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Postprandial Hepatic Lipid Metabolism Requires Signaling through Akt2 Independent of the Transcription Factors FoxA2, FoxO1, and SREBP1c

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SUMMARY

Under conditions of obesity and insulin resistance, the serine/threonine protein kinase Akt/PKB is required for lipid accumulation in liver. Two forkhead transcription factors, FoxA2 and FoxO1, have been suggested to function downstream of and to be negatively regulated by Akt and are proposed as key determinants of hepatic triglyceride content. In this study, we utilize genetic loss of function experiments to show that constitutive activation of neither FoxA2 nor FoxO1 can account for the protection from steatosis afforded by deletion of Akt2 in liver. Rather, another downstream target positively regulated by Akt, the mTORC1 complex, is required in vivo for de novo lipogenesis and Srebp1c expression. Nonetheless, activation of mTORC1 and SREBP1c is not sufficient to drive postprandial lipogenesis in the absence of Akt2. These data show that insulin signaling through Akt2 promotes anabolic lipid metabolism independent of Foxa2 or FoxO1 and through pathways additional to the mTORC1-dependent activation of SREBP1c.

INTRODUCTION

Insulin is the primary hormone responsible for coordinating the metabolic response to nutrient intake. Following a meal, insulin suppresses the hepatic glucose production that provides the brain substrate during fasting and directs ingested nutrients into long-term energy stores. In particular, insulin promotes triglyceride (TG) synthesis and storage utilizing dietary fatty acids and induces de novo lipogenesis in liver and adipose tissue. Much of this synchronized metabolic response is thought to be mediated by hormone-dependent reduction in the transcription of gluconeogenic genes, such as glucose-6-phosphatase, catalytic subunit (G6pc), and cytosolic phosphoenolpyruvate carboxykinase 1 (Pck1), and activation of lipogenic gene expression via the master regulator sterol regulatory element-binding transcription factor 1c (Srebp1c) (Argaud et al., 1996; Horton et al., 2002; O’Brien et al., 2001). Thus, the transcriptional targets of the insulin signaling pathway are of considerable interest.

The serine/threonine kinase Akt (also known as protein kinase B, PKB) is now well-established as a major mediator of the metabolic actions of insulin (Whiteman et al., 2002). Akt2 is the predominant isoform in the insulin-responsive tissues like liver, muscle, and adipose tissue (Cho et al., 2001). Akt2 knockout mice are “diabetic,” whereas liver-specific disruption of the Akt2 gene has mild effects on glycemia but more dramatic effects on lipogenic gene expression and steatosis in obese, insulin-resistant mice (Cho et al., 2001; Leavens et al., 2009). The canonical nuclear target of Akt, established in worms and flies as well as vertebrates, is the transcription factor Foxhead box O (FoxO) (Gross et al., 2009). Akt phosphorylates and inhibits the transcriptional activity of FoxO at least in part by promoting its exclusion from the nucleus (Durham et al., 1999; Guo et al., 1999; Nakae et al., 1999; Rena et al., 1999; Tang et al., 1999). There are now considerable data implicating FoxO1 as an intermediate in the pathway by which insulin through Akt suppresses hepatic gluconeogenic gene expression and glucose output (Haeusler et al., 2010b; Li et al., 2007; Liu et al., 2008; Matsumoto et al., 2007; Nakae et al., 2001; Puigserver et al., 2003; Qu et al., 2006).

In contrast to the consensus of opinion regarding the role of FoxO1 in the control of gluconeogenic gene expression, its contribution to regulation of hepatic lipid metabolism remains unclear. In some studies, expression of a constitutively active FoxO1 in liver induces Srebp1c expression and hepatic TG accumulation, though others have not found this response; in addition, FoxO1 inhibits TG secretion (Matsumoto et al., 2006; Zhang et al., 2006). In contrast, Kamagate et al. reported that FoxO1 is necessary and sufficient to promote hepatic very-low-density lipoprotein-associated TG (VLDL-TG) production and hypertriglyceridemia via its regulation of microsomal TG transfer protein (MTP), and further proposed that this pathway accounts for...
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Akt2 Regulates Hepatic TG Accumulation Independent of FoxA2

