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Lipid-induced insulin resistance: unravelling the mechanism

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Abstract

Insulin resistance has long been associated with obesity. More than 40 years ago, Randle and colleagues postulated that lipids impaired insulin-stimulated glucose use by muscles through inhibition of glycolysis at key points. However, work over the past two decades has shown that lipid-induced insulin resistance in skeletal muscle stems from defects in insulin-stimulated glucose transport activity. The steatotic liver is also resistant to insulin in terms of inhibition of hepatic glucose production and stimulation of glycogen synthesis. In muscle and liver, the intracellular accumulation of lipids—namely, diacylglycerol—triggers activation of novel protein kinases C with subsequent impairments in insulin signalling. This unifying hypothesis accounts for the mechanism of insulin resistance in obesity, type 2 diabetes, lipodystrophy, and ageing; and the insulin-sensitising effects of thiazolidinediones.

Introduction

Obesity is now a pandemic that is largely caused by a combination of our genetics, evolutionary pressures that favour metabolic efficiency,¹ and a modern environment in which highly palatable, calorie-dense food is widely available and inexpensive.² There are now more overweight than underweight people worldwide, and children are increasingly at risk of becoming obese.^{3–5} In tandem with the obesity epidemic, the prevalence of related disorders, such as metabolic syndrome, non-alcoholic fatty liver disease, and type 2 diabetes mellitus, is also rising. Insulin resistance plays a crucial part in the pathogenesis of all these disorders, yet the cellular mechanisms are still poorly understood. Here, we review studies in human beings and rodents that have informed our current understanding of the mechanistic links between lipid accumulation and insulin resistance. We first discuss some of the pioneering studies in this specialty.

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All authors contributed to writing this Seminar, and have approved the final version.

Conflicts of interest

We declare that we have no conflicts of interest.

Glucose-fatty-acid cycle

Randle and colleagues⁶ postulated a mechanism more than 40 years ago by which fatty acids could impair insulin-stimulated glucose oxidation in muscle. They reported that incubation of preparations of the rat heart with fatty acids increased intracellular concentrations of glucose-6-phosphate (G6P) and glucose, and incubation of preparations of diaphragm increased intracellular concentrations of glycogen (figure 1). According to Randle and colleagues' theory, fat oxidation increased the ratios of acetyl coenzyme A to coenzyme A and NADH to NAD⁺ in the mitochondria, which in turn resulted in the inactivation of pyruvate dehydrogenase. Accumulation of citrate inhibits phosphofructokinase and thus increases intracellular concentrations of G6P, promoting glycogen synthesis and inhibiting hexokinase. The resulting intracellular accumulation of glucose prevents further glucose uptake. Thus, in their model, the availability of lipids as a source of fuel generated metabolic signals that impaired the use of glucose through inhibition of the key glycolytic enzymes.

Testing Randle and colleagues' hypothesis

Investigation of the association between fatty acids and insulin resistance is difficult in individuals who are already obese or diabetic because of the confounding effects of other co-morbidities. These effects are avoided by investigation of the mechanisms of insulin resistance in the offspring of patients with type 2 diabetes mellitus, who are young, lean, and insulin resistant. When Perseghin and colleagues⁷ compared such individuals with controls matched for age and body-mass index (BMI), they noted an inverse correlation between plasma concentrations of fatty acids and insulin sensitivity. However, insulin sensitivity was more tightly associated with intramyocellular lipid content (assessed non-invasively by use of proton [1H]-magnetic resonance spectroscopy [MRS]) in lean offspring of patients with type 2 diabetes mellitus,^{8,9} and with intramuscular triglyceride content in muscle biopsy samples from non-diabetic male Pima Indians.¹⁰

Does insulin resistance alter the intramyocellular concentration of G6P and glycogen in human beings? In Randle and colleagues' proposed theory about the glucose-fatty-acid cycle, muscle glycogen content was predicted to be high as a result of the accumulation of intracellular glucose and G6P. Measurement of these metabolites in vivo is difficult; repeated muscle biopsies are needed to measure a rate of change, and metabolite concentrations are affected by even brief periods of ex-vivo hypoxia.¹¹ Use of MRS circumvents these difficulties, allowing non-invasive and real-time, sequential measurement of these metabolites in situ and thus avoiding the confounding effects of hypoxia. 13-carbon [13C]-MRS and 31-phosphorus [31P]-MRS were used to non-invasively measure insulin-stimulated changes in muscle glycogen, and in intramyocellular G6P in patients with type 2 diabetes mellitus and non-diabetic first-degree relatives.¹²⁻¹⁴ When measured by use of 13C-MRS, the rates of insulin-stimulated glucose uptake and glycogen synthesis were more than 50% lower in the patients with diabetes than in the control individuals,¹³ and was associated with a reduction in concentrations of G6P in muscle.¹⁴ Similarly, insulin-stimulated concentrations of G6P and rates of muscle glycogen synthesis were reduced in lean, normoglycaemic, insulin-resistant first-degree relatives of patients with type 2 diabetes mellitus.¹²

