenesis of NASH⁶⁻⁸. Recent studies indicate that the unfolded protein response (UPR), an adaptive response that regulates ER function during ER stress through transcriptional and translational modulation of factors involved in ER homeostasis, also plays a critical role in free fatty acid (FFA) and free cholesterol mediated lipotoxicity⁸⁻¹⁴. Under conditions of nutrient overload, excess FFA and cholesterol induce activation of the UPR^{9, 14}. Failure of the UPR to maintain ER homeostasis in the setting of metabolic stress, resulting from nutrient overload, leads to ER stress-mediated hepatocyte apoptosis^{1, 8, 10, 11, 15-18}. Therefore, strategies to prevent ER stress-related hepatocyte death are highly desirable and would likely halt progression of NAFL to NASH.

Accumulating evidence suggests that C/EBP homologous protein (CHOP) is a major transcriptional regulator of ER stress-mediated apoptosis^{19, 20}. Studies implicate CHOP in a wide-array of common diseases including neurodegenerative, cardiovascular, and metabolic disorders, including NAFLD²¹⁻²⁷. Apart from modulating ER stress-mediated apoptosis, CHOP also plays a role in transcriptional regulation of cellular lipid metabolism^{20, 28} suggesting that increased hepatic expression of CHOP in NAFLD patients may protect hepatocytes from FFA and cholesterol induced lipotoxicity. However, a mechanism whereby CHOP mediates protection against FFA-induced ER stress is not entirely clear.

We and others have demonstrated a beneficial role for glucagon-like peptide-1 (GLP-1) in *in vivo* and *in vitro* for NAFLD treatment²⁹⁻³⁴. GLP-1 is an incretin hormone secreted by the L-cells of the distal small intestine and proximal colon³⁵. Prior data indicate that GLP-1 analogues can suppress FAA-induced steatosis in isolated hepatocytes, and in animal models of NASH fed a diet high in carbohydrate, saturated fat, and fructose. A recent small-scale double-blind placebo-controlled clinical trial in which the long-acting GLP-1 analogue, liraglutide was tested demonstrated liraglutide reduced metabolic dysfunction, increased hepatic insulin sensitivity, and reduced hepatic de novo lipogenesis (DNL) *in vivo*³⁶. Studies in mice and cell culture models of steatosis demonstrate that GLP-1 agonists, liraglutide or exendin-4, protect steatotic hepatocytes from apoptosis by improved handling of FFA-induced ER stress^{31, 34}. Here, by using global knock out mice for the critical ER stress protein, CHOP, we provide *in vivo* evidence that liraglutide mediates its protective effects through CHOP to ultimately prevent hepatocyte apoptosis and NAFLD progression. Our results demonstrate that in the absence of CHOP, liraglutide administration fails to

ameliorate steatohepatitis in mice fed a HFCD rendering hepatocytes far more susceptible to death.

MATERIALS AND METHODS

Mice

CHOP^{-/-} mice on a C57BL/6 background were purchased from Jackson Laboratories (Bar Harbor, ME). Animals were housed in standard micro-isolator cages and maintained on a 12 h: 12 h light/dark cycle. All animals received humane care and all procedures were approved by the Institutional Animal Care and Use Committee of the Veterans' Administration Hospital in Decatur, GA.

Diet

Five week old age and weight matched adult male mice were fed a high fat, high cholesterol diet and high fructose diet (HFCD) containing 0.2% cholesterol, 20% protein, 43% CHO, 23% fat (6.6% trans-fat) and 2.31% fructose (TD.130885; Harlan Laboratories)^{37, 38}; or the standard diet (ND) containing 16% protein, 61% carbohydrate and 7.2% fat. All cohorts were fed *ad libitum* for 16 weeks. After 12 weeks of feeding, mice received daily intraperitoneal injections of saline or liraglutide (200 μ g/Kg body weight) for 4 weeks. Mice were weighed weekly and food intake per cage was also measured weekly. At the end of the 16 weeks, mice were fasted for 8 hours prior to euthanasia.

Histopathology and immunohistochemistry

Formalin-fixed liver tissues were paraffin-embedded, sectioned, and stained with H&E, Sirius Red, and Oil Red O as described previously³⁷. Sirius Red stained areas were quantified by ImageJ software³⁹. NASH scoring was performed by a liver pathologist using metrics for the NASH-CRN⁴⁰. Apoptosis was detected using a terminal deoxynucleotidyl transferase dUTP nick end-labeling (TUNEL) assay as per the manufacturer's instructions (R&D Systems, Minneapolis, MN). For immunohistochemistry, paraffin-embedded liver tissue sections were probed with F4/80 antibody (eBiosciences, San Diego, Ca), followed by detection with HRP-conjugated secondary antibody and DAB substrate kit (Cell Signaling, Danvers, MA) following manufacturer's guidelines. Photomicrographs of histologic sections were obtained using a Zeiss Light Microscope (Zeiss, Jena, Germany).

