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Stearoyl-CoA Desaturase and its Relation to High-Carbohydrate Diets and Obesity

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Abstract

Obesity is currently a worldwide epidemic and public health burden that increases the risk for developing insulin resistance and several chronic diseases such as diabetes, cardiovascular diseases and non-alcoholic fatty liver disease. The multifactorial causes of obesity include several genetic, dietary and lifestyle variables that together result in an imbalance between energy intake and energy expenditure. Dietary approaches to limit fat intake are commonly prescribed to achieve the hypocaloric conditions necessary for weight loss. But dietary fat restriction is often accompanied by increased carbohydrate intake, which can dramatically increase endogenous fatty acid synthesis depending upon carbohydrate composition. Since both dietary and endogenously synthesized fatty acids contribute to the whole-body fatty acid pool, obesity can therefore result from excessive fat or carbohydrate consumption. Stearoyl-Coenzyme A desaturase-1 (SCD1) is a delta-9 fatty acid desaturase that converts saturated fatty acids into monounsaturated fatty acids (MUFA) and this activity is elevated by dietary carbohydrate. Mice lacking *Scd1* are protected from obesity and insulin resistance and are characterized by decreased fatty acid synthesis and increased fatty acid oxidation. In this review, we address the association of high-carbohydrate diets with increased SCD activity and summarize the current literature on the subject of SCD1 and body weight regulation.

Keywords

Stearoyl-CoA Desaturase; SCD1; Obesity; Insulin; Carbohydrate; Lipogenesis

1. Introduction

Obesity is currently a worldwide epidemic prevalent in both adults and children that is caused by an imbalance of high energy consumption with low energy expenditure [1]. Increased weight gain is associated with an insulin resistance syndrome that elevates the risk for several chronic diseases including diabetes, cardiovascular diseases and non-alcoholic fatty liver disease [2, 3]. In response to positive energy balance, the adipose tissue stores this surplus energy primarily in triglyceride-rich lipid droplets. Sustained nutrient overload is hypothesized to impair normal adipocyte function leading to abnormal adipokine production and leakage of nutrients from the adipocyte into insulin-sensitive tissues such as muscle and liver [4]. These events contribute

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to both local and systemic insulin resistance resulting in hyperinsulinemia, increased hepatic glucose production, reduced peripheral glucose uptake and dyslipidemia [2,5,6].

Alterations in dietary macronutrient composition and caloric intake can be effective treatments for weight loss and plasma lipid management. Most food-based recommendations disseminated to the general public share a consistent message to consume at least 55% of calories from carbohydrate, increase consumption of fruits, vegetables, legumes and whole grains, but minimize intake of sugar, cholesterol, saturated and trans fatty acids [7,8]. Many popular weight-loss strategies share some of these recommendations, but their target dietary carbohydrate intakes span a wide range between 22.2% and 81.0% [9]. While dietary adherence is an important factor in the efficacy of any intervention, individual variation in response to dietary changes also influences the outcome [10]. For example, interactive effects between genetic modifiers and low-fat, high-carbohydrate diets can effect levels of plasma very low-density lipoprotein (VLDL) triglyceride levels, low-density lipoprotein (LDL) subclass distribution and insulin secretion response [11–13].

One of the gene-nutrient interactions that may be responsible for the differential response to nutritional intervention is the dietary modulation of stearoyl-Coenzyme A desaturase (SCD). SCD is a delta-9 fatty acid desaturase that converts saturated fatty acids into monounsaturated fatty acids (MUFA) [14]. These MUFA are key substrates for the formation of complex lipids such as triglycerides, cholesterol esters, wax esters and diacylglycerols. Additionally, increased cellular SCD1 activity has been suggested to influence fatty acid partitioning by promoting fatty acid synthesis but decreasing oxidation [15–17]. Previous studies have shown that the reduced MUFA synthesis that occurs in *Scd1*-deficient mice or those treated with *Scd1*-targeted antisense oligonucleotides is associated with several metabolic changes that elicit protection from obesity, cellular lipid accumulation and insulin resistance [14,17–21]. *Scd1* levels are elevated by both dietary and hormonal factors, such as glucose, fructose, saturated fatty acids and insulin, but repressed by polyunsaturated fatty acids (PUFA) and leptin [14,16,22–25]. Therefore, high levels of *Scd1* elicited by both dietary carbohydrate and certain fatty acids may be a significant contributor to the development of obesity and other chronic diseases.