Could these changes be induced in healthy, otherwise insulin-sensitive individuals? Lipid infusions combined with heparin to activate lipoprotein lipase, raised plasma concentrations of fatty acids, promoted accumulation of muscle lipids,¹⁵ and impaired oral glucose tolerance¹⁶ and insulin-stimulated glucose disposal in healthy individuals.^{17,18} Roden and colleagues¹⁷ combined lipid infusions with the normoglycaemic-hyperinsulinaemic clamp in healthy individuals while monitoring synthesis of muscle glycogen with 13C-MRS, and

concentrations of G6P with 31P–MRS. After about 3 h of hyperlipidaemia, muscle G6P concentrations were actually reduced, followed by a decrease in insulin-stimulated rates of glucose disposal. These findings contrasted with Randle and colleagues' hypothesis and results of previous studies in which an increase was reported in intramyocellular concentrations of G6P in muscle biopsy samples during similar conditions.¹⁹ Thus, in healthy individuals, exposure to high concentrations of plasma fatty acids caused insulin resistance associated with an induced defect in either glucose transport or phosphorylation activity, not an impairment in glycolysis.

To discern which of these two possible effects takes place, Dresner and colleagues²⁰ used a novel 13C–MRS method to measure intramyocellular concentrations of free glucose in healthy people under similar conditions of high and low plasma concentrations of fatty acids. If there was a block at the hexokinase step, the concentration of intramyocellular glucose would be expected to increase. Instead, raising plasma fatty acid concentration attenuated the accumulation of intracellular glucose, implying that insulin-stimulated glucose transport activity was blunted. Cline and colleagues²¹ studying individuals with type 2 diabetes mellitus and healthy controls during matched hyperinsulinaemic-normoglycaemic conditions, reported that intramyocellular concentrations of glucose were much lower than predicted had hexokinase been controlling the rate of insulin-stimulated glucose uptake by muscle. This result suggested that in people with type 2 diabetes mellitus, impairment of insulin-stimulated glucose use by muscle was also largely due to reductions in insulin-stimulated glucose transport.

In parallel with these studies, there were complementary advances in knowledge about glucose transporter (GLUT) proteins. The GLUT family of proteins (also called solute carriers 2A [SLC2A]) is diverse, with 13 isoforms.²² However, GLUT4 is distinguished from the others as a high-affinity, insulin-responsive transporter that is highly expressed in muscle and adipose tissue.²³ Insulin-mediated translocation of GLUT4 to the muscle sarcolemmal membrane was impaired in patients with type 2 diabetes mellitus.^{24,25} Thus, a defect in glucose transport, not glycolysis, was implicated as the cause of reduced insulin-mediated glucose metabolism in patients with type 2 diabetes mellitus in two independent studies.^{24,25} Moreover, the results of these studies suggested that ectopic accumulation of lipid within the muscle might be the cause of insulin resistance.

Diacylglycerol-induced insulin resistance

The coordinated intracellular response to insulin requires an intricate relay of signals. In skeletal muscle, insulin binds to its receptor, activating the receptor tyrosine kinase activity, with subsequent phosphorylation and activation of insulin-receptor substrate 1 (IRS1; figure 2). When phosphorylated, IRS1 activates 1-phosphatidylinositol 3-kinase (PI3K). This enzyme, through signalling intermediates, activates Akt2, which phosphorylates and inactivates AS160, a protein that prevents translocation of GLUT4 through its interaction with Rab proteins.²⁶ Thus, insulin promotes the docking and fusion of GLUT4-containing vesicles to the plasma membrane. The tyrosine phosphorylation of IRS1 and associated activation of PI3K were impaired in rodent models of insulin resistance.^{27,28} Similarly, IRS1-associated PI3K activity was greatly reduced in the muscles of individuals being given lipid infusions, indicating that the lipid-induced reduction in insulin-stimulated glucose transport was attributable to a defect in insulin signalling.²⁰ But what connects lipid accumulation with impaired insulin action in muscle? This answer is related to the family of protein kinase C (PKC) serine-threonine kinases.

