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Hepatic Diacylglycerol-Associated Protein Kinase C ϵ **Translocation Links Hepatic Steatosis to Hepatic Insulin Resistance in Humans**

Graphical Abstract



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In Brief

ter Horst et al. show that simple hepatic steatosis in the context of NAFLD does not fully explain hepatic insulin resistance in obese humans. Impaired hepatic insulin action was characterized by accumulation of diacylglycerol subspecies in the cytosol of hepatocytes and translocation of PKC ϵ from the cytosol to the membrane.

Highlights

- The presence of hepatic steatosis is associated with insulin resistance
- Intrahepatic triglycerides are not sufficient for hepatic insulin resistance
- Diacylglycerol in hepatic cytosol predicts insulin inhibition of glucose production
- Diacylglycerol-associated insulin resistance is characterized by PKC_E translocation



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Hepatic Diacylglycerol-Associated Protein Kinase C ϵ Translocation Links Hepatic Steatosis to Hepatic Insulin Resistance in Humans

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SUMMARY

Hepatic lipid accumulation has been implicated in the development of insulin resistance, but translational evidence in humans is limited. We investigated the relationship between liver fat and tissue-specific insulin sensitivity in 133 obese subjects. Although the presence of hepatic steatosis in obese subjects was associated with hepatic, adipose tissue, and peripheral insulin resistance, we found that intrahepatic triglycerides were not strictly sufficient or essential for hepatic insulin resistance. Thus, to examine the molecular mechanisms that link hepatic steatosis to hepatic insulin resistance, we comprehensively analyzed liver biopsies from a subset of 29 subjects. Here, hepatic cytosolic diacylglycerol content, but not hepatic ceramide content, was increased in subjects with hepatic insulin resistance. Moreover, cytosolic diacylglycerols were strongly associated with hepatic PKC_E activation, as reflected by PKC_E translocation to the plasma membrane. These results demonstrate the relevance of hepatic diacylglycerol-induced PKC ε activation in the pathogenesis of NAFLD-associated hepatic insulin resistance in humans.

INTRODUCTION

Insulin resistance and non-alcoholic fatty liver disease (NAFLD) are common and consequential complications of obesity (Grundy, 2000). Ectopic triglyceride accumulation in the liver has clearly been linked to non-alcoholic steatohepatitis (NASH), liver cirrhosis, and hepatocellular carcinoma (Wong et al., 2014), but it is less clear whether there is a causal relationship between intrahepatic triglyceride (IHTG) accumulation and hepatic insulin resistance (Farese et al., 2012; Shulman, 2000).

Elevated IHTG content is associated with hepatic and/or muscle insulin resistance in most, but not in all, human studies (Bugianesi et al., 2005; Deivanayagam et al., 2008; Gastaldelli et al., 2007; Korenblat et al., 2008; Kotronen et al., 2008; Petersen et al., 2002, 2005; Sanyal et al., 2001; Seppälä-Lindroos et al., 2002). Moreover, in addition to triglycerides, it is now established that other (possibly more bioactive) lipid species may also accumulate in hepatocytes in the context of NAFLD. In several animal models, buildup of diacylglycerol (DAG) has been linked to hepatic insulin resistance through activation of protein kinase C_{ε} (PKC ε), which may directly interfere with hepatic insulin signaling (Jornayvaz et al., 2011; Samuel et al., 2004, 2007; Samuel and Shulman, 2016; Xia et al., 2015). In this regard, PKC_E has recently been shown to phosphorylate insulin receptor (INSR) threonine¹¹⁶⁰, which led to inhibition of INSR kinase activity (Petersen et al., 2016). Furthermore, mice with a threonineto-alanine mutation at the homologous residue threonine $^{\rm 1150}$ (Insr^{T1150A} mice) were protected from high-fat-diet-induced