This review addresses the association of high-carbohydrate diets with obesity and increased SCD activity. The first part of this review clarifies several nutritional variables that comprise a high-carbohydrate diet that can impact its metabolic effect. In particular, we emphasize the distinction between low- and high-glycemic index carbohydrate sources in relation to obesity. Next, we discuss how these nutritional components and the associated hormonal changes regulate the expression of SCD1. The final part of this review will summarize the current literature on the subject of SCD1 and body weight regulation.

2. Defining the high-carbohydrate diet

2.1 The origins of the low-fat, high-carbohydrate diet

The recommendation to substitute carbohydrate calories for fat calories is partly rooted in studies from both human and animal models that showed a plasma LDL-cholesterol raising effect of the dietary saturated fats lauric (12:0), myristic (14:0), and palmitic acid (16:0) compared to unsaturated fats [26–28]. One strategy to limit saturated fat intake involves the isocaloric substitution of dietary unsaturated fat for saturated fat. The low-fat, high-carbohydrate diet is an extension of this approach that limits total fat intake as opposed to targeting saturated fat. This is achieved by exchanging dietary fat with carbohydrate or protein. Since fat is more calorie dense (~9 kcal/g) than carbohydrate or protein (~4 kcal/g), this can be a useful strategy to reduce caloric intake to promote negative energy balance and weight loss. For the purpose of this review, changes in carbohydrate and fat calories are assumed to occur in the absence of altered protein intake. Thus, the low-fat, high-carbohydrate approach

results in a substantial increase in the percent calories coming from carbohydrate at the expense of fat. However, it is important to note that increased dietary protein intake at the expense of carbohydrate or fat calories has been associated with increased satiety and compliance with weight loss and weight maintenance strategies [29,30].

2.2 Not all carbohydrates are created equal

The metabolic response to carbohydrate intake is influenced by both carbohydrate mass as well as carbohydrate composition [31,32]. Carbohydrates exist as monosaccharides (i.e. glucose, fructose and galactose), disaccharides (i.e. sucrose and lactose), or longer polymers of glucose (i.e. maltodextrins and starches). Dietary monosaccharides are readily absorbed by enterocytes in the upper small intestine and rapidly transported out the cell into the portal circulation. Disaccharides must first be cleaved into monosaccharides by brush-border glucosidases prior to absorption. For example, dietary sucrose is cleaved by sucrase to yield equal amounts of glucose and fructose. High-fructose corn syrup is a processed sugar derived from corn starch that contains approximately equal amounts of glucose and fructose. Increased consumption of simple sugars such as sucrose and high-fructose corn syrup that rapidly enter the bloodstream after ingestion has been suggested to be a large contributor to the increased incidence of obesity and the metabolic syndrome [33].

In contrast, plant starches are complex carbohydrates consisting of hundreds to several thousands of glucose units linked by α -1,4-glycosidic bonds (amylose), or a branched form linked by both α -1,4- and α -1,6-glycosidic bonds (amylopectin). These starches require salivary and pancreatic-derived amylase enzymes to cleave the glycosidic bonds and convert starch into maltose, maltotriose, and α -dextrins (or maltodextrins), which are further hydrolyzed to glucose by brush border enzymes. The physical accessibility of the carbohydrate in foods to hydrolytic enzymes is an important factor in controlling the rate of monosaccharide entry into the bloodstream. The amylose:amylopectin ratio of starch can influence carbohydrate absorption due to the ability of amylose-rich starches to form more tightly packed, hydrolysis-resistant granules [34]. The physical form (liquid vs. solid), degree of processing (whole vs. refined), method of preparation (raw vs. cooked) and fiber content can also have a significant impact on the rate of carbohydrate absorption [34]. These variables are often expressed as the glycemic index of a food, which is defined as the blood glucose response elicited by a test food relative to a glucose standard containing an equivalent mass of carbohydrate [35]. Both the glycemic index as well as the total carbohydrate content of a food will influence postprandial glycemia and elicit a proportional insulin secretory response (Figure 1) [32].

