



Published in final edited form as:

Cell. 2016 November 03; 167(4): 1052–1066.e18. doi:10.1016/j.cell.2016.10.015.

## Inflammation Improves Glucose Homeostasis Through IKK $\beta$ -XBP1s Interaction

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### SUMMARY

It is widely believed that inflammation associated with obesity has an important role in the development of type 2 diabetes. I $\kappa$ B kinase beta (IKK $\beta$ ) is a crucial kinase that responds to inflammatory stimuli such as Tumor Necrosis Factor  $\alpha$  (TNF $\alpha$ ), by initiating a variety of intracellular signaling cascades, and is considered to be a key element in the inflammation-mediated development of insulin resistance. We show here, contrary to expectation, that IKK $\beta$ -mediated inflammation is a positive regulator of hepatic glucose homeostasis. IKK $\beta$  phosphorylates the spliced form of X-Box Binding Protein 1 (XBP1s) and increases the activity of XBP1s. We have used three experimental approaches to enhance the IKK $\beta$  activity in the liver of obese mice, and observed increased XBP1s activity, reduced ER stress, and a significant improvement in insulin sensitivity and consequently in glucose homeostasis. Our results reveal a beneficial role of IKK $\beta$ -mediated hepatic inflammation in glucose homeostasis.

### ETOC blurb

Inflammatory signaling via IKK $\beta$  in the liver is beneficial for glucose homeostasis, running counter to the prevailing view that inflammation caused by obesity leads to insulin resistance

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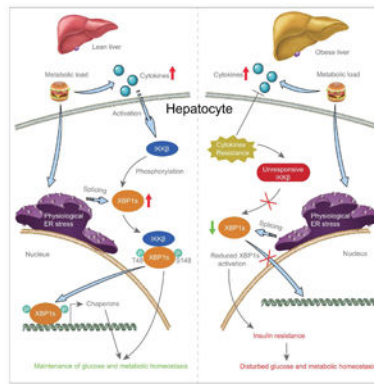
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### AUTHOR CONTRIBUTION

UO came up with the idea to investigate role of IKK $\beta$  in regulation of ER stress and XBP1s activity. UO and JL designed the experiments. JL has performed most of the experiments with help from DI, JL, HH, BA, PM, MASH, MFU and SWP. The LIKK $\beta$ <sup>Tg(+/+)</sup> mouse model was created by KT and MK. Data were analyzed by UO, MK and JL. The manuscript was written by UO, JL, SWP and MK.

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## INTRODUCTION

According to projections of the World Health Organization, type 2 diabetes will affect 300 million people or more by the year 2025 (Wild et al., 2004; Xu et al., 2013). The total cost of diabetes and its complications to the global economy in 2010 was estimated at around \$376 billion; this is expected to increase to \$490 billion in 2030 (Hu, 2011). Despite enormous research efforts in the study of obesity and diabetes, the molecular mechanisms that lead to insulin resistance and type 2 diabetes associated with obesity is not fully understood.

A leading notion in the fields of obesity and type 2 diabetes is that obesity leads to increased inflammation in a variety of metabolically active tissues locally, initiates inflammatory stimuli such as Tumor Necrosis Factor  $\alpha$  (TNF $\alpha$ ) and other cytokines either locally or systemically, and that these stimuli in turn lead to the development of insulin resistance and type 2 diabetes (Gregor and Hotamisligil, 2011; Olefsky and Glass, 2010; Shoelson et al., 2006). To date no pharmacologic treatment has emerged that is based on blocking inflammation or the activity of inflammatory molecules in obesity.

Certain conditions, such as accumulation of misfolded proteins in the endoplasmic reticulum (ER) lumen, and uncontrolled nutrient and energy homeostasis, perturb ER homeostasis and lead to a condition defined as ER stress (Lee and Ozcan, 2014; Park and Ozcan, 2013; Walter and Ron, 2011). ER stress is marked by activation of a complex intracellular signaling network that is referred to as the unfolded protein response (UPR) (Gardner et al., 2013; Lee and Ozcan, 2014; Park and Ozcan, 2013; Walter and Ron, 2011).

IRE1, which is one of the main UPR signaling molecules, is activated by autophosphorylation, and it cleaves the full-length mRNA of the X-box binding protein 1 (XBP1), leading to the translation of the spliced form of XBP1 (XBP1s), a master regulator for ER homeostasis. Nuclear translocation and activity of XBP1s are substantially reduced in the liver of obese versus lean mice (Park et al., 2010). Decreased XBP1s activity plays a key role in the development of insulin resistance and type 2 diabetes in obese mice (Lee et al., 2011; Park et al., 2010; Zhou et al., 2011), and that reinstatement of XBP1s activity in the liver of obese and diabetic mice greatly increases insulin sensitivity, reduces blood glucose levels, and restores glucose homeostasis (Deng et al., 2013; Zhou et al., 2011).

Increased inflammation in obesity is also believed to contribute to the development of ER stress (Gregor and Hotamisligil, 2011; Zhang et al., 2008). However, we recently showed that the inflammatory signaling molecule p38 mitogen-activated protein kinase (p38 MAPK) - which is also activated by TNF $\alpha$  - in fact increases the activity of XBP1s and reduces ER stress; also that activation of p38 MAPK in the liver of obese and diabetic mice significantly enhances insulin sensitivity and improves glucose homeostasis.

I $\kappa$ B kinase beta (IKK $\beta$ ) is one of the most important kinases that mediates the effects of general inflammatory stimuli inside the cell, and is a major downstream regulator of TNF signaling (Hacker and Karin, 2006; O'Dea and Hoffmann, 2009; Senftleben and Karin, 2002). TNF activates p38 MAPK and IKK $\beta$  simultaneously (Hacker and Karin, 2006; Karin, 2005b; O'Dea and Hoffmann, 2009). IKK $\beta$  was previously believed to cause ER stress. However, recent unexpected findings on the positive effects of TNF-p38 MAPK-XBP1s axis signaling on ER stress (Lee et al., 2011) led us to investigate whether IKK $\beta$  has any effect on XBP1s, and thus on ER stress signaling.

## RESULTS

### IKK $\beta$ Upregulates XBP1s Activity by Increasing Its Stability

To investigate whether IKK $\beta$  influences the activity of XBP1s, we co-expressed IKK $\beta$  and XBP1s in human embryonic kidney 293 (HEK293) cells by infecting the cells with an XBP1s-expressing adenovirus (Ad-XBP1s) and different doses of an adenovirus that expresses constitutively active IKK $\beta$  (Ad-ca-IKK $\beta$ ). Overexpression of ca-IKK $\beta$  increased protein levels of XBP1s in a dose dependent manner (Figure 1A). Total IKK $\beta$  levels were up-regulated in response to Ad-ca-IKK $\beta$  infection (Figure 1A). Consequently, the nuclear levels of p65 (Figure S1A) and NF- $\kappa$ B target genes were also significantly up-regulated (Figures S1B).

We also performed the same experiment in four other cell lines, which all yielded similar results (Figures S1C-F). Co-expression of ca-IKK $\beta$  with human XBP1s (hXBP1s) has also increased the protein levels of hXBP1s (Figure S1G).

Next, we infected HEK293 cells with Ad-XBP1s alone or with Ad-ca-IKK $\beta$ , and analyzed the levels of endogenous XBP1s mRNA. IKK $\beta$  did not enhance endogenous XBP1s mRNA levels (Figure 1B). Analysis of mRNA levels of both exogenously and endogenously expressed XBP1s in other cells lines also did not show any alterations (Figure S1H and S1I). These results provided support that IKK $\beta$  increased protein levels of XBP1s without effecting the mRNA levels and thus without creating ER stress. Activating transcription factor-6 (ATF6) is another UPR transcription factor and cleaved form (ATF6n) is up-regulated in response to ER stress to re-establish ER homeostasis (Lee and Ozcan, 2014; Park and Ozcan, 2013). We next co-expressed ca-IKK $\beta$  with ATF6n. Unlike XBP1s, IKK $\beta$  did not alter ATF6n levels (Figure S1J). These results indicate that IKK $\beta$  has specific activity on XBP1s.

We also infected HEK293 cells with Ad-XBP1s alone, or with Ad-XBP1s plus Ad-ca-IKK $\beta$ , and analyzed level of XBP1s target genes and nuclear translocation of XBP1s. XBP1s target

gene expression, including homocysteine-inducible ER stress protein (*HERPUDI*), heat shock protein 5 (*HSPA5*), and protein disulphide isomerase family A member 3 (*PDIA3*) was significantly increased at the presence of XBP1s (Figure 1C), which is further increased by ca-IKK $\beta$  co-expression (Figure 1C). IKK $\beta$ -expression also increased nuclear translocation of XBP1s (Figure 1D).

Next we sought to investigate whether other isoforms of IKK, IKK $\alpha$  and IKK $\gamma$ , also has effects on XBP1s' activity. Similar to IKK $\beta$ , IKK $\alpha$  also increased the XBP1s protein levels (Figure S1K). However, IKK $\gamma$  did not lead to an increase in XBP1s protein levels (Figure S1L). Considering the major reported roles of IKK $\beta$  in regulation of metabolic homeostasis, we focused our research on further investigating the IKK $\beta$ .

Given that IKK $\beta$  increases the level of XBP1s protein without altering its mRNA levels (Figures 1A and 1B), we examined whether IKK $\beta$  affects the stability of the XBP1s protein. We first infected HEK293 cells with Ad-XBP1s alone, or with Ad-XBP1s plus Ad-ca-IKK $\beta$ , and then treated the cells with cycloheximide to inhibit global translation. In the absence of IKK $\beta$  expression, XBP1s started to degrade within 5 minutes of treatment, and was undetectable 20 minutes after adding the cycloheximide (Figure 1E); however, this degradation rate was significantly reduced when IKK $\beta$  and XBP1s were expressed together (Figure 1E). Densitometric analysis of western blot results from three independent experiments (Figure 1E, S1M and S1N) confirmed that XBP1s degradation was significantly inhibited by IKK $\beta$  (Figure 1F). The half-life of XBP1s was around 10 minutes in the absence of IKK $\beta$ ; but when IKK $\beta$  was co-expressed with XBP1s, the half-life of XBP1s was prolonged to 20 minutes (Figure 1F). These results indicate that IKK $\beta$  increases the amount of XBP1s protein by enhancing its stability. Considering that XBP1s is recycled in a proteasome-dependent manner (Zhang et al., 2014), we also tested whether IKK $\beta$  affects the level of ubiquitination of XBP1s. A myc-tagged ubiquitin (Ub-myc)-expressing vector was transfected into cells with a flag-tagged XBP1s (XBP1s-flag), in the presence or absence of an HA-tagged constitutively active IKK $\beta$  (IKK $\beta$ -HA). We found that co-expression of IKK $\beta$  and XBP1s robustly reduced the ubiquitination of XBP1s, without affecting global ubiquitination levels in the cell (Figure 1G).

To investigate whether p65 signaling is involved in IKK $\beta$ -mediated XBP1s stabilization, we used a highly efficient p65 siRNA. (Figure 1H) and co-transfected XBP1s and IKK $\beta$  in p65 siRNA transfected cells; depletion of p65 did not block IKK $\beta$ -mediated upregulation of XBP1s protein levels (Figure 1H).

### IKK $\beta$ Interacts with and Phosphorylates XBP1s

The increased protein level of XBP1s caused by IKK $\beta$  prompted us to hypothesize that IKK $\beta$  stabilizes XBP1s by direct modification, through a physical interaction. To test this hypothesis, we transfected cells ca-IKK $\beta$ -HA and XBP1s-flag expressing vectors and immunoprecipitated XBP1s from cell lysates; IKK $\beta$  was indeed co-immunoprecipitated with XBP1s (Figure 2A), which indicates that XBP1s and IKK $\beta$  interact. We also immunoprecipitated IKK $\beta$  and examined the precipitates for the presence of XBP1s. XBP1s was also co-immunoprecipitated with IKK $\beta$  (Figure 2B).

We next investigated whether the interaction between XBP1s and IKK $\beta$  that we observed *in vitro*, exists between endogenous XBP1s and endogenous IKK $\beta$  *in vivo*. We have previously reported that re-feeding after fasting greatly increases XBP1s protein levels in the liver of lean mice (Park et al., 2010). To investigate whether endogenous XBP1s interacts with IKK $\beta$  during normal physiological processes, lean healthy mice were fasted for 24 hours, then re-fed for one hour, at which point liver lysates were processed for IKK $\beta$  immunoprecipitation; we found that IKK $\beta$  immunoprecipitation pulled down XBP1s (Figure 2C), which confirmed that endogenous XBP1s and endogenous IKK $\beta$  interact in the liver.

To determine whether IKK $\beta$ 's kinase activity is responsible for the increased stability and consequent enhanced protein levels of XBP1s, HEK293 cells were transfected with XBP1s-flag in the presence or absence of ca-IKK $\beta$ -HA; the cells were then treated with [(aminocarbonyl)amino]-5-(4-fluorophenyl)-3-thiophenecarboxamide (TPCA-1), an IKK $\beta$  inhibitor. Inhibition of IKK $\beta$ 's kinase activity greatly blocked the IKK $\beta$ -mediated increase in total protein levels of XBP1s (Figure 2D). To also investigate whether endogenous IKK $\beta$  can also affect XBP1s levels, we expressed XBP1s in cells and then treated with vehicle or with TNF $\alpha$  at the absence or presence of TPCA-1. TNF $\alpha$  stimulation increased XBP1s protein levels. However, pre-treatment with IKK $\beta$  inhibitor TPCA-1 blocked TNF $\alpha$ -induced up-regulation of XBP1s (Figure 2E). Next, we sought to investigate whether the IKK $\beta$ 's effect on XBP1s is mediated through its kinase activity. For this purpose, we compared the effects of ca-IKK $\beta$  and a mutant IKK $\beta$  that lacks kinase activity (kd-IKK $\beta$ ). Expression of ca-IKK $\beta$  greatly upregulated XBP1s protein levels. However, kd-IKK $\beta$ , despite expression of more protein compared to ca-IKK $\beta$ , did not have any effect on XBP1s (Figure 2F). Furthermore, we also stimulated the cells with IL-1 $\beta$ , another cytokine that have ability to increase IKK activity. As shown in Figure S10, stimulation of cells, which are transfected with XBP1s, led to a robust increase in XBP1s protein levels and IKK $\beta$  phosphorylation.

These findings then led us to ask whether IKK $\beta$  phosphorylates XBP1s: we transfected HEK293 cells with an XBP1s-flag in the presence or absence of IKK $\beta$ -HA, and immunoprecipitated XBP1s from the cell lysates. Use of a general anti-phosphoserine/threonine antibody revealed a robust, IKK $\beta$ -mediated increase in levels of XBP1s phosphorylation (Figure 2G).

Together, the above results indicate that elevated IKK $\beta$  activity enhances Ser/Thr phosphorylation of XBP1s. However to address whether IKK $\beta$  has a direct physical interaction with XBP1s, and whether it directly phosphorylates XBP1s, we transfected HEK293 cells with a flag-ca-IKK $\beta$ , then immunoprecipitated the ca-IKK $\beta$  protein from the cell lysates, and performed an *in vitro* IKK $\beta$  kinase assay, using a His-TF-XBP1s fusion protein as the substrate; an IKK $\beta$  substrate and TPCA-1 (20  $\mu$ g ml<sup>-1</sup>) were used as positive and negative controls, respectively. Immunoprecipitated IKK $\beta$  successfully phosphorylated its substrate in *in vitro* setting (**line 5**, Figure 2H); but when TPCA-1 was present during the assay, the phosphorylation was blocked (**line 6**, Figure 2H). Analysis of XBP1s phosphorylation confirmed that XBP1s was robustly phosphorylated by IKK $\beta$  (**line 2**, Figure 2H) and that this phosphorylation was inhibited by TPCA-1 (**line 3**, Figure 2H).

To determine the sites on XBP1s that are phosphorylated by IKK $\beta$ , we performed the same *in vitro* kinase assay as described above with His-TF-XBP1s protein, in the presence and absence of IKK $\beta$ , then immunoprecipitated XBP1s from the reaction buffer. Subsequently, possible phosphorylation sites on XBP1s were analyzed with tandem mass spectrometry (MS/MS). Results showed that IKK $\beta$  directly phosphorylates XBP1s at Thr48 and Ser148 residues (Figures S2A and S2B).

We have previously produced an antibody that is specific for p-XBP1<sup>Thr48</sup> (Lee et al., 2011). We next infected HEK293 cells with Ad-XBP1s alone, or with Ad-XBP1s plus Ad-ca-IKK $\beta$ ; immunoblotting with p-XBP1<sup>Thr48</sup> antibody demonstrated that XBP1s<sup>Thr48</sup> phosphorylation was indeed increased by IKK $\beta$  (Figure 2I and 2J).