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Table 1. Characteristics of Study Participants According to Extent of Hepatic Steatosis, n = 133						
	No Steatosis (n = 52)	Mild Steatosis (n = 41)	Severe Steatosis (n = 40)	р		
Female sex	27 (51)	14 (34)	18 (45)	0.229		
Age (years)	49 ± 2	51 ± 2	49 ± 2	0.634		
BMI (kg/m²)	39 ± 1	41 ± 1	41 ± 1	0.442		
Body fat (%)	45 ± 1	45 ± 2	46 ± 1	0.695		
Fasting glucose (mmol/L)	5.2 ± 0.1	5.2 ± 0.1	5.4 ± 0.1	0.301		
Triglycerides (mmol/L)	1.3 ± 0.1	1.6 ± 0.1	1.7 ± 0.2^{a}	0.039		
Cholesterol (mmol/L)	4.9 ± 0.1	4.9 ± 0.2	5.0 ± 0.2	0.671		
High-density lipoprotein (mmol/L)	1.2 ± 0.0	1.1 ± 0.1	1.1 ± 0.0	0.287		
Low-density lipoprotein (mmol/L)	3.0 ± 0.1	3.1 ± 0.2	3.2 ± 0.1	0.620		
Alanine aminotransferase (U/L)	26 ± 2	38 ± 3^{b}	$40 \pm 3^{\circ}$	<0.001		
Aspartate aminotransferase (U/L)	22 ± 1	30 ± 3	32 ± 3^{b}	0.004		
Gamma-glutamyl transpeptidase (U/L)	34 ± 4	42 ± 7	42 ± 4	0.320		
C-reactive protein (mg/L)	4.9 ± 0.9	6.0 ± 1.0	7.6 ± 1.3	0.197		
Resting energy expenditure (kcal/day)	1,862 ± 48	2,000 ± 72	1,931 ± 54	0.256		
IHTG (%)	2.9 ± 0.2	10.4 ± 0.4 ^c	$23.3 \pm 1.0,^{c},^{d}$	<0.001		

Data are count (%) or mean ± SEM. No, mild, and severe steatosis were defined as intrahepatic triglycerides (IHTG) <5.56%, 5.56%–15%, or >15%, respectively.

^ap < 0.05 versus no steatosis.

^bp < 0.01 versus no steatosis.

^cp < 0.001 versus no steatosis.

^dp < 0.001 versus mild steatosis.

hepatic insulin resistance and displayed increased hepatic insulin signaling, suppression of hepatic glucose production, and hepatic glycogen synthesis compared with wild-type (WT) controls during hyperinsulinemic clamp studies. These results provide a strong mechanistic link for DAG-PKCε-mediated hepatic insulin resistance in the context of NAFLD. In contrast, other studies have suggested that an increase in hepatic ceramide content is the major mediator of lipid-induced hepatic insulin resistance in the context of NAFLD (Chavez and Summers, 2012; Holland et al., 2011; Luukkonen et al., 2016), although the mechanism by which ceramides cause hepatic insulin resistance is less clear. In the few available translational studies, hepatic DAG content, but not hepatic ceramide content, correlated with indices of hepatic insulin resistance in obese humans (Kumashiro et al., 2011; Magkos et al., 2012).

There are also circumstances, in both animals and humans, in which hepatic insulin resistance is dissociated from hepatic steatosis (Amaro et al., 2010; Benhamed et al., 2012; Brown et al., 2010; Buettner et al., 2004; Monetti et al., 2007; Semple et al., 2009; Visser et al., 2011). Some, but not all, of these apparently conflicting observations may be the result of differences among studies in diagnosis and definition of fatty liver and/or insulin resistance (Farese et al., 2012). Moreover, a better understanding of the subcellular compartmentalization of hepatic DAG species and PKC_E may explain some of the apparent dissociations between hepatic steatosis and hepatic insulin resistance (Samuel and Shulman, 2016). For instance, short-term high-fat feeding promotes hepatic insulin resistance, which is associated with increases in cytosolic and membrane DAG content and activation of PKC_E as reflected by translocation of PKC_E from the cytosol to the plasma membrane (Samuel et al., 2004, 2007). In contrast,

antisense oligonucleotide knockdown of hepatic CGI-58 leads to hepatic steatosis, which is associated with considerable accumulation of DAGs in lipid droplets (Cantley et al., 2013). Contrary to the high-fat-fed mice, this increase in lipid droplet DAGs is not associated with PKC ϵ activation or hepatic insulin resistance. It is currently unknown how these paradigms translate to humans.

Therefore, we here report a clinical study in obese non-diabetic humans aimed to (1) determine the relationship between liver fat content and insulin resistance using gold-standard metabolic flux measurements and (2) determine whether the subcellular distribution of individual hepatic lipid species and PKC ε in liver biopsies is relevant for hepatic insulin resistance in humans.

RESULTS AND DISCUSSION

The Presence of NAFLD, but Not the Extent of Hepatic Steatosis, Is Associated with Insulin Resistance in Obese Humans

To determine whether the accumulation of IHTG in the context of NAFLD relates to abnormalities in glucose metabolism and/or tissue-specific parameters of insulin action, we measured these metabolic fluxes using the two-step hyperinsulinemic-euglycemic clamp technique in a cohort of 133 obese adults. Fifty-two subjects had normal IHTG by proton magnetic resonance spectroscopy (¹H-MRS) (no steatosis, IHTG <5.56% [Szczepaniak et al., 2005]). Of the 81 subjects with ¹H-MRS-defined hepatic steatosis, 41 subjects had mild steatosis (IHTG 5.56%–15%), and 40 subjects had severe steatosis (IHTG >15%). These groups did not differ in most baseline characteristics, but subjects with hepatic steatosis had higher plasma triglycerides and aminotransferases than those without steatosis (Table 1).