It has been proposed that insulin inhibits FoxA2 via Akt-dependent phosphorylation and nuclear exclusion, controlling lipid export, fatty acid oxidation, and accumulation of TG in the liver (Wolfrum et al., 2004, 2008; Wolfrum and Stoffel, 2006). We generated mice lacking both Akt2 and FoxA2 in liver by breeding Akt2(loxP/loxP);FoxA2(loxP/loxP) mice to AFP-Cre, which is expressed exclusively in liver (Zhang et al., 2005). As shown in Figure 1A, we achieved essentially complete deletion of Akt2 in both Akt2(loxP/loxP);FoxA2(loxP/loxP) and Akt2(loxP/loxP);FoxA2(loxP/loxP) mice. FoxA2 protein levels were also greatly reduced in both FoxA2 single-knockout and Akt2, FoxA2 double-knockout hepatocytes (Figure 1A). Mice of all four genotypes were submitted to glucose tolerance tests, which showed indistinguishable rates of glucose disposal (see Figure S1A available online). In addition, there were no significant changes in serum insulin or TG levels from either fed or overnight-fasted mice lacking Akt2, FoxA2, or both in liver (Figures S1B and S1C). Adipose tissue mass was the same among the mice of all four genotypes (Figure S1D).

The TG content in livers deficient in Akt2, FoxA2, or both was unchanged compared to wild-type controls when assayed in mice under chow-fed conditions (Figure 1B). Akt2 is required for accumulation of liver TG under some situations that induce hepatic steatosis, and expression of a constitutively active FoxA2 reduces hepatic TG accumulation in leptin-deficient (ob/ob) mice (Leavens et al., 2009; Wolfrum et al., 2004; Wolfrum and Stoffel, 2006). To address the question of whether endogenous FoxA2 is responsible for the reduction in hepatic TG in livers deficient in Akt2, mice lacking Akt2, FoxA2, or both in liver were fed a high-fat diet (HFD) for 3 months. Glucose tolerance tests were indistinguishable among the four genotypes (Figure S2A), as were serum insulin and TG levels under both fasting and fed conditions (Figures S2B and S2C). Loss of Akt2 in livers trended to a decrease of hepatic TG accumulation relative to wild-type controls after HFD feeding, and this was unaffected by deletion
of FoxA2 (Figure 1C). These data show that, at least under conditions of diet-induced obesity (DIO), FoxA2 activity is unlikely to be the cause of the protection from steatosis rendered by removal of Akt2 from liver. These data are also consistent with previous findings challenging the notion of FoxA2 as a target of insulin signaling (Zhang et al., 2005).

**Loss of Akt2 Suppresses Hepatic TG Levels in FoxO1-Deficient Liver**

FoxO1, a transcription factor negatively regulated by the insulin/Akt signaling pathway, promotes expression of gluconeogenic genes such as G6pc and Pck1 (Gross et al., 2009). To assess the function of FoxO1 as an independent target of Akt2, we derived Akt2, FoxO1 and combined liver-specific double-knockout mice. Akt2 and FoxO1 protein levels were significantly reduced in Akt2 and FoxO1 single-knockout livers, respectively, and in Akt2, FoxO1 double-knockout livers (Figure 2A, left), and real-time PCR from either FoxO1 knockout livers or Akt2, FoxO1 double-knockout livers showed reduction of FoxO1 mRNA levels compared to Akt2 liver under normal chow feeding (Figure 2B). Under conditions of normal chow feeding, both Akt2 and FoxO1 mice showed lower levels of blood glucose, as did Akt2 and FoxO1 mice (Figure 2B). These data show that, at least under conditions of diet-induced obesity (DIO), FoxA2 activity is unlikely to be the cause of the protection from steatosis rendered by removal of Akt2 from liver.
from wild-type controls (Figures 2B and 2C). These data are consistent with previously published results that support a model in which FoxO1 regulates glucose output and is downstream of Akt (Matsumoto et al., 2007).

After an overnight fast, AFP > Cre;Akt2(loxP/loxP), AFP > Cre;FoxO1(loxP/loxP) and AFP > Cre;Akt2(loxP/loxP);FoxO1(loxP/loxP) mice had hepatic TG levels indistinguishable from that in wild-type controls (Leavens et al., 2009; Matsumoto et al., 2007) (Figure 2D). Fed AFP > Cre;FoxO1(loxP/loxP) mice displayed increased hepatic TG levels compared to controls and AFP > Cre;Akt2(loxP/loxP);FoxO1(loxP/loxP) mice, though TG secretion was unaltered (Figure 2E and Figure S4). Surprisingly, the accumulation of liver TG in the AFP > Cre;FoxO1(loxP/loxP) mice was suppressed by concomitant deletion of Akt2 (Figure 2E). Though deletion of FoxO1 leads to a phenotype opposite to that of deletion of Akt2, these data are inconsistent with a model in which Akt2 regulates lipid accumulation through suppression of FoxO1 activity.