We also made attempts to produce an antibody against XBP1s (which is phosphorylated on the Ser<sup>148</sup> residue) but we failed. Thus, to determine the role of IKK $\beta$ -mediated phosphorylation of the Thr48 and Ser148 residues of XBP1s, we replaced Thr48 and Ser148 of XBP1s with alanines (XBP1s(T48A) and XBP1s(S148A)). We also created a double mutant XBP1s (XBP1s(DM)) bearing both substitutions. HEK293 cells were transfected with XBP1s, XBP1s<sup>T48A</sup>, XBP1s<sup>S148A</sup>, or XBP1s<sup>DM</sup> constructs, with or without a ca-IKK $\beta$ -expressing vector. The T48A or S148A single mutations substantially blocked the IKK $\beta$ -mediated up-regulation of nuclear, cytoplasmic and total I XBP1s (Figure S2C), whereas double Thr48A and Ser148A mutation greatly reduced the effects of IKK $\beta$  on XBP1s (Figure S2C).

Next, to determine whether Thr48 and Ser148 are the main sites phosphorylated by IKK $\beta$ , we transfected HEK293 cells with flag-tagged XBP1s or flag-tagged XBP1s(DM), along with an IKK $\beta$ -HA, and subsequently immunoprecipitated XBP1s. Upon probing the XBP1s immunoprecipitates with an anti-phosphoserine/threonine antibody, we observed that IKK $\beta$ -mediated phosphorylation of XBP1s was greatly diminished when the Thr48 and Ser148 residues were mutated to alanine (Figure S2D).

To assess how IKK $\beta$ -mediated phosphorylation of XBP1s affects its expression, we transfected HEK293 cells with XBP1s or XBP1s(DM), along with one that expresses IKK $\beta$ ; the cells were then treated with cycloheximide (20  $\mu\text{g ml}^{-1}$ ). The rate of degradation of XBP1s(DM) protein was significantly higher than that of XBP1s (Figures S2E and S2H). Combined densitometric analysis of three independent experiments (Figure S2E-S2G) further demonstrated that the Thr48A/Ser148A double mutation reduced the half-life of XBP1s by more than 50% (Figure S2H).

### The Interaction between IKK $\beta$ and XBP1s is Lost in Obesity

We next asked whether the interaction between IKK $\beta$  and XBP1s is altered in obesity. Wild-type (WT) and *ob/ob* mice were fasted for 24 hours, then given food *ad libitum* for 1 hour. Immunoprecipitation experiments showed that XBP1s was present in the IKK $\beta$  immunoprecipitates from the liver lysates of lean mice, in which XBP1s expression had been induced by re-feeding (Figure 3A); however, this interaction between XBP1s and IKK $\beta$  was not detected in *ob/ob* mice under the same re-feeding condition (Figure 3A). Meanwhile, as per our previous findings (Park et al., 2010), total XBP1s levels were lower in



the liver of *ob/ob* mice than in those of lean mice (Figure 3A and 3B). The total levels of IKK $\beta$  in both mice (lean and *ob/ob*) were comparable (Figure 3A). These results indicate that the interaction between XBP1s and IKK $\beta$  is reduced in the liver of *ob/ob* mice and also raised the question whether IKK $\beta$  is activated during the re-feeding. Indeed, there was a significant up-regulation in IKK $\beta$  phosphorylation in response to re-feeding in lean mice (Figure 3C and 3D and S3A). The basal phosphorylation status of IKK $\beta$  in the obese mice was higher in fed state than the lean mice. However, re-feeding did not lead to any further increase (Figure 3C and 3D and S3A). In other words, obese mice were un-responsive to the stimuli created by re-feeding in increasing IKK $\beta$  phosphorylation. An IKK $\beta$  kinase assay also documented that re-feeding significantly up-regulated the kinase activity of IKK $\beta$ , but similar to the phosphorylation status, there was no up-regulation of IKK $\beta$  kinase activity in obese mice following the re-feeding (Figure 3E). Finally we investigated whether NF- $\kappa$ B activity is also regulated by re-feeding in the lean and obese mice. In parallel to the IKK $\beta$  activity, p65 nuclear levels were significantly upregulated after re-feeding in the lean mice, but this response was blunted in the obese mice (Figure 3F and 3G). Figure 3G was created by combination of three independent experiments (Figure 3F, S3B and S3C). Finally analysis of NF- $\kappa$ B target genes following the re-feeding has shown a similar pattern namely that various target gene expression such as Chemokine (C-X-C motif) ligand 10 (*Cxcl10*), Chemokine (C-C motif) ligand 2 (*Ccl2*), Chemokine (C-C motif) ligand 4 (*Ccl4*), Chemokine (C-C motif) ligand 5 (*Ccl5*), Hepatocyte growth factor (*Hgf*) and interferon gamma (*Ifng*) were significantly up-regulated in the lean mice but there were no response in the obese mice to refeeding (Figure 3H).

Contrary to previous reports (Cai et al., 2005), the above results collectively suggest a positive involvement of IKK $\beta$  in the regulation of glucose homeostasis and normal physiological responses. We thus examined what effect up-regulated IKK $\beta$  activity has in livers of obese mice by using different experimental approaches.

### Adenovirus-Mediated Hepatic IKK $\beta$ Gain-Of-Function Improves Glucose Homeostasis

To increase IKK $\beta$  activity in liver, we injected Ad-ca-IKK $\beta$  (or Ad-LacZ, as a control) into *ob/ob* mice through tail vein. Ad-ca-IKK $\beta$  injection resulted in significant increase in total IKK $\beta$  and nuclear levels of p65 in the liver of *ob/ob* mice relative to the control group (Figures S3D and S3E).

Analysis of blood glucose of *ob/ob* mice showed that ca-IKK $\beta$  expression significantly reduced the blood glucose levels at fed and fasted states (Figures S3F and S3G). Glucose tolerance test (GTT) done on day 4 post-injection revealed a major improvement in glucose disposal rate in ca-IKK $\beta$  expressing mice (Figure S3H). Results of an insulin tolerance test (ITT), done on day 6 post injection, also showed a significant improvement in whole-body insulin sensitivity in IKK $\beta$ -expressing *ob/ob* mice, when absolute blood glucose levels were taken into account (Figure S3I). However, analysis of ITT results as percent values of the initial resting blood glucose values had shown no difference between the groups (Figure S3J), which lead us to conclude that whole body insulin sensitivity is not altered in the Ad-ca-IKK $\beta$  group. Furthermore, circulating insulin levels were significantly reduced in the Ad-ca-IKK $\beta$ -injected mice when compared to controls (Figure S3K); and plasma aspartate

transaminase (AST) and alanine transaminase (ALT) levels were also significantly reduced in Ad-ca-IKK $\beta$ -injected mice (Figures S3L and S3M). Food intake (Figures S3N) and body weight (Figures S3O) were not altered.

We next investigated whether total and nuclear levels of XBP1s are up-regulated, and consequently whether ER stress was reduced, when IKK $\beta$  activity was increased. As shown in Figure S3P, total as well as nuclear levels of XBP1s were increased in the ca-IKK $\beta$ -expressing group, along with a marked reduction in PERK phosphorylation and expression of DNA damage inducible transcript 3 (*DDIT3*) (Figure S3Q). Collectively, these observations indicate that ER stress was reduced. Phosphorylation levels of c-Jun amino terminal kinase (JNK), p38 mitogen activated protein kinase (p38) and its downstream target activating transcription factor-2 (ATF2) were not altered by ca-IKK $\beta$  expression (Figure S3R).

### AAV-Mediated Expression of IKK $\beta$ in Livers of *Ob/ob* Mice Improves Glucose Metabolism

We next used adeno-associated virus (AAV), which cannot induce immune responses by itself *in vivo* (Daya and Berns, 2008; Kotterman and Schaffer, 2014; Rehman et al., 2008), to increase activation of IKK $\beta$  in the liver. AAV-ca-IKK $\beta$  injection increased both total IKK $\beta$  and nuclear p65 levels in liver of *ob/ob* mice (Figure 4A). In parallel, expression levels of IKK $\beta$  target genes (e.g., chemokine (C-X-C motif) receptor 2 (*Cxcr2*); CD14 antigen (*Cd14*); superoxide dismutase 2 (*Sod2*); CD40 antigen (*Cd40*); and nitric oxide synthase 2 (*Nos2*)) were increased when mice were injected with AAV-ca-IKK $\beta$  (Figure S4A). Expression of ca-IKK $\beta$  also up-regulated the mRNA amounts of cytokines such as interleukin 1 (*Il-1*), interleukin 4 (*Il-4*), and interleukin 6 (*Il-6*) (Figure S4B), but not TNF $\alpha$  mRNA levels in the liver (Figure S4B). IL-6 plasma levels were also significantly up-regulated (Figure S4C). The AAV-ca-IKK $\beta$  used in this study employed a liver-specific thyroxine-binding globulin (TBG) promoter. Analysis of IKK $\beta$  levels in the epididymal white adipose tissue (WAT), in brown adipose tissue (BAT), and in muscle and kidney tissues documented that the increase in IKK $\beta$  levels was confined to the liver (Figure S4D).

Once liver-specific pattern of expression of IKK $\beta$  was successfully established for AAV-ca-IKK $\beta$ , we injected the vector (or AAV-GFP as a control) into the tail vein of *ob/ob* mice. On day 5 post injection, measurement of blood glucose at fed and fasted state showed that the ca-IKK $\beta$ -expressing mice had significantly lower blood glucose levels compared to control mice (Figures 4B and 4C); in addition, circulating insulin levels in the ca-IKK $\beta$ -expressing mice were also markedly reduced (Figure 4D). GTT, performed on day 8 after injection, revealed a significant improvement in the AAV-ca-IKK $\beta$ -injected group (Figures 4E and 4F). ITT done on day 13 post injection also showed a significantly enhanced whole-body insulin sensitivity in the ca-IKK $\beta$ -expressing group (Figures 4G and 4H).

Next, we investigated whether hepatic insulin receptor signaling is altered in *ob/ob* mice as a result of ca-IKK $\beta$  expression. Insulin-stimulated IR tyrosine phosphorylation, AKT<sup>Ser473</sup>, AKT<sup>Thr308</sup>, and GSK3 $\beta$  phosphorylations were significantly increased in AAV-ca-IKK $\beta$ -injected *ob/ob* mice when compared with the GFP-expressing control mice (Figure 5A-E). These results indicate that increased hepatic IKK $\beta$  activity improves IR signaling.



Increased activity of IKK $\beta$  also led to substantial upregulation of XBP1s protein amounts, and increased XBP1s<sup>Thr48</sup> phosphorylation (Figures 5F and 5G). Expression of XBP1s target genes - such as DnaJ homolog, subfamily B, member 9 (*Dnajb9*); heat shock protein 5 (*Hspa5*); homocysteine-inducible, ER stress-inducible, ubiquitin-like domain member 1 (*Herpud1*); protein disulfide isomerase associated 3 (*Pdia3*) - were also significantly increased in the IKK $\beta$ -expressing group, pointing to increased XBP1s activity (Figure 5H). Finally, increased IKK $\beta$  activity reduced PERK<sup>Thr980</sup> phosphorylation (Figure 5I), expression of *Ddit3* (Figure 5J) and activating transcription factor-4 (ATF4) protein levels (Figure S4E). Phosphorylation of JNK, P38 and ATF2 were not altered (Figure S4E). Additionally, ALT and AST levels were significantly reduced (Figures 5K and 5L). No differences in food intake, bodyweight and body temperature were detected (Figure S4F-S4H).

Five weeks post injection, no necrosis or liver damage was detected in AAV-GFP- or AAV-ca-IKK $\beta$ -injected mice (Figure S4I). Immunostaining for active caspase-3 also did not reveal increased apoptosis in the liver of AAV-ca-IKK $\beta$ -injected mice (Figure S4I). And Trichrome staining (to evaluate the presence of any fibrosis caused by IKK $\beta$  activity) revealed no difference between the groups (Figure S4I).

### AAV-Mediated Hepatic IKK $\beta$ Expression is Beneficial in DIO Mice

We next examined the effects of hepatic overexpression of IKK $\beta$  in DIO mice. As with results obtained from the *ob/ob* models, DIO mice injected with AAV-ca-IKK $\beta$  displayed significantly lower blood glucose and insulin levels compared to the control group (Figure 6A and 6B). GTT, performed on post-injection day 19, revealed a significant improvement in the IKK $\beta$ -expressing group (Figures 6C and 6D). ITT, performed on post-injection day 21, showed that whole-body insulin sensitivity was also significantly enhanced in the IKK $\beta$  group (Figures 6E and 6F). As shown in Figure 6G, analysis of blood glucose levels 10 weeks after the injections still showed a significant decline in the IKK $\beta$ -expressing group. GTT was performed again eight weeks after the injection and it showed improved glucose tolerance in the IKK $\beta$ -expressing group (Figures 6H and 6I).

At the time of sacrifice, IKK $\beta$  and XBP1s levels were analyzed. Expression of IKK $\beta$  was maintained (Figure 6J) and its target genes were significantly elevated (Figure S5A). XBP1s protein levels were also markedly upregulated, in parallel with the high expression levels of IKK $\beta$  (Figure 6J), without any changes in the corresponding mRNA levels (Figure 6K); and in addition, expression of XBP1s target genes was significantly increased (Figure 6K). AAV-mediated expression of IKK $\beta$  led to a marked decrease in PERK phosphorylation, ATF4 protein levels (Figure 6L and S5B) and a significant reduction in *Ddit3* expression levels (Figure 6M) even 14 weeks post injection.

Histological analysis showed that expression of IKK $\beta$  in the liver for 14 weeks almost completely abolished hepatosteatosis (Figure 6N); and consistent with this, liver triglyceride levels were also significantly reduced (Figure 6O). q-PCR analysis of lipogenic genes (such as acetyl-Coenzyme A carboxylase alpha (*Acaca*), acetyl-Coenzyme A carboxylase beta (*Acacb*), diacylglycerol O-acyltransferase 2 (*Dgat2*) and fatty acid synthase (*Fasn*)) revealed a significantly decreased expression in the IKK $\beta$ -expressing mice when compared to

controls (Figure S5C). Furthermore, IKK $\beta$ -expression in the liver did not alter in body temperature (Figure S5D), body weight (Figure S5E) or food intake (Figure S5F). Finally, ALT and AST levels were significantly reduced in the IKK $\beta$  group, confirming a beneficial role of IKK $\beta$  in the liver (Figure S5G and S5H).

To investigate the role of IKK $\beta$  over expression on glucose homeostasis in more detail, we performed hyperinsulinemic euglycemic clamp (HEC) studies in the DIO mice. As shown in supplementary Figure S5I, hepatic glucose production was significantly reduced during the HEC procedure in the ca-IKK $\beta$  expressing group, whereas, glucose infusion rate was significantly increased (Figure S5J and S5K). Glucose uptake rate was not different between the groups (Figure S5L). Finally analysis of blood glucose levels during the whole HEC experiment documented that euglycemic state were reached in both groups (Figure S5M).

### Glucose homeostasis is improved in Liver-specific IKK $\beta$ transgenic mice

Earlier studies report that glucose homeostasis is disturbed in liver-specific constitutively active IKK $\beta$  transgenic mice (Cai et al., 2005). Based on our findings from experiments on Adenovirus and AAV-mediated constitutively active IKK $\beta$  expression, we created a transgenic mouse model in which the expression of constitutively active human IKK $\beta$ <sup>S177E/S181E</sup> was driven under the albumin promoter (LIKK $\beta$ <sup>+/+</sup>) to investigate the phenotype of liver-specific IKK $\beta$  transgenic mice. Expression of constitutively active human IKK $\beta$  was increased about 3-fold, compared to endogenous IKK $\beta$  (Figure 7A and 7B). In parallel, amounts of nuclear p65 were also markedly increased (Figure 7A) and NF- $\kappa$ B target genes, such as *Cd14*, *Cd40*, *Ccl2*, *Il-10*, *Il-1r1* and *Il-6* were up-regulated (Figure S6A).

We placed LIKK $\beta$ <sup>Tg(+/+)</sup> and control mice (LIKK $\beta$ <sup>Tg(-/-)</sup>) on a HFD and investigated the phenotype of this model for a period of 14 weeks in three independent cohorts. Analysis of blood glucose levels throughout the experimental period revealed that LIKK $\beta$ <sup>Tg(+/+)</sup> mice have significantly reduced blood glucose levels (Figure 7C); consistent with this, plasma insulin levels in the LIKK $\beta$ <sup>Tg(+/+)</sup> mice were also significantly lower than in WT mice (Figure 7D). GTTs were performed in all three independent cohorts (Figures S6B-S6D): the combined AUC analysis (Figure 7E) documented a significant improvement in the glucose tolerance of LIKK $\beta$ <sup>Tg(+/+)</sup> mice. With ITTs, when the data points were analyzed as absolute blood glucose levels measured during the experiments in each cohort (Figures S6E-S6G), two of them (Figures S6E and S6F) displayed significant differences, and one did not (Figure S6G). However, when we analyzed percent changes as a function of the starting blood glucose levels of these three independent ITTs (Figure 7F), no differences were detected, leading us to conclude that general insulin sensitivity in the whole body of LIKK $\beta$ <sup>Tg(+/+)</sup> mice is not altered.