The PKC family consists of three main groups: conventional (α , β I, β II, and γ), novel (δ , ϵ , η , and θ), and atypical (ζ and λ).²⁹ Classic PKCs become activated when calcium binds to

the C2 domain, increasing the affinity of the C1 domain for diacylglycerol, which then removes a pseudosubstrate from the catalytic domain. By contrast, the C1 domains of novel PKCs intrinsically have much greater affinity for diacylglycerol,³⁰ though subtle structural differences alter this affinity and suggest different thresholds for activation.³¹ The potential role of PKCs in regulating insulin action has long been recognised. Phorbol esters potently and non-selectively activate PKCs, and impair activation of the insulin receptor in vitro.^{32–35} Thus, the accumulation of diacylglycerol within the cells might similarly activate PKCs.

In rodents, a high-fat diet increased the concentration of diacylglycerol in muscles and activated novel PKCs.³⁶ Similarly, infusion of lipid and heparin or heparin for 5 h caused insulin resistance in muscles that was associated with accumulation of intracellular diacylglycerol and specific activation of PKC θ .³⁷ In this model, insulin resistance was attributed to lipid-induced defects in the insulin signalling pathway that originated from a reduction in tyrosine phosphorylation of IRS1.³⁸ By contrast, phosphorylation of IRS1 at the serine-307 residue was increased, preventing IRS1 from interacting with the insulin receptor.³⁹ The importance of activation of novel PKCs and serine phosphorylation of IRS1 for the development of insulin resistance was shown in mice without PKC θ ,⁴⁰ and in those with serine-to-alanine mutations in key residues of IRS1 (thereby preventing serine phosphorylation).⁴¹ Though lipid accumulation in muscle was similar to that in wild-type mice, those with the serine-to-alanine mutations were protected from fat-induced insulin resistance in muscle. Consistent with the results of these studies, Itani and colleagues reported activation of both muscle PKC β 2 and PKC δ after lipid or heparin infusions,⁴² and PKC θ in patients with type 2 diabetes mellitus.⁴³ Increased serine phosphorylation of IRS1 has also been noted in the muscles of individuals who are insulin-resistant.^{44,45} However, still not known is how the activation of novel PKCs might relate to serine phosphorylation of IRS1, and which kinases (ie, jun-N terminal kinase, I κ kinase β) might have a role in the pathway.

Diacylglycerol hypothesis

Insulin resistance develops with the accumulation of fatty-acid metabolites (namely diacylglycerols) within insulin-responsive tissues.⁴⁶ Genetic murine models have been invaluable in establishing this theory—eg, tissue-specific overexpression of lipoprotein lipase promotes tissue-specific lipid accumulation and selective insulin resistance.⁴⁷ By contrast, prevention of lipid entry into muscle by removal of lipoprotein lipase,⁴⁸ or other proteins involved in fat transport (CD36/49/50 or FATP151) prevents lipid accumulation in muscles and protects against insulin resistance. Increasing energy expenditure also protects against lipid accumulation, as shown in mice overexpressing muscle-specific uncoupling protein 352 or not expressing acetyl coenzyme A carboxylase 2.⁵³ Hoehn and colleagues⁵⁴ described a different ACC2 knockout mouse that, despite the expected increases in fat oxidation, did not have an increased energy expenditure, and gained similar amounts of weight when given a high-fat diet. Together, these data suggest that a shift in substrate preference, without an increase in total energy expenditure, will not protect against fat-induced insulin resistance.

Further insights were gained with animal models in which obesity and triglyceride content were disassociated from insulin action—eg, the *ob/ob* adiponectin transgenic mouse is leptin deficient (*ob/ob*) and overexpresses adiponectin, an adipokine that improves insulin sensitivity. These mice are much heavier than *ob/ob* mice, but have lower liver content of triglycerides and diacylglycerol, and as a result are more insulin sensitive, which is consistent with the diacylglycerol hypothesis.⁵⁵ In human beings, plasma adiponectin concentrations are directly related to insulin sensitivity and inversely related to ectopic lipid

accumulation.^{56–60} Thus, adiponectin might play an important part in mediating ectopic fat accumulation, and the development of insulin resistance.