Akt2 Regulates Hepatic TG Levels Independent of FoxO1 upon HFD Feeding

Suppression of FoxO1 by shRNA in livers from DIO mice decreases blood glucose and insulin levels and improves insulin sensitivity as assessed by the euglycemic, hyperinsulinemic clamp (Samuel et al., 2006). To investigate the role of FoxO1 as a downstream mediator of Akt’s role in metabolism under pathological conditions, we placed the wild-type controls, AFP > Cre;Akt2(loxP/loxP), AFP > Cre;FoxO1(loxP/loxP) and AFP > Cre;Akt2(loxP/loxP);FoxO1(loxP/loxP) mice, on a HFD for 3 months. Glucose tolerance tests were indistinguishable in mice of all four genotypes (Figure 3A). Fasting and fed serum insulin and TG levels were also similar among these same mice (Figures 3B and 3C). However, under fed conditions, loss of hepatic Akt2 in HFD-fed mice trended to a lower TG level in liver, whereas liver TG upon ablation of FoxO1 did not differ from wild-type controls (Figure 3D). Deletion of FoxO1 concomitant with Akt2 did not reverse the protection from steatosis in AFP > Cre;Akt2(loxP/loxP) mice (Figure 3D), again militating against the idea that FoxO1 regulates lipid metabolism downstream of Akt2.

Loss of Akt2 Decreases Hepatic TG Levels in FoxO1 Knockout Mice after Aurothioglucose Treatment

In DIO, deletion of Akt2 in liver reduces hepatic TG without altering lipogenesis, whereas in ob/ob mice protection from steatosis is accompanied by a concomitant reduction in lipogenesis and lipogenic gene expression (Leavens et al., 2009). To study the AFP > Cre;Akt2(loxP/loxP);FoxO1(loxP/loxP) mice under conditions that mimic ob/ob mice, we injected mice with aurothioglucose (goldthioglucose, GTG), which causes hypothalamic damage, hyperphagia, and weight gain (Brecher et al., 1965; Marshall and Mayer, 1954). As shown in Figure 4A, 8 weeks after GTG injection, mice of all genotypes displayed significantly increased body weight and hepatic TG levels compared to age-matched controls (compare Figure 4B to Figure 2E). AFP > Cre;Akt2(loxP/loxP) mice trended to lower hepatic TG, while AFP > Cre;FoxO1(loxP/loxP) mice had significantly elevated TG in liver (Figure 4B). Interestingly, concomitant deletion of Akt2 and...
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In order to address the pathways downstream of Akt2 that mediate lipid homeostasis, we sought to establish a model more physiological than hypothalamic obesity that would test the dependency of hepatic lipogenesis on Akt2. To accomplish this, mice were fasted overnight and then refed a lipogenic high-carbohydrate diet (HCD). As expected, blood glucose and serum insulin levels increased by 1.5 hr after HCD refeeding (Figure 5A and 5B). Liver weights normalized to compared to wild-type controls, indicating the minimal requirement for Akt2 in hepatic lipogenesis versus glycemia (Figure 5D). These data point to a discrepancy between the requirements for Akt2 in hepatic lipogenesis versus glycemia that in primary hepatocytes, insulin utilizes a pathway independent of FoxO1 (Nakae et al., 2001) into Akt2-null livers (data not shown). These data are consistent with those presented above in which Akt2 is epistatic to FoxO1 regarding the control of liver TG content.