HFD feeding led to a blunted insulin-stimulated IR tyrosine phosphorylation in the liver of LIKK $\beta$ <sup>Tg(-/-)</sup> mice, while IKK $\beta$  overexpression significantly enhanced it (Figures 7G and 7H). Analysis of downstream IR signaling, reflected by AKT<sup>Ser473</sup>, AKT<sup>Thr308</sup> and GSK3 $\beta$ <sup>Ser9</sup> phosphorylations (Figures 7G, and 7I-K), also documented significant up-regulation of insulin signaling in the LIKK $\beta$ <sup>+/+</sup> liver, when compared to the LIKK $\beta$ <sup>-/-</sup> group.

XBP1s levels in these mice were consistent with the results of adenovirus- and AAV-mediated IKK $\beta$  gain-of-function experiments, with a marked increase in the levels of XBP1s protein (Figure 7L), but no change in the corresponding mRNA levels (Figure 7M). XBP1s target genes were also significantly upregulated (Figure 7M). Analysis of PERK<sup>Thr980</sup> phosphorylation (Figure 7N) and *Ddit3* expression levels (Figure 7O) revealed significant downregulation in both, indicating a reduction in ER stress.

After 14 weeks of HFD feeding, mice were sacrificed and liver pathology was examined histologically by H&E staining. Hepatosteatosis induced by 14 weeks of HFD feeding in the LIKK $\beta$ <sup>Tg(-/-)</sup> group was completely blocked in LIKK $\beta$ <sup>Tg(+/+)</sup> mice (Figure 7P), indicating that IKK $\beta$  has strong anti-lipogenic properties in the liver. Indeed, the levels of expression of lipogenic genes such as, Acetyl-coenzyme A carboxylase alpha (*Acaca*), Acetyl-coenzyme A carboxylase beta (*Acacb*), Fatty acid synthase (*Fasn*), stearoyl-coenzyme A desaturase (*Scd1*) were dramatically reduced in the LIKK $\beta$ <sup>Tg(+/+)</sup> mice (Figure 7Q) and hepatic triglyceride levels were also significantly depressed (Figure 7R). In parallel, the ALT and AST levels were significantly lower in LIKK $\beta$ <sup>Tg(+/+)</sup> mice (Figures S6H and S6I). There were no increases in JNK, p38 MAPK and ATF2 activity (Figures S6J). Food intake (Figure S6K), body weight (Figure S6L) and body temperature (Figure S6M) were unaltered.

### Depletion of XBP1s diminishes IKK $\beta$ 's beneficial effect in the liver

To address whether IKK $\beta$ 's effect on improvement of glucose homeostasis is mainly mediated by increased XBP1s activity, we depleted XBP1s while simultaneously overexpressing ca-IKK $\beta$  in the liver. For this reason, we first generated an adenovirus that expresses XBP1 shRNA (Ad-shXBP1). To test whether Ad-shXBP1 can effectively deplete XBP1s protein and mRNA, we first infected the MEFs with Ad-shXBP1 or with an adenovirus that expresses LacZ shRNA (Ad-shLacZ) as a control. Subsequently, cells were either incubated with vehicle or tunicamycin. As shown in Figure S7A, stimulation of cells with tunicamycin in Ad-shLacZ-infected cells led to robust production of XBP1s. However, this increase was completely abolished in the Ad-shXBP1-infected cells. Next, we infected the cells with Ad-shLacZ and Ad-shXBP1 and subsequently divided each group into two subgroups, one of which was stimulated with vehicle (DMSO) or tunicamycin. In both vehicle- and tunicamycin-treated cells, Ad-shXBP1 infection led to a highly significant reduction in XBP1s mRNA levels (Figure S7B). These results indicate that the Ad-shXBP1 that we produced efficiently silences XBP1s expression.

Next we injected AAV-GFP or AAV-ca-IKK $\beta$  together with Ad-shLacZ or Ad-shXBP1 into the tail vein of obese and diabetic *db/db* mice. Three days after the injections, AAV-ca-IKK $\beta$  + Ad-shLacZ-injected group had significantly lower blood glucose levels when compared with AAV-GFP- + Ad-shLacZ-injected group. However, the mice injected with AAV-ca-IKK $\beta$  + Ad-shXBP1 had significantly higher levels of blood glucose starting from the third day of injections (Figure S7C). Subsequently we have performed a GTT on post injection day six. AAV-ca-IKK $\beta$  + Ad-shLacZ- injected group had a significantly higher rate of glucose clearance from circulation when compared to the AAV-GFP+ Ad-shLacZ-injected group (Figure S7D). However, depletion of XBP1s in ca-IKK $\beta$  overexpressing conditions by injection of AAV-ca-IKK $\beta$  + Ad-shXBP1 significantly lowered the glucose clearance rate

from circulation when compared to the AAV-ca-IKK $\beta$ +Ad-shLacZ-injected group, indicating that improved glucose tolerance with ca-IKK $\beta$  overexpression significantly diminishes when the XBP1s' expression is silenced (Figure S7D and S7E). Analysis of blood glucose levels on post injection day 6 showed significant difference between the AAV-ca-IKK $\beta$  + Ad-shXBP1 and the AAV-ca-IKK $\beta$  + Ad-shLacZ groups (Figure S7F). In parallel, insulin levels were significantly reduced in the AAV-ca-IKK $\beta$  + Ad-shLacZ group and returned back to control levels in the AAV-ca-IKK $\beta$  + Ad-shXBP1 group (Figure S7G). All of these alterations happened in the absence of any change in the bodyweight or food intake of the groups (Figure S7H and S7I).

On day 10 post injections, mice were sacrificed and liver was collected for further analysis. Analysis of total XBP1s protein levels from the liver of three groups showed that ca-IKK $\beta$  overexpression increased XBP1s protein levels, and use of Ad-shXBP1 have blocked this up-regulation (Figure S7J and S7K), which confirms that XBP1shRNA has successfully depleted XBP1s protein. Ca-IKK $\beta$  overexpression mediated increase in XBP1s nuclear levels was also reduced with the expression of XBP1 shRNA (Figure S7L and S7M). Next, we analyzed PERK<sup>Thr980</sup> phosphorylation. In parallel to the results obtained from previous experiments ca-IKK $\beta$  overexpression reduced PERK<sup>Thr980</sup> phosphorylation (Figure S7N and S7O). However, depleting XBP1s in this setting significantly increased PERK phosphorylation, indicating that reduced ER stress after ca-IKK $\beta$  overexpression is mainly mediated by up-regulation of XBP1s protein levels (Figure S7N and S7O).

## DISCUSSION

IKK $\beta$  is one of the most highly studied kinases in the intracellular inflammatory pathways (Hacker and Karin, 2006; Karin, 2005a; O'Dea and Hoffmann, 2009; Senftleben and Karin, 2002), and is regarded as a major intracellular mediator of inflammation-related hepatic insulin resistance.

In this report, we have uncovered an unexpected signaling network between IKK $\beta$  and XBP1s, and documented that IKK $\beta$  increases the stability and activity of XBP1s both in vitro and in vivo. Further analysis has shown that hepatic IKK $\beta$  is phosphorylated and activated at postprandial states and re-feeding induced increase in IKK $\beta$  activity leads to its binding to XBP1s. It is interesting to note that basal phosphorylation levels of hepatic IKK $\beta$  at fasted states are high, but it does not respond to the stimuli created by re-feeding in obesity. The majority of signaling systems work by ligand-dependent increase in the activity of the elements of those signaling systems. For example, in obese and insulin resistant conditions, *basal* phosphorylation of IRS proteins or down stream signaling molecules such as AKT are high compared to the lean states. However, insulin-stimulated up-regulation of IRS activity at postprandial conditions is severely suppressed in obese insulin resistant states and does not respond to stimuli, and therefore cannot activate downstream signaling. What we see in the case of IKK $\beta$  is very similar to the phenomenon observed in insulin resistant conditions for the insulin receptor signaling molecules.

Based on the initial biochemical findings, we used three gain-of-function approaches to explore whether IKK $\beta$  could have beneficial effects on hepatic glucose homeostasis. These

three approaches yielded similar results, and established with certainty that IKK $\beta$  increases XBP1s activity, reduces ER stress, enhances hepatic IR signaling, and improves glucose homeostasis, blocks the development of hepatosteatosis or reduces already existing fatty liver disease.

An earlier study used an animal model similar to the LIKK $\beta$ <sup>Tg(+/+)</sup> mice reported that glucose homeostasis was disturbed in these mice. It is not uncommon to see different phenotypes from the same genetic animal models in different animal facilities of different institutes, or even on different backgrounds within the same animal facility. However, the findings we present here are based on three different experimental approaches, and the results from all three are consistent showing that hepatic IKK $\beta$  activity is beneficial for maintaining glucose homeostasis.

One question remains to be answered is that why both deletion (Arkan et al., 2005) and re-activation of IKK $\beta$  in the liver improves glucose homeostasis. One possible explanation is that embryonic deletion of IKK $\beta$  in the liver could elicit a compensatory response through hyper-activation of IKK $\alpha$  and creating the outcome observed in liver specific IKK $\beta$  knockout mice. Indeed, we have shown that IKK $\alpha$  has also a strong activity towards XBP1s.

For the last two decades, research on obesity and type 2 diabetes has viewed increased local or systemic inflammation as a major contributor to the development of type 2 diabetes in obesity (Gregor and Hotamisligil, 2011; Hotamisligil, 2006; Olefsky and Glass, 2010; Shoelson et al., 2006). The proposed role for inflammation in type 2 diabetes has shifted from involvement of systemic inflammation to local inflammation, sub-acute inflammation to chronic inflammation, and finally to a condition called metaflammation (Hotamisligil, 2006). Emerging evidence points to a contrasting, or at least much more complicated role, for inflammation in type 2 diabetes. Increased inflammatory stimuli are believed to block insulin receptor signaling through inhibition of insulin receptor substrates (IRS), by increasing their serine phosphorylation (Hotamisligil et al., 1996). In contrast, Copps et al. have shown that the main phosphorylation site (serine 307) in the IRS1 protein (that is responsive to inflammatory stimuli) (Copps et al., 2010) is a beneficial phosphorylation site *in vivo*; also, that phosphorylation of this site enhances insulin action, rather than blocking it (Copps et al., 2010). Several groups also report that interleukin 6 (IL-6), a cytokine that mediates inflammation, is beneficial for glucose homeostasis in obesity (Awazawa et al., 2011; Sadagurski et al., 2010), and that depletion of IL1R1 in these mice leads to a more severe obesity phenotype and glucose intolerance (Garcia et al., 2006; McGillicuddy et al., 2013). Consistent with these results, local inactivation of the main inflammatory cytokine TNF in WAT, reportedly *worsens* glucose homeostasis (Wernstedt Asterholm et al., 2014); this landmark study directly contradicts and challenges the central hypothesis (Hirosumi et al., 2002; Hotamisligil et al., 1993) that increased local TNF-mediated inflammation in WAT creates glucose intolerance and insulin resistance. Activation of IKK $\beta$  in adipose tissue also improves glucose homeostasis (Jiao et al., 2012). In addition, a recent study (Lancaster et al., 2016) has shown that double-stranded RNA-dependent protein kinase (PKR), which was claimed as a critical mediator and critical component of an inflammatory complex leading to insulin resistance (Nakamura et al., 2010) does not cause insulin resistance. Moreover, a number of anti-inflammatory approaches in human clinical trials have failed to improve

insulin sensitivity: for example, statins, which have anti-inflammatory activity, reduce the activity of AP-1 and NF- $\kappa$ B. A recent report analyzed five trials involving 32,752 patients, and found that 12 percent of these patients were more likely to have diabetes (Preiss et al., 2011). Another study employed TNF $\alpha$  antagonism (e.g., a TNF $\alpha$ -specific antibody, or Fc-fusion TNF $\alpha$  receptors) to treat type 2 diabetes: these trials also failed to improve patients' insulin sensitivity (Gao and Ye, 2012).

The convergence of the p38 MAPK (Lee et al., 2011) and IKK $\beta$  signaling pathways on XBP1s suggests that XBP1s is an important inflammatory signaling node, which has a reduced activity in obesity (Lee et al., 2011; Park et al., 2010). Given that p38 MAPK and IKK $\beta$  are mainly activated by cytokines (Hacker and Karin, 2006; Karin, 2005b; Senftleben and Karin, 2002), and that they are unresponsive to the stimuli in obesity conditions, we propose that obesity could be a condition of generalized cytokine resistance, wherein cytokines are unable to act on target tissues to activate relevant pathways to keep glucose homeostasis under control. Cytokine resistance may be the cause of up-regulated circulating cytokines in obesity.

In conclusion, our results elucidate the relationship between obesity and inflammation, and question the concept that hepatic inflammation or hepatic activation of IKK $\beta$  is detrimental for glucose homeostasis. Recent advances in our understanding of the obesity-inflammation-type 2 diabetes axis underscore the beneficial role of inflammation, and suggest that therapeutic developments for obesity and type 2 diabetes may, in fact, arise from a shift in our thinking about the mechanisms that underlie these disorders.

## STAR Methods

### CONTACT FOR REAGENT AND RESOURCE SHARING

Further information and requests for reagents may be directed to, and will be fulfilled by the corresponding author Umut Ozcan (umut.ozcan@childrens.harvard.edu).

### EXPERIMENTAL MODEL AND SUBJECT DETAILS

**Mice**—To generate the Diet-induced obese (DIO) mice, wild-type C57BL/6J male mice were fed with a high-fat diet (HFD, 45 kcal% from fat) for 16-20 weeks. Liver-specific IKK $\beta$  transgenic hemizygous mice (LIKK $\beta$ <sup>tg(+/-)</sup>) were kindly provided by Dr. Michael Karin. Homozygous (LIKK $\beta$ <sup>tg(+++)</sup>) and control (LIKK $\beta$ <sup>tg(-/-)</sup>) mice were obtained by interbreeding IKK $\beta$  transgenic hemizygous (LIKK $\beta$ <sup>tg(+/-)</sup>) mice. BKS.Cg-Dock7<sup>m(+/-)</sup> *Lep<sup>db</sup>/J* mice and BKS.Cg-*Lep<sup>ob</sup>/J* mice were obtained from the Jackson Laboratory. All animal experiments were approved by Institutional Animal Care and Use Committee (IACUC) at Boston Children's Hospital.

To increase the protein levels and nuclear translocation of XBP1s in the livers of lean mice, C57BL/6J were fasted for 24 hours, then re-fed for one hour, at which point the mice were sacrificed.

**Cell Lines**—Human embryonic kidney 293 (HEK293), Chinese hamster ovary (CHO), Hepa-1c1c7 (murine hepatoma cell), HepG2 (human hepatoma cell) and MDA-MB-231



(breast adenocarcinoma cell) cells were maintained in DMEM with 10% fetal bovine serum (FBS), 10 U ml<sup>-1</sup> penicillin, and 1 mg ml<sup>-1</sup> streptomycin at 37°C in a 5% CO<sub>2</sub> humidified atmosphere.

## METHOD DETAILS

**Biochemical Reagents**—Anti-XBP1s, anti-insulin receptor (IR), anti-phosphotyrosine (PY99), anti-myc, anti-HA antibodies, anti-ATF4, anti-ATF6, and HRP-conjugated goat anti-mouse and goat anti-rabbit antibodies were from Santa Cruz Biotechnology (Santa Cruz, CA). Rabbit anti-IKKβ (Rabbit), anti-IKKα, anti-IKKγ, anti-p-JNK, anti-JNK2, anti-p-P38, anti-P38, anti-p-c-Jun, anti-c-Jun, anti-ATF2, anti-p-ATF2, anti-ubiquitin, anti-phospho-AKT (Thr308), anti-phospho-AKT (Ser473), anti-AKT, anti-phospho-GSK3α/β(Ser21/9), anti-GSK3β, anti-phospho-PERK (Thr980), PERK antibodies, anti-p65, anti-Nup98, anti-Lamin A/C and anti-α-tubulin antibodies and mouse anti-ubiquitin antibody were purchased from Cell Signaling Technology (Beverly, MA). Anti-Flag and anti-His antibody and chemical compound [(aminocarbonylamino)-5-(4-fluorophenyl)-3-thiophenecarboxamide (TPCA-1) and cycloheximide (CHX) were purchased from Sigma Aldrich (St. Louis, MO). Anti-phosphoserine/threonine antibody was purchased from ECM Bioscience (Versailles, KY). Rabbit polyclonal anti-phospho-XBP1s<sup>Thr48</sup> was raised against CRAAGSEASGT(p)PQARKRQR by Covance (Denver, PA) as described previously (Lee et al., 2011). Mouse anti-XBP1s was purchased from Biolegend (San Diego, CA). Mouse anti-IKKβ was purchased from Novus (Littleton, CO). Anti-Phosphoserine/threonine antibody was from Abcam (Cambridge, MA). Recombinant Human TNF-α protein was purchased from R&D systems (Minneapolis, MN).