Serological analysis

Blood obtained by cardiac puncture was collected for measurements of serum alanine aminotransferase (ALT), and aspartate aminotransferase (AST) concentrations using an AST and ALT Activity Assay Kit (Sigma-Aldrich, St. Louis, MO).

Quantitative real-time PCR

Isolation of total RNA from liver, cDNA synthesis and qRT-PCR were performed as previously described⁴¹. Expression data were normalized to 18S rRNA and data are presented as fold change in gene expression compared to WT- standard chow (ND) fed controls.

Glucose tolerance test (GTT) and insulin tolerance test (ITT)

GTTs and ITTs were performed at baseline, and after 16 weeks of HFCD feeding as described previously³⁰. Blood glucose concentration was measured using a hand held glucometer (Freestyle Flash, Abbott Laboratories, Abbott Park, IL) as described elsewhere³⁰.

Immunoblotting

Liver tissues were homogenized and sonicated in RIPA lysis buffer (Sigma-Aldrich, St. Louis, MO) containing protease inhibitor cocktail and phosphostop (Roche Diagnostics, Indianapolis, IN), and total protein was extracted by centrifugation. A total of 30 μ g protein was resolved on 4–12% precast SDS-PAGE gel (Invitrogen, Grand Island, NY) and transblotted onto a polyvinylidene fluoride membrane (Bio-Rad, Hercules, CA). Membranes were probed with antisera against p-JNK, JNK, p-PDK, PDK, p-AKT, AKT, p-PKC ζ , PKC ζ , p-PERK, PERK, p-elF2 α , elF2 α , p-IRE-1, IRE-1, GRP78, CHOP, caspase-3, cleaved caspase-3, ATF-6 α , XBP-1, BCL-2, Bax, LC-3B and β -actin (Cell Signaling Technology, Danvers, MA). Isotype-matched horseradish peroxidase conjugated secondary antibodies, enhanced chemiluminescence substrate (Pierce, Rockford, IL) and a FluorChem 8900 digital imaging system (AlphaInnotech, San Leandro, CA) were used to visualize protein bands. Densitometric analyses was performed with VisionWorks® Software, version 6.8 (UVP, Upland, CA).

Hepatic hydroxyproline quantification:

Hepatic 4-hydroxyproline concentration was quantified using the Hydroxyproline Assay kit (Sigma-Aldrich, St. Louis, MO) following manufacturer's guidelines.

Primary mouse hepatocyte isolation and in vivo fatty acid and exendin-4 treatment

Primary mouse hepatocytes from 8 weeks old WT and $CHOP^{-/-}$ mice were isolated as previously described⁴². Isolated hepatocytes were cultured in William's medium E Formulation (Sigma-Aldrich, St. Louis, MO) supplemented with 10% fetal bovine serum (FBS; Atlanta Biologicals, Flowery Branch, GA) and 5% Penicillin-Streptomycin (Sigma-Aldrich, St. Louis, MO). The *in vitro* fatty acid and exendin-4 treatments were performed as described previously³¹. Hepatocytes were pretreated with 400 µM palmitic acid (PA) in ethanol (EtOH) for 24 h, after which hepatocytes were treated with 10 nM exendin-4 and PA for an additional 24 hr. Hepatocytes treated with EtOH (Control), or EtOH plus exendin-4 (Control + Ex4), or PA alone were used as controls. At the end of the experiment, cells were subjected to RNA isolation or Oil Red O staining as described above. Gene expression data were normalized to GAPDH and data are presented as fold change in gene expression compared to WT + EtOH controls.

Statistical analysis

Values were reported as means \pm SEM and each treatment group included 5-7 mice. All experiments were repeated at least two times on two separate occasions. Data were analyzed by analysis of variance (ANOVA) with Bonferroni's post hoc test for determination of statistical significance (p < 0.05).