3. Nutritional control of lipogenesis and the role of SCD1

3.1 Cellular sensing and fate of dietary carbohydrate

The transport of glucose into the cell and subsequent phosphorylation to glucose-6-phosphate are the primary rate-limiting steps for glucose metabolism. Pancreatic β -cells and liver hepatocytes, as well as some cells of the hypothalamus, kidney, and small intestine, express the high K_m plasma membrane GLUT2 glucose transporter and glucokinase glucose phosphorylating enzyme. This allows these cells to act as “glucose-sensors” by coupling millimolar changes in blood glucose concentrations to corresponding signal-generating metabolic flux rates. For example, dietary modifications resulting in an increased blood glucose load elicit a greater β -cell insulin secretory response [32,36] (Figure 1). It is important to note that amino acids and fatty acids are relatively weak insulin secretagogues alone, but can also acutely augment glucose-stimulated insulin secretion [37–40]. In contrast to β -cells and hepatocytes, tissues such as muscle and adipose require insulin to stimulate glucose uptake by promoting the translocation of a large pool of intracellular GLUT4 glucose transporters to the cell surface for glucose transport and subsequent phosphorylation by the low K_m hexokinase

[41]. One fate of glucose-6-phosphate is to be utilized to replenish cellular glycogen stores in tissues such as muscle and liver. However, adipose tissue and liver also have a large capacity for lipogenesis and an alternative fate of glucose-6-phosphate is the conversion to acetyl-CoA for fatty acid synthesis and storage as triglycerides.

3.2 Stimulation of hepatic de novo lipogenesis by dietary carbohydrate

While both adipose and liver may perform de novo lipogenesis (DNL), the liver is unique in its ability to dramatically increase fatty acid synthesis in response to changes in dietary macronutrient intake. Of particular interest is the effect of dietary carbohydrate, especially sugars, in humans to increase hepatic DNL [42]. Hepatic DNL is lowest after fasting and peaks in the postprandial period [43], but is also strongly influenced by dietary composition. Whereas minimal DNL occurred on a high-fat diet (40% of calories as fat, 45% as glucose polymers), hepatic fatty acid synthesis was markedly increased upon low-fat feeding (10% of calories as fat and 75% as glucose polymers) [44]. Furthermore, eucaloric substitution of dietary starch for sugar reduced hepatic DNL [45]. Additionally, fructose has been shown to be a more potent inducer of hepatic lipogenesis than glucose in mice, rats and humans [24,46–48]. This is partly due to the remarkable ability of the liver to clear approximately 70% of blood fructose [49]. Also, fructose metabolism is not inhibited at the level of phosphofructokinase as is glucose, allowing for fructose to bypass this regulator step and serve as an unregulated source of precursors for hepatic lipogenesis [33].

The rate of intestinal carbohydrate absorption into the circulation also influences the physiological response to ingested carbohydrate (Figure 1). Behall *et al.* investigated the effect of slowly absorbed amylose starch compared to rapidly absorbed amylopectin starch in humans and found that consumption of an amylose containing meal elicited a lower postprandial glucose and insulin response compared to those fed an equivalent amount of amylopectin [50–52]. Furthermore, a meta-analysis of 37 prospective human cohort studies found that both high-glycemic index and high-glucose load diets are independently associated with increased risk for several chronic diseases, including type 2 diabetes and coronary heart disease, suggesting that postprandial glycemia and insulin secretion contribute to disease progression [31]. This relationship is also exemplified by studies in mice and rats, in which feeding amylopectin starch promoted more adiposity, hepatic steatosis, hyperinsulinemia, and increased plasma triglycerides compared to feeding amylose starch [53–55].

3.3 Transcriptional control of *Scd1* by dietary carbohydrate and fatty acids

The amenability of the rodent animal model to both dietary and genetic manipulation has allowed for the elucidation of several molecular mechanisms involved in carbohydrate-induced lipogenesis. One mechanism for elevated hepatic lipogenesis in response to dietary carbohydrate is increased insulin secretion, which promotes the activation of the lipogenic transcription factors liver-X-receptor (LXR) and sterol regulatory element binding protein 1c (SREBP-1c) [56,57]. LXR activation has also been shown to increase the expression of mRNAs encoding both SREBP-1c and carbohydrate responsive element binding protein (ChREBP) [58]. Increased hepatic glucose flux promotes the formation of xylose-5-phosphate via the hexose monophosphate shunt to promote activation of protein phosphatase 2A and subsequent dephosphorylation of ChREBP to an active lipogenic transcription factor [48,59,60]. Together, SREBP, LXR and ChREBP act to couple increased carbohydrate intake with induction of lipogenic genes including liver pyruvate kinase (L-PK; *Pklr*), acetyl-CoA carboxylase (*ACC α* *Acaca*, *ACC β* ; *Acacb*), fatty acid synthase (FAS; *Fasn*) and *Scd1* (Figure 1).