**Plasmids**—Mouse IKKβ plasmid, pCR-Flag-IKKβ was purchased from Addgene (Cambridge, MA). Plasmids expressing constitutively active IKKβ (pcDNA3.1-IKKβ) with amino acid substitution mutations (S177E/S181E) were generated by PCR-based mutagenesis using pCR-Flag-IKKβ as a template and QuikChange II site-directed mutagenesis kit (Agilent Technologies), following manufacturer's instruction. Constitutively active IKKβ (S177E/S181E) was subsequently cloned into pcDNA3.1 (Life Technologies).

Primer sequences for amino acid substitution of IKKβ (S177E/S181E) were as follows:

(forward), 5' -  
GGAGCTGGATCAGGGCGAGCTGTGCACGGAATTTGTGGGGACTCTGC-3';  
(reverse),  
5' -GCAGAGTCCCCACAAATTCGGTGCACAGCTCGCCCTGATCCAGCTCC-3'.

Human XBP1s plasmid and mouse plasmid expressing kinase death IKKβ (pCR-Flag-IKKβ-KM) were purchased from Addgene. Vectors expressing mouse XBP1s (no tagged or flag-tagged XBP1s) and NH<sub>2</sub>-terminal (1–373 aa) of activating transcription factor 6a (ATF6n) were constructed previously (Ozcan et al. 2009). The XBP1s plasmids with amino acid substitution mutations (T48A, S148A or double mutations T48A/S148A) were generated by PCR-based mutagenesis using pcDNA3.1-Flag-XBP1s as a template and QuikChange II site-directed mutagenesis kit.

Primer sequences for T48A were as following:

(forward), 5'-GGGTCGGAGGCGAGCGGGGCACCGCAGGCTCGCAAGCGG-3';  
(reverse), 5'-CCGCTTGCAGCCTGCGGTGCCCGCTCGCCTCCGACCC-3'.

Primer sequences for S148A were as following:

(forward), 5'-CACGCTTGGGAATGGACGCGCTGGATCCTGACGAGG-3';  
(reverse), 5'-CCTCGTCAGGATCCAGCGCGTCCATTCCCAAGCGTG-3'.

**Total Protein Extraction from Cells**—Cells were lysed in lysis buffer (25 mM Tris-HCl (pH 7.4), 10 mM NaF, 10 mM Na<sub>4</sub>P<sub>2</sub>O<sub>7</sub>, 2 mM Na<sub>3</sub>VO<sub>4</sub>, 1 mM EGTA, 1 mM EDTA, 1% NP-40, protease inhibitor (Roche) and phosphatase inhibitor cocktails (Roche)). After rotation at 4°C for 20 minutes, cell lysates were centrifuged at 16,000 × g for 20 minutes at 4°C and supernatants were collected. Protein concentrations were measured by using Lowry protein assay (Bio-Rad) and normalized with lysis buffer for each sample to have an equal concentration. Proteins were denatured by boiling at 100°C for 5 minutes in 1X Laemmli buffer. The lysates were cooled to room temperature before loading to western blot gel.

**Total Protein Extraction from Tissue**—Liver, adipose, muscle and kidney tissues (ranging 50-100 mg) were homogenized with TissueLyser II (QIAGEN) in 1 mL of ice-cold tissue lysis buffer (25 mM TrisHCl, pH 7.4; 10 mM Na<sub>3</sub>VO<sub>4</sub>, 100 mM NaF, 50 mM Na<sub>4</sub>P<sub>2</sub>O<sub>7</sub>, 10 mM EGTA, 10 mM EDTA, 1% NP-40, protease inhibitor (Roche) and phosphatase inhibitor cocktails (Roche)). After homogenization, lysates were rotated at 4°C for 20 minutes and centrifuged at 16,000 × g for 20 minutes. The top lipid layer was carefully removed and the remaining supernatant was transferred to 1.5 mL Eppendorf tubes for further centrifugation. This step was repeated twice to completely remove the lipid. Protein concentrations were determined by using Lowry protein assay (Bio-Rad) and were normalized with lysis buffer for each sample. Proteins were denatured in 1X Laemmli buffer by boiling at 100°C for 5 minutes. The lysates were cooled to room temperature before loading for western blot analysis.

**Western Blotting**—Cell or tissue lysate samples, prepared as described above, were resolved on SDS-PAGE. Proteins were then transferred to polyvinylidene fluoride (PVDF) membrane. The membrane was blocked in Tris-buffered saline solution with 0.05 % Tween 20 (TBST; pH 7.4) with 10% blocking reagent provided with BM Chemiluminescence Western blotting substrate (POD) assay system for 1 h, and then incubated with primary antibody in TBST with 10% blocking reagent at 4°C overnight. After the incubation, the membrane was washed three times in TBST and incubated with secondary antibody in TBST with 10% blocking reagent for 1 h at room temperature. After subsequent three-time washing in TBST, membrane was developed using a chemiluminescence assay system (Roche) and exposed to X-ray films (Denville Scientific). Relative protein levels were quantified by Image J program (NIH, Bethesda, MD).

For stripping, membrane was vigorously shaken in stripping buffer (62.5 mM Tris-HCl, pH 6.7; 2% SDS; 100 mM 2-mercaptoethanol) at 55°C for 15 minutes. After stripping, membrane was washed three times in TBST (3 times for 20 minutes).

**Preparation anti-IKK $\beta$ -coated Dynabeads**—Mouse anti-IKK $\beta$  antibody (8  $\mu$ g per sample) (Novus, Littleton, CO) was conjugated to Dynabeads Protein G (50  $\mu$ L per sample) (Life Technologies) with disuccinimidyl suberate (DSS) and bis(sulfosuccinimidyl) suberate (BS3) crosslinker (Thermo Scientific #21585) following its instructions with modifications, which is as following: For each sample, 50  $\mu$ L of dynabeads were spun down and the supernatant was discarded, and then dynabeads were incubated with 8  $\mu$ g of antibody and 160  $\mu$ L of conjugation buffer (PBS + 0.05% Tween-20) for 10 minutes at room temperature. The beads-antibody complex was washed with conjugation buffer (20 mM sodium phosphate, 0.15 M NaCl, 0.05% Tween-20, pH 8.5) twice by gentle pipetting. 2 mg of Bis (sulfosuccinimidyl) suberate (BS3) was dissolved in 0.69 mL conjugation buffer and 250  $\mu$ L of BS3 solutions was immediately added to the beads-antibody complex.

The mixture was rotated for 30 minutes at room temperature, and then the reaction was stopped by adding 12.5  $\mu$ L of quenching buffer (1 M Tris HCl, pH 7.5). After four times washing with immunoprecipitation buffer (PBST: 0.02% Tween, 0.1% sodium azide), the beads were stored in immunoprecipitation buffer at 4°C, until used for immunoprecipitation later.

**Immunoprecipitation**—Cell lysates (1-4 mg per sample) or tissue lysates (2-8 mg per sample) were incubated with antibody (1-8  $\mu$ g per sample) overnight at 4°C with a gentle rotation. Protein A Sepharose CL-4B beads (for rabbit immunoglobulin G [IgG], GE Healthcare Life Sciences) or protein G Sepharose CL-4B beads (for mouse IgG, GE Healthcare Life Sciences) (10-30  $\mu$ L per sample) were added to the tubes and rotated at 4°C for 1 h. Beads were precipitated by centrifugation at 800  $\times$  g for 30 seconds and washed three times with cold lysis buffer. The pellet was resuspended in 2X Laemmli buffer and incubated at 100°C for 5 minutes. The supernatants were collected and used for western blot.

For the immunoprecipitation experiments in figure 2E, we immunoprecipitated the total lysates with anti-XBP1 antibody, then loaded three times more immunoprecipitated samples from XBP1s-transfected cells than the samples from the co-transfected cells with XBP1s and IKK $\beta$ .

For the immunoprecipitation experiments in figure 3B, we used three times more immunoprecipitation samples from XBP1s-transfected cells, XBP1s(DM)-transfected cells and XBP1s(DM)-together with- IKK $\beta$ -transfected cells than that of XBP1s-together with- IKK $\beta$ -transfected cells when loading to SDS-PAGE gels to perform western blot.

For the immunoprecipitation experiments in figure 4A, after immunoprecipitation of IKK $\beta$  from the liver lysates, we loaded two times more immunoprecipitates (IP) samples from *ob/ob* mice liver than that of lean mice liver to SDS-PAGE gels, and XBP1s and IKK $\beta$  protein levels were detected by immunoblotting.

**Nuclear Protein Extraction**—Cells were lysed with 400  $\mu$ L of cytoplasmic lysis buffer (10 mM HEPES (pH 7.5), 2 mM  $MgCl_2$ , 1 mM EDTA, 1 mM EGTA, 10 mM KCl, 10 mM NaF, 0.1 mM  $Na_3VO_4$ , 1 mM dithiothreitol [DTT], protease inhibitor (Roche) and phosphatase inhibitor cocktails (Roche)). After 15 minutes of incubation on ice, 25  $\mu$ L of 10% NP-40 was added to the cell lysates. The lysates were then incubated for 5 minutes on ice and vortexed 2 times for 10 sec. The lysates were centrifuged for 30 seconds at  $16,000 \times g$ , and supernatants were collected as cytoplasmic fractions. The pellets were re-suspended in 50  $\mu$ L of nuclear extraction buffer (25 mM HEPES (pH 7.5], 500 mM NaCl, 10 mM NaF, 10% glycerol, 0.2% NP-40, 5 mM  $MgCl_2$ , and 1 mM DTT, protease inhibitor (Roche) and phosphatase inhibitor cocktails (Roche)) and sonicated 3 times on ice. The nuclear fractions were centrifuged for 5 minutes at  $16,000 \times g$  and the supernatant was collected to obtain nuclear proteins. Both cytoplasmic and nuclear proteins were quantified using Bio Rad protein assay and normalized with lysis buffer to have an equal concentration. The proteins were denatured by boiling at  $100^\circ C$  following mixing with 5X Laemmli buffer.

For nuclear extraction from liver tissues, 80 mg of liver tissue was cut into small pieces and washed with PBS. Nuclear proteins were isolated using a commercially available kit from Thermo according to their instructions.

100 mg liver tissues was cut into small pieces and then homogenized using a Dounce homogenizer or a tissue grinder in 800  $\mu$ L CER I. And then the tissue mixture was vortexed vigorously on the highest setting for 15 seconds to fully suspend the tissue samples, followed by 10 minutes incubation on ice. 44  $\mu$ L ice-cold CER II was added to the mixture and the tubes were vortexed for 5 seconds on the highest setting. The samples were incubated on ice for one minute, vortexed for another 5 seconds, and centrifuged for 5 minutes at the speed of  $16,000 \times g$ . The supernatant which contains cytoplasmic extract, was immediately transferred to a new pre-chilled tube and further lysed with Laemmli buffer. The pellet (nuclear fraction) was suspended in 200  $\mu$ L ice-cold NER buffer by vortexing, followed by ultrasound for 20 sec. The lysate was centrifuged at maximum speed ( $\sim 16,000 \times g$ ) for 10 minutes. The supernatant (nuclear extract) was immediately transferred to a new pre-chilled tube, and further lysed by adding Laemmli buffer.

**Glucose Tolerance Test**—Mice were fasted overnight (6 p.m.–9 a.m.), and D-glucose ( $0.3 \text{ g kg}^{-1}$  for *ob/ob*,  $1.25 \text{ g kg}^{-1}$  for DIO and  $1.5 \text{ g kg}^{-1}$  for  $LIKK\beta^{tg(+/+)}$  mice) was administered intraperitoneally. Blood glucose levels were measured from the tail using a glucose meter (Contour, Bayer) at 0, 15, 30, 60, 90, and 120 minutes after D-glucose administration.

**Insulin Tolerance Test**—Mice were fasted for 6 h (8 a.m.–2 p.m.), and recombinant human insulin from Eli Lilly was administered intraperitoneally ( $2 \text{ IU kg}^{-1}$  for *ob/ob*,  $1.5 \text{ IU kg}^{-1}$  for DIO and  $1.4 \text{ IU kg}^{-1}$  for  $LIKK\beta^{tg(+/+)}$  mice. Blood glucose levels were measured from the tail using a glucose meter (Contour, Bayer) at 0, 15, 30, 60, 90, and 120 minutes after insulin administration.

**Blood Glucose, Plasma Insulin and Plasma IL-6 Measurements**—Mice were fasted for 6 h and their blood glucose levels were measured with a glucose meter (Bayer)

after the blood was collected from tail vein. For enzyme-linked immunosorbent assay (ELISA), blood was collected from the tail vein and plasma was separated by centrifugation at  $2,000 \times g$ ,  $4^{\circ}\text{C}$  for 30 minutes. Plasma insulin levels were determined by an Ultra-Sensitive Mouse Insulin ELISA kit from Crystal Chem (Downers Grove, IL). Plasma IL-6 levels were measured with a Mouse IL-6 ELISA kit from Boston Biochem, Inc (Cambridge, MA).

**Analysis of in vivo insulin signaling**—For in vivo insulin signaling analysis, mice were anaesthetized with xylazine (100 mg per kg) / ketamine (20mg per kg) (Pfizer) after 6 h of fasting. Insulin (1 IU  $\text{kg}^{-1}$ ) or saline was infused into the liver via the portal vein. Five minutes after infusion, livers were harvested and frozen in liquid nitrogen immediately and stored at  $-80^{\circ}\text{C}$  until use.

### **Blood Alanine Transaminase (ALT) and Aspartate Transaminase (AST) Measurements**

**Serum sample preparation:** The fresh blood were collected with heparin sodium coated capillary and the sample were centrifugated at 3,000 rpm at  $4^{\circ}\text{C}$  for 30 minutes. The supernatant blood plasma were collected and the ALT and AST levels in the plasma were measured with ALT Color Endpoint Assay and AST Color Endpoint Assay kits (Bio Scientific, Austin, TX), respectively, according to manufacturer's instructions with no modifications.

**AST Assay:** Firstly, 6 serial dilutions of the Oxaloacetate Control were prepared by using the Pyruvate Dilution Buffer and the concentration of these Control Dilutions were as follows: 800 IU/L, 400 IU/L, 200 IU/L, 100 IU/L, 50 IU/L and 0 IU/L. 5  $\mu\text{L}$  of each sample or standard (in duplicate) was added to the bottom of the microplate (96 well-plate) wells and 50  $\mu\text{L}$  of AST Reagent Solution was added to the wells. The plate were covered with adhesive film and incubated the plate at  $37^{\circ}\text{C}$  for 10 minutes. The adhesive film were carefully removed and 50  $\mu\text{L}$  AST Color Reagent was added to the wells. The plate was covered with adhesive film again and incubated at  $37^{\circ}\text{C}$  for 10 minutes. Then, 200  $\mu\text{L}$  0.1 M HCl was added to each well and the absorbance of each wells were measured at 510 nm. The AST concentration in each sample was calculated by the equation:

$$\text{AST concentration} = (\text{mean absorbance} - y - \text{intercept})/\text{slope}$$

Use the mean absorbance values for each serum sample to determine the corresponding concentration of AST from the standard curve.

**ALT Assay:** Firstly, we made 6 serial dilutions of the Pyruvate Control using the Pyruvate Dilution Buffer and the concentration of these Pyruvate Control Dilutions are as follows:

$$150 \text{ IU/L}, 70 \text{ IU/L}, 32 \text{ IU/L}, 15 \text{ IU/L}, 7 \text{ IU/L} \text{ and } 0 \text{ IU/L}.$$

10  $\mu\text{L}$  of each sample or standard (in duplicate) was added to the bottom of the microplate (96 well-plate) wells and 50  $\mu\text{L}$  of ALT Reagent Mix was added to the wells. The plate were covered with adhesive film and incubated the plate at  $37^{\circ}\text{C}$  for 30 minutes. The adhesive

film were carefully removed and 50  $\mu$ L DNPH Color Solution was added to the wells. The plate was covered with adhesive film again and incubated at 37°C for 10 minutes. Then, 200  $\mu$ L 0.5 M NaOH was added to each well and the plate was incubated at 37°C for 5 minutes. The absorbance of each wells were measured at 510 nm.

**RNA Preparation and Real-Time Quantitative PCR**—Total RNA was extracted from cells or the liver of mice using Trizol reagent (Life Technologies) following its instruction. Purified total RNA was used to obtain cDNA using a cDNA synthesis kit (Bio-Rad) following these conditions: 25°C for 5 minutes, 42°C for 30 minutes, and 85°C for 5 minutes. The gene expression was analyzed with iQ5 Multicolor Real-Time PCR Detection System (BioRad, Hercules, California) with SYBR Green Supermix (BioRad, Hercules, California). The mRNA levels were normalized to 18S as a house keeping gene.