Figure 1. The Presence of Hepatic Steatosis, but Not Its Extent, Is Associated with Insulin Resistance in Obese Humans

(A) The basal rate of EGP was assessed after an overnight fast.

(B) Hepatic insulin sensitivity is expressed as the insulin-mediated suppression of basal EGP.

(C) Adipose tissue insulin sensitivity is expressed as the insulin-mediated suppression of circulating FA.

(D) Peripheral insulin sensitivity is expressed as the insulin-stimulated Rd of glucose.

Data are mean \pm SEM (n = 125–133). ***p \leq 0.001.

Hepatic steatosis was not associated with differences in the basal rate of endogenous glucose production (EGP) (Figure 1A). However, subjects with mild or severe hepatic steatosis had higher fasting plasma insulin levels than subjects without hepatic steatosis (Table S1). Since fasting insulin × basal EGP is an index of hepatic insulin resistance (Korenblat et al., 2008), this suggests that basal hepatic responsiveness to insulin may be impaired in obese subjects with steatosis (Figure S1A). Indeed, during the low-dose insulin step of the hyperinsulinemiceuglycemic clamp, insulin-mediated suppression of EGP was impaired in subjects with mild or severe hepatic steatosis, indicating hepatic insulin resistance (Figure 1B). In both steatosis groups, insulin suppression of plasma fatty acids (FA) and insulin stimulation of the glucose rate of disappearance (Rd) were also reduced, indicating adipose tissue and peripheral/muscle insulin resistance, respectively (Figures 1C, 1D, S1B, and S1C). Notably, there were no differences in any of these parameters between subjects with mild or severe hepatic steatosis (Figures 1B-1D). Finally, these results were not caused by differences in insulin clearance (Figures S1D and S1E) or levels of glucoregulatory hormones during the clamps (Table S1). Therefore, data from our large cohort of obese subjects show that the presence of liver steatosis is associated with impaired insulin sensitivity in the liver, adipose tissue, and muscle, but that the extent of hepatic steatosis does not predict the severity of insulin resistance.

IHTG Are Not Sufficient or Essential for the Development of Hepatic Insulin Resistance in Obese Humans

In agreement with the previous data, regression analysis confirmed that the relationship between IHTG and insulin suppression of EGP was best described by a nonlinear model: IHTG and hepatic insulin action were negatively correlated in the lower range of steatosis but dissociated in the higher range of steatosis (Figure 2A). More importantly, obese subjects with hepatic steatosis displayed nearly the same variation in hepatic insulin sensitivity as subjects without hepatic steatosis (suppression of EGP ranging from 38%–99% and 51%–100%, respectively). In fact, the distribution of both variables clearly shows that hepatic steatosis can occur without hepatic insulin resistance, and vice versa (Figure 2A). Therefore, IHTG are not strictly sufficient or necessary for hepatic insulin resistance (Farese

et al., 2012), at least in the conditions of our study. We observed similar relationships between IHTG and parameters of extrahepatic insulin sensitivity (Figures 2B and 2C). These observations are consistent with previous human studies that have demonstrated IHTG accumulation without insulin resistance (Amaro et al., 2010; Visser et al., 2011) or insulin resistance in the absence of IHTG (Semple et al., 2009). Therefore, the available data do not implicate that IHTGs are a cause for hepatic insulin resistance, and this indicates that (an)other factor(s) must be responsible for NAFLD-associated hepatic insulin resistance.

Hepatic Insulin Resistance Is Associated with DAG Accumulation In Cytosol

Non-invasive measurement methods such as ¹H-MRS cannot distinguish between inert lipids such as triglycerides (Yu et al., 2002) and bioactive lipids such as DAGs or ceramides. The subcellular distribution of these lipids may also be critical to their biological effects. Therefore, to determine whether these parameters are important for hepatic insulin resistance in obese humans, we obtained liver biopsies from a subset of subjects that underwent bariatric surgery <2 weeks after the clinical experiments (n = 29).