RESULTS

Liraglutide treatment attenuates steatohepatitis in HFCD-fed WT mice

We have previously demonstrated that liraglutide treatment significantly reduced steatosis in WT mice fed a high fat and high sugar diet (HFD)³⁰⁻³⁴. We recently reported that mice fed 0.2% cholesterol in addition to HFD (HFCD) for 16 weeks resulted in mild steatohepatitis characterized by steatosis, inflammation, and mild fibrosis³⁷. Here, we first determined whether liraglutide treatment would also be effective in suppressing steatohepatitis in HFCD-fed mice. As shown in Fig. 1A-M, WT mice fed the HFCD for 16 weeks developed histological features of steatohepatitis indicated by ballooning degeneration of hepatocytes, inflammatory cell infiltration, and sinusoidal fibrosis. In agreement with previous reports³⁰⁻³⁴, four weeks of liraglutide treatment significantly reduced hepatic steatosis, body weight, and both liver and visceral fat weight expressed as percent of body weight (Fig. 1A-F). Liraglutide treatment also significantly reduced the NASH-CRN histology score⁴³, hepatic hydroxyproline levels and serum AST and ALT levels (Fig 1G-J).

CHOP modulates liraglutide-mediated suppression of steatohepatitis

As anticipated, CHOP expression was markedly increased in HFCD-fed WT mice, and the daily administration of liraglutide suppressed CHOP expression (Fig. 6A,H). Therefore, we investigated whether CHOP was pivotal in liraglutide-mediated suppression of steatohepatitis. Consistent with previous observations⁴⁴, HFCD-fed *CHOP^{-/-}* mice developed more severe histological features of NASH compared with HFCD-fed WT mice (Fig. 1A-C, G & 2A-C, G). HFCD-fed CHOP-/- mice gained greater body weight, and had greater liver weight and visceral fat weight compared with HFCD-fed WT mice (Fig. 1D-F & 2D-F). HFCD-fed CHOP^{-/-} mice had higher hepatic fibrosis compared with HFCD-fed WT mice (Fig. 1K-M & 2K-M). In addition, hepatic inflammation in HFCD-fed CHOP-/mice was higher than HFCD-fed WT mice as evidenced by increased infiltration of macrophages, increase in tumor necrosis factor (TNF)-a and monocyte chemoattractant protein (MCP)-1 transcript levels, and higher serum AST and ATL levels (Fig. 3A-C & 2A, G. M). Liraglutide treatment attenuated steatosis, and hepatic inflammation in the HFCD-fed CHOP-/- mice, but this effect was far less robust compared to the liraglutide treated HFCDfed WT mouse cohort (Fig. 1A-M; 2A-M & 3A-C). Taken together, these data suggest that CHOP serves to protect mouse liver from HFCD-mediated hepatic inflammation. Also, that CHOP plays an essential role in liraglutide-mediated protection from NASH-associated liver injury.

HFCD-fed CHOP^{-/-} mice are resistant to liraglutide-mediated improvement in insulin sensitivity and glucose homeostasis

We and others have previously demonstrated that liraglutide treatment attenuates steatosis by improving hepatic insulin sensitivity in both *in vitro* and *in vivo* models of hepatic steatosis³⁰⁻³⁴. Since CHOP plays a role in suppressing metabolic genes during ER stress^{20, 28}, we tested whether liraglutide improves insulin sensitivity in the absence of CHOP. As seen in Fig 4A-D, liraglutide treatment significantly improved hyperglycemia and insulin sensitivity in HFCD-fed WT mice. However, HFCD-fed *CHOP*^{-/-} mice remained insulin resistant and hyperglycemic despite 4 weeks of liraglutide treatment (Fig. 4A-D). To

determine the molecular mechanisms that could account for persistent insulin resistance in the *CHOP*^{-/-} mice, we performed a detailed analysis of the downstream mediators of the insulin-signaling pathway. As shown in Fig. 4E-J, we demonstrated that activation of PDK, AKT and PKC ζ were significantly attenuated in both WT and *CHOP*^{-/-} mice fed HFCD. Although we did not observe any significant differences in JNK activation between ND and HFCD fed WT mice, HFCD significantly increased JNK activation in *CHOP*^{-/-} mice, which also correlated with impaired insulin sensitivity in these mice (Fig. 4E,G). In agreement with our previous *in vitro* results³⁴, liraglutide treatment significantly activated PDK, AKT, and PKC ζ and reduced JNK activation in HFCD-fed WT mice (Fig. 4E-J). Interestingly, liraglutide treatment also induced activation of PDK, AKT and PKC ζ in HFCD-fed *CHOP*^{-/-} mice. However liraglutide administration in *CHOP*^{-/-} mice failed to suppress JNK activation; and, while insulin signaling elements were activated, glucose handling in the *CHOP*^{-/-} mice fed the HFCD was not improved.