The effect of dietary carbohydrates, such as glucose, fructose and sucrose, to robustly increase hepatic *Scd1* is due to both SREBP-1c-dependent and independent mechanisms [21,24,61]. Both fructose feeding [24] and LXR agonists [62,63] have been shown to increase hepatic

Scd1 in both SREBP-1c^{+/+} and SREBP-1c^{-/-} mice. ChREBP deficiency also abrogates the increase in *Scd1* elicited by both dietary carbohydrate [64] and LXR agonists [58], although the presence of carbohydrate responsive element in the promoter of *Scd1* has yet to be identified. *Scd1* expression has been shown to be positively regulated by direct LXR binding to an LXR-response element in the *Scd1* promoter, as well as LXR-mediated activation of SREBP-1c transcription [61,62]. Consistent with fructose being more lipogenic than glucose [46–48], fructose is also a more potent inducer of hepatic *Scd1* [24]. As opposed to hepatic carbohydrate metabolism, intracerebroventricular glucose administration strongly decreases hepatic *Scd1* [65]. However, the physiological significance of this brain-liver circuit in the context of concomitant direct insulin and glucose effect on the liver remains to be established.

In addition to metabolic effects elicited by dietary carbohydrate, fatty acids also modulate the transcriptional activation of *Scd1* and other lipogenic genes [23,25]. The binding of SREBP-1 to the SREBP response element of the *Scd1* promoter is decreased by dietary *n-3* and *n-6* PUFA, partly due to post-translational repression of SREBP-1c maturation [61]. PUFA also act to decrease mRNA abundance of SREBP-1c potentially by accelerating mRNA decay [66,67]. Additionally, *n-3* and *n-6* PUFA decrease the nuclear abundance of ChREBP; however, the mechanism is still unresolved [68,69]. In contrast to the anti-lipogenic effects of PUFA, dietary saturated fat strongly induces *Scd1* expression in a mechanism that may involve fatty acid upregulation of PGC1- β and subsequent coactivation of SREBP-1c and LXR [16,70]. Thus, the lipogenic potential of a low-fat, high-carbohydrate diet may be increased or decreased by relative amounts of dietary saturated fats and PUFA, respectively.

Recently, Cao *et al.* have suggested that one product of SCD1, palmitoleate, produced and released by the adipose tissue suppresses hepatic lipogenic rates through a specific inhibition of *Scd1* in liver [71]. In their genetic model, mice lacking two major adipocyte fatty acid binding proteins (ap2 and mal1) showed a diminished fatty acid binding capacity in the adipose tissue [72] associated with a 2- to 3-fold increase in palmitoleate content of adipose lipids and plasma free fatty acids and robust repression of hepatic *Scd1* [71]. This study also reported reduced hepatic *Scd1* in mice after tripalmitoleate infusion compared to tripalmitate infusion [71]. However, the abundant dietary MUFA oleic acid has been previously shown to have no repressive effect on hepatic *Scd1* expression relative to the well-documented effects of *n-3* and *n-6* PUFA linoleic, linolenic, arachidonic, eicosapentaenoic, and docosahexaenoic acid [61, 73–75]. Additionally, increased plasma palmitoleate in humans has been independently associated with both hypertriglyceridemia and abdominal adiposity [76]. Therefore, further *in vivo* validation is necessary to directly evaluate the ability of altered circulating palmitoleate levels to elicit metabolic changes and repress hepatic *Scd1* compared to both oleate and *n-3* and *n-6* PUFA.

4. The relationship of SCD1 to carbohydrate-induced adiposity

4.1 Lessons from *Scd1*-deficient mice

Mice with a targeted deletion of *Scd1* or with spontaneous *asebia* mutations in *Scd1* have provided several novel insights into the metabolic role of SCD1 [19,77]. *Scd1*-deficient mice have reduced hepatic fatty acid and triglyceride synthesis in response to high-carbohydrate feeding [18,21,24]. This is partly due to decreased maturation of SREBP-1 protein and expression of mRNAs encoding fatty acid synthesis genes [16,24]. Normal SREBP-1c activation in *Scd1*^{-/-} mice is restored by feeding high levels of dietary MUFA, but not saturated fat, indicating that induction of the hepatic lipogenic program requires adequate levels of MUFA derived from the diet or from endogenous synthesis [16,24]. We have recently shown that liver-specific deletion of *Scd1* using the Cre-loxP system is also sufficient to block this carbohydrate-induced lipogenic program [78].

The lean phenotype of *Scd1*-deficient mice also results from increased energy expenditure and oxygen consumption due to enhanced fatty acid oxidation and thermogenesis in liver, muscle, and brown adipose tissue, through both transcriptional and post-transcriptional mechanisms [20,79–82]. *Scd1*-deficient mice also have elevated AMP-activated protein kinase (AMPK) activity in muscle and liver, which results in inhibitory phosphorylation of acetyl-CoA carboxylase, reduced malonyl-CoA levels and enhanced import of fatty acids into the mitochondria for oxidation [16,80,81,83]. The mechanism for how cellular MUFA homeostasis regulates the SREBP-1c maturation and AMPK activation are currently unknown.