Akt2 Is Required for Postprandial De Novo Lipogenesis

We next turned to the question: since FoxO1 and FoxA2 do not appear to function downstream of Akt2 in lipid homeostasis, what pathway could mediate Akt2’s effect on lipogenic gene expression? The most likely answer lies in the recent observation that in primary hepatocytes, insulin utilizes a pathway dependent on mTORC1 to stimulate the processing and expression of SREBP1c, a transcription factor that drives coordinated expression of lipogenic genes (Azzout-Marniche et al., 2000; Chakrabarti et al., 2010; Duvel et al., 2010; Fleischmann and Iyedjian, 2000; Li et al., 2010; Porstmann et al., 2008). To test the requirement for this pathway in vivo, we injected raptorKoxPloxP mice with an adeno-associated virus expressing liver-specific Cre recombinase (AAV > Cre) or green fluorescent protein (AAV > GFP), thus generating hepatic raptor knockout mice or controls. As shown in Figure 7A, raptor protein levels were substantially reduced in AAV > Cre; raptorKoxPloxP mice, and the phosphorylation of an mTORC1 target, ribosomal protein S6 (S6), was not detected, as assayed by reactivity with a phosphospecific antibody (Figure 7A). Expression of the lipogenic genes Srebp1c, Fas, Acly, and Gpat were reduced in AAV > Cre; raptorKoxPloxP mice, though Acc and Gck were unchanged under fed conditions (Figure 7B). Most notably, de novo lipogenesis following a high-carbohydrate meal was largely abrogated in liver from AAV > Cre; raptorKoxPloxP mice (Figure 7C).

Having established the critical role of mTORC1 in de novo lipogenesis, we then asked whether defects in its expression and posttranslational processing of SREBP1c explained the disturbed lipid metabolism in AAFP > Cre; Akt2FloxPloxP mice. Following a high-carbohydrate meal, there was an increase in feeding suppressed insulin-induced gene 2a (INSIG2a) expression for at least 6 hr independent of the presence of hepatic Akt2 (Figure 5E).

To test whether unrestrained activity of FoxO1 is the major factor responsible for the lack of postprandial de novo lipogenesis and hepatic weight gain in AAFP > Cre; Akt2FloxPloxP mice, we measured fatty acid synthesis in livers from AAFP > Cre; Akt2FloxPloxP; FoxO1FloxPloxP mice. Elimination of FoxO1 in liver was without effect on the increase in blood glucose evident after ingestion of a HCD (Figure 6A). However, deletion of FoxO1 in liver appeared to reverse the prevention in hepatic weight gain evident in the AAFP > Cre; Akt2FloxPloxP livers, much like the suppression of a similar defect in livers lacking IRS1 and IRS2 (Figure 6B) (Dong et al., 2008). Deletion of FoxO1 alone in liver led to a trend for increased de novo lipogenesis following a meal, consistent with the increase in hepatic TG under several conditions (Figures 2E, 4B, and 6C). Unlike liver size, the decrease in lipid synthesis in livers lacking Akt2 was not reversed by concomitant removal of FoxO1; instead, AAFP > Cre; Akt2FloxPloxP; FoxO1FloxPloxP livers demonstrated rates of de novo lipogenesis virtually identical to those in AAFP > Cre; Akt2FloxPloxP livers (Figure 6C). Delivery by adenovirus of a dominant-negative form of FoxO1 (Nakae et al., 2001) into AAFP > Cre; Akt2FloxPloxP; FoxO1FloxPloxP livers also failed to reverse the decreased lipogenesis in Akt2 null livers (data not shown). These data are consistent with those presented above in which Akt2 is epistatic to FoxO1 regarding the control of liver TG content.

mTORC1 Is Required for De Novo Lipogenesis in Mouse Liver

Cell Metabolism

AFP > Cre; Akt2(loxP/loxP) wild-type mice, but this was blunted in body weight also increased significantly after 6 hr refeeding in (Figures 5 A and 5B). Liver weights normalized to compared to wild-type controls, indicating the minimal requirement for Akt2 in hepatic lipogenesis versus glycemia (Figure 5D). These data point to a discrepancy between the requirements for Akt2 in hepatic lipogenesis versus glycemia following a high-carbohydrate meal. Hepatic mRNAs encoding lipogenic enzymes increased after refeeding a high-carbohydrate meal (Figure 5 E). Surprisingly, the expression of Srebp1c and its targets Acly and Acc were only minimally if at all reduced in AAFP > Cre; Akt2(loxP/loxP) mice, though there was a blunting in the induction of glucokinase (Gck) and Fasn (Figure 5E). HCD

Figure 4. Loss of Hepatic Akt2 Decreases TG Accumulation in Liver Independent of FoxO1 after Aurothioglucose Treatment

(A) Mice with aurothioglucose (GTG) injection gained more weight than age-matched not-treated (NT) controls. Male mice were fasted overnight and then injected with 0.3 g/kg bodyweight GTG at 4–5 weeks old, and were maintained on normal chow for 8 weeks before analysis; n = 4–8 for each genotype; *p < 0.05 by two-way ANOVA followed by Bonferroni’s post test. (B) Hepatic TG levels under fed conditions in mice injected with GTG as described above, n = 6–10 for each group. *p < 0.05, **p < 0.01 by one-way ANOVA followed by Tukey’s post test; error bars represent SEM. Abbreviations are as indicated in Figure 2.