**Genomic DNA Purification for Genotyping**—Mouse tails were lysed in 100  $\mu$ L 1X lysis buffer (4.5 mL of 1.5M Tris pH 8.8; 1.7 mL of 1M (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>; 0.3 mL of 0.5M MgCl<sub>2</sub>; 0.5 mL of Triton X100; 92 mL autoclaved H<sub>2</sub>O and 1 mL betamercaptoethanol) with 2.5  $\mu$ L of Proteinase K and 0.5  $\mu$ L of betamercaptoethanol, at 55°C for 15 hours. Samples were then boiled at 100°C for 20 minutes and cooled to room temperature, followed by a 20-sec centrifugation at 800  $\times$  g. The supernatant was removed to new tubes, and equal volume of phenol chloroform solution was added. Samples were vortexed for 20 seconds and centrifuged at 16,000  $\times$  g at 4°C for 20 minutes. The top layer was transferred to new tubes and two volumes of isopropanol were added to each tube. Samples were quickly vortexed and then centrifuged at 16,000  $\times$  g at 4°C for 10 minutes. The supernatant was discarded and the DNA pellet was washed with 500  $\mu$ L of 70% ethanol by centrifugation at 16,000  $\times$  g at 4°C for 10 minutes. Same step was repeated with 100% ethanol. Then, the supernatant was discarded and the DNA pellet was let dry at room temperature. Finally, 100  $\mu$ L of distilled H<sub>2</sub>O was added to the purified DNA, and the tubes were heated at 55°C for 10 minutes. After vortexing, the samples were centrifuged at 16,000  $\times$  g at 4°C for 10 minutes and the purified genomic DNA was used for genotyping.

**Adenovirus Production**—Adenovirus-producing plasmids containing a gene of constitutively active IKK $\beta$  or XBP1s were constructed using adenoviral expression kit from Life Technologies. The adenovirus vectors were digested with Pac I, and then transfected into 293A producer cells in 6-well-plates. The media was replaced with DMEM containing 10% FBS and 1% penicillin/streptomycin the next day. The cells were transferred to 10 cm tissue culture dishes 48 h after the transfection. The culture media was replaced with fresh media every 2-3 days until cytopathic effect (CPE) was observed. The cells were collected when 80% CPE was observed and adenovirus was harvested by freezing at -80°C and thawing at 37°C and repeating it for four times. Finally, cell lysates were removed by centrifugation at 2,000  $\times$  g for 30 minutes at 25°C and the supernatant containing adenovirus particles was stored at -80°C.

**Adenovirus Transduction**—To transduce cells in 10 cm dishes with adenovirus, cells were washed with culture medium containing 1% FBS and incubated with 2.5 mL of media containing 1% FBS and adenovirus. The dishes were gently rocked every 15 minutes for 1 h



and 7.5 mL of media containing 1% FBS was added to each dish. The cells were incubated for additional 16 h.

**Adenovirus or Adeno-Associated Virus (AAV) Injection to Mice**—Virus (either adenovirus or AAV) was introduced to mice through the tail vein injection. Virus was thawed at 25°C before injection and the desired amount of virus was diluted with saline to a final volume of 100  $\mu$ L per mouse. Mice were restrained and their tails were heated mildly with a heating lamp to achieve vasodilatation. Virus was injected through the tail vein. To prevent the back flow of virus solution, mild pressure was applied at the spot of injection immediately after injection until no bleeding was achieved.

**Production of AAV**—The constitutively active IKK $\beta$  coding sequence was generated by PCR-based method using pcDNA3.1-IKK $\beta$ (S177E/S181E) as a template. The following primer pair was used in this PCR:

(forward), 5'-TACTTACGCGTACCATGGACTACAAGGACGACGAT-3';

(reverse), 5'-ACAAGTCGACCTACTAATCGCAGGCCTGCTCC-3'.

The resulting constitutively active IKK $\beta$  coding sequence was cloned into an AAV plasmid (TBG-AAV2.1) to create TBG-IKK $\beta$  vector, in which thyroid hormone-binding globulin (TBG) promoter controls hepatocyte-specific expression of IKK $\beta$ . The subsequent AAV virus production and purification were finished by Gene Therapy Center and Vector Core at University of Massachusetts Medical School.

**Recombinant XBP1s Protein**—Recombinant XBP1s proteins were produced and purified as described previously (Lee et al., 2011). Briefly, XBP1s coding sequence with 6 $\times$  His-TF tag (N terminus) sequence were cloned into a pGSC1 plasmid. The final protein sequence of the construction (6 $\times$  His +TF tag + 3C protease cleavage site) was as follows:

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MNHKVHHHHHHMQVSVETTQGLGRRVTITIAADSIETAVKSELVNAVKKVRI
DGFRKKGKVP MNIVAQRYGASVRQDVLGDLMSRNFIDAIKEKINPAGAPTYVP
GEYKLGEDFTYSVEFEVYPEVELQGLEAIEVEKPIVEVTDADVDGMLDTRLK
QQATWKEKDGAVEAEDRVTIDFTGSVDGEEFEGGKASDFVLAMGQGRMIPG
FEDGIKGHKAGEEFTIDVTFPEEYHAENLKGKAAKFAINLKKVEERELPELTA
EFIKRFGVEDGSVEGLRAEVRK
NMERELKSAIRNRVKSQAIEGLVKANDIDVPAALIDSEIDLRRQAAQRFGGN
EKQALELPRELFEEQAKRRVVGLLLGEVIRTNELKADEERVKGLIEEMASAY
EDPKEVIEFYSKNKELMDNMRNVALEEQA VEAVLAKAKVTEKETTFNELMN
QQASAGLEVLFGQP.
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To produce recombinant XBP1s protein, 5 ng of pGSC1 plasmid containing XBP1s with His-TF tag was transformed into ArcticExpress<sup>TM</sup> (DE3) RP host strain *E. coli* (Agilent Technologies) to produce recombinant XBP1s protein. The total protein was extracted and purified by Ni-affinity resin (Life Technologies).

**Protein Degradation Assay**—HEK293 cells were transfected with XBP1s vector alone or together with IKK $\beta$  vector. After 24 h of transfection, the cells were treated with

cycloheximide (CHX 20  $\mu\text{g ml}^{-1}$ , Sigma), a translation inhibitor. Cells without treatment and at 5, 10, 15, 25 and 45 minutes following CHX incubation were flash frozen in liquid nitrogen. Protein levels were determined by western blotting and the subsequent quantification was performed from scanned immunoblotting results with ImageJ software.

### **Genotyping of Liver-Specific IKK $\beta$ Transgenic Mice Using TaqMan-based Real-Time PCR**

Litters from breedings of IKK $\beta$  transgenic hemizygous parents were genotyped using TaqMan-based real-time PCR in order to distinguish between the homozygous and hemizygous transgenic mice. Real-time PCR was conducted in triplicate in total 10  $\mu\text{L}$  of reaction solution with 5  $\mu\text{L}$  of 2X Taqman Genotyping Master Mix (Applied Biosystems, Life Technologies), 0.5  $\mu\text{L}$  of 20X IKK $\beta$  gene expression assay, 0.5  $\mu\text{L}$  of 20X GAPDH gene expression assay, and 10 ng genomic DNA (purified as described above) in each well of a 384-well plate. Real-time PCR was conducted on the ABI 6 (Applied Biosystems) using the following program: hold stage (50°C for 2 minutes, 95°C for 10 minutes) and PCR stage (40 cycles of 95°C for 15 sec, 60°C for 1 min). SDS Analysis (version 2.3, a software from ABI) was used to determine the Ct values of each reaction. Primer probe combinations were purchased as proprietary reagents from ABI-Life Technologies Inc:

Gapdh, Life Technologies, Cat. # 4400291; Assay ID: Mm00186822\_cn;

IKK $\beta$ , Life Technologies, Cat. # 4351372; Assay ID: Hs01559467\_m1.

**Histological Analysis**—The liver was collected immediately from AAV-injected *ob/ob* mice, AAV-injected DIO and LIKK $\beta^{\text{tg}(+/+)}$  and LIKK $\beta^{\text{tg}(-/-)}$  mice and stored in 10% formalin in phosphate buffer. After two weeks of incubation, the samples were embedded in paraffin and sectioned. The tissue sections were used for Hematoxylin and Eosin (H&E) Staining, and immunohistochemical staining for cleaved caspase-3 (active caspase-3) and Trichrome staining (Rodent Histopathology core, Harvard Medical School; Specialized Histopathology Services-Longwood, Brigham and Women's Hospital).

**IKK $\beta$  Kinase Activity Assays**—IKK complexes were immunoprecipitated from mouse liver lysates using anti-IKK $\gamma$  antibodies (BD PharMingen, San Jose, CA) and protein A-Sepharose beads (GE Healthcare Life Sciences, Pittsburgh, PA) as described above. In vitro IKK $\beta$  Kinase Activity assay was conducted with the IKK complexes harvested, using Cyclex IKK Assay kit from MBL International (Woburn, MA), following its instructions, with no modifications.

To assay partially purified recombinant IKK $\beta$ , add 10  $\mu\text{L}$  of each sample or standard to the wells of the assay plate on ice. Duplicate wells containing 10 m units/10  $\mu\text{L}$  IKK $\beta$  positive control (Cat# CY-E1176-2) has been included in each assay as a positive control for phosphorylation. Begin kinase reaction by addition of 90  $\mu\text{L}$  Kinase Reaction Buffer (Kinase Buffer plus 20X ATP) and incubate for 30 minutes at 30°C. Wash the wells four times with Wash Buffer (containing 2% Tween-20) making sure each well is filled completely. Remove residual Wash Buffer by gentle tapping or aspiration. Pipette 100  $\mu\text{L}$  of Anti-phospho-I $\kappa$ B $\alpha$  serine32 monoclonal antibody (AS-2E8) into each well and incubate for 30 minutes at RT. Wash the wells five times. Pipette 100  $\mu\text{L}$  of HRP conjugated anti-mouse IgG into each well

and incubate for 30 minutes at RT. Wash the wells five times. Add 100  $\mu$ L of Substrate Reagent (containing chromogenic substrate, tetra-methylbenzidine (TMD)) to each well and incubate for 5-15 minutes at RT. Add 100  $\mu$ L of Stop Solution (containing 1 N  $H_2SO_4$ ) to each well. The absorbant signal in each well is measured with a spectrophotometric plate reader at dual wavelengths of 450/540 nm. All samples and standards have been assayed in duplicate.

**IKK $\beta$  Kinase Assay with XBP1s as Substrates**—The in vitro IKK $\beta$  kinase assay was conducted as described previously (Hu et al., 2005; Shinohara et al., 2005). HEK293 cells were transfected with flag-IKK $\beta$  (S177E/S181E) vector, and the constitutively active IKK $\beta$  protein was purified through immunoprecipitation with an anti-flag antibody. Equal amount of purified IKK $\beta$  protein was incubated with His-TF-XBP1s fusion protein produced as described above or with commercial IKK $\beta$  substrate (Millipore, Darmstadt, Germany), in a kinase assay buffer (20 mM Hepes, pH 7.6; 25 mM  $\beta$ -glycerophosphate; 0.1 mM  $Na_3VO_4$ ; 4mM NaF; 2 mM  $MnCl_2$ ; 10 mM  $MgCl_2$ ; 10  $\mu$ M ATP) at 37°C for 30 minutes. For a negative control reaction, TPCA-1 (20  $\mu$ g  $mL^{-1}$ ), an IKK $\beta$  inhibitor, was added to the reaction solution of IKK $\beta$  together with His-TF-XBP1s or IKK $\beta$  substrate at 37°C for 30 minutes. The kinase reaction was stopped by adding 1X Laemmli buffer and boiling at 100°C for 10 minutes. IKK $\beta$ -mediated phosphorylation of XBP1s or IKK $\beta$  substrate was analyzed by western blotting with anti-serine/threonine antibody.

**Mass Spectrometric Analysis by LC-MS/MS**—His-TF-XBP1s and purified constitutively active IKK $\beta$  protein (described above) were incubated in a kinase assay buffer (20 mM Hepes, pH 7.4; 25 mM  $\beta$ -glycerophosphate; 0.1 mM  $Na_3VO_4$ ; 4 mM NaF; 2 mM  $MnCl_2$ ; 10 mM  $MgCl_2$ ; 10  $\mu$ M ATP) at 37°C for 30 minutes. His-TF-XBP1s was pulled down with anti-His antibody, and resolved in SDS-PAGE and stained with Coomassie blue (Bio-Rad). Excised His-TF-XBP1s from Coomassie-stained gel was extracted and digested with trypsin and digested XBP1s peptide mixtures were subjected to microcapillary liquid chromatography–tandem mass spectrometry (LC-MS/MS) analysis (Taplin Biological Mass Spectrometry Facility, Harvard Medical School). Resulting MS/MS spectra of XBP1s fragments were analyzed by the SEQUEST (Thermo Scientific) algorithm.

**Tissue Lipid Extraction**—Around 75 mg of liver tissue (exact weight was recorded and used to normalize lipid amount) was homogenized with Homogenizer (Thermo Fisher Scientific) in 1 mL 50 mM NaCl on ice, following the addition of 5 mL of chloroform/methanol mixture (chloroform : methanol = 2:1). The homogenized tissue solution was vortexed for 10 seconds and centrifuged at 16,000  $\times$  g for 10 minutes. The aqueous phase was carefully removed, and the left oleic phase was mixed with 1.5 mL of methanol. The resulting mixture was vortexed for 10 seconds and centrifuged at 16,000 X g for 10 minutes. The lipid extract (top phase) was carefully moved to a new tube and mixed with 75  $\mu$ L of 10% Triton X100 in acetone, and dried in fume hood. The resulting pellet was used for cholesterol or triglyceride determination with cholesterol assay kit (Wako, Richmond, VA) or Triglyceride assay kit (Wako, Richmond, VA) according to manufacture instructions.

**Hyperinsulinemic euglycemic clamp**—Prior to the clamp experiment, one catheter was inserted into the jugular vein for infusions. After 5–7 days of recovery, mice that had lost less than 10% of their preoperative weight were subjected to the clamp. On the day of experimentation, each mice was deprived of food for 3.5 h in the morning and then infused continuously with D-[3-3H]-glucose (PerkinElmer) (0.05  $\mu$ Ci/min) at a rate of 1 $\mu$ L/min for 1.5 h. After basal sampling, a primed continuous infusion of human regular insulin (Humulin, Eli Lilly) is infused at a rate of 4 mU/kg/min. The insulin solutions were prepared with 3% BSA. 20% glucose was infused at variable rates as needed to maintain plasma glucose at ~120 mg/dl. All infusions were done using micro infusion pumps (KD Scientific). Blood glucose concentrations were monitored regularly according to a fixed scheme from tail vein. Steady state was considered achieved during 75-120 minutes, when a fixed glucose-infusion rate maintained the glucose concentration in blood constantly for 45 minutes. The [3-3H] glucose content in serum during basal conditions and at steady state was measured in a liquid scintillation counter.

## QUANTIFICATION AND STATISTICAL ANALYSIS

Error bars are represented as mean  $\pm$  SEM. Statistical significance was calculated by Student's *t* test, one-way analysis of variance (ANOVA) or two-way analysis of variance (ANOVA). When ANOVA indicated a significant difference among the groups, the statistical difference between two groups were compared using a stricter criterion for statistical significance according to the Bonferroni rule (corrected P value = pairwise P value  $\times$  no. of comparisons). Significance was accepted at \*P < 0.05, \*\*P < 0.01 or \*\*\*P < 0.001.

## DATA AND SOFTWARE AVAILABILITY

Image J software was downloaded from the website: <https://imagej.nih.gov/ij/>. GraphPad Prism 5 software was used for two-way analysis of variance (ANOVA) with Bonferroni multiple-comparison analysis

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgments

We thank Morris F. White and Rongya Tao for helping us to perform the hyperinsulinemic-euglycemic clamp experiments. This work was supported by funds provided to UO from Department of Medicine, Boston Children's Hospital, by NIH grants R01DK081009 and R01DK098496 to UO, by the American Diabetes Association's Career Development grant #7-09-CD-10, and by NIH grant AI043477 to MK. UO is a scientific founder, shareholder, and member of the scientific advisory board and board of directors of ERX Pharmaceuticals, Inc.