4.2 Liver-specific loss of *Scd1* reveals a role in carbohydrate-induced adiposity

The decreased fatty synthesis and increased fatty acid oxidation phenotypes elicit resistance to obesity and hepatic steatosis in whole-body *Scd1*-deficient mice when stressed with a high-fat (60% calories from fat) or high-carbohydrate diet (10% calories from fat) [16,20,78]. However, liver-specific *Scd1*-deficient mice are protected from high-carbohydrate, but not high-fat diet-induced obesity and hepatic steatosis [78]. We can conclude from this study that SCD1 activity in extrahepatic tissues is involved in obesity resistance from high-fat diets. Hepatic *Scd1* deficiency is apparently insufficient to induce the hypermetabolism phenotype that occurs in whole-body *Scd1*-deficient mice. Binczek *et al.* have proposed that disruption of the epidermal lipid barrier in whole-body *Scd1*-deficient mice is responsible for the obesity resistance [84]. However, metabolic effects derived from inhibition of *Scd1* in the adipose or other tissues are also possible. This is supported by the studies of Jiang *et al.* that reported antisense oligonucleotide-mediated inhibition of *Scd1* in the liver and adipose protected mice from high-fat diet-induced obesity without concomitant skin abnormalities [17]. Interestingly, *ap2/mal1* knockout mice are protected from high-fat diet induced obesity, insulin resistance, and hepatic triglyceride accumulation, coincident with elevated plasma palmitoleate levels and transcriptional repression of hepatic *Scd1* [71]. However, these findings are not in accordance with the lack of high-fat diet-induced obesity resistance observed in liver-specific *Scd1*-deficient mice [78]. Furthermore, whole-body *Scd1*-deficient mice have decreased plasma palmitoleate levels despite obesity resistance and reduced hepatic lipogenesis [20,85]. This suggests that the *ap2/mal1* knockout model is causing obesity resistance independent of altered plasma palmitoleate levels and hepatic *Scd1* repression.

The reduced carbohydrate-induced adiposity in liver-specific *Scd1*-deficient mice suggests that the hepatic conversion of carbohydrate to fatty acids is requisite for not only hepatic triglyceride synthesis but also expansion of adipose stores (Figure 2). The liver packages its lipid stores into triglyceride-rich VLDL particles, which are secreted into the circulation for hydrolysis by lipoprotein lipase in peripheral tissues [86]. Relative to control mice, liver *Scd1*-deficient mice fed the high-carbohydrate diet (10% calories from fat) had lower hepatic lipogenic induction, hepatic triglyceride accumulation, and fasting plasma triglycerides [78]. These phenotypes all corresponded with smaller white adipose tissue depots. Similar results were also observed in whole-body *Scd1*-deficient mice fed a very low-fat, high-carbohydrate diet. This is reminiscent of studies by Scribner *et al.* in which long-term feeding of a high-glycemic index diet to mice promoted higher hepatic triglycerides, plasma triglycerides and adiposity relative to mice fed a low-glycemic index diet, presumably due to differential induction of hepatic lipogenesis [54,55]. This data suggests that the transport of de novo synthesized hepatic triglycerides to the adipose tissue must occur in order for high-carbohydrate diets to promote weight gain (Figure 2). However, a dysregulation of hepatic carbohydrate metabolism or insulin signaling may also contribute to decreased adiposity observed in these models.

4.3 Metabolic effects of a very-low fat, high-carbohydrate diet in mice

Most rodent standard or purified low-fat, high-carbohydrate diets contain at least 10% of calories from fat and supply sufficient MUFA and PUFA for survival. We have recently