Liquid chromatography/tandem mass spectrometry (LC-MS/ MS) analysis of snap-frozen liver biopsies revealed that neither hepatic ceramide (r = 0.32, p = 0.117) nor total DAG (r = 0.31, p = 0.109) content predicted hepatic insulin sensitivity in morbidly obese subjects (Figures S2A and S2B). However, DAG may be present in the cell membrane, cytosol, and lipid droplets (Cantley et al., 2013; Kumashiro et al., 2011). Using a subcellular fractionation method (Cantley et al., 2013), we separated liver biopsies into membrane and cytosolic fractions. Here, we observed that DAG accumulation in the cytosolic fraction (r = 0.43, p = 0.024), but not DAG accumulation in the membrane (r = 0.19, p = 0.342), predicted hepatic insulin resistance assessed by clamp (Figures S2C and S2D). Cytosolic DAG also strongly correlated with other parameters of hepatic insulin action including the Hepatic Insulin Sensitivity Index, HOMA-IR, and fasting plasma insulin levels (data not shown). Finally, we analyzed individual hepatic DAG species in relation to hepatic insulin sensitivity (Table S2). In the cytosolic fraction, hepatic DAGs composed of C18:1-C16:0, C16:0-C16:0, and C18:1-C18:1 were most abundant and also showed strong negative



correlations with suppression of EGP. In contrast, hepatic DAG composed of C20:4–C20:5 was highly abundant in the membrane fraction and positively correlated with hepatic insulin sensitivity. Individual hepatic ceramide species were not related to hepatic insulin action (Table S3).

Hepatic DAG-Associated Insulin Resistance and $\text{PKC}\epsilon$ Translocation

Hepatic DAGs are proposed to impair insulin signaling through activation of novel PKC (nPKC) (Farese et al., 2012; Nagle et al., 2009; Samuel and Shulman, 2016). In our prior study, we found that PKC ε was the major nPKC isoform expressed in human liver, and activation of PKC ε correlated with hepatic DAG content in the cytosol and HOMA-IR (Kumashiro et al., 2011). However, the association between PKC ε activation and insulin suppression of hepatic glucose output has not been reported in humans. Therefore, we measured PKC ε translocation to the cell membrane as an index of PKC ε activation (Kumashiro et al., 2011; Schmitz-Peiffer and Biden, 2008).

We compared liver biopsies from insulin-sensitive subjects (insulin suppression of EGP >80%, n = 10) with biopsies from insulin-resistant subjects (insulin suppression of EGP <65%, n = 8). The insulin-resistant subjects displayed both impaired fasting glucose concentrations and hyperinsulinemia (Table S4). In addition, insulin-resistant livers had a marked, 2.5-fold increase in cytosolic DAG content, but no difference in total DAG or ceramide content (Figures 3A-3D). In fact, all individual DAG species were increased in the hepatic cytosol of insulin-resistant subjects (Figure 3E). Notably, hepatic DAG-associated hepatic insulin resistance was characterized by a 2-fold increase in translocation of PKC ε from the cytosol to the membrane (Figures 3F and 3G). Taken together, this suggests that hepatic DAGs in the cytosol promote PKC ε translocation from the cytosol to the membrane where it can bind INSR and phosphorylate threonine¹¹⁶⁰, which, in turn, inhibits INSR kinase activity (Petersen et al., 2016).

Sources for Hepatic DAG Accumulation In Obese Humans

Hepatic DAG may accumulate from a variety of sources including circulating FA derived from adipose tissue lipolysis (Donnelly et al., 2005; Finck and Hall, 2015). To explore the potential role of adipose tissue lipolysis in promoting increased hepatic DAG content, we assessed rates of lipolysis measured by tracer dilution of $[1,1,2,3,3-^2H_5]$ glycerol during hyperinsulinemic

(B) Adipose tissue insulin sensitivity.
(C) Peripheral insulin sensitivity.
Lines are best fit and 95% Cl (n = 125–133).

conditions. Using this approach, we found that insulin-mediated suppression of the glycerol rate of appearance (Ra)

and circulating FA levels during insulin infusion both strongly correlated with hepatic DAG accumulation, indicating that higher release of FA from lipolysis during hyperinsulinemic (postprandial) conditions is associated with increased cytosolic DAG content (Figures S3A and S3B). In line, hepatic lipid synthesis in rats is primarily driven by substrate (FA) delivery to the liver, which occurs in an insulin-independent manner (Vatner et al., 2015). Thus, these data are consistent with the possibility that most of the hepatic DAG derives from FA from increased lipolysis in insulin-resistant adipose tissue, underlining the importance of healthy insulin-sensitive adipose tissue in the prevention of ectopic lipid disposition and hepatic insulin resistance.