## References

- Arkan MC, Hevener AL, Greten FR, Maeda S, Li Z-W, Long JM, Wynshaw-Boris A, Poli G, Olefsky J, Karin M. IKK- $\beta$  links inflammation to obesity-induced insulin resistance. *Nat Med.* 2005; 11:191–198. [PubMed: 15685170]
- Awazawa M, Ueki K, Inabe K, Yamauchi T, Kubota N, Kaneko K, Kobayashi M, Iwane A, Sasako T, Okazaki Y, et al. Adiponectin enhances insulin sensitivity by increasing hepatic IRS-2 expression via a macrophage-derived IL-6-dependent pathway. *Cell Metab.* 2011; 13:401–412. [PubMed: 21459325]

- Cai D, Yuan M, Frantz DF, Melendez PA, Hansen L, Lee J, Shoelson SE. Local and systemic insulin resistance resulting from hepatic activation of IKK-beta and NF-kappaB. *Nature medicine*. 2005; 11:183–190.
- Copps KD, Hancer NJ, Opore-Ado L, Qiu W, Walsh C, White MF. Irs1 Serine 307 Promotes Insulin Sensitivity in Mice. *Cell Metabolism*. 2010; 11:84–92. [PubMed: 20074531]
- Daya S, Berns KI. Gene therapy using adeno-associated virus vectors. *Clinical microbiology reviews*. 2008; 21:583–593. [PubMed: 18854481]
- Deng Y, Wang ZV, Tao C, Gao N, Holland WL, Ferdous A, Repa JJ, Liang G, Ye J, Lehrman MA, et al. The Xbp1s/GalE axis links ER stress to postprandial hepatic metabolism. *The Journal of clinical investigation*. 2013; 123:455–468. [PubMed: 23257357]
- Gao ZG, Ye JP. Why do anti-inflammatory therapies fail to improve insulin sensitivity? *Acta pharmacologica Sinica*. 2012; 33:182–188. [PubMed: 22036866]
- Garcia MC, Wernstedt I, Berndtsson A, Enge M, Bell M, Hultgren O, Horn M, Ahren B, Enerback S, Ohlsson C, et al. Mature-onset obesity in interleukin-1 receptor I knockout mice. *Diabetes*. 2006; 55:1205–1213. [PubMed: 16644674]
- Gardner BM, Pincus D, Gotthardt K, Gallagher CM, Walter P. Endoplasmic reticulum stress sensing in the unfolded protein response. *Cold Spring Harbor perspectives in biology*. 2013; 5:a013169. [PubMed: 23388626]
- Gregor MF, Hotamisligil GS. Inflammatory Mechanisms in Obesity. *Annual Review of Immunology*. 2011; 29:415–445.
- Hacker H, Karin M. Regulation and function of IKK and IKK-related kinases. *Science's STKE : signal transduction knowledge environment*. 2006; 2006:re13.
- Hirosumi J, Tuneman G, Chang L, Gorgun CZ, Uysal KT, Maeda K, Karin M, Hotamisligil GS. A central role for JNK in obesity and insulin resistance. *Nature*. 2002; 420:333–336. [PubMed: 12447443]
- Hotamisligil GS. Inflammation and metabolic disorders. *Nature*. 2006; 444:860–867. [PubMed: 17167474]
- Hotamisligil GS, Peraldi P, Budavari A, Ellis R, White MF, Spiegelman BM. IRS-1-mediated inhibition of insulin receptor tyrosine kinase activity in TNF-alpha- and obesity-induced insulin resistance. *Science*. 1996; 271:665–668. [PubMed: 8571133]
- Hotamisligil GS, Shargill NS, Spiegelman BM. Adipose expression of tumor necrosis factor-alpha: direct role in obesity-linked insulin resistance. *Science*. 1993; 259:87–91. [PubMed: 7678183]
- Hu FB. Globalization of Diabetes: The role of diet, lifestyle, and genes. *Diabetes Care*. 2011; 34:1249–1257. [PubMed: 21617109]
- Jiao P, Feng B, Ma J, Nie Y, Paul E, Li Y, Xu H. Constitutive activation of IKKbeta in adipose tissue prevents diet-induced obesity in mice. *Endocrinology*. 2012; 153:154–165. [PubMed: 22067324]
- Karin M. Inflammation-activated Protein Kinases as Targets for Drug Development. *Proceedings of the American Thoracic Society*. 2005a; 2:386–390. [PubMed: 16267367]
- Karin M. Inflammation-activated protein kinases as targets for drug development. *Proceedings of the American Thoracic Society*. 2005b; 2:386–390. discussion 394–385. [PubMed: 16267367]
- Kotterman MA, Schaffer DV. Engineering adeno-associated viruses for clinical gene therapy. *Nature reviews Genetics*. 2014; 15:445–451.
- Lancaster GI, Kammoun HL, Kraakman MJ, Kowalski GM, Bruce CR, Febbraio MA. PKR is not obligatory for high-fat diet-induced obesity and its associated metabolic and inflammatory complications. *Nature communications*. 2016; 7:10626.
- Lee J, Ozcan U. Unfolded protein response signaling and metabolic diseases. *The Journal of biological chemistry*. 2014; 289:1203–1211. [PubMed: 24324257]
- Lee J, Sun C, Zhou Y, Lee J, Gokalp D, Herrema H, Park SW, Davis RJ, Ozcan U. p38 MAPK-mediated regulation of Xbp1s is crucial for glucose homeostasis. *Nat Med*. 2011; 17:1251–1260. [PubMed: 21892182]
- McGillcuddy FC, Reynolds CM, Finucane O, Coleman E, Harford KA, Grant C, Sergi D, Williams LM, Mills KH, Roche HM. Long-term exposure to a high-fat diet results in the development of glucose intolerance and insulin resistance in interleukin-1 receptor I-deficient mice. *American journal of physiology Endocrinology and metabolism*. 2013; 305:E834–844. [PubMed: 23921145]

- Nakamura T, Furuhashi M, Li P, Cao H, Tuncman G, Sonenberg N, Gorgun CZ, Hotamisligil GS. Double-stranded RNA-dependent protein kinase links pathogen sensing with stress and metabolic homeostasis. *Cell*. 2010; 140:338–348. [PubMed: 20144759]
- O’Dea E, Hoffmann A. NF- $\kappa$ B signaling. *Wiley interdisciplinary reviews Systems biology and medicine*. 2009; 1:107–107. [PubMed: 20151024]
- Olefsky JM, Glass CK. Macrophages, inflammation, and insulin resistance. *Annual review of physiology*. 2010; 72:219–246.
- Park SW, Ozcan U. Potential for therapeutic manipulation of the UPR in disease. *Seminars in immunopathology*. 2013; 35:351–373. [PubMed: 23572207]
- Park SW, Zhou Y, Lee J, Lu A, Sun C, Chung J, Ueki K, Ozcan U. The regulatory subunits of PI3K, p85alpha and p85beta, interact with XBP-1 and increase its nuclear translocation. *Nature medicine*. 2010; 16:429–437.
- Preiss D, Seshasai SR, Welsh P, Murphy SA, Ho JE, Waters DD, DeMicco DA, Barter P, Cannon CP, Sabatine MS, et al. Risk of incident diabetes with intensive-dose compared with moderate-dose statin therapy: a meta-analysis. *Jama*. 2011; 305:2556–2564. [PubMed: 21693744]
- Rehman KK, Trucco M, Wang Z, Xiao X, Robbins PD. AAV8-mediated gene transfer of interleukin-4 to endogenous beta-cells prevents the onset of diabetes in NOD mice. *Molecular therapy : the journal of the American Society of Gene Therapy*. 2008; 16:1409–1416. [PubMed: 18560422]
- Sadagurski M, Norquay L, Farhang J, D’Aquino K, Copps K, White MF. Human IL6 enhances leptin action in mice. *Diabetologia*. 2010; 53:525–535. [PubMed: 19902173]
- Senftleben U, Karin M. The IKK/NF-kappa B pathway. *Critical care medicine*. 2002; 30:S18–26.
- Shoelson SE, Lee J, Goldfine AB. Inflammation and insulin resistance. *The Journal of clinical investigation*. 2006; 116:1793–1801. [PubMed: 16823477]
- Walter P, Ron D. The unfolded protein response: from stress pathway to homeostatic regulation. *Science*. 2011; 334:1081–1086. [PubMed: 22116877]
- Wernstedt Asterholm I, Tao C, Morley TS, Wang QA, Delgado-Lopez F, Wang ZV, Scherer PE. Adipocyte inflammation is essential for healthy adipose tissue expansion and remodeling. *Cell Metab*. 2014; 20:103–118. [PubMed: 24930973]
- Wild S, Roglic G, Green A, Sicree R, King H. Global prevalence of diabetes: estimates for the year 2000 and projections for 2030. *Diabetes Care*. 2004; 27:1047–1053. [PubMed: 15111519]
- Xu Y, Wang L, He J, Bi Y, Li M, Wang T, Wang L, Jiang Y, Dai M, Lu J, et al. Prevalence and control of diabetes in Chinese adults. *Jama*. 2013; 310:948–959. [PubMed: 24002281]
- Zhang X, Zhang G, Zhang H, Karin M, Bai H, Cai D. Hypothalamic IKKbeta/NF-kappaB and ER stress link overnutrition to energy imbalance and obesity. *Cell*. 2008; 135:61–73. [PubMed: 18854155]
- Zhang Y, Liu CM, Cao XC, Zang Y, Zhou YB, Li J. Involvement of transcription factor XBP1s in the resistance of HDAC6 inhibitor Tubastatin A to superoxidation via acetylation-mediated proteasomal degradation. *Biochemical and biophysical research communications*. 2014; 450:433–439. [PubMed: 24909686]
- Zhou Y, Lee J, Reno CM, Sun C, Park SW, Chung J, Lee J, Fisher SJ, White MF, Biddinger SB, et al. Regulation of glucose homeostasis through a XBP-1-FoxO1 interaction. *Nature medicine*. 2011; 17:356–365.



**HIGHLIGHTS**

- IKK $\beta$  phosphorylates XBP1s on at least on two residues.
- Phosphorylation of XBP1s increases its stability.
- Increased hepatic activity of IKK $\beta$  does not cause insulin resistance.
- IKK $\beta$  is beneficial for glucose homeostasis.



(E) HEK293 cells transfected with XBP1s alone or together with ca-IKK $\beta$  and further treated with cycloheximide (CHX, 20  $\mu\text{g ml}^{-1}$ ) for the indicated time periods. XBP1s and tubulin protein levels were analyzed with immunoblotting. The blots from the cells transfected with XBP1s only were exposed for a longer time (left bands) and the blots from cells transfected with XBP1s and ca-IKK $\beta$  together were exposed for a shorter time (right bands) to achieve comparable autoradiographic signals for total XBP1s from both groups at the zero time point.

The results represented in A-E were reproduced in three independent experiments.

(F) The graph depicts the averaged ratio of the autoradiographic signals of XBP1s to tubulin levels in HEK293 cells from three independent experiments. These three independent experiments are shown in **Figure 1E**, Figure S1M and S1N. The dotted lines show the half-life of XBP1s protein.

(G) 293T cells were transfected with XBP1s-flag and Ub-myc with or without ca-IKK $\beta$ -HA. Ubiquitinated XBP1s levels in HEK293 (left); The global ubiquitination levels in HEK293 cells were detected by western blotting using anti-Ub antibody (right). These results were reproduced in four independent experiments.

(H) XBP1s, IKK $\beta$ , p65 and tubulin protein levels in HEK293 cells, which were transfected with XBP1s alone or together with ca-IKK $\beta$ , after transfection with p65 and control siRNA. These results were reproduced in two independent experiments.

Error bars are represented as mean  $\pm$  SEM. Significance was determined by student's *t* test (B), one-way analysis of variance (ANOVA) (C), and two-way analysis of variance (ANOVA) with Bonferroni multiple-comparison analysis (F). \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001, NS: Nonsignificant.



(E) XBP1s and tubulin protein levels in HEK293 cells transfected with XBP1s and further treated with DMSO or TPCA-1 (20  $\mu$ M) for 2 hours following incubating with vehicle or TNF $\alpha$  (100 ng ml<sup>-1</sup>) for 30 minutes.

(F) XBP1s, IKK $\beta$  and tubulin protein levels in HEK293 cells transfected with the indicated vectors.

(G) Phosphorylated XBP1s levels in HEK293 transfected with the indicated vectors. We specifically loaded three times more amounts of IP samples from XBP1s-transfected cells when compared to the XBP1s-ca-IKK $\beta$  co-transfected cells to achieve comparable total XBP1s autoradiographical signal from each sample.

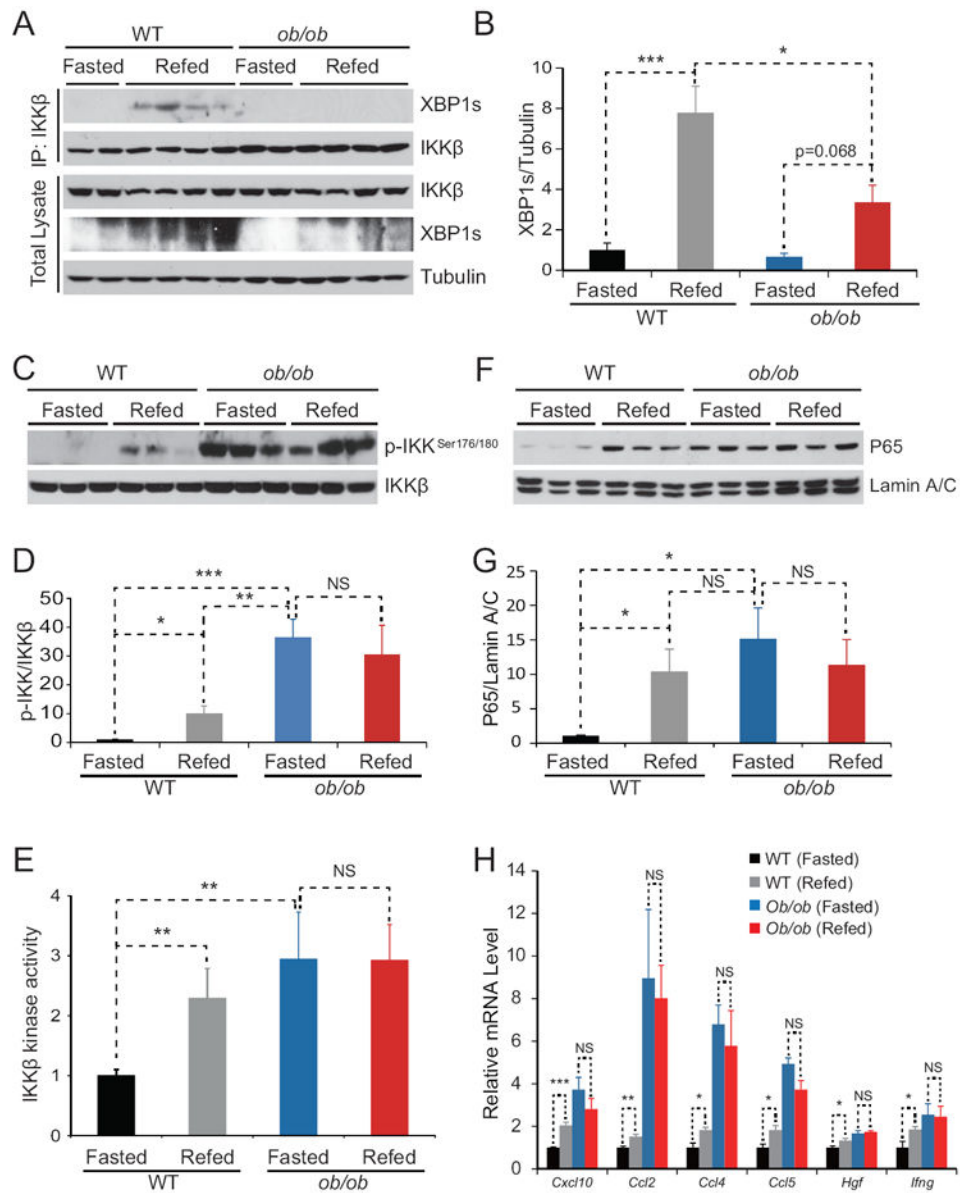
(H) Phospho-XBP1s levels after IP of XBP1s from IKK $\beta$  kinase assay.

(I) Phospho-XBP1s<sup>Thr48</sup> and total XBP1s protein levels in HEK293 cells infected with the indicated adenovirus.

(J) Densitometric quantification of ratio of Phospho-XBP1s<sup>Thr48</sup> to total XBP1s protein in **Figure 2I**.

Experiments in A, B, D and I were reproduced in three and in C, E, F, G and H were reproduced in two independent experiments.

Error bars are represented as mean  $\pm$  SEM. Significance was determined by student's *t* test. \*\*P < 0.01, \*\*\*P < 0.001.



**Figure 3. IKK $\beta$  Activation Response to Refeeding is Impaired in *Ob/Ob* Mice**

Lean and age-matched *ob/ob* mice were fasted for 24 hours and refeed for 1 hour.

(A) XBP1s Immunoblotting in the IKK $\beta$  immunoprecipitates from the liver lysates (top). We specifically loaded two times more amounts of IP samples from the liver of *ob/ob* mice liver when compared to those of lean mice. XBP1s and tubulin protein levels in total liver lysates (bottom).

(B) Densitometric quantification of ratio of XBP1s protein level to tubulin protein in **Figure 3A**.

(C) Phospho-IKK $\beta^{\text{Ser176/180}}$  and IKK $\beta$  protein levels in total liver lysates.

(D) Densitometric quantification of ratio of Phospho-IKK $\beta^{\text{Ser176/180}}$  level to IKK $\beta$  protein level in **Figure 3B** and Figure S3A (n=6 for each group).