Liraglutide treatment fails to improve lipid, cholesterol and glucose metabolism in the absence of CHOP

Next, to understand the discrepancy between insulin signaling pathway activation and failure to improve glucose handling following liraglutide administration in *CHOP*^{-/-} mice, we investigated the effect of liraglutide on key mediators of lipid, cholesterol and glucose metabolism. As seen in Fig. 5A-D, HFCD significantly increased peroxisomal β -oxidation and uptake of long chain fatty acids as indicated by increased expression of Carnitine palmitoyltransferase (CPT)-1a, Acyl-CoA oxidase (ACOX)-1 and ACOX-2, key mediators of peroxisomal β -oxidation, in WT HFCD-fed mice. HFCD also increased CPT-1a and ACOX-2 expression in *CHOP*^{-/-} mice, however the expression level was significantly lower than the WT mice (Fig. 5A-D). Liraglutide treatment suppressed β -oxidation in HFCD-fed WT mice as indicated by increased peroxisome proliferator-activated receptor (PPAR)-a expression and decrease in CPT-1a, ACOX-1 and ACOX-2 expression. None of these key transcription factors and enzymes associated with β -oxidation were suppressed in the liraglutide administered HFCD-fed *CHOP*^{-/-} mice, nor was PPARa increased, suggesting that CHOP is required for liraglutide-mediated suppression of β -oxidation (Fig. 5A-D).

In contrast to β -oxidation, expression levels of sterol regulatory element-binding protein (SREBP)-1c, PPAR- γ , fatty acid synthase (FAS) and carbohydrate-responsive elementbinding protein (ChREBP), key mediators of *de novo* lipogenesis and carbohydrate metabolism, were significantly higher in HFCD-fed *CHOP*^{-/-} mice suggesting a role of CHOP in regulating carbohydrate and fat metabolism (Fig. 5E-H). Liraglutide treatment suppressed *de novo* lipogenesis in both WT and *CHOP*^{-/-} mice fed HFCD as indicated by reduced expression of PPAR- γ and FASN (Fig. 5E-H). Interestingly, liraglutide failed to suppress SREBP1c and ChREBP in HFCD-fed *CHOP*^{-/-} mice suggesting that CHOP may play a role in Liraglutide-mediated suppression of glucose metabolism. Taken together, these data suggest that CHOP is required for liraglutide-mediated improvement in lipid, cholesterol, and glucose metabolism even though key players of the hepatocyte insulin signaling pathway were activated in both WT and the *CHOP*^{-/-} mice fed the HFCD.

Liraglutide treatment attenuates ER stress in HFCD-fed WT mice but not in CHOP-/- mice

Since hepatic ER stress contributes to the development of steatosis^{10, 11, 15, 17, 45}, we investigated whether liraglutide treatment attenuates steatohepatitis by suppressing ER stress in the absence of CHOP. As shown in Fig 6A-H, HFCD significantly activated PKR like ER kinase (PERK) and inositol requiring 1 (IRE-1) pathways, as indicated by increased phosphorylation of PERK and its downstream effector E74-like factor 2a (elF2a), and IRE-1 and its downstream effector X-box binding protein (XBP)-1, respectively, in WT mice. HFCD also increased activation of activating transcription factor (ATF)-6α in WT mice. Similarly all three key UPR pathways were also significantly activated in the HFCD-fed *CHOP*^{-/-} mice (Fig. 6A-H). However, in the absence of CHOP, HFCD induced significantly higher activation of PERK, IRE-1 and ATF-6a pathways compared to WT control (Fig. 6A-H). Liraglutide treatment attenuated HFCD-induced activation of PERK and IRE-1 pathways in WT mice suggesting that liraglutide treatment improved ER homeostasis. However, in the HFCD-fed CHOP^{-/-} mice, liraglutide treatment was ineffective in suppressing elF2 and IRE-1 activation, which could account for why liraglutide failed to suppress ER stress in the absence of CHOP. It should be noted that increased activation of IRE-1 in HFCD-fed *CHOP*^{-/-} mice did not increase XBP-1 activation in these mice. Furthermore, liraglutide treatment significantly increased XBP-1 activation in both WT and *CHOP*^{-/-} mice fed HFCD; however, this increase did not correlate with liraglutide-mediated decrease in IRE-1 activation in these mice. We also did not observe any significant differences in GRP78 expression between WT and CHOP-/- mice fed HFCD; nor did liraglutide treatment affect GRP78 expression in WT mice fed HFCD (Fig. 6A,E). However, liraglutide treatment significantly increased GRP78 expression in the HFCD-fed CHOP-/mice. HFCD-induced ER stress in WT mice correlated with increased expression of CHOP, and liraglutide treatment significantly reduced CHOP expression (Fig. 6A,H). Taken together, these data suggest that liraglutide treatment attenuated HFCD-induced hepatic injury by suppressing ER stress, and implicate a central role for CHOP in protecting hepatocytes from HFCD-induced ER stress.