FoxO1 led to livers that accumulated TG levels comparable to those in AAFP > Cre; Akt2(loxP/loxP) rather than AAFP > Cre; FoxO1(loxP/loxP) mice (Figure 4B). This showed that the Akt2 mutant phenotype was dominant to that of FoxO1, which, again, did not support Akt2 as upstream of FoxO1, but rather suggested that Akt2 regulates hepatic TG levels downstream of or in parallel to FoxO1.
the phosphorylated form of protein S6 and eukaryotic translation initiation factor 4E-binding protein 1 (4EBP1), which reflects activation of mTORC1 by nutrients and insulin. Surprisingly, this was unchanged in $\text{AFP} > \text{Cre};\text{Akt2}^{\text{loxP/loxP}}$ mice (Figure 7D). Both the precursor and the processed form of SREBP1c increased 6 hr after HCD feeding; induction of these
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Figure 6. Loss of Hepatic Akt2 Decreases De Novo Lipogenesis Independent of FoxO1
(A) Fasting and refed blood glucose levels. Two-month-old male mice were placed on HCD for 3 days, fasted overnight, and refed with HCD for 6 hr. Blood glucose levels were taken at fasting, 3 hr refeding, and 6 hr refeding time points, n = 5–7 for each genotype, “p < 0.01 by two-way ANOVA followed by Bonferroni’s post test, error bars represent SEM.
(B) Liver weight to body weight ratio. Mice were as described above. Six hours after refedting, mice were sacrificed, and body weight and wet liver weight were determined; n = 5–7 for each condition, “p < 0.05, **p < 0.01 by two-way ANOVA followed by Bonferroni’s post test, error bars represent SEM.
(C) De novo lipogenesis, n = 4–7 for each genotype, **p < 0.01 by one-way ANOVA followed by Tukey’s post test; error bars represent SEM. Abbreviations are as indicated in Figure 2.

In Figure 2E, we assessed hepatic lipogenesis by assessing [3H]-palmitate incorporation into fatty acids. We also measured liver weight to body weight ratio to determine whether Akt2 regulated hepatic lipogenesis. As shown in Figure 2E, Akt2-deficient mice had increased hepatic lipogenesis after refeeding with HCD. By contrast, Akt2 deficiency did not affect the ratio of liver weight to body weight or liver weight to body weight ratio. These results demonstrate that Akt2 is required for the regulation of hepatic lipogenesis.

The metabolic disturbances associated with type 2 diabetes mellitus and the metabolic syndrome have been perceived as being due to an inability of insulin to promote effectively its anabolic actions, a condition long referred to as insulin resistance. However, this view has been modified recently with the increasing appreciation that, while the “insulin-resistant” liver displays some features consistent with an inability of insulin to act, most notably an inappropriately high glucose output, its characteristic increased de novo lipogenesis and TG synthesis is increased, leading to high liver weight to body weight ratio. The latter is a precursor for fatty acid synthesis. The activity of ACLY is regulated transcriptionally as a target of SREBP1c as well as at the posttranslational level by Akt and protein kinase A phosphorylation (Berwick et al., 2002; Horton et al., 2002; Potapova et al., 2000). Insulin injection stimulated phosphorylation of Akt and ACLY, and phosphorylation of both was blunted in livers from AFP > Cre;Akt2^{loxP/loxP} mice (Figure 7E). To study this in more detail, hepatocytes were isolated from control and Akt2 KO livers. Again, insulin promoted Akt and ACLY phosphorylation.