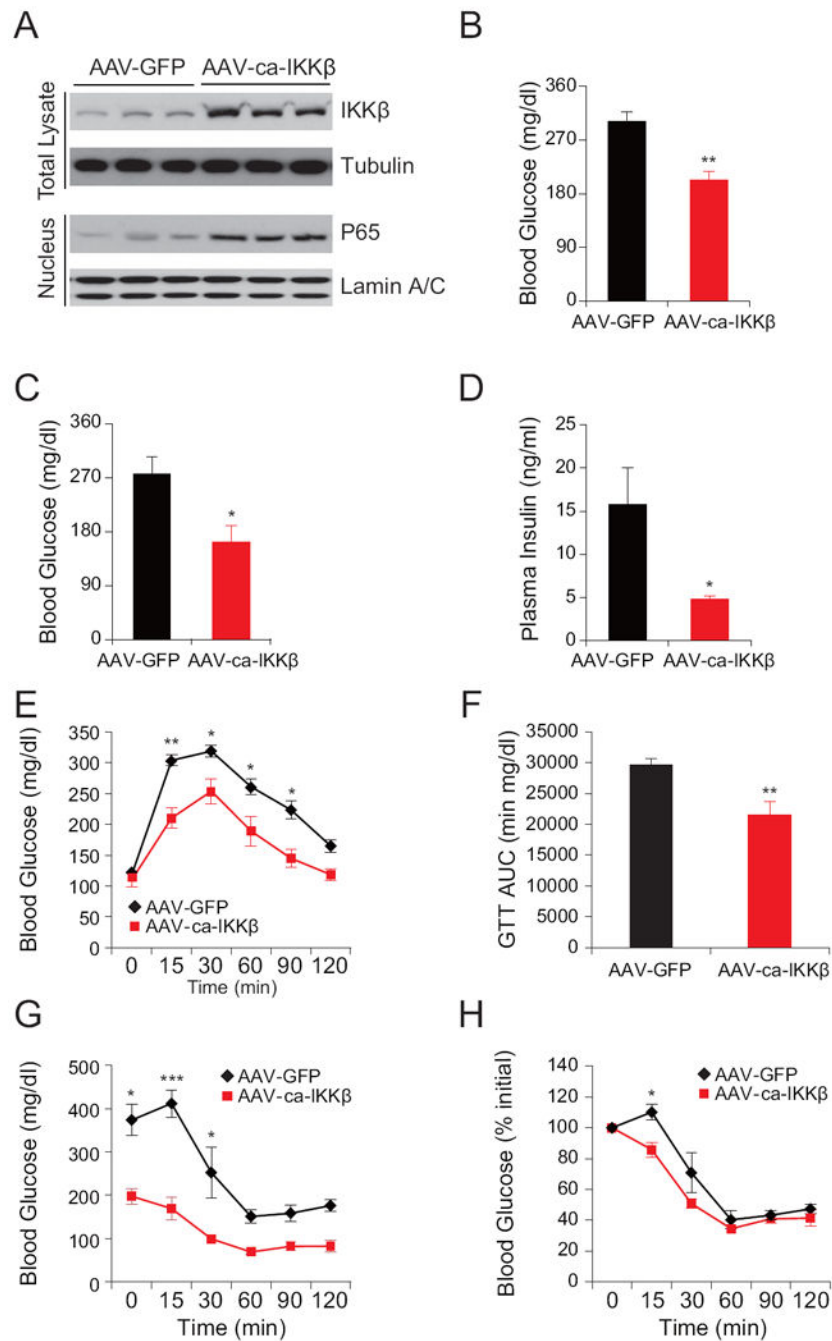


**(E)** IKK $\beta$  activity in mice livers (n=8 for each group). The average of the signals of IKK $\beta$  activities in the liver of fasted WT mice were used for normalization of IKK $\beta$  activities in the liver in other groups. The IKK $\beta$  activities in the four groups are expressed as fold difference.

**(F)** p65 and Lamin A/C protein levels in nuclear extracts of the liver of WT and *ob/ob* mice.

**(G)** Densitometric quantification of the ratio of p65 protein level to Lamin A/C protein in **Figure 3E**, Figure S3B and S3C (n=9 for each group).

**(H)** Gene expression levels of *Cxcl10*, *Ccl2*, *Ccl4*, *Ccl5*, *Hgf* and *Ifng* in the liver of WT and *ob/ob* mice (n=5 for each groups). 18S was used for normalization of gene expression. Error bars are represented as mean  $\pm$  SEM. Significance was determined by student's *t* test. \*P < 0.05 \*\*P < 0.01, \*\*\*P < 0.001, NS: Non-significance.



#### Figure 4. IKKβ Gain of Function in the Liver of *Ob/ob* Mice is Beneficial

Male *ob/ob* mice (8 weeks old) were injected with AAV-ca-IKKβ or AAV-GFP at a dose of  $2.4 \times 10^9$  genome copies (GC)  $g^{-1}$  ( $n=5$  in each group) through the tail vein.

(A) IKKβ and tubulin protein levels in total lysates and p65 and Lamin A/C protein levels in nuclear extracts of the liver on post-injection day 30.

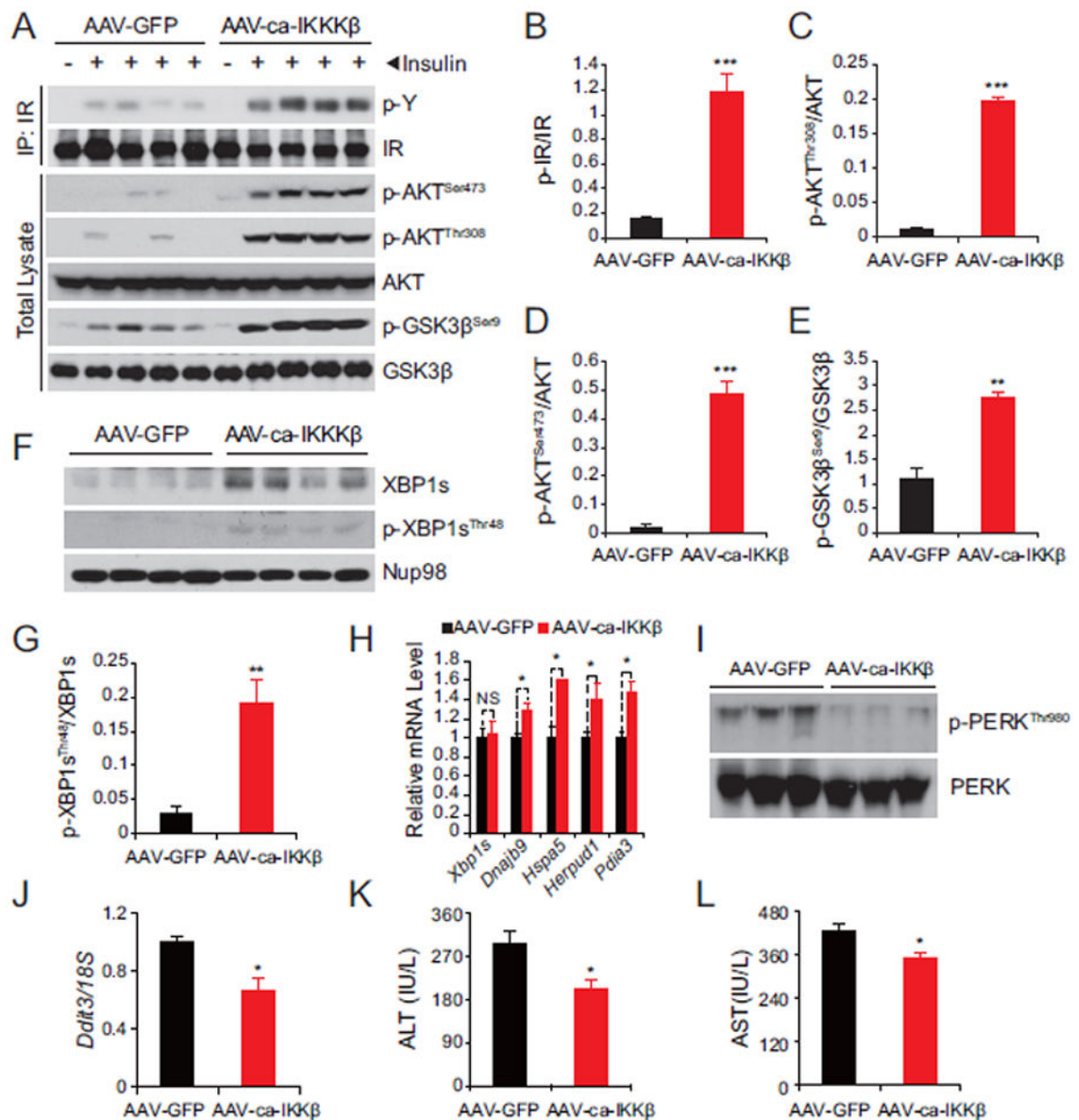
(B-C) Blood glucose levels (mg  $dl^{-1}$ ) (B) at fed state and (C) after 6 hours of fasting on day 5 post injection.

**(D)** Plasma insulin levels ( $\text{ng ml}^{-1}$ ) on post injection day 24. Each experiment was repeated in two independent groups.

**(E)** GTT on post injection day 11. **(F)** Area under curve (AUC) analysis of GTT from **Figure 4E**.

**(G-H)** ITT performed 13 day after the AAV injection. **(G)** The curve depicts the absolute blood glucose ( $\text{mg dl}^{-1}$ ) levels, **(H)** Percent values of initial blood glucose concentration during ITT.

Each experiment was reproduced in four independent groups. Error bars are represented as mean  $\pm$  SEM. Significance was determined by student's *t* test (B-D and F) or two-way ANOVA with Bonferroni multiple-comparison analysis (E, G and H). \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



**Figure 5. IKK $\beta$  Reduces ER Stress in the Liver of *Ob/ob* Mice**  
 Male *ob/ob* mice (8 weeks old) were injected with AAV-ca-IKK $\beta$  and AAV-GFP at a dose of  $2.4 \times 10^9$  GC  $g^{-1}$  (n=5 in each group) through the tail vein.  
 (A) Protein lysates were subjected to IR-IP and subsequently phospho-tyrosine and IR immunoblotting. Direct lysates were used to analyze phospho AKT<sup>Ser473</sup>, AKT<sup>Thr308</sup> and GSK3 $\beta$ <sup>Ser9</sup> and the total protein levels of AKT and GSK3 $\beta$ .  
 (B-E) Densitometric quantifications of the western blots bands in (A). Graphs depict the ratio of the autoradiographical signals of (B) phospho-IR/total IR, (C) phospho-AKT<sup>Thr308</sup>/total AKT, (D) phospho-AKT<sup>Ser473</sup>/total AKT and (E) phospho-GSK3 $\beta$ <sup>Ser9</sup>/total GSK3 $\beta$ .  
 (F) Nuclear phospho-XBP1s<sup>Thr48</sup> and total XBP1s protein levels in the liver of *ob/ob* mice. Nup98 was used as a loading control.

**(G)** The graph depicts the ratio of the signals of phospho-XBP1s<sup>Thr48</sup> protein to XBP1s protein shown in **(F)**.

**(H)** Gene expression levels of *Xbp1s*, *Dnajb9*, *Hspa5*, *Herpud1* and *Pdia3* in livers of *ob/ob* mice.

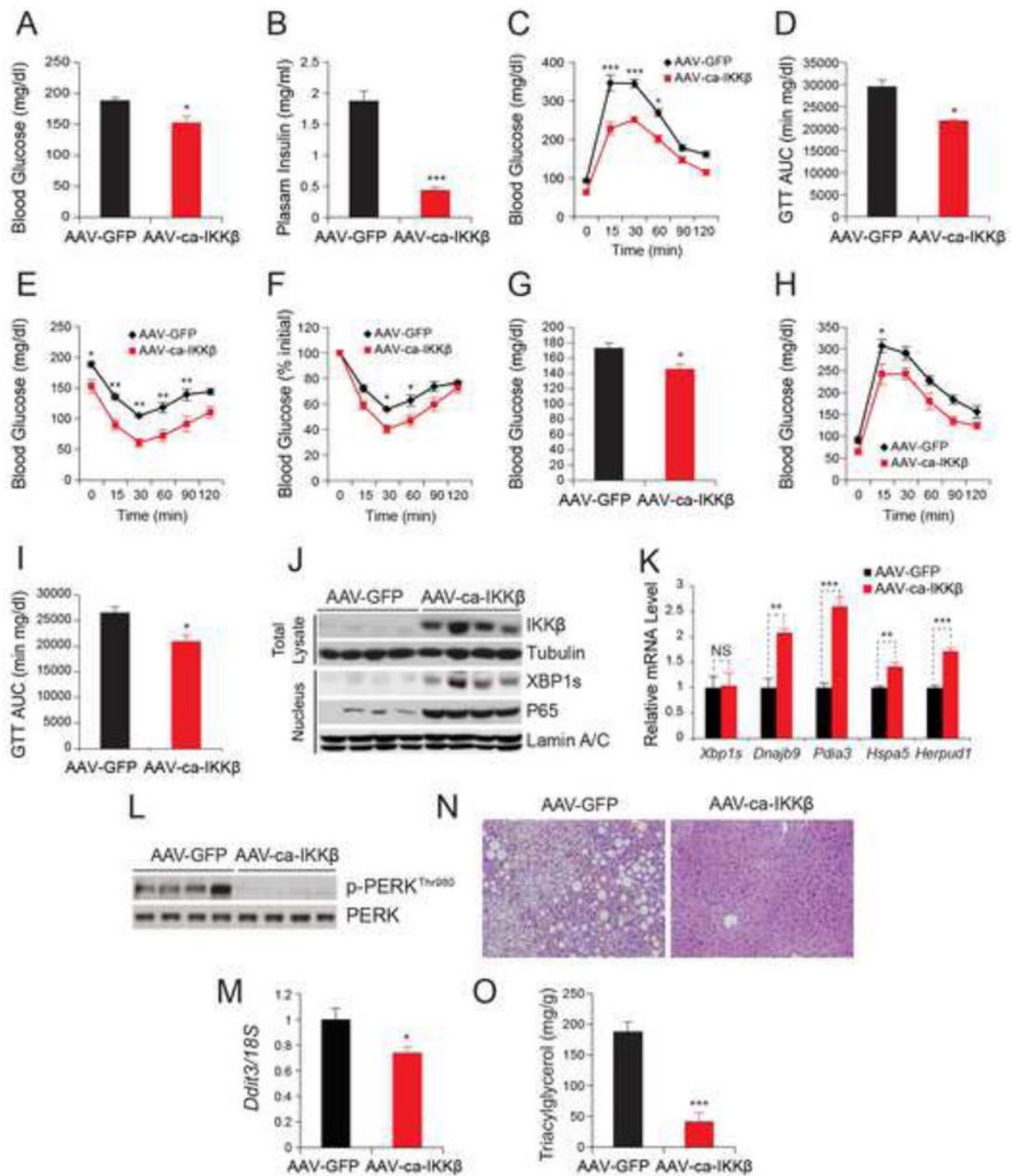
**(I)** PERK<sup>Thr980</sup> phosphorylation and total PERK protein levels in livers of *ob/ob* mice.

**(J)** *Ddit3* mRNA levels in livers of *ob/ob* mice. *18S* was used for normalization of gene expression.

**(K-L)**, **(K)** Plasma alanine transaminase (ALT; IU L<sup>-1</sup>) and **(L)** Aspartate transaminase (AST; IU L<sup>-1</sup>) of *ob/ob* mice 12 days after injection.

Error bars are represented as mean ± SEM. Significance was determined by student's *t* test.

\*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001, NS: Non-significance.



### Figure 6. IKK $\beta$ Improves Glucose Homeostasis in DIO Mice

The diet induced obese (DIO) mice were injected with AAV-ca-IKK $\beta$  or AAV-GFP at a dose of  $2.4 \times 10^9$  GC g $^{-1}$  (n=5 in each group) via the tail vein.

(A) Blood glucose levels (mg dl $^{-1}$ ) after 6 hours fasting on post injection day 21.

(B) Plasma insulin concentrations (ng ml $^{-1}$ ) on day 22 post injection.

(C) GTT on day 19 post injection.