Liraglutide treatment suppresses hepatic apoptosis in HFCD-fed WT mice but not in $CHOP^{-/-}$ mice

Since CHOP plays a role in ER stress-induced apoptosis, and liraglutide treatment has been shown to suppress lipotoxicity-induced hepatocyte apoptosis^{15, 17}, we next investigated whether liraglutide treatment was equally effective in suppressing apoptosis in the absence of CHOP. As seen in Fig 7A-F, the HFCD increased hepatocyte apoptosis in both WT and *CHOP*^{-/-} mice indicated by increased number of TUNEL positive cells and caspase-3 activation in the liver. However, the absence of CHOP resulted in significantly higher apoptosis in HFCD-fed *CHOP*^{-/-} mice compared with WT controls (Fig. 7A). Higher apoptosis in HFCD-fed CHOP^{-/-} mice correlated with increased Bax and LC-3B expression, and decreased BCL-2 expression (Fig. 7B-F). Liraglutide treatment reduced apoptosis in both WT and *CHOP*^{-/-} mice fed HFCD, however apoptosis remained significantly higher in liraglutide treated HFCD-fed *CHOP*^{-/-} mice compared with WT mice fed HFCD (Fig. 7A). While liraglutide treatment significantly lowered caspase-3 activation in *CHOP*^{-/-} mice, it did not reduce Bax and LC-3B expression in this cohort (Fig. 7B-F).

Taken together, these data suggest that CHOP is critical for liraglutide-mediated suppression of diet-induced hepatocyte death.

GLP-1 analog exendin-4 attenuates fatty acid deposition in primary hepatocytes from WT mice but not in primary hepatocytes from CHOP^{-/-} mice.

To investigate whether the role of CHOP in modulating liraglutide-mediated protection from NASH in vivo is hepatocyte specific, primary liver hepatocytes isolated from WT and *CHOP*^{-/-} mice were cultured in 400 µM PA palmitic acid (PA) and treated with 10 nM exendin-4. As seen in Fig. 8A, PA treatment significantly increased fat deposition in both WT and CHOP^{-/-} hepatocytes indicated by increased Oil Red O stained fat droplets in the hepatocytes. It should be noted that fat deposition in the CHOP^{-/-} hepatocytes treated with ethanol alone was higher compared with WT hepatocytes treated with ethanol (Fig. 8A). Exendin-4 reduced fat deposition in the PA treated WT hepatocytes, but failed to reduce fat deposition in PA treated CHOP-/- hepatocytes (Fig. 8A). Further analysis revealed that PA treatment significantly increased peroxisomal β-oxidation and uptake of long chain fatty acids as indicated by increased expression of CPT-1a, and ACOX-1 transcripts in both WT and *CHOP*^{-/-} hepatocytes (Fig. 8B-C). Exendin-4 treatment suppressed β -oxidation in PA treated WT hepatocytes, however did not suppress β -oxidation and fatty acid uptake in the PA treated *CHOP*^{-/-} hepatocytes (Fig. 8B-C). Additionally, exendin-4 also failed to suppress de novo lipogenesis and glucose metabolism in the absence of CHOP as indicated by significantly higher expression of FAS and SREBP1c transcripts in PA + Ex4 treated CHOP^{-/-} hepatocytes (Fig. 8D-E). Together these data support our *in vivo* observations and demonstrate that CHOP plays a role in modulating GLP-1 mediated protection from fat deposition in hepatocytes.

Discussion

In the present study, we demonstrate a critical role for CHOP in regulating diet-induced ER stress in the setting of administration of the long-acting GLP-1 analogue liraglutide *in vivo*, along with confirmatory *in vitro* studies in primary hepatocytes derived from $CHOP^{-/-}$ mice. Here, we show that CHOP plays a protective role in reducing ER stress-induced hepatocyte apoptosis, and is essential for the action of liraglutide in mediating these events during nutrient excess. Furthermore, our results demonstrate the contribution of liraglutide mediation of cell carbohydrate and lipid metabolism in preventing disease progression.

CHOP is a bZIP-containing transcription factor, and is a common point of molecular convergence for all three canonical ER stress transducers, IRE1, PERK, and ATF6^{46, 47}. CHOP is among the highly inducible genes during ER stress and its expression is primarily regulated at the transcriptional level by various components of the UPR pathway^{20, 48}. CHOP is primarily considered a pro-apoptotic transcription factor that mediates ER stress-induced cell death through the regulation of the Bcl-2 family of proteins⁴⁶. Consequently, CHOP deficiency provides partial resistance to ER stress-mediated apoptosis in *in vivo* models of chemically induced ER stress^{46, 47}. However, paradoxically, in our model of nutrient excess, the results indicate that CHOP plays a protective role, and not a deleterious one, in restoring ER homeostasis.