Insulin-Stimulated Phosphorylation of ATP Citrate Lyase
Lyase Is Blunted in Akt2-Deficient Livers

To explore other potential mechanisms utilized by Akt2 to regulate lipid metabolism, we examined two other pathways known to modulate hepatic lipogenesis activity responding to insulin signaling. ATP-citrate lyase (ACLY) catalyzes the cleavage of cytosolic citrate to oxaloacetate and acetyl-CoA,
accumulation more closely resemble a state of augmented insulin sensitivity. The simplest, though by no means only, explanation for this phenomenon is that insulin resistance occurs at a step distal to a bifurcation in insulin signaling, resulting in selective inhibition of the hormone’s effect on glucose output, while permitting signaling to lipid metabolism. Work from our laboratory as well as others suggests that this divergence occurs downstream of Akt, and thus the identification of the specific pathways that Akt2 utilizes to regulate metabolism has achieved particular importance (He, et al., 2010; Leavens, et al., 2009). In this study, we investigated the roles of two proposed Akt target transcription factors, FoxO1 and FoxA2, and an indirect route for control of a third, SREBP1c, which is regulated by mTORC1. Our data show that, whereas suppression of FoxO1 activity is downstream of Akt in the inhibition of hepatic glucose output, phosphorylation of neither FoxO1 nor FoxA2 is likely to mediate insulin’s effects on de novo lipogenesis and TG accumulation in liver. Our data also support the notion that mTORC1 is required for maintenance of Srebp1c expression and de novo lipogenesis, but its activation is not sufficient to reverse the reduction of de novo lipogenesis coincident with deletion of Akt2. Moreover, it is likely that the mechanism by which Akt2 induces lipogenesis after a high-carbohydrate meal is not based on regulation of gene expression by any of the three transcription factors assessed in this study.

Considerable data support a role for FoxO1 in the control of hepatic gluconeogenic genes expression, though there is much less clarity regarding its importance for glycemia and hepatic lipid metabolism. As has been reported, we found that inhibition of FoxO1 in the nuclear localization of FoxA2, even in the extreme case of elevated insulin levels in ob/ob mice (Chen, et al., 2005). In loss-of-function studies, liver-specific FoxA2 knockout mice displayed no abnormality in hepatic TG levels, consistent with the current results (Bochkis, et al., 2008). Under normal chow feeding or in mice with DIO, loss of hepatic FoxA2 did not affect serum glucose and insulin or hepatic TG levels (Figures 1B and 1C and Figures S1 and S2). These data are compatible with a model in which inhibition of FoxA2 is not required for Akt2 to regulate hepatic TG accumulation. These results are qualified by the possibility that, in other mouse strains, the requirements for hepatic steatosis might be different.

Recently, mTORC1 has been shown to enhance de novo lipogenesis by regulating Srebp1c expression and processing (Chakrabarti, et al., 2010; Duvel, et al., 2010; Li, et al., 2010; Porstmann, et al., 2008). The full activation of mTORC1 requires both nutrients and growth factor signaling, which determine cellular location and Akt-dependent activation of the mTORC1 complex, respectively (Sancak, et al., 2008, 2010; Sengupta, et al., 2010b). In tissue culture cells and primary hepatocytes, insulin/Akt regulates Srebp1c expression and processing and de novo lipogenesis in an mTORC1-dependent manner (Azzout-Marniche, et al., 2000; Chakrabarti, et al., 2010; Duvel, et al., 2010; Fleischmann and Iynedjian, 2000; Li, et al., 2010; Porstmann, et al., 2008). However, the model of an obligate role for mTORC1 in hepatic lipogenesis has not been tested genetically in vivo. Herein we report that conditional deletion of raptor in liver abolished mTORC1 activity, leading to a reduction in the expression of Srebp1c and its targets and decreased de novo lipogenesis (Figures 7A–7C). Interestingly, even though AFP > Cre;Akt2(loxP/loxP)
and AFP > Cre;Akt2(loxP/loxP), FoxO1(loxP/loxP) mice exhibit significantly decreased de novo lipogenesis following a high-carbohydrate meal, activation of mTORC1 and SREBP1c were largely unaffected (Figures 5D, 7D, and 7E). These data indicate that mTORC1 activation is not sufficient to induce lipogenesis under these conditions. During the revision of this paper, Yecies et al. reported an mTORC1-independent pathway by which Akt2 signals through INSIGI2a to promote SREBP1c processing (Yecies et al., 2011). We found no change in the normal decrease in hepatic INSIGI2a mRNA in AFP > Cre;Akt2(loxP/loxP) mice after HCD refeeding and only a modest reduction in the nuclear form of SREBP1c or expression of its target genes (Figures 5E and 7E). Thus, the mTORC1-independent pathway revealed in this study by liver-specific deletion of Akt2 does not represent induction of INSIGI2a and is unlikely to be related to SREBP1c processing or expression. Rather, we propose that the requirement for Akt2 is linked to its role in the posttranslational control of other targets such as ACLY.