explored the metabolic effect of feeding a high-sucrose, very low-fat (HSVLF) diet for 2 weeks in whole-body and liver-specific *Scd1*-deficient mice [78,87,88]. This diet contributes ~2.5% calories from fat (derived from corn oil), which limits total fat as well as dietary MUFA and PUFA. In response to the HSVLF diet, control mice are able to upregulate de novo lipogenesis in order to synthesize sufficient MUFA to maintain metabolic homeostasis. However, whole-body *Scd1*-deficient mice develop a complex phenotype involving loss of body weight, hypoglycemia, depleted hepatic glycogen, hypercholesterolemia, and cholestasis [87]. Subsequent hepatic gene expression analysis revealed significant endoplasmic reticulum stress and inflammation that are associated with the metabolic dysfunction [88]. Supplementation of the HSVLF diet with dietary unsaturated fat, but not saturated fat, ameliorated most of the HSVLF-induced metabolic perturbations [87]. However, dietary MUFA alone was not as effective as a fat source containing both MUFA and PUFA to reduce the hypercholesterolemic effects of the HSVLF diet. Liver-specific *Scd1*-deficient mice fed the HSVLF diet also developed loss of body weight, hypoglycemia and depleted hepatic glycogen, but only modest hypercholesterolemia and none of the cholestasis-like phenotypes observed in whole-body *Scd1*-deficient mice [78].

These findings emphasize two important roles for SCD1 during very low-fat, high-carbohydrate feeding conditions. 1) Lack of SCD1 results in an increased requirement for dietary unsaturated fat to compensate for reduced de novo MUFA synthesis. 2) Endogenous MUFA synthesis is essential during dietary unsaturated fat insufficiency and influences the dietary requirement of PUFA. This model is consistent with the observation that hepatic *Scd1* and MUFA synthesis are repressed by dietary PUFA intake, which may have evolved as a mechanism to maintain cellular unsaturated fatty acid balance [61]. Studies by Chakravarthy *et al.* in mice with liver-specific deletion of FAS have provided additional evidence for the importance of de novo fatty acid synthesis [89]. When fed a high-carbohydrate diet without fat or subjected to prolonged fasting, these mice developed hypoglycemia and fatty liver coincident with reduced expression of PPAR α target genes. These phenotypes did not occur in mice fed a standard chow diet and were corrected by administration of a potent PPAR α agonist, suggesting that new fat synthesized by FAS is the preferred endogenous activator of PPAR α [89]. Since SCD1 is downstream of FAS and converts FAS-derived saturated fatty acids into MUFA, it is possible that impaired hepatic MUFA synthesis is responsible for many of the metabolic phenotypes elicited by hepatic deficiency of either SCD1 or FAS.

4.4 The relation of human SCD1 to carbohydrate-induced obesity

Similar to mouse SCD1, human SCD1 is expressed in several tissues, is highest in liver and adipose, and shares 94.1% amino acid identity with mouse SCD1 [90]. The promoter of human *SCD1* has been cloned and characterized [91]. Transcription factor binding sites for SREBP, NF-Y, and the PUFA response element were present in the human *SCD1* promoter similar to the mouse *Scd1* promoter. Most human studies have inferred SCD activity via the ratio of monounsaturated to saturated fatty acids in plasma lipids, as discussed below. However, Chong *et al.* have recently shown that a short-term feeding of a high-carbohydrate diet elicits a parallel increase in both DNL and hepatic SCD activity, as measured by conversion of [2 H]-palmitic acid to [2 H]-palmitoleic acid in VLDL-triglyceride [94]. These data strongly suggest that human SCD1 expression and hepatic lipid synthesis respond similarly to dietary carbohydrate as found in rodent models.

Sampath and Ntambi recently reviewed the use of the SCD plasma desaturation index (18:1/18:0 or 16:1/16:0) in humans as a surrogate marker for SCD activity and its association with metabolic outcomes [92]. Although this plasma index does not allow for the distinction between liver and adipose SCD1 activity, it has been shown to correlate well with plasma triglyceride levels [76,85,93,94] and body weight [76,95]. A 4–6 week-long high-carbohydrate

diet increased the 18:1/18:0 ratio and this explained 44% of the variance in the plasma triglyceride response [85]. Additionally, a short-term 3-day low-fat, high-carbohydrate feeding elicited a parallel increase in the desaturation index, hepatic lipogenesis, and plasma triglycerides [94]. An elevated SCD desaturation index has also been shown to correlate well with the dyslipidemia observed in familial combined hyperlipidemia, potentially due to genetic variations that affect SCD1 activity [93]. Mangravite *et al.* have recently shown that adipose levels of *SCD1* mRNA are lowered independently by both acute weight loss as well as isocaloric reduction in carbohydrate intake [96]. Furthermore, adipose *SCD1* levels correlated with plasma triglycerides, but this was independent of carbohydrate intake and the plasma SCD desaturation index [96]. This suggests that modulation of both hepatic and adipose SCD1 levels may influence the plasma triglyceride response via different mechanisms.