Perspective

In this study, we demonstrated that the presence of hepatic steatosis in obese subjects is associated with insulin resistance in the liver, adipose tissue, and peripheral tissues. In subjects with hepatic steatosis, however, the amount of IHTG per se did not additionally affect tissue-specific parameters of insulin action. Noteworthy, although all subjects were obese, there was a wide range in insulin sensitivity. The distribution of the data further showed that IHTG were neither sufficient nor necessary for the development of hepatic insulin resistance, indicating that another underlying mechanism must be responsible for impaired insulin suppression of EGP in the context of NAFLD. Recent animal studies showed that the major underlying mechanism for hepatic insulin resistance involves hepatic DAG-induced PKC ε activation (Kumashiro et al., 2011; Petersen et al., 2016; Samuel and Shulman, 2016), but efforts to translate this paradigm to humans have been limited. Therefore, we comprehensively analyzed liver samples from subjects with normal or impaired insulin suppression of EGP and found that hepatic DAG accumulation in the cytosol was increased in obese humans with hepatic insulin resistance. Moreover, this was strongly associated with hepatic PKC_E activation as reflected by its translocation to the cell membrane. In contrast, there was no relationship between hepatic ceramide content and hepatic insulin resistance in these subjects. Taken together, these results support a model for the pathogenesis of hepatic insulin resistance in NAFLD, in which DAGs in the cytosol promote translocation of PKC ε to the membrane, where it inhibits hepatocellular insulin signaling. Here, we substantiate the relevance of this pathway in human metabolic disease.

Some nuances with respect to our findings should be made. First, the above-mentioned mechanism for lipid-induced hepatic insulin resistance should be appreciated in light of



(A) Hepatic ceramide content in obese subjects with normal hepatic insulin sensitivity or hepatic insulin resistance.

(B) Total hepatic DAG content.

(C) Hepatic DAG content in cytosolic fraction.

(D) Hepatic DAG content in membrane fraction.

(E) Individual DAG species in hepatic cytosol. Note the segmented y axis.

(F) Representative bands show PKC ε translocation from cytosol to membrane.

(G) Translocation of $\mathsf{PKC}\epsilon$ from cytosol (c) to membrane (m) as an index of activation.

Data are mean \pm SEM (n = 18–29). *p < 0.05. **p < 0.01. ***p = 0.001.

perform measurements in subcellular compartments but conclude that their data do not exclude the possibility that DAGs contribute to hepatic insulin resistance in humans, indicating that results are not strictly conflicting. In addition, the I148M gene variant in PNPLA3 was associated with hepatic steatosis without insulin resistance, whereas metabolic NAFLD was associated with a harmful hepatic lipidome (Luukkonen et al., 2016). We have also reported that hepatic steatosis in the context of familial hypobetalipoproteinemia is not associated with hepatic insulin resistance (Visser et al., 2011), suggesting that genetics may, in part, explain why some people with hepatic steatosis develop insulin resistance, whereas others do not. Un-

interconnected defects in other organ systems. For instance, extrahepatic regulation of adipose tissue lipolysis and FA availability by insulin is another important regulator of hepatic gluconeogenesis through substrate-driven regulation of pyruvate carboxylase activity/flux by hepatic acetyl CoA, which is independent of canonical hepatic insulin signaling (Perry et al., 2015). Thus, PKC ε inhibition of hepatic insulin signaling may mainly affect insulin regulation of net hepatic glycogen synthesis. Second, previous studies have suggested that increased hepatic ceramide synthesis may be causal in lipid-induced hepatic insulin resistance (Chavez and Summers, 2012; Holland et al., 2011; Luukkonen et al., 2016), but ceramides were unrelated to hepatic insulin resistance in the present study. Others have also reported such dissociations (Kumashiro et al., 2011; Magkos et al., 2012; Perry et al., 2013). Some of these conflicting results may be due to differences in experimental techniques or limited sample size/power. In one study, "metabolic NAFLD" was associated with an increase in hepatic ceramides and elevated HOMA-IR, but also with an increase in four out of five DAG species (Luukkonen et al., 2016). The authors did not fortunately, we do not have genotype data in this study, but it will be of interest to determine whether the I148M and other variants play a role in the susceptibility to lipid mediated-insulin resistance. Third, there is ample (reverse causal) evidence suggesting that extrahepatic insulin resistance contributes to the development of hepatic steatosis. In individuals with muscle and adipose tissue insulin resistance, nutrients are diverted to the liver. Accompanied by a compensatory hyperinsulinemia, these conditions provide both substrate (glucose and FA) and stimulus (insulin signaling through sterol regulatory elementbinding protein 1c) for hepatic triglyceride synthesis and de novo lipogenesis (Cohen et al., 2011; Perry et al., 2014; Petersen et al., 2007; Rabøl et al., 2011; Vatner et al., 2015). Finally, an important limitation in the interpretation of these data is the cross-sectional design of the study. However, we believe that obtaining serial liver biopsies in humans would be unethical in the absence of urgent medical indications, and numerous animal studies support a causal role for hepatic DAG accumulation and PKC_E activation in hepatic insulin resistance (Jornayvaz et al., 2011; Petersen et al., 2016; Samuel et al., 2007; Samuel and Shulman, 2016). These data offer further support for the DAG-PKC ε hypothesis for lipid-induced hepatic insulin resistance in NAFLD and support the development of interventions that target hepatic DAG-induced PKC ε activation for the prevention and/or treatment of type 2 diabetes.

