

The regulation of ApoB metabolism by insulin

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The leading cause of death in diabetic patients is cardiovascular disease. Apolipoprotein B (ApoB)-containing lipoprotein particles, which are secreted and cleared by the liver, are essential for the development of atherosclerosis. Insulin plays a key role in the regulation of ApoB. Insulin decreases ApoB secretion by promoting ApoB degradation in the hepatocyte. In parallel, insulin promotes clearance of circulating ApoB particles by the liver via the low-density lipoprotein receptor (LDLR), LDLR-related protein 1 (LRP1), and heparan sulfate proteoglycans (HSPGs). Consequently, the insulin-resistant state of type 2 diabetes (T2D) is associated with increased secretion and decreased clearance of ApoB. Here, we review the mechanisms by which insulin controls the secretion and uptake of ApoB in normal and diabetic livers.

Cardiovascular disease in diabetes

Most individuals with diabetes ultimately die of cardiovascular disease (CVD). Hyperglycemia is a central feature of the diabetic state, and it has been suggested that hyperglycemia promotes cardiovascular disease by increasing inflammation, oxidative stress, and the formation of advanced glycosylation end products (for a review, see [1]). However, the normalization of serum glucose levels does not consistently reduce the risk of CVD in diabetic patients [2,3]. A more important factor may be derangements in lipoprotein metabolism. Although diabetic patients are not usually hypercholesterolemic, they do show abnormalities in lipoprotein kinetics and composition. Moreover, statin drugs, which reduce serum cholesterol, effectively reduce CVD risk in diabetic patients [4]. We review here the effects of insulin on the hepatic metabolism of ApoB-containing lipoprotein particles, as these are essential for the development of atherosclerosis.

Overview of lipoprotein metabolism

The lipoprotein system allows the intercellular transport of cholesterol and triglycerides, which serve distinct roles in the body. Triglycerides are an important source of energy. Cholesterol, by contrast, is a structural element of membranes as well as a precursor for steroids and bile acids. In contrast to triglycerides, most mammalian tissues

are not capable of catabolizing cholesterol. They must therefore return excess cholesterol to the liver, which can secrete cholesterol into the bile, either directly or after conversion to bile acids [5].

Lipoproteins can be divided into classes that vary in their apolipoproteins, lipid composition and, therefore, function. Apolipoprotein B is a structural protein that is an integral component of chylomicrons, as well as very low-density lipoprotein (VLDL), intermediate-density lipoprotein (IDL), and low-density lipoprotein (LDL) particles. Chylomicrons are secreted by the intestine and enable the transport of dietary triglycerides to other tissues after feeding. They contain a truncated form of ApoB, referred to as ApoB48 because it is 48% of the full-length protein [6]. Chylomicrons are lipolyzed within the vasculature, releasing their fatty acids to peripheral cells, particularly the adipose tissue, for storage. When most of the triglycerides have been removed, the chylomicron remnant is cleared by the liver [5].

VLDL, which is secreted by the liver, also transports triglycerides. In humans, VLDL contains only ApoB100, the full-length protein, whereas either ApoB100 or ApoB48 is present in the VLDL of mice and some other rodents [6]. As the VLDL particle delivers triglycerides to other tissues it is successively lipolyzed into an IDL and then LDL particle. LDL-cholesterol can be taken up by cells that require exogenous cholesterol. However, most LDL is cleared by the liver [5].

The other major class of lipoproteins is high-density lipoprotein (HDL). HDL does not contain ApoB, and high HDL levels are correlated with decreased risk of atherosclerosis. HDL can remove excess cholesterol from peripheral tissues, such as endothelial cells and macrophages, and return it to the liver for excretion from the body, a process known as reverse cholesterol transport [7].

Insulin plays a central role in coordinating lipoprotein metabolism. In the fed state, triglycerides and other nutrients are available from the gut. In this case, high insulin levels act on the adipocyte to promote triglyceride uptake and inhibit free fatty acid release. This promotes the storage of triglycerides for later use. In fasting, this process is reversed: insulin levels fall, and free fatty acids are released from the adipocyte and delivered to the liver. The liver re-esterifies the fatty acids to triglycerides and secretes them as VLDL [7]. Moreover, the ability of the adipose to take up triglycerides is impaired in the absence of insulin, and the triglycerides are therefore preferentially utilized by other tissues, such as skeletal muscle and liver [7,8].