Despite the ongoing controversy surrounding the expression of GLP-1 receptor in the liver, studies conducted in our laboratory, and others, substantiate a direct role for GLP-1 receptor agonists in suppressing hepatic steatosis in both cell culture and animal models^{29-31, 34}. It is now well-established that GLP-1 receptor agonists suppress hepatic accumulation of triglycerides not only by promoting insulin sensitivity in fat-loaded hepatocytes, but also by restoring ER homeostasis and reducing subsequent lipotoxicity-mediated hepatocyte death^{29-31, 34}. In agreement with these previous reports, liraglutide administration protected WT mice from HFCD-induced steatohepatitis. However, liraglutide treatment did not have a significant effect on HFCD-induced steatohepatitis in *CHOP*^{-/-} mice. Liraglutide treatment, in the absence of CHOP failed to protect the deleterious effects of HFCD-induced liver injury and cell death, strongly suggesting that CHOP plays a direct role in liraglutide-mediated protection from HFCD-induced steatohepatitis. Our results are substantiated by *in vitro* findings of primary hepatocytes derived from *CHOP*^{-/-} mice, which demonstrate that GLP-1 analogues must play a role in modulating carbohydrate and lipid metabolism in the hepatocyte.

Apart from modulating ER stress associated apoptosis, CHOP also regulates carbohydrate and lipid metabolism by modulating transcriptional control of master regulators of glucose metabolism, fatty acid synthesis, and cholesterol metabolism^{20, 28}. Our data demonstrate a relationship between liraglutide administration and CHOP mediated regulation of key genes associated with hepatocyte metabolism in the setting of a Western diet (HFCD). In the HFCD-fed *CHOP*^{-/-} cohort, liraglutide administration failed to repress β -oxidation, and transcriptional regulators of *de novo* lipogenesis and cholesterol metabolism suggesting CHOP functions as a transcriptional repressor of metabolic genes during HFCD-induced ER stress. Taken together, these data suggest that CHOP plays a major role in the liraglutidemediated transcriptional regulation of hepatic metabolism during nutrient excess. Our findings provide a molecular framework in support of the recently reported small-scale clinical trial, which demonstrated liraglutide reduced hepatic insulin resistance and *de novo* lipogenesis¹⁶.

Apart from its insulinotropic effect, GLP-1 analogs also increase insulin sensitivity and glucose uptake by skeletal muscle, hepatocytes, and adipose tissue^{30, 34, 49}. Our results confirm liraglutide- mediated improvement in insulin sensitivity in HFCD-fed WT mice, but this was not true in the CHOP^{-/-} mice fed the HFCD. These data provide additional evidence for a role of CHOP in regulating glucose and more importantly fatty acid metabolism. While the insulin signaling pathway was activated in mice given liraglutide in the setting of the Western diet, only WT mice had restored glucose tolerance, not the *CHOP*^{-/-} mice. As anticipated the WT mice fed the Western diet had a reduction in JNK. That alone could explain why there is less caspase-3 activation and reduced TUNEL staining. This was not the case for the CHOP-/- mice. In both HFCD-fed CHOP-/- cohorts, JNK activity was unchanged, yet caspase-3 activity and TUNEL staining were suppressed in the CHOP^{-/-} mice fed the HFCD upon liraglutide administration. This would imply that either CHOP regulates the insulin pathway downstream of PDK-1, AKT and PKC- ζ , or that CHOP functions to suppress de novo lipogenesis. A reduction in de novo lipogenesis would explain reduced lipotoxicity, reduced hepatic inflammation, and ultimately reduced hepatocyte apoptosis.

The implication that CHOP is involved in ER stress-induced lipotoxicity stems from strong correlational data in NAFLD⁵⁰. However, rather than attenuating disease progression, genetic ablation of CHOP accelerates NAFLD progression in both HFD and methionine and choline deficient models of NASH^{44, 51}. The ability of CHOP to transcriptionally modulate fatty acid metabolism may explain its apparent paradoxical protective role during chronic nutrient excess conditions^{20, 28}. This conclusion is consistent with our data showing that liraglutide fails to promote insulin sensitivity, reduce steatosis, restore ER homeostasis and reduce subsequent lipotoxicity-mediated hepatocyte death in the absence of CHOP. Increased hepatocyte death in HFCD-fed CHOP-/- mice can be attributed to lower expression of anti-apoptotic protein BCL-2 and higher expression of Bax and LC-3B in these mice. In the CHOP^{-/-} mice, liraglutide administration reduced ER stress-related proteins, but not to the level of the WT mice fed the HFCD. Liraglutide treatment also reduced caspase-3 activation in HFCD-fed CHOP^{-/-} mice but this did not correlate with a decrease in TUNEL positive cells suggesting that another pathway in vivo must account for higher hepatocyte apoptosis following liraglutide administration. The increased expression of Bax and LC-3B in HFCD-fed CHOP^{-/-} mice and inability of liraglutide to attenuate apoptosis by reducing Bax and LC-3B levels in the absence of caspase-3 activation implies that both caspase dependent and independent cell death pathways are at play in the HFCDfed $CHOP^{-/-}$ mice. Future studies will be required to clarify the mechanisms of caspase independent cell death pathways.