Taking these data in aggregate, we suggest that Akt2 promotes hepatic lipid accumulation and de novo lipogenesis independent of FoxA2 and FoxO1. Moreover, activated mTORC1 and SREBP1c and increased lipogenic gene expression is required but not sufficient for postprandial lipogenesis. Thus, our data provide strong support for the notion that nutrients have a direct role in the liver to promote anabolic lipid metabolism by a process dependent on the presence of a functional mTORC1 complex as well as other Akt-dependent signaling pathways.

**EXPERIMENTAL PROCEDURES**

**Animals**

The AFP > Cre;Akt2(loxP/loxP), AFP > Cre;FoxO1(loxP/loxP), and RaptorloxP/loxP mice have been described previously (Leavens et al., 2009; Matsumoto et al., 2007; Sengupta et al., 2010a; Zhang et al., 2005). All animals were raised and treated with approval from the University of Pennsylvania IACUC in accordance with National Institutes of Health (NIH) guidelines. Mice were of a 129/FVB/C57BL6 mixed background, and male littermates were used for all studies. In the FoxA2 experiments, the wild-type control group included mice of genotypes AFP > Cre;Akt2loxP/+, FoxA2loxP/loxP, and Akt2loxP/loxP; FoxA2loxP/loxP. Mice labeled as AFP > Cre;Akt2loxP/loxP included the genotypes AFP > Cre;Akt2loxP/loxP, FoxA2loxP/loxP, and AFP > Cre;Akt2loxP/loxP, FoxA2loxP/loxP. Mice labeled as AFP > Cre;FoxA2loxP/loxP included the genotypes AFP > Cre;Akt2loxP/loxP, FoxA2loxP/loxP, and AFP > Cre;Akt2loxP/loxP, FoxA2loxP/loxP. In the FoxO1 experiments, wild-type controls include genotypes AFP > Cre;Akt2loxP/loxP, FoxO1loxP/loxP and Akt2loxP/loxP; FoxO1loxP/loxP. Mice indicated as AFP > Cre;Akt2loxP/loxP, FoxA2loxP/loxP and AFP > Cre; Akt2loxP/loxP, FoxA1loxP/loxP. In all cases there were no metabolic phenotypes among the pooled genotypes including Cre expressed alone. Mice were main-
tained on a normal Chow diet (laboratory rodent diet 5001, LabDiet) unless otherwise noted under 12 hr light/dark cycle (7 a.m./7 p.m.) in a barrier facility. When fed with HFD (D123311, Research Diets), mice were started at an age of 1 month and remained for 3 months.

For aurothioglucose (GTG, A0606, Sigma Aldrich) treatment, mice were maintained on normal chow. At 5–6 weeks old, mice were fasted for 24 hr and then injected intraperitoneally (i.p.) with one dose of 0.3 g/kg GTG in PBS. After injection, mice were kept fasted for another 20 hr before normal Chow diet was freely accessible for the following 2 months. This protocol has been suggested to reduce mortality caused by GTG injection in mice (Brecher et al., 1965).

For acute excision of Raptor in liver, the adeno-associated virus was generated by Viral Vector Core at University of Pennsylvania. GFP or Cre recombinase was driven by a liver-specific promoter, thyroxine-binding globulin (TBG) promoter. Mice were injected postorbitaly at a dosage of 1.0 × 1011 gc virus diluted in PBS at 6 weeks of age and maintained on normal Chow for another 2 weeks before analysis.

**SGE Secretion Assay**

TG secretion assay was performed as described previously (Millar et al., 2005). Mice were fasted overnight and then injected i.p. with 1 g/kg poloxamer 407 (kindly provided by Dr. John S. Millar, University of Pennsylvania). Blood samples were collected at indicated time points and assayed for serum TG.

**Statistics**

All data are presented as mean ± SEM. As noted in the figure legends, data were analyzed by using unpaired Student’s t test with two-tailed analysis, one-way ANOVA followed by Tukey’s post test, or two-way ANOVA followed by Bonferroni’s post test.

**SUPPLEMENTAL INFORMATION**

Supplemental Information includes seven figures, one table, Supplemental Experimental Procedures, and Supplemental References and can be found with this article online at doi:10.1016/j.cmet.2011.09.001.
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Akt2 Targets Regulate Hepatic Lipid Metabolism