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Regulation of VLDL production

Within the hepatocyte, insulin exerts its effects via a complex signaling cascade [9]. Some of the major branches of this cascade are shown in Figure 1. Upon binding insulin, the insulin receptor recruits the insulin receptor substrate (IRS) proteins, leading to activation of phosphatidylinositide 3-kinase (PI3K), which mediates many of the metabolic effects of insulin. PI3K activates Akt, which inhibits the transcription factor FoxO1 by phosphorylating it. Phosphorylated FoxO1 is excluded from the nucleus, and is therefore unable to activate transcription of its targets, including the gluconeogenic genes. Akt also stimulates glycogen synthesis by inhibiting glycogen synthase kinase 3 (GSK3). Furthermore, Akt activates lipogenesis, in part via the mammalian target of rapamycin complex 1 (mTORC1) signaling complex, which is necessary for the induction of the lipogenic transcription factor sterol regulatory element binding protein (SREBP)-1c [10]. Consequently, in the hepatocyte, insulin suppresses glucose production and stimulates the production of glycogen and triglycerides. It also acutely inhibits the secretion of ApoB [11].

The synthesis and secretion of ApoB are complex. Surprisingly, most ApoB protein is degraded before secretion, and the amount of ApoB secreted is determined largely by the proportion of newly synthesized polypeptide that escapes degradation [6]. There are multiple mechanisms

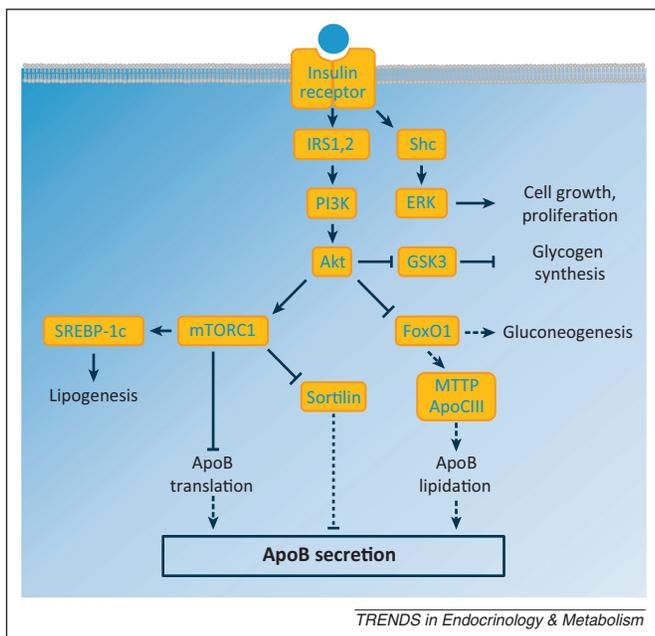


Figure 1. Insulin signaling cascade. Insulin binds to its receptor and recruits the insulin receptor substrate (IRS) proteins, leading to the activation of phosphatidylinositide 3-kinase (PI3K) and subsequently Akt. Akt serves as a major branch point in the signaling cascade: its inactivation of glycogen synthase kinase 3 (GSK3) de-represses and therefore stimulates glycogen synthesis, whereas its activation of mammalian target of rapamycin complex 1 (mTORC1) stimulates lipogenesis via the transcription factor sterol regulatory element binding protein 1c (SREBP-1c). Activation of mTORC1 also has dual effects on ApoB synthesis: it inhibits ApoB secretion by decreasing ApoB translation, but promotes ApoB secretion by suppressing sortilin. The inhibition of FoxO1 by Akt represses expression of gluconeogenic genes and thus decreases gluconeogenesis; it also inhibits the expression of microsomal triglyceride transfer protein (MTTP) and ApoCIII, thereby inhibiting lipitation of ApoB and decreasing ApoB secretion. Finally, insulin can also activate extracellular signal-regulated kinase (ERK) via Shc to stimulate cell growth and proliferation. Solid arrows denote pathways that are active during insulin signaling; dashed arrows indicate pathways that are inactive during insulin signaling.