EXPERIMENTAL PROCEDURES

Subjects

Subjects participated in metabolic studies at the Academic Medical Center (Amsterdam, the Netherlands). Treatment-naive subjects were included in this study if they were obese (BMI \geq 30 kg/m²) with stable weight (<5% weight change) for >3 months prior to examinations, and if they had undergone a two-step hyperinsulinemic-euglycemic clamp study according to standard operating procedures and liver ¹H-MRS before any intervention. Subjects were excluded in case of a history of diabetes, use of alcohol (>2 units/day) or recreational drugs, use of antipsychotic or antidepressant medication, or any somatic disorder except for stable obesity-related conditions (i.e., NAFLD/NASH, dyslipidemia, hypertension, obstructive sleep apnea).

All subjects completed a medical evaluation including history, physical examination, and blood tests. Body composition was determined by bioelectrical impedance (Maltron BF-906). Resting energy expenditure was assessed by indirect calorimetry (V_{max} Encore 29n; CareFusion).

All procedures were approved by the Academic Medical Center medical ethics committee, and all subjects provided written informed consent in accordance with the Declaration of Helsinki.

Liver Fat Content

We assessed IHTG content by ¹H-MRS as described (van der Valk et al., 2014). This method has high diagnostic accuracy and high precision with low variability for assessment of hepatic steatosis in NAFLD (Dulai et al., 2016).

Hyperinsulinemic-Euglycemic Clamps

Glucose kinetics and tissue-specific parameters of insulin sensitivity were assessed during a two-step hyperinsulinemic-euglycemic clamp study as described (ter Horst et al., 2015). Briefly, subjects were admitted to the metabolic unit after an overnight fast. A primed continuous infusion of the stable isotope-labeled glucose tracer [6,6-2H2]glucose (>99% enriched; Cambridge Isotopes) was started. In a subset of subjects (n = 29), we also infused the stable isotope-labeled glycerol tracer [1,1,2,3,3-2H5]glycerol (>99% enriched; Cambridge Isotopes). Basal rates of EGP and lipolysis were determined after 2 hr of tracer equilibration. Next, hepatic insulin sensitivity (expressed as suppression of basal EGP by insulin) and adipose tissue insulin sensitivity (expressed as suppression of circulating FA levels or suppression of glycerol Ra by insulin) were assessed after 2 hr of low-dose insulin infusion (Actrapid 20 mU · m⁻² · min⁻¹; Novo Nordisk Farma). Finally, peripheral insulin sensitivity (expressed as insulin-stimulated Rd of glucose) was assessed after 2 hr of high-dose insulin infusion (60 mU · m⁻² · min⁻¹). During hyperinsulinemia, plasma glucose was maintained at 5.0 mmol/L by frequent bedside monitoring and variable infusion of exogenous glucose (enriched with 1% [6,6-2H2] glucose).

Liver Biopsies

A subset of subjects (n = 29) underwent bariatric surgery shortly (<2 weeks) after the clinical assessments. Subjects were instructed to maintain stable weight by consumption of a weight-maintenance diet in the preoperative period. During surgery, an experienced surgeon obtained a wedge biopsy from the right liver lobe. Biopsies were immediately snap-frozen in liquid nitrogen and stored at -80° C.

Plasma Hormones and Metabolites

Glucose, insulin, glucagon, cortisol, and FA were measured as described (ter Horst et al., 2015). Enrichments of $[6,6^2H_2]$ glucose and $[1,1,2,3,3^{-2}H_5]$ glycerol (tracer-to-tracee ratios) were measured by gas chromatography-MS (Ackermans et al., 2001).

Hepatic Lipid Metabolites

Extraction, purification, and assessment of DAGs and ceramides from liver was performed by LC-MS/MS in a blinded fashion as described (Kumashiro et al., 2011). Subcellular fractionation into membrane and cytosolic compartments was performed as reported (Cantley et al., 2013).

PKC_E Translocation Assay

Membrane translocation of PKC ε was assayed as described (Kumashiro et al., 2011) and expressed as the ratio of membrane protein band density over cytosol protein band density.

Calculations

Fluxes (EGP, glucose Rd, and glycerol Ra) were calculated using modified versions of the Steele equations for the steady state (basal EGP and glycerol Ra) or non-steady state (during insulin infusion) and expressed as μ mol \cdot (kg body weight)⁻¹ \cdot min⁻¹ (Steele, 1959).