In summary, our data indicate that *in vivo* the administration of liraglutide may be protective against NAFLD disease progression by initially reducing *de novo* lipogenesis and thus reducing lipotoxicity. As clinical trials continue, further clarification for the molecular mechanisms accounting for the pleotropic role of long-acting GLP-1 analogues in the liver is warranted.

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Abbreviations

NAFLD	non-alcoholic fatty liver disease
NASH	non-alcoholic steatohepatitis
СНОР	CCAAT/enhancer-binding protein homologous protein

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Figure 1. Liraglutide treatment attenuates steatohepatitis in HFCD-fed WT mice

Photomicrographs of (A) Hematoxylin and Eosin, (B) Oil Red O, and (C) Sirius Red stained liver sections of WT mice fed a ND or HFCD for 16 weeks. After 12 weeks of feeding, mice received daily injections of saline or liraglutide (Lira) for 4 weeks. Black arrowheads, inflammatory immune cells; blue arrowheads, fat droplets; black arrows, collagen deposition. (D) Body, (E) liver, and (F) visceral fat weight as well as (G) NASH-CRN score, serum (H) ALT and (I) AST levels, and (J) hepatic hydroxyproline levels measured at the end of the 16 week experiments. Changes in liver and visceral fat weights are reported as percentages of body weight. (K) Quantitative analysis of Sirius Red stained liver tissue sections. Quantitative RT-PCR analysis of hepatic (L) α SMA and (M) TNF- α transcript levels measured at the end of the 16 week experiments. Data are presented as mean \pm SEM; n = 5-7 mice per group. Hashtags indicate significant differences (p < 0.05) between ND- or HFCD-fed WT mice. "o" indicates significant differences (p < 0.05) between saline or liraglutide-treated WT mice. Scale 20 µm.



Figure 2. CHOP modulates liraglutide-mediated suppression of steatohepatitis

Photomicrogaphs of (A) Hematoxylin and Eosin, (B) Oil Red O, and (C) Sirius Red stained liver tissue sections of *CHOP*^{-/-} mice fed a ND or HFCD for 16 weeks. After 12 weeks of feeding, mice received daily injections of saline or liraglutide (Lira) for 4 weeks. Black arrowheads, inflammatory immune cells; blue arrowheads, fat droplets; black arrows, collagen deposition. (D) Body, (E) liver, and (F) visceral fat weight as well as (G) NASH-CRN score, serum (H) ALT and (I) AST levels, and (J) hepatic hydroxyproline levels measured at the end of the 16 week experiments. Changes in liver and visceral fat weights are reported as percentages of body weight. (K) Quantitative analysis of Sirius Red stained liver tissue sections. Quantitative RT-PCR analysis of hepatic (L) α SMA and (M) TNF- α transcript levels measured at the end of the 16 week experiments. Data are presented as mean \pm SEM; n = 5-7 mice per group. Hashtags indicate significant differences (p < 0.05) between ND- or HFCD-fed WT mice. "o" indicates significant differences (p < 0.05)



Figure 3. Hepatic macrophage infiltration is higher in HFCD-fed *CHOP^{-/-}* mice

(A) Immunohistochemical staining of F4/80⁺ macrophages in the liver. Black arrows, F4/80⁺ macrophages. Quantitative RT-PCR analysis of (B) F4/80 and (C) MCP-1 mRNA transcript levels in the liver of WT or *CHOP*^{-/-} mice fed a ND or HFCD for 16 weeks. After 12 weeks of feeding, mice received daily injections of saline or liraglutide (Lira) for 4 weeks. Data are presented as mean \pm SEM; n = 5-7 mice per group. Hashtags indicate significant differences (p < 0.05) between ND- or HFCD-fed *CHOP*^{-/-} mice. Stars indicate significant differences (p < 0.05) between WT and *CHOP*^{-/-} mice. "o" indicate significant differences (p < 0.05) between saline or liraglutide treated *CHOP*^{-/-} mice. Scale 20 µm.