As discussed in sections 2 and 3, dietary conditions that promote increased insulin secretion will result in high SCD1 activity, which is associated with a metabolic state favoring hepatic triglyceride accumulation and an expansion of adipose triglyceride stores. Changes in SCD1 activity may influence obesity in two ways. In both human and animal models, elevated levels of SCD1 have been suggested to influence the partitioning of fatty acids towards storage and away from oxidation [15–17]. However, elevated hepatic SCD1 activity and MUFA production may also lead to increased synthesis and transport of VLDL triglycerides to peripheral tissues (Figure 2). In human studies, dietary carbohydrates (primarily sugars) have consistently been shown to increase both hepatic de novo lipogenesis and fasting plasma triglycerides, but the relative contribution of increased VLDL triglyceride production and clearance to the elevated plasma triglyceride phenotype is unresolved (reviewed in [11,42]). Interestingly, Chaput *et al.* recently reported a novel interaction between dietary composition and insulin secretion in which patients predisposed to acute hypersecretion of insulin gained the most weight, especially among those consuming low-fat diets [13]. Since insulin has many physiological effects occurring in a variety of tissues, the effects of hyperinsulinemia on the development of obesity are likely not limited to altered hepatic triglyceride production.

5. Conclusion

High-carbohydrate diets, especially those enriched in simple carbohydrates, increase hepatic de novo lipogenesis coincident with elevated hepatic *Scd1* expression. This involves pro-lipogenic effects of increased circulating glucose, fructose and insulin to promote activation of SREBP-1, ChREBP, and LXR. Whole-body or liver-specific deletion of *Scd1* in mice dramatically reduces the lipogenic effects of dietary carbohydrate by decreasing MUFA production and preventing the upregulation of lipogenic gene expression. However, several key questions remain unanswered in relation to the role of SCD1 in metabolism. First, does oleate or another product of SCD1 directly activate SREBP processing? If so, would increased hepatic *Scd1* expression in the absence of elevated blood glucose and insulin be sufficient to promote SREBP processing and de novo lipogenesis? Alternatively, the impaired SREBP processing and lipogenesis may be the indirect result of energy deficiency or altered membrane composition. The role of extrahepatic SCD1 in carbohydrate-induced adiposity also needs to be investigated. Since liver-specific *Scd1*-deficient mice are protected from high-carbohydrate, but not high-fat diet-induced obesity, *Scd1* expression in extrahepatic tissues such as the adipose, muscle, brain, and skin may also play a pivotal metabolic role. However, the blunted carbohydrate-induced lipogenesis in liver-specific *Scd1*-deficient mice suggest that these tissues cannot compensate for local hepatic depletion of MUFA. Together, the current body of literature suggests that high-carbohydrate diets act through modulation of hepatic SCD1 activity to increase the conversion of carbohydrates to MUFA, which enhance hepatic triglyceride synthesis and secretion. Presumably, this rise in hepatic triglyceride secretion promotes the storage of carbohydrates as triglycerides in the adipose tissue, contributing to the development of obesity.

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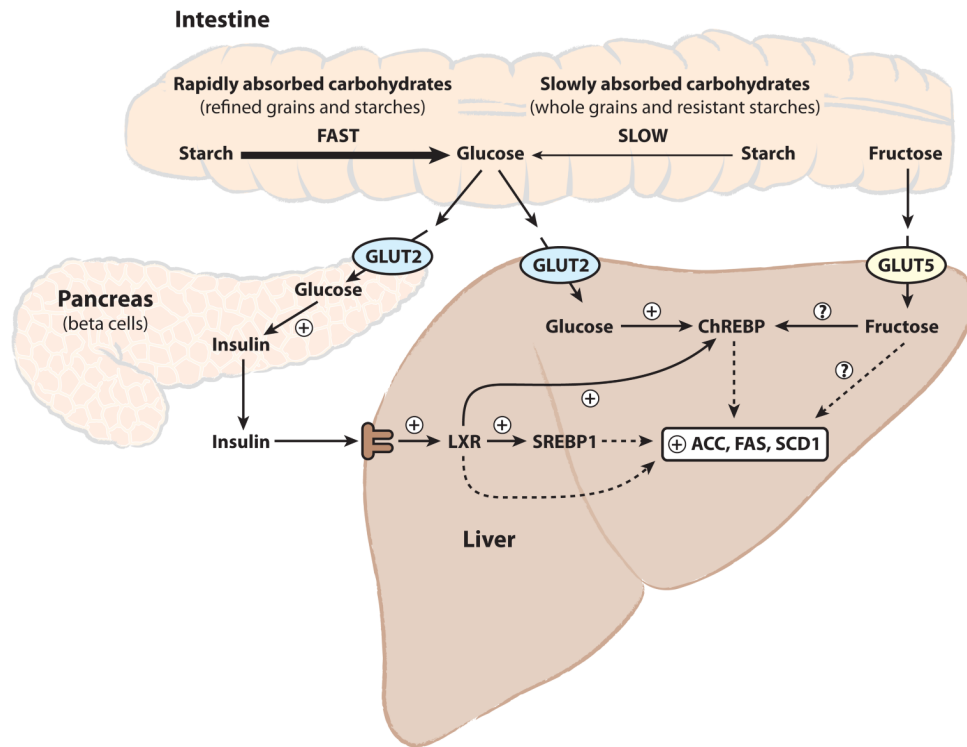