Statistical Analyses

Groups were compared by two-tailed t test (two groups) or one-way ANOVA with Bonferroni correction (more than two groups). Correlations between continuous variables were evaluated by Pearson's correlation and regression analyses (linear and nonlinear). Findings were considered significant if p < 0.05. Statistical analyses were performed using IBM SPSS Statistics v.23 and GraphPad Prism v.6 (La Jolla).

SUPPLEMENTAL INFORMATION

Supplemental Information includes three figures and four tables and can be found with this article online at http://dx.doi.org/10.1016/j.celrep.2017.05.035.

AUTHOR CONTRIBUTIONS

K.W.t.H. researched data, performed the analysis, contributed to discussions about the results, and wrote the manuscript. P.W.G. and R.I.V. researched data. M.T.A. and D.Z. were responsible for laboratory analyses. A.J.N. was responsible for ¹H-MRS experiments. S.E.I.F., J.A.R., M.N., D.F.V., V.T.S., K.F.P., G.I.S., and M.J.S. contributed to discussions about the results. All authors critically reviewed and approved the final manuscript.

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Supplemental Information

Hepatic Diacylglycerol-Associated Protein

Kinase C^{ϵ} Translocation Links Hepatic

Steatosis to Hepatic Insulin Resistance in Humans

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	No steatosis	Mild steatosis	Severe steatosis	p ^a
	(n = 52)	(n = 41)	(n = 40)	
Basal (fasting)				
Insulin (pmol/l)	84 ± 5	131 ± 9**	145 ± 16***	< 0.001
Glucagon (ng/l)	94 ± 5	98 ± 6	81 ± 5	0.129
Cortisol (nmol/l)	228 ± 13	214 ± 14	213 ± 22	0.754
FA (mmol/l)	0.66 ± 0.03	0.65 ± 0.03	0.63 ± 0.03	0.702
Low-dose insulin infusion (step 1)				
Insulin (pmol/l)	301 ± 13	343 ± 18	326 ± 24	0.249
Glucagon (ng/l)	78 ± 4	80 ± 5	76 ± 4	0.858
Cortisol (nmol/l)	271 ± 16	252 ± 20	259 ± 15	0.562
FA (mmol/l)	0.11 ± 0.01	0.16 ± 0.01*	0.17 ± 0.01**	0.004
High-dose insulin infusion (step 2)				
Insulin (pmol/l)	765 ± 30	811 ± 47	777 ± 35	0.664
Glucagon (ng/l)	67 ± 4	70 ± 6	65 ± 4	0.778
Cortisol (nmol/l)	226 ± 16	218 ± 20	213 ± 15	0.842
FA (mmol/l)	0.03 ± 0.00	0.04 ± 0.00	0.05 ± 0.00*	0.009

Table S1. Hormones and Metabolites during Hyperinsulinemic-Euglycemic Clamp Studies. Related to Figures 1 and 2.

Data are mean ± SEM. No, mild and severe hepatic steatosis were defined as IHTG < 5.56%, 5.56

- 15%, or > 15%, respectively. Fatty acids (FA), intrahepatic triglycerides (IHTG).

^a Groups were compared by 1-way ANOVA with Bonferroni correction for multiple testing.

* p < 0.05 vs no steatosis. ** p < 0.01 vs no steatosis. *** p < 0.001 vs no steatosis.



Figure S1. Additional Metabolic Flux Data. Related to Figure 1.

(A) The Hepatic Insulin Sensitivity Index was reduced in subjects with hepatic steatosis, but did not differ between subjects with mild or severe steatosis. The index was calculated as 100 / (fasting plasma insulin x basal endogenous glucose production).

(B) Expressing peripheral insulin sensitivity as percentage increase relative to basal glucose uptake yielded similar results.

(C) Expressing peripheral insulin sensitivity relative to lean body mass yielded similar results.

(D) Insulin clearance during low-dose insulin infusion (step 1 of the clamp) did not differ between subjects with varying extent of steatosis. Insulin clearance was calculated as insulin infusion rate / plasma insulin concentration during the clamp.

(E) Insulin clearance during high-dose insulin infusion (step 2) did not differ between subjects with varying extent of steatosis.

Data are mean ± SEM (n = 133). Rate of disappearance (Rd), fat-free mass (FFM).

** p < 0.01 vs no steatosis. *** p < 0.001 vs no steatosis.





(A) Hepatic ceramide content and insulin suppression of EGP.

(B) Total hepatic DAG content and insulin suppression of EGP.

(C) Hepatic DAG content in the cytosolic fraction and insulin suppression of EGP.

(D) Hepatic DAG content in the membrane fraction and insulin suppression of EGP.

Lines were fitted by least squares method (n = 25-29). Diacylglycerol (DAG), endogenous glucose production (EGP).