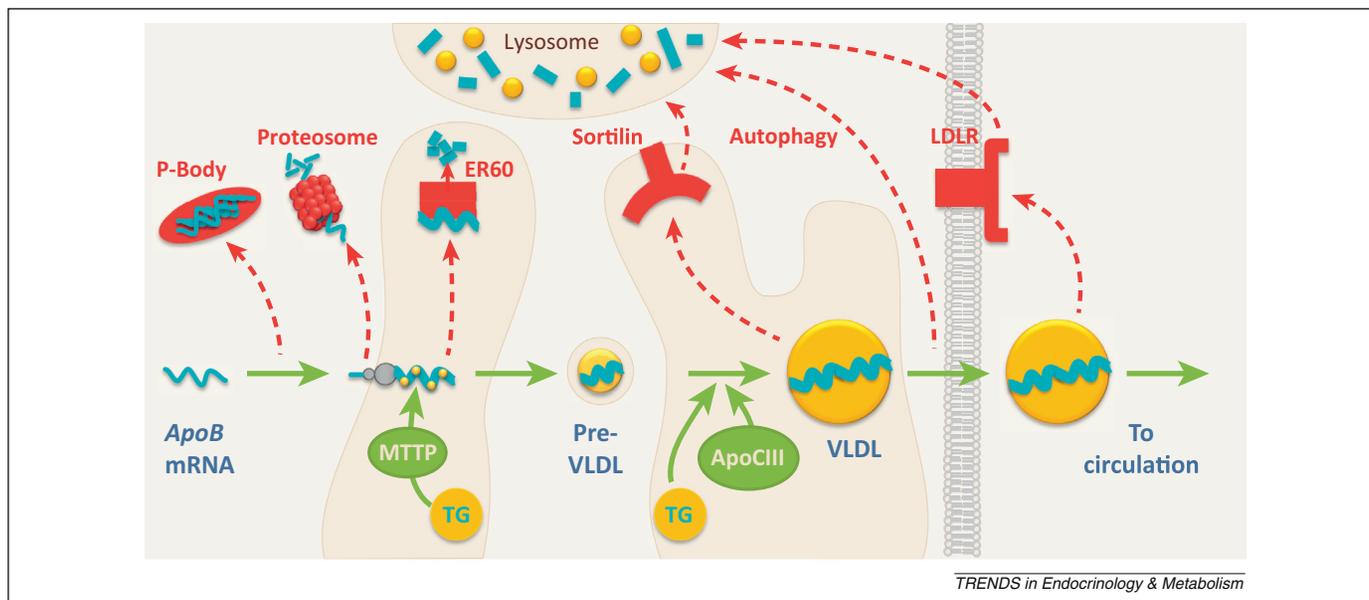
by which insulin regulates ApoB (Figure 2). Although insulin does not alter *ApoB* mRNA levels [12], it inhibits ApoB translation by promoting the trafficking of *ApoB* mRNA into P-bodies, aggregates of translationally repressed mRNAs [13,14]. The effects of insulin on ApoB translation require signaling through PI3K and mTORC1, and may involve the association of RNA-binding proteins with the *ApoB* mRNA 5'-untranslated region [15].

As ApoB is translated, it is lipidated by microsomal triglyceride transfer protein (MTTP). MTTP adds triglycerides to the nascent ApoB during its cotranslational translocation into the lumen of the endoplasmic reticulum [16]. This produces a dense, lipid-poor, pre-VLDL particle. The expression of MTTP is driven by FoxO1, and is thus inhibited by insulin [17]. Because ApoB must be lipidated by MTTP to escape proteosomal degradation [16], insulin may promote ApoB degradation by decreasing expression of MTTP. Consistent with this, insulin-resistant rodents show increased amounts of nuclear FoxO1, *Mttp* mRNA, and MTTP protein, as well as increased ApoB secretion [17,18]. However, acute insulin treatment in rodents does not decrease either *Mttp* mRNA or MTTP activity [19]; this may due to the long half-life of *Mttp* mRNA (~4.4 days) [20]. ApoB may also be regulated by ER60, an endoplasmic reticulum resident chaperone and protease that binds to and degrades ApoB when overexpressed [21]; interestingly, ER60 is decreased and ApoB secretion is increased in the insulin-resistant fructose-fed hamster [22].