Figure 1. High-carbohydrate diets induce hepatic lipogenesis via direct and indirect liver nutrient sensing mechanisms. After carbohydrate ingestion, starches are broken down in the intestine into glucose and transported into the bloodstream. The blood glucose load is influenced by both the total carbohydrate intake as well as the rate of starch digestion. Simple sugars such as glucose, fructose, and sucrose require minimal digestion, unlike starches, and are rapidly absorbed. Glucose-sensing by the pancreatic β -cells results in a proportional increase in insulin secretion. Hepatic sensing of glucose and fructose together with increased hepatic insulin-signaling promotes the activation of ChREBP, LXR, and SREBP to induce hepatic lipogenic gene expression, including SCD1.

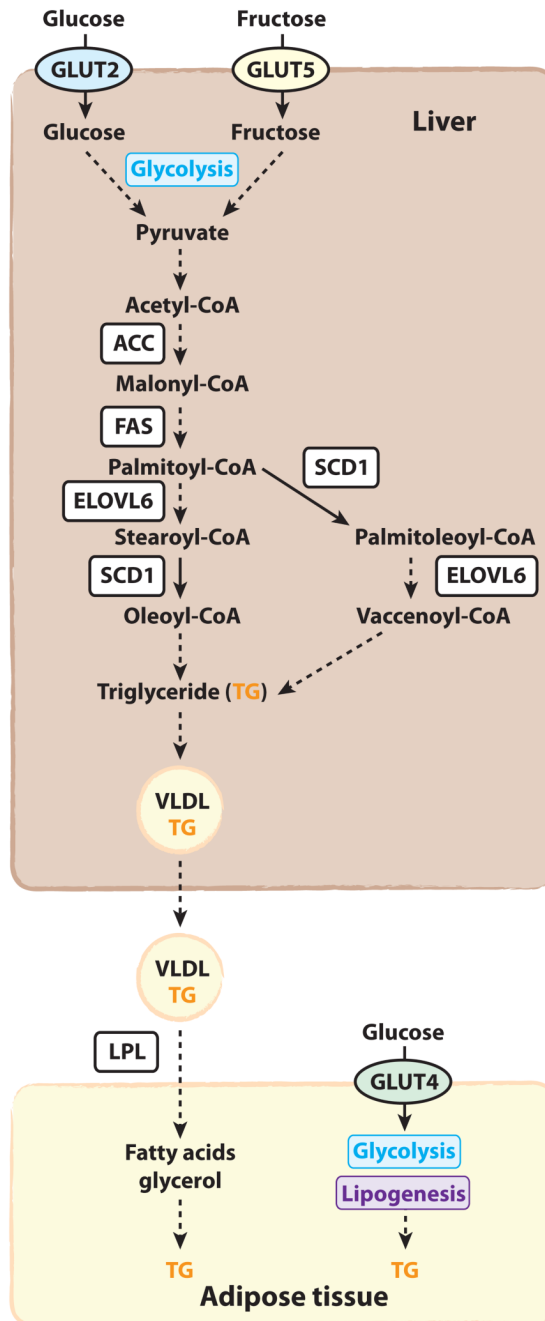


Figure 2.

In response to increased carbohydrates and insulin, the hepatic levels of lipogenic enzymes are increased (Figure 1). The augmented lipogenic capacity of the liver allows for the large load of dietary carbohydrate to be converted to fatty acids for storage as triglyceride (TG) and secretion as VLDL. SCD1 catalyzes the conversion of the saturated fatty acid products of FAS into MUFA, which are the preferred substrate for triglyceride synthesis. Peripheral tissues such as adipose tissue take up these triglycerides from VLDL in a lipoprotein-lipase (LPL) dependent mechanism. Adipocytes can also uptake glucose in a GLUT4-dependent mechanism for conversion to triglycerides.