Diacylglycerol acyl transferase (DGAT) 1 transfers a fatty acid from a fatty acyl coenzyme A to diacylglycerol to make a triglyceride. Mice overexpressing DGAT1 in skeletal muscles (MCK-DGAT1)⁶¹ accumulate tri glycerides in their muscles, but are protected from fat-induced muscle insulin resistance. These mice model the paradox of elite endurance athletes who are insulin sensitive but who have increased triglyceride content in their muscles.^{62,63} In the MCK-DGAT1 mice, though muscle triglyceride content was increased, diacylglycerol content was low, consistent with protection from fat-induced insulin resistance. In man, a single bout of exercise was sufficient to induce similar changes, in which increased DGAT1 expression in muscle was associated with increased concentrations of triglyceride, reduced concentrations of diacylglycerol, and improved insulin sensitivity.⁶⁴ Diacylglycerol can also be converted into phosphatidic acid, a major membrane lipid, by diacylglycerol kinases.⁶⁵ Expression of diacylglycerol kinase δ was decreased in skeletal muscles of rodents with hyperglycaemia and patients with poorly controlled diabetes.⁶⁵ In mice, haploinsufficiency of this enzyme increased the muscle content of diacylglycerol, but not of triglycerides, and resulted specifically in muscle insulin resistance. By disassociation of diacylglycerol and triglyceride content, the results of these studies support the hypothesis that diacylglycerol also plays a part in causing muscle insulin resistance in individuals with poorly controlled type 1 diabetes.

Diacylglycerol-induced insulin resistance in muscle and liver can readily be explained in most forms of obesity, in which increased delivery of fatty acids overwhelms the capacity of cells to oxidise fat or convert diacylglycerols to triacylglycerols. The increases in intramyocellular lipid content in healthy, lean, insulin-resistant offspring of parents with type 2 diabetes mellitus,⁹ and in healthy lean, elderly individuals have not been completely accounted for.⁶⁶ One possibility is a reduction in lipid oxidation.⁴⁴ The oxidative capacity of muscles can be quantified in vivo by measurement of the rates of flux in the citric acid cycle (with ¹³C–MRS in combination with an infusion of ¹³C–acetate), and the rates of ATP synthesis (with ³¹P–MRS).⁶⁷ In lean, insulin-resistant offspring of parents with type 2 diabetes mellitus, the rate of ATP synthesis was reduced by 30% compared with controls matched for age and BMI.⁶⁸ Befroy and colleagues⁶⁹ noted that the flux of the citric acid cycle in muscle was reduced by 30% in a similar group of individuals. This decrease in flux in the citric acid cycle and in ATP synthesis was similar to the 38% reduction in mitochondrial density.⁴⁴ Thus, at least in this cohort, a reduction in mitochondrial content probably accounted for the reduced rate of oxidative phosphorylation in mitochondria. Whether this reduction in mitochondrial density is a primary cause of lipid oxidation or acquired as a result of the lipid oxidation is unknown. Although results of some microarray studies have indicated a reduction in expression of the peroxisome proliferator-activated receptor (PPAR) γ coactivator 1 α in the muscles of patients with type 2 diabetes mellitus,^{70,71} they were not replicated in lean, insulin-resistant first-degree relatives of patients with type 2 diabetes mellitus, which suggests that other factors bring about the reduction in mitochondrial content in these individuals.⁴⁴ Irrespective of whether the reduction in mitochondrial content in the muscles is a primary defect, at the very least it reduces the mitochondrial fatty acid oxidation that promotes the accumulation of diacylglycerol within the muscle and contributes to the development of skeletal muscle insulin resistance in young, lean, insulin-resistant offspring of parents with type 2 diabetes mellitus.

Although the offspring of patients with type 2 diabetes mellitus have an inherited predisposition towards intramyocellular lipid accumulation and insulin resistance, almost all people manifest similar changes with increasing age. Results of population-based studies have clearly shown an association between ageing and insulin resistance.^{72–74} As in the

young insulin-resistant individuals, the presence of peripheral insulin resistance in the elderly was associated with accumulation of intramyocellular lipids.^{66,75,76} To investigate the potential role of decreased oxidative and phosphorylation activities in this process, Petersen and colleagues⁶⁶ used ¹³C-MRS and ³¹P-MRS to measure muscle mitochondrial function in a group of young and old people carefully matched not only for BMI, but also for body composition and activity. They noted that age-related decreases in the oxidative and phosphorylation activities were associated with increases in intramyocellular content of triglycerides and insulin resistance. These findings along with those from the studies of the insulin-resistant offspring of patients with type 2 diabetes mellitus, suggest that impaired mitochondrial function, with an impaired capacity to oxidise fatty acids, predisposes muscle to intramyocellular fat accumulation and reduced insulin sensitivity. By use of similar ¹³C-MRS techniques, Boumezbeur and colleagues⁷⁷ reported similar reductions in neuronal mitochondrial activity in healthy elderly individuals, which suggests that these age-associated reductions in mitochondrial activity might occur in several organs. Because mitochondria have a key role in the regulation of glucose-stimulated β -cell insulin secretion, age-associated reductions in mitochondrial function might also play a part in the impairment of β -cell function associated with ageing, and promote progression to impaired glucose tolerance and type 2 diabetes mellitus in elderly people.⁷⁸