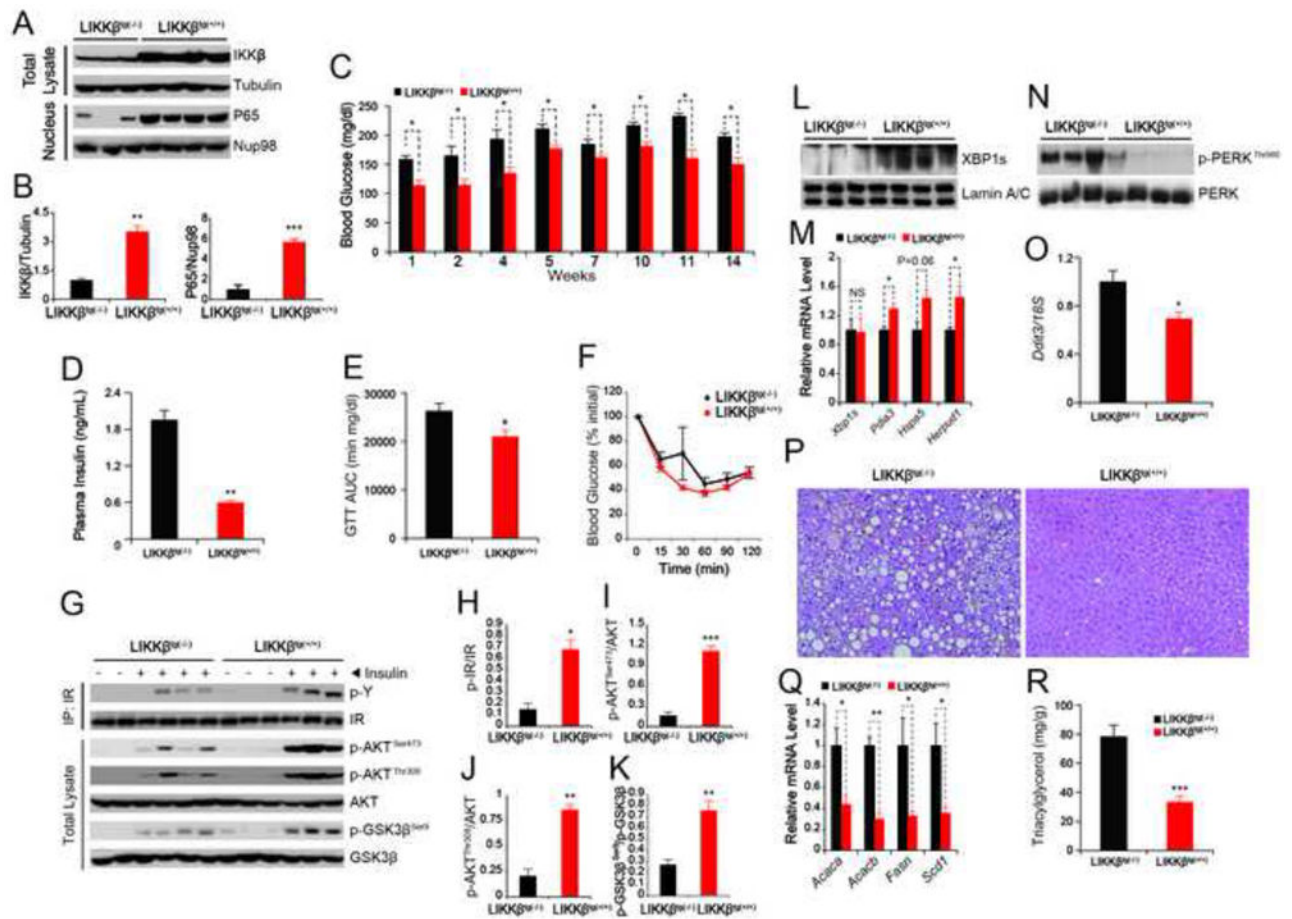
(D) Area under the curve of GTT in **Figure 6C**.

(E-F) ITT on day 21 post injection. The curves depict (E) the absolute blood glucose concentration and (F) the percentage of initial glucose concentration. Each experiment was reproduced in two independent groups.



- (G) Blood glucose levels ( $\text{mg dl}^{-1}$ ) at 6-hour fasted state 10 weeks after injection.
- (H) GTT 11 weeks after injection.
- (I) AUC analysis of GTT in **Figure 6H**.
- (J) IKK $\beta$  and tubulin protein levels in total lysates and XBP1s, p65 and Lamin A/C protein levels in nuclear extracts of the liver 14 weeks after injection.
- (K) Gene expression levels of *Xbp1s*, *Dnajb9*, *Hspa5*, *Herpud1* and *Pdia3* in livers of DIO mice 24 days after injection.
- (L) PERK<sup>Thr980</sup> phosphorylation and total PERK protein levels in livers of DIO mice 4 weeks after injection.
- (M) *Ddit3* mRNA levels in the liver of DIO mice 14 weeks after injection. 18S was used for normalization of gene expression.
- (N) H&E staining of liver sections from DIO mice 14 weeks after injection.
- (O) Triglyceride contents ( $\text{mg g}^{-1}$ ) in the liver 14 weeks after injection.

All Error bars are represented as mean  $\pm$  SEM. Significance was determined by student's *t* test (A, B, D, G, I, K, M and O) or two-way ANOVA with Bonferroni multiple-comparison analysis (C, E, F and H). \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001, NS: Non-significance.



### Figure 7. $LIKK\beta^{Tg(+/+)}$ Mice Have Improved Metabolic Homeostasis

$LIKK\beta^{Tg(+/+)}$  mice and control mice  $LIKK\beta^{Tg(-/-)}$  were fed with a HFD (n=4 in  $LIKK\beta^{Tg(+/+)}$  group and n=5 in  $LIKK\beta^{Tg(-/-)}$  group) for 14 weeks.

(A) IKK $\beta$  and tubulin protein levels in total liver lysates, and p65 and Lamin A/C protein levels in nuclear extracts of the liver.

(B) Densitometric quantifications of the western blot showing the ratio of IKK $\beta$  to tubulin and p65 to Nup98 in **Figure 7A**.

(C) Blood glucose levels (mg dl<sup>-1</sup>) at 6-hour fasting after at indicated weeks of HFD-feeding.

(D) Plasma insulin concentrations (ng ml<sup>-1</sup>) after 12 weeks of HFD-feeding.

(E) The graph depicts AUC analysis of the GTTs from three independent groups of  $LIKK\beta^{Tg(+/+)}$  and their corresponding controls after 8 weeks of HFD-feeding (n=14 in  $LIKK\beta^{Tg(+/+)}$  group and n=18 in  $LIKK\beta^{Tg(-/-)}$  group). For each GTT data and analysis of the three independent groups, see Figure S6B-S6D.

(F) The graph depicts ITT (percent values) results from three independent groups of  $LIKK\beta^{Tg(+/+)}$  and their corresponding controls after 6 weeks of HFD-feeding (n=14 in  $LIKK\beta^{Tg(+/+)}$  group and n=18 in  $LIKK\beta^{Tg(-/-)}$  group). For each ITT data and analysis of the absolute blood glucose levels of the three independent groups, see Figure S6E-S6G.

(G) Phospho-tyrosine and IR levels in IR-IP samples. Phospho AKT<sup>Ser473</sup>, AKT<sup>Thr308</sup> and GSK3 $\beta$ <sup>Ser9</sup> and the total protein levels of AKT and GSK3 $\beta$  were determined from the direct lysates of the liver of mice, which was infused with insulin through portal vein.

(H-K) Densitometric quantifications of the western blot bands in **Figure 7G**. Graphs depict the ratios of the autoradiographic signals of (H) phospho-IR/total IR, (I) phospho-AKT<sup>Thr308</sup>/total AKT, (J) phospho-AKT<sup>Ser473</sup>/total AKT and (K) phospho-GSK3 $\beta$ <sup>Ser9</sup>/total GSK3 $\beta$ .

(L) XBP1s and Lamin A/C protein levels in the nuclear extracts of the liver of LIKK $\beta$ <sup>Tg(+/+)</sup> and LIKK $\beta$ <sup>Tg(-/-)</sup> mice.

(M) Gene expression levels of *Xbp1s*, *Pdia3*, *Hspa5* and *Herpud1* in the liver of LIKK $\beta$ <sup>Tg(+/+)</sup> and LIKK $\beta$ <sup>Tg(-/-)</sup> mice.

(N) PERK<sup>Thr980</sup> phosphorylation and total PERK protein levels in the liver of LIKK $\beta$ <sup>Tg(+/+)</sup> and LIKK $\beta$ <sup>Tg(-/-)</sup> mice.

(O) *Ddit3* mRNA levels in livers of LIKK $\beta$ <sup>Tg(+/+)</sup> and LIKK $\beta$ <sup>Tg(-/-)</sup> mice after 14 weeks of HFD-feeding. 18S was used for normalization of gene expression.

(P) H&E staining of liver sections from LIKK $\beta$ <sup>Tg(+/+)</sup> and LIKK $\beta$ <sup>Tg(-/-)</sup> mice.

(Q) Gene expression levels of *Acaca*, *Acacb*, *Fasn* and *Scd1* in the liver of LIKK $\beta$ <sup>Tg(+/+)</sup> and LIKK $\beta$ <sup>Tg(-/-)</sup> mice.

(R) Triglyceride level in the liver of LIKK $\beta$ <sup>Tg(+/+)</sup> and LIKK $\beta$ <sup>Tg(-/-)</sup> mice.

Error bars are represented as mean  $\pm$  SEM. Significance was determined by student's *t* test (B, C, D, E, F, H, I, J, K, M, O, Q and R) or two-way ANOVA with Bonferroni multiple-comparison analysis (E). \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001, NS: Non-significance.

## KEY RESOURCES TABLE

REAGENT or RESOURCE	SOURCE	IDENTIFIER
<i>Antibodies</i>		
Rabbit polyclonal anti-XBPIs	Santa Cruz Biotechnology	Cat#sc-7160
Rabbit polyclonal anti-insulin receptor (IR)	Santa Cruz Biotechnology	Cat#sc-711
Mouse monoclonal anti-phosphotyrosine (PY99)	Santa Cruz Biotechnology	Cat#sc-7020
Mouse monoclonal anti-myc	Santa Cruz Biotechnology	Cat#sc-40
Mouse monoclonal anti-HA antibodies	Santa Cruz Biotechnology	Cat#sc-7392
Rabbit polyclonal anti-ATF4	Santa Cruz Biotechnology	Cat#sc-200
Rabbit polyclonal anti-ATF6	Santa Cruz Biotechnology	Cat#sc-22799
Goat anti-mouse IgG-HRP	Santa Cruz Biotechnology	Cat#sc-2005
Goat anti-rabbit IgG-HRP	Santa Cruz Biotechnology	Cat#sc-2030
Rabbit polyclonal anti-IKK $\beta$	Cell Signaling Technology	Cat#2684
Rabbit polyclonal anti-IKK $\alpha$	Cell Signaling Technology	Cat#2682
Rabbit polyclonal anti-IKK $\gamma$	Cell Signaling Technology	Cat#2685
Rabbit polyclonal anti-p-JNK	Cell Signaling Technology	Cat#9251
Rabbit polyclonal anti-JNK2	Cell Signaling Technology	Cat#4672
Rabbit monoclonal anti-p-P38	Cell Signaling Technology	Cat#4631
Rabbit polyclonal anti-P38	Cell Signaling Technology	Cat#9212
Rabbit polyclonal anti-p-c-Jun	Cell Signaling Technology	Cat#9164
Rabbit monoclonal anti-c-Jun	Cell Signaling Technology	Cat#9165
Rabbit monoclonal anti-ATF2	Cell Signaling Technology	Cat#9226
Rabbit polyclonal anti-p-ATF2	Cell Signaling Technology	Cat#9225
Mouse monoclonal anti-ubiquitin	Cell Signaling Technology	Cat#3936
Rabbit monoclonal anti-phospho-AKT (Thr308)	Cell Signaling Technology	Cat#4056
Rabbit monoclonal anti-phospho-AKT (Ser473)	Cell Signaling Technology	Cat#4058
Rabbit polyclonal anti-AKT	Cell Signaling Technology	Cat#9272
Rabbit polyclonal anti-phospho-GSK3 $\alpha/\beta$ (Ser21/9)	Cell Signaling Technology	Cat#9331
Rabbit monoclonal anti-GSK3 $\beta$	Cell Signaling Technology	Cat#9315

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Rabbit monoclonal anti-phospho-PERK (Thr980)	Cell Signaling Technology	Cat#3179
Rabbit monoclonal PERK antibodies	Cell Signaling Technology	Cat#3192
Rabbit monoclonal anti-p65	Cell Signaling Technology	Cat#4764
Rabbit monoclonal anti-Nup98	Cell Signaling Technology	Cat#4283
Rabbit polyclonal anti-Lamin A/C	Cell Signaling Technology	Cat#2032
Rabbit monoclonal anti- $\alpha$ -tubulin	Cell Signaling Technology	Cat#2125
Mouse monoclonal Anti-Flag anti-His	Sigma Aldrich	Cat#F3165
Rabbit polyclonal anti-phosphoserine/threonine	Sigma Aldrich	SAB1305538
Rabbit polyclonal anti-phospho-XBP1s <sup>Thrs</sup>	ECM Biosciences	Cat#PP2551
Mouse monoclonal anti-XBP1s	Covance	Raised against CRAAGSEASGT(p)PQARRKQR
Mouse monoclonal anti-IRK $\beta$	Biologend	Cat#647502
Rabbit polyclonal anti-Phosphoserine/threonine	Novus	Cat#10AG2
Mouse monoclonal anti-IRK $\gamma$	Abcam	Cat#Ab17464
	BD Pharmingen	Cat#559675
<b>Chemicals, Peptides, and Recombinant Proteins</b>		
(aminocarboxylamino)-5-(4-fluorophenyl)-3-thiophenecarboxamide (TPCA-1)	Sigma Aldrich	T1452
Cycloheximide (CHX)	Sigma Aldrich	C1988
Recombinant Human TNF- $\alpha$ protein	R&D systems	210-TA
Recombinant human insulin	Eli Lilly	HI-210
Recombinant XBP1s	This paper	N/A
D-[3-3H]-glucose	PerkinElmer	NET331C250UC
<b>Deposited Data</b>		
<b>Experimental Models: Cell Lines</b>		
Human: Human embryonic kidney 293 (HEK293)	ATCC	CRL-1573
Hamster: CHO cells	ATCC	CRL-2092
Mouse: Hepa-1c1c7 (hepatoma cell)	ATCC	CRL-2026
Human: HepG2 (hepatoma cell)	ATCC	HB-8065
Human: MDA-MB-231 (breast adenocarcinoma cell)	ATCC	HTB-26
<b>Experimental Models: Organisms/Strains</b>		

REAGENT or RESOURCE	SOURCE	IDENTIFIER
<b>Mouse:</b>		
B6.Cg- <i>Lep<sup>fl/j</sup></i>	The Jackson Laboratory	Cat# JAX:000632
IKK $\beta^{fl/+}$	Michael Karin	N/A
B6.BKS(D)- <i>Lep<sup>fl/j</sup></i>	The Jackson Laboratory	Cat# JAX:000697
<b>Recombinant DNA</b>		
Plasmid: pCR-Flag-IKK $\beta$ (mouse)	Addgene	Cat#15465
Plasmid: pcDNA3.1-IKK $\beta$ (S177E/S181E)-pcw107	Addgene	Cat#64609
Plasmid: Human XBPIs plasmid	Addgene	Cat#63680
Plasmid: pcDNA3.1-Flag-XBPIs(mouse)	Jaemin L., et al., 2011	N/A
Plasmid: pCR-Flag-IKK $\beta$ -KM(mouse)	Michael Karin	N/A
Plasmid: Vector expressing NH2-terminal (1-373 aa) of activating transcription factor 6n (ATF6n)	Ozcan L., et al., 2009	N/A
Plasmid: pcDNA3.1-Flag-XBPIs(T48A)	This paper	N/A
Plasmid: pcDNA3.1-Flag-XBPIs(S148A)	This paper	N/A
Plasmid: pcDNA3.1-Flag-XBPIs(T48A/S148A)	This paper	N/A
Ad-ca-IKK $\beta$	Michael Karin	N/A
Ad-XBPIs	This paper	N/A
AAV-GFP	Gene Therapy Center and Vector Core, University of Massachusetts Medical School	N/A
AAV-ca-IKK $\beta$	This paper	N/A
<b>Sequence-Based Reagents</b>		
Primers for Realtime analysis, see Table S1	This paper	N/A
Primer: sequences for amino acid substitution of IKK $\beta$ (S177E/S181E): (forward), 5'-GGAGCTGGATCAGGGCGAGCTGTGCACGGAATTTGTGGGACTCTGC-3'; (reverse), 5'-GCAGAGTCCCCACAAATCCGTGCACAGCTCGCCCTGATCCAGCTCC-3'.	This paper	N/A
Primer: sequences for amino acid substitution of XBPI(T48A): (forward), 5'-GGGTCGGAGGGGAGCGGGGACCCGAGGCTCGCAAGCGG-3'; (reverse), 5'-CCGCTTGCGAGCCTGCGGTGCCCCCGCTCGCCTCCGACCC-3'.	This paper	N/A
Primer: sequences for amino acid substitution of XBPI (S148A):	This paper	N/A



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REAGENT or RESOURCE	SOURCE	IDENTIFIER
(forward), 5'-CACGCTTGGGAATGGACGCGCTGGATCCCTGACGAGG-3'; (reverse), 5'-CCTCGTCAGGATCCAGCGCGTCCATTCCCAAGCGTG-3'.		
<i>Software and Algorithms</i>		
Image J	<a href="https://imagej.nih.gov/ij/">https://imagej.nih.gov/ij/</a>	N/A
GraphPad Prism 6	Purchased by Boston Children Hospital	N/A
<i>Other</i>		