Figure 4. HFCD-fed *CHOP^{-/-}* mice are resistant to liraglutide-mediated improvement in insulin sensitivity and glucose homeostasis

(A, C) Glucose and (B, D) insulin tolerance in (A, B) WT and (C, D) $CHOP^{-/-}$ mice fed a ND or HFCD for 16 weeks. After 12 weeks of feeding, mice received daily injections of saline (sal) or liraglutide (Lira) for 4 weeks. (E, F) Representative western blot images and (G-J) densitometry measurements of expression level of various effector molecules of the insulin-signaling pathway in the liver. Hashtags indicate significant differences (p < 0.05) between ND- or HFCD-fed mice. Stars indicate significant differences (p < 0.05) between saline or Liraglutide treated mice. Data are presented as mean ± SEM; n = 5-7 mice per group.



Figure 5. HFCD-fed $CHOP^{-/-}$ mice are resistant to liraglutide-mediated improvement in lipid, cholesterol and glucose metabolism

Quantitative RT-PCR analysis of (A) *PPAR-a*, (B) *CPT-1a*, (C) ACOX-1, (D) ACOX-2, (E) *SREBP1c*, (F) *PPAR-* γ , (G) FAS, and (H) ChREBP transcript levels in the liver of WT or *CHOP*^{-/-} mice fed a ND or HFCD for 16 weeks. After 12 weeks of feeding, mice received daily injections of saline or liraglutide for 4 weeks. Data are presented as mean ± SEM; *n* = 5-7 mice per group. Hashtags indicate significant differences (*p* < 0.05) between ND- or HFCD-fed mice. Stars indicate significant differences (*p* < 0.05) between WT and *CHOP*^{-/-} mice. "o" indicate significant differences (*p* < 0.05) between saline or Liraglutide treated mice.



Figure 6. Liraglutide treatment attenuates ER stress in HFCD-fed WT mice but not in *CHOP*^{-/-} mice

(A) Representative western blot images and (B-H) densitometry measurements of various ER stress parameters in the liver of WT and $CHOP^{-/-}$ mice fed a ND or HFCD for 16 weeks. After 12 weeks of feeding, mice received daily injections of saline (sal) or liraglutide (Lira) for 4 weeks. Data are presented as mean \pm SEM; n = 5-7 mice per group. Hashtags indicate significant differences (p < 0.05) between ND- or HFCD-fed mice. Stars indicate significant differences (p < 0.05) between WT and $CHOP^{-/-}$ mice. "o" indicate significant differences (p < 0.05) between WT and $CHOP^{-/-}$ mice.



Figure 7. Liraglutide treatment suppresses apoptosis in HFCD-fed WT mice but not in *CHOP*^{-/-} mice

(A) Representative immunofluorescence images of liver tissue sections stained with terminal deoxynucleotidyl transferase (TUNEL; green) to detect apoptotic cells. (B) Representative western blot images and (C-F) densitometry measurements of hepatic cleaved caspase 3, BCL-2, Bax and LC-3B expressions in WT and *CHOP*^{-/-} mice fed a ND or HFCD for 16 weeks. After 12 weeks of feeding, mice received daily injections of saline (sal) or Liraglutide (Lira) for 4 weeks. Data are presented as mean \pm SEM; n = 5-7 mice per group. Hashtags indicate significant differences (p < 0.05) between ND- or HFCD-fed mice. Stars indicate significant differences (p < 0.05) between WT and *CHOP*^{-/-} mice. "o" indicate significant differences (p < 0.05) between saline or Liraglutide treated mice. Scale 20 µm.



Figure 8. GLP-1 analog exendin-4 attenuates fatty acid deposition in primary hepatocytes from WT mice but not in primary hepatocytes from $CHOP^{-/-}$ mice

(A) Oil Red O stained micrographs of primary hepatocytes isolated from WT and $CHOP^{-/-}$ mice. Quantitative RT-PCR analysis of (B) ACOX-1, (C) CPT-1a, (D) FAS and (E) SREBP1c transcript levels in the primary hepatocytes isolated from WT and $CHOP^{-/-}$ mice. Hepatocytes were pretreated with 400 µM palmitic acid (PA) in ethanol for 24 h after which hepatocytes were treated with 10 nM exendin-4 and 400 µM PA for an additional 24 hr. Hepatocytes treated with ethanol (EtOH), or EtOH plus exendin-4 (Control + Ex4), or PA alone served as controls. Data presented are representative of 2 independent experiments with four replicates per treatment group. Hashtags indicate significant differences (p < 0.05) between EtOH or PA treated hepatocytes. Stars indicate significant differences (p < 0.05) between WT and $CHOP^{-/-}$ hepatocytes. "o" indicate significant differences (p < 0.05) between untreated or liraglutide treated mice. Scale 10 µm.