A second lipidation step, in which the bulk of lipids are added to ApoB, occurs independently of MTTP [11,23]. This maturation step can be inhibited by insulin/PI3K signaling [24]. Although the exact mechanisms are unknown, one potential mediator is ApoCIII, an apolipoprotein secreted by the liver and to a lesser extent by the intestine. Individuals with a null allele of APOCIII manifest a 45% reduction in serum triglycerides and a 27% decrease in coronary heart disease risk [25]. ApoCIII is best known for its ability to inhibit lipoprotein lipase and hepatic lipase, and therefore lipolysis of triglycerides carried on chylomicrons and VLDL [26], but it has also been suggested to act intracellularly to promote ApoB secretion. Expression of ApoCIII in McA-RH7777 rat hepatoma cells, which lack endogenous ApoCIII, increases ApoB and triglyceride secretion by promoting the second step of ApoB lipidation [27]. However, ApoB secretion *in vivo* is not increased by ApoCIII expression and VLDL triglyceride secretion is increased only upon expression of very high levels of ApoCIII [28].

Transcription of *ApocIII*, like *Mttp*, is induced by FoxO1, and therefore inhibited by insulin. *ApocIII* transcription is also induced by the carbohydrate response element binding protein (ChREBP), which is activated by glucose [29], and peroxisome proliferator-activated receptor gamma coactivator 1 β (PGC-1 β), which is activated by fatty acids [30]. In the diabetic state, the presence of insulin resistance, hyperglycemia, and elevated free fatty acids would all be expected to drive *ApocIII* transcription. Indeed, ApoCIII levels are increased in the serum of patients with T2D [31].

The triglycerides used for ApoB lipidation are derived from three sources: free fatty acids released from the



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Figure 2. Synthesis of ApoB and secretion of very low-density lipoprotein (VLDL). During translation, ApoB is lipidated with triglycerides (TG) by microsomal triglyceride transfer protein (MTTP) in the endoplasmic reticulum (ER) to form a lipid-poor pre-VLDL particle. The pre-VLDL particle is trafficked through the ER and Golgi compartments, where additional triglycerides are added in an MTTP-independent manner promoted by ApoCIII, before secretion of the mature VLDL particle into the circulation. The synthesis and secretion of ApoB can be inhibited by multiple mechanisms, including proteasomal, ER60-mediated, autophagic, and lysosomal degradation. Green arrows indicate ApoB–VLDL synthesis pathways; red dashed arrows indicate ApoB degradation pathways.

adipocyte, hepatic uptake of chylomicron and VLDL remnants, and *de novo* lipogenesis. Interestingly, insulin has opposing effects on the different sources of hepatic triglycerides. Although insulin decreases the free fatty acid supply to the liver by suppressing adipose tissue lipolysis, it promotes the uptake of remnant particles (see below) and lipogenesis (for a review, see [11]).

The lipidated ApoB particle can also be degraded before being secreted, and this may be a more important site of insulin action, at least *in vitro*. Although the exact mechanisms are again unclear, insulin stimulation of ApoB degradation is blocked by brefeldin A, which inhibits intracellular transport between the endoplasmic reticulum and the Golgi [32], as well as by knockout of Atg5, a protein necessary for autophagy [33].

Insulin may also regulate the lipidated ApoB particle via sortilin [34,35]. Sortilin is a member of the Vps10p-domain family of transmembrane receptors, which bind and traffic extracellular ligands to the lysosome; sortilin can also traffic Golgi-localized ligands to the lysosome [36]. Sortilin appears to target ApoB to the lysosome for degradation because the ability of sortilin to inhibit ApoB secretion is blocked by mutations that interfere with sortilin trafficking to the lysosome and by disruptors of lysosome function [34]. Consistent with this, sortilin overexpression decreases VLDL secretion in most, although not all, studies [34,37,38]. Surprisingly, recent work has shown that sortilin is suppressed by mTORC1 [38], a downstream target of insulin. This suggests that insulin may actually promote ApoB secretion via sortilin, although overall it inhibits ApoB secretion via other pathways.

Finally, insulin can antagonize ApoB secretion by promoting the re-uptake of secreted VLDL back into the hepatocyte before it exits the space of Disse, located between the hepatocyte and endothelial cell. This re-uptake is mediated by the LDL receptor and potentially HSPGs

[39,40]. If the VLDL particle escapes re-uptake, it can enter the circulation and deliver its triglyceride cargo to other tissues.

Regulation of lipoprotein clearance

ApoB particles are removed from the circulation primarily by the liver; this process is promoted by insulin and impaired in diabetes [41–43]. Here, we examine the effects of insulin on the hepatic receptor systems that mediate ApoB clearance: the LDLR, LRP1, and HSPGs. However, it is important to note that other factors in the diabetic state could also contribute to delayed ApoB clearance. First, there is a defect in triglyceride lipolysis. Second, hyperglycemia and increased oxidative stress promote glycosylation and oxidation of lipoprotein particles, and this could interfere with their uptake and clearance [44,45]. Finally, the overproduction of lipoproteins from the liver as well as the intestine (for a review, see [46]) could indirectly impair lipoprotein clearance by competing for the clearance apparatus.

LDL receptor (LDLR)

LDLR, which binds to ApoB100 and ApoE, is a key determinant of serum cholesterol levels and cardiovascular disease risk. Over 30 years ago insulin was shown to promote LDL uptake in fibroblasts [47,48]. Subsequent studies in cultured hepatoma cells showed that insulin increases *LDLR* mRNA [49]. The effects of insulin on *LDLR* mRNA appear to be mediated by the transcription factor SREBP-1c, which also mediates the effects of insulin on lipogenesis. Thus, in the livers of rodents with decreased insulin levels due to fasting [50] and type 1 diabetes (T1D) [51], *Ldlr* mRNA levels are decreased.

The effects of insulin on LDLR protein, however, are less clear. The livers of mice with liver-specific knockdown [52] or knockout [53] of the insulin receptor show reduced levels

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of Ldlr protein. By contrast, Ldlr protein levels in the liver do not change with fasting or T1D [50,51], and reports of Ldlr protein levels in obesity/T2D are conflicting [52,54].

The discordance between LDLR mRNA and protein levels under particular circumstances could be due to proprotein convertase subtilisin/kexin type 9 (PCSK9). PCSK9 is a secreted protein that binds to the LDLR and promotes its degradation. *Pcsk9* knockout mice have increased levels of Ldlr protein and therefore increased LDL clearance [55]. Humans with gain of function mutations in *PCSK9* show increased levels of LDL-cholesterol and increased risk of cardiovascular disease, whereas individuals with loss of function mutations in *PCSK9* show decreased levels of LDL-cholesterol and reduced risk of cardiovascular disease [55]. Moreover, therapies to inhibit PCSK9 have recently been shown to be effective in reducing LDL-cholesterol levels [56].

Interestingly, insulin increases *PCSK9* mRNA *in vitro* [50]. *In vivo*, fasting and T1D decrease insulin and *Pcsk9* mRNA [50,51]. In these models, the reduced expression of *Pcsk9* has been invoked to account for the fact that the abundance of Ldlr protein does not change even as *Ldlr* mRNA is decreased. Taken together, these data illustrate the fact that the effects of insulin and diabetes on LDLR are complex, because the ability of insulin to induce PCSK9 could reduce LDLR expression, even as insulin action on SREBP-1c and other pathways promotes LDLR expression.

LDLR-related protein 1 (LRP1)

LRP1 is another member of the LDLR family. It contains a very large extracellular domain capable of binding more than 30 ligands, including lipoproteins, growth factors, and protease inhibitors [57]. Although LRP1 is unable to bind to ApoB and therefore LDL, it is able to bind remnant particles via molecules such as ApoE [57]. However, the role of LRP1 in lipoprotein clearance remains unclear. In *Ldlr*-deficient mice, liver-specific knockout of *Lrp1* leads to impaired removal of chylomicron remnants and a 2- to 3-fold increase in plasma cholesterol and triglycerides [58]. In mice that are wild type for *Ldlr*, however, liver-specific knockout of *Lrp1* has little effect on plasma lipoproteins [58]. Thus, LRP1 is not necessary for lipoprotein clearance in the presence of functional LDLR.

Insulin has been shown to promote the translocation of LRP1 from intracellular vesicles to the cell surface [59,60]. Thus, treatment with glucose, which induces insulin secretion, increases LRP1 protein in plasma membrane preparations [59]. It also increases uptake of the LRP1 ligands, α 2-macroglobulin and chylomicron remnants. The ability of insulin to promote α 2-macroglobulin uptake is blunted in *ob/ob* mice, a model of T2D, suggesting that obesity and insulin resistance could cause impaired LRP1 translocation [59]. In the adipocyte, LRP1 exists in the same vesicles as the Glut4 transporter, the major mediator of insulin-sensitive glucose transport [61], suggesting that the same signaling mechanisms utilized by insulin to translocate Glut4 may be used to translocate LRP1.

Interestingly, LRP1 also mediates the clearance of activated factor VIII, which promotes clotting: mice deficient in *Lrp1* show increased levels of circulating factor VIII and decreased factor VIII clearance [62]. LRP1 also appears to

increase HDL-cholesterol, as hepatic knockout of *Lrp1* decreases HDL synthesis and lowers HDL levels [63]. Thus, defects in LRP1 could potentially account for multiple features of obesity/T2D: a pro-coagulant state, low HDL-cholesterol, and, if there are concomitant defects in the LDLR, post-prandial lipemia.

Heparan sulfate proteoglycans (HSPGs)

HSPGs consist of a core polypeptide onto which numerous negatively charged sugar moieties, heparan sulfates, are attached [64]. The result is a large, highly negatively charged, flexible structure. In the liver, HSPGs are positioned to trap lipoproteins within the space of Disse, rapidly clearing them from the circulation; they may also promote the uptake of these particles into hepatocytes by internalizing them directly, or facilitating their interaction with other receptors [64,65]. The liver expresses multiple HSPGs, including perlecan, syndecans 1, 2, and 4, and glypicans 1 and 4 [66]. The role of HSPGs, particularly syndecan-1, in lipoprotein clearance has been suggested by the findings that liver-specific knockout of either syndecan-1 (*Sdc1*) or GlcNAc *N*-deacetylase/*N*-sulfotransferase-1 (*Ndst1*), an enzyme that participates in the assembly of heparan sulfate moieties, can reduce clearance of remnant lipoproteins [66,67]. The reduction in remnant clearance occurs even in mice wild type for *Ldlr*, indicating that deficiencies in the HSPGs, unlike *Lrp1*, cannot be compensated for by Ldlr.

Over 30 years ago, insulin deficiency was found to produce defects in the sulfation of proteoglycans [68], although this has not been found in all studies [69]. Moreover, perlecan protein and *Ndst1* mRNA are decreased in insulin-deficient rodents [70,71]. In addition, *db/db* mice, a model of T2D, show increased expression of heparan sulfate glucosamine 6-*O*-endosulfatase 2 (*Sulf2*), an enzyme that promotes HSPG degradation [72]. Knockdown of *Sulf2* in *db/db* mice increased the clearance of chylomicron remnants and lowered plasma triglyceride levels [73].

Other players

The lipolysis-stimulated lipoprotein receptor (LSR), which is expressed primarily in hepatocytes, promotes the uptake of ApoB- and ApoE-containing lipoproteins in the presence of high concentrations of fatty acids [74]. Interestingly, *Lsr* is decreased in mouse models of T1D [75] and T2D [76]. Knockdown of *Lsr* using adenoviruses encoding shRNA against *Lsr* increased serum triglycerides, whereas adenoviral reconstitution of *Lsr* in *ob/ob* mice lowered serum triglyceride and cholesterol levels [76]. Second, ApoCIII interferes with the binding of ApoB particles to the LDL receptor, preventing their removal from the circulation [77]; this is in addition to its effects on lipolysis and ApoB secretion. Finally, sortilin not only inhibits ApoB secretion but also promotes the clearance of ApoB-containing lipoproteins, because overexpression of sortilin in the liver increases, and knockout of sortilin decreases, the fractional catabolic rate of LDL [34]. Consistent with this, genome-wide association studies in humans have shown that polymorphisms predicted to increase sortilin expression are strongly associated with lower LDL-cholesterol levels [78].

The diabetic dyslipidemia

Given the central role of insulin in the control of ApoB lipoprotein metabolism, it is not surprising that derangements in insulin signaling produce dyslipidemia. Indeed, liver-specific knockout of the insulin receptor increases secretion of ApoB, decreases LDL clearance, and markedly increases susceptibility to atherosclerosis in mice [53].

T2D is the most common form of diabetes and is associated with obesity. It is primarily due to insulin resistance – that is, the failure of insulin to maintain glucose homeostasis – resulting in hyperglycemia and a compensatory hyperinsulinemia. For example, the failure of insulin to suppress FoxO1 promotes gluconeogenic gene expression and hyperglycemia. Interestingly, however, not all components of the insulin signaling cascade become impaired in T2D, a phenomenon known as ‘selective insulin resistance’. For example, mTORC1 remains active in T2D, presumably because it either remains sensitive to insulin and is driven by the hyperinsulinemia present [79], or is activated directly by excess nutrients [80].

In normal physiology, insulin suppresses FoxO1 in the liver, decreasing MTP and ApoCIII expression, and therefore, ApoB secretion. Insulin also suppresses lipolysis in adipose tissue, decreasing fatty acid flux to the liver. Although insulin also stimulates lipogenesis and suppresses sortilin, the net effect of insulin is to suppress ApoB secretion. With the development of T2D, however, FoxO1 becomes disinhibited, leading to increased expression of MTP and ApoCIII (Figure 3). At the same time, mTORC1 remains activated, suppressing sortilin. Consistent with this, the ability of insulin to suppress ApoB secretion is lost in humans with T2D [81,82] and ApoB secretion is increased [81,83–86].

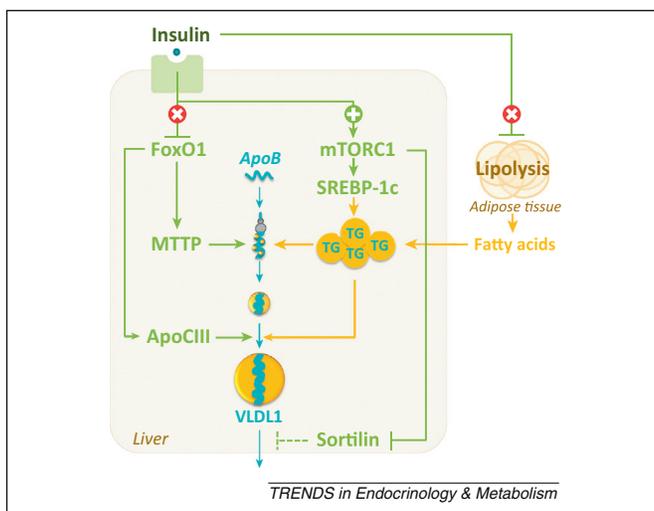


Figure 3. Type 2 diabetes (T2D) drives very low-density lipoprotein (VLDL) secretion through multiple pathways. T2D is characterized by selective insulin-resistance: insulin fails to suppress lipolysis in the adipose tissue and FoxO1 in the liver, but is still able to activate mammalian target of rapamycin complex 1 (mTORC1). The disinhibition of FoxO1 leads to increased expression of microsomal triglyceride transfer protein (MTP) and ApoCIII, thereby promoting ApoB secretion. At the same time, the stimulation of mTORC1 in the liver leads to activation of sterol regulatory element binding protein 1c (SREBP-1c) and increased lipogenesis, as well as suppression of sortilin. Because sortilin increases ApoB degradation, the suppression of sortilin further promotes ApoB secretion. Finally, the increase in lipogenesis, coupled with the increased flux of fatty acids to the liver, enlarges the pool of triglycerides (TG) available to lipidate ApoB, driving the formation of lipid-rich VLDL1 particles. The overall result is an increase in the secretion of ApoB in the form of VLDL1 particles in T2D.

T2D also promotes the accumulation of hepatic triglycerides, which fuel the lipidation of ApoB. First, the ability of insulin to suppress lipolysis in the adipocyte is lost in T2D, leading to increased free fatty acid flux to the liver. Second, the ability of insulin to stimulate mTORC1, SREBP-1c, and hepatic lipogenesis appears to remain intact [87]. The resulting excess of triglycerides promotes the formation of the VLDL1 subclass of particles, which are larger and more triglyceride-rich than the VLDL2 subclass of particles [11,88,89]. Consistent with this, patients with T2D show an increase in the secretion of VLDL1, rather than VLDL2, particles [83,90]. Finally, T2D is associated with a decrease in triglyceride lipolysis [91] as well as ApoB clearance [42,43].

Hypertriglyceridemia is therefore common in T2D. Hypertriglyceridemia promotes the development of small, dense LDL, and decreased HDL via the enzyme cholesterol ester transfer protein (CETP). CETP, which is not present in mice and rats, catalyzes the transfer of triglycerides in VLDL for cholesterol esters in LDL and HDL, resulting in triglyceride-rich LDL and HDL particles. Triglyceride-rich LDL particles can be lipolyzed to small, dense LDL, which are thought to be highly atherogenic because of their greater susceptibility to oxidation, reduced affinity for the LDL receptor, and increased uptake by arterial tissue [45]. Triglyceride-rich HDL particles are more susceptible to degradation, leading to lower HDL-cholesterol levels. Thus, increased VLDL1 secretion is central to the pathogenesis of all of the features of the dyslipidemia of T2D: hypertriglyceridemia, decreased HDL-cholesterol, and increased small dense LDL-cholesterol. Note that hypercholesterolemia, although an established risk factor for cardiovascular disease, is not usually present in patients with T2D [92].

T1D, by contrast, is due to autoimmune destruction of the β cells of the pancreas, leading to hypoinsulinemia and decreased insulin action on all pathways. Consequently, lipogenesis is decreased in rodent models of T1D [93]. Nonetheless, patients with untreated T1D have decreased lipoprotein clearance and hypertriglyceridemia [94,95]; those with poorly controlled T1D also have hypercholesterolemia, and increased VLDL-cholesterol and LDL-cholesterol [96,97]. Well-controlled T1D patients, who are made insulin-replete by injections of exogenous insulin, however, generally show normal plasma triglycerides and cholesterol, normal or decreased LDL-cholesterol, and normal or increased HDL-cholesterol [96,97], indicating that exogenous insulin is sufficient to restore lipoprotein homeostasis.

Concluding remarks

In summary, insulin plays a central role in the secretion and clearance of ApoB lipoproteins, which play a key role in promoting atherosclerosis. Although diabetes is not usually associated with hypercholesterolemia, there are important changes in lipoprotein kinetics, including increased secretion and decreased clearance of ApoB particles. Consequently, the number of LDL particles, which is closely tied to atherosclerosis risk, is elevated [98]. Moreover, qualitative differences in LDL – such as the formation of small dense LDL, which are potentially intrinsically more

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atherogenic – are present. Thus, even though the development of cardiovascular disease is undoubtedly multifactorial, defining and reversing the changes in ApoB metabolism that occur in diabetes could prove most effective in reducing mortality.

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