Mechanisms of hepatic insulin resistance

Ectopic lipid accumulation in the liver is now widely known as non-alcoholic fatty liver disease. Formerly thought of as benign steatosis, this liver disease is now the most common chronic cause of raised serum concentrations of liver-derived enzymes in adults and children,⁷⁹ and it is closely associated with obesity, insulin resistance, and type 2 diabetes mellitus.^{80–82} Insulin action in the liver has many similarities with insulin action in muscle. In the liver, insulin activates the insulin receptor kinase, which phosphorylates IRS1 and IRS2, leading to activation of PI3K and ultimately Akt2 (figure 2).⁸³ At this point, Akt2 activation promotes glycogen synthesis and inhibits gluconeogenesis.

Many investigators have suggested that non-alcoholic fatty liver disease develops in the setting of insulin resistance.^{84–88} Increased hepatic steatosis was reported in genetic mouse models of muscle insulin resistance, specifically mice with muscle-specific deletions of the insulin receptor (MIRKO)⁸⁷ and the GLUT4 transporter.⁸⁸ Petersen and colleagues⁸⁹ compared postprandial de-novo lipogenesis in lean healthy, young insulin-sensitive and insulin-resistant individuals matched for age, BMI, percentage of fat mass, blood pressure, and physical activity. They postulated that insulin resistance in skeletal muscles, by altering the distribution of postprandial energy storage, would lead to an atherogenic dyslipidaemia that arises in metabolic syndrome. Specifically, insulin resistance in muscle would impair storage of carbohydrate as glycogen in muscle. Instead, carbohydrates would be redirected to the liver, and become substrates for hepatic de-novo lipogenesis. After a carbohydrate challenge, the insulin-resistant group had substantial elevations in plasma insulin but glycogen synthesis in muscle was 61% lower than in the controls, as measured with ¹³C-MRS. Furthermore, insulin-resistant individuals had a large increase in liver triglyceride content, attributable to a roughly two-fold increase in hepatic de-novo lipogenesis. These changes were associated with a 60% increase in concentration of fasting plasma triglycerides and a roughly 20% decrease in plasma concentration of HDL. Notably, visceral fat mass, as measured with abdominal MRI, was identical in these cohorts of normal-weight, insulin-sensitive and insulin-resistant individuals, which suggests these features of the metabolic syndrome can develop independently of increased visceral adiposity. Stefan and colleagues⁹⁰ also noted that, in a cohort of obese people, insulin-sensitive and insulin-resistant individuals were distinguished on the basis of lipid accumulation in the muscles and livers, but not subcutaneous or visceral adiposity. Fabbrini and colleagues⁹¹ similarly

reported that intrahepatic triglyceride content, not visceral adiposity, was associated with insulin resistance and increased triglyceride secretion.

Individuals of Asian-Indian ancestry are a population at risk of developing non-alcoholic fatty liver disease. Comparison of young, normal-weight men of Asian-Indian descent with eastern Asian, white, black, and Hispanic men showed that the Asian-Indian individuals had a greatly increased prevalence of insulin resistance.⁹² The most striking difference was a near doubling in the average liver triglyceride content in the Asian-Indian men when compared with matched white men. Though still low compared with individuals with marked obesity and type 2 diabetes mellitus, young Asian-Indian, normal-weight men were highly susceptible to hepatic steatosis associated with insulin resistance, increasing their risk of developing type 2 diabetes mellitus, steatohepatitis, and liver cirrhosis.⁹² Two polymorphisms (rs2854116 and rs2854117) in the apolipoprotein C3 (*ApoC3*) gene that seem to predispose individuals to the development of non-alcoholic fatty liver disease and insulin resistance have been identified.⁹³ This polymorphism leads to a roughly 30% higher plasma concentration of ApoC3, and postprandial hypertriglyceridaemia. As a result, the livers of carriers of these polymorphisms can take up increased amounts of lipid from the chylomicron remnant, leading to non-alcoholic fatty liver disease and hepatic insulin resistance. Transgenic mice that overexpress *ApoC3* provide genetic evidence in support of this hypothesis. When given a high-fat diet, these mice had greater accumulation of liver diacylglycerol than did wild-type mice, and the accumulation of diacylglycerol was associated with activation of PKC ϵ and substantial hepatic insulin resistance (Lee H-Y, Yale University School of