	Cytosolic fraction		Membrane fraction			
Species	r ^a	р	Abundance	r ^a	р	Abundance
			[mean ± SEM (nmol/g)]			[mean ± SEM (nmol/g)]
C18:1-C16:0	-0.44	0.019	222.6 ± 33.9	0.01	0.948	215.2 ± 14.6
C20:4-C20:5	-0.12	0.551	17.1 ± 2.3	0.41	0.028	84.7 ± 6.9
C16:0-C20:4	-0.36	0.061	0.7 ± 0.1	0.39	0.041	1.6 ± 0.1
C18:0-C20:4	-0.27	0.172	0.6 ± 0.1	0.41	0.032	2.0 ± 0.1
C18:0-C18:0	-0.47	0.012	15.2 ± 2.1	-0.12	0.559	15.3 ± 0.9
C18:2-C18:0	-0.33	0.083	7.9 ± 1.2	0.34	0.078	12.9 ± 0.8
C18:1-C18:0	-0.42	0.027	27.6 ± 4.2	0.06	0.760	32.8 ± 2.4
C16:0-C16:0	-0.47	0.013	73.5 ± 12.0	-0.21	0.296	71.0 ± 4.5
C18:0-C16:0	-0.45	0.015	23.5 ± 3.5	-0.10	0.630	20.6 ± 1.3
C18:1-C18:1	-0.41	0.030	326.4 ± 51.2	0.17	0.377	377.4 ± 34.2
C18:1-C18:2	-0.33	0.091	59.6 ± 9.5	0.34	0.076	65.8 ± 6.8
C18:2-C18:2	-0.23	0.233	9.8 ± 1.6	0.38	0.044	10.4 ± 1.1
C16:0-C18:2	-0.33	0.085	79.0 ± 12.4	0.29	0.130	100.9 ± 5.9

Table S2. Correlations Between Individual Hepatic DAG Species and Insulin Sensitivity of EGP Suppression (n = 29). Related to Figure 3.

Highly abundant species that were also related to hepatic insulin sensitivity are highlighted. Diacylglycerol (DAG), endogenous glucose production (EGP).

^a Correlations were evaluated by Pearson's correlation coefficient.

Species	r ^a	р	Abundance
			[mean ± SEM (nmol/g)]
C16:0	-0.27	0.187	58.8 ± 2.1
C18:0	-0.24	0.254	16.6 ± 0.6
C20:0	-0.28	0.180	43.8 ± 6.6
C22:0	-0.27	0.190	65.1 ± 2.4
C24:1	-0.16	0.447	40.9 ± 1.2
C24:0	-0.06	0.779	52.4 ± 2.1

Table S3. Correlations Between Individual Hepatic Ceramide Species and Insulin Sensitivity of EGP Suppression (n = 25). Related to Figure 3.

Endogenous glucose production (EGP).

^a Correlations were evaluated by Pearson's correlation coefficient.

	Sensitive	Resistant	p ^a
	(n = 10)	(n = 8)	
Female sex	6 (60)	3 (38)	0.343
Age (years)	51 ± 2	45 ± 4	0.135
Body mass index (kg/m ²)	41 ± 1	47 ± 3	0.055
Body fat content (%)	45 ± 1	47 ± 3	0.603
Fasting glucose (mmol/l)	4.8 ± 0.1	5.5 ± 0.3	0.013
Fasting insulin (pmol/l)	115 ± 20	183 ± 20	0.029
Basal EGP (µmol·kg⁻¹·min⁻¹)	6.9 ± 0.3	7.2 ± 0.3	0.518
Insulin suppression of EGP (%)	87 ± 2	56 ± 1	< 0.001
IHTG content (%)	9.2 ± 3.0	14.2 ± 2.6	0.209

 Table S4. Characteristics of Subjects with Normal Hepatic Insulin Sensitivity or with Hepatic

 Insulin Resistance, Who Had Liver Biopsies Taken during Bariatric Surgery. Related to Figure 3.

Data are count (%) or mean \pm SEM. Hepatic insulin sensitivity and resistance were defined as insulin suppression of EGP > 80% and < 65%, respectively. Endogenous glucose production (EGP), intrahepatic triglyceride (IHTG).

^a Groups were compared by 2-tailed independent samples t test.



Figure S3. Release of FA from Lipolysis during Hyperinsulinemic Conditions Predicts Hepatic DAG Accumulation. Related to Figure 3.

(A) The rate of lipolysis during hyperinsulinemic-euglycemic clamps was assessed by tracerdilution of $[1,1,2,3,3-^{2}H_{5}]$ glycerol.

(B) Plasma FA concentrations during hyperinsulinemia.

Data were fitted by least squares method (n = 28). Diacylglycerol (DAG), fatty acids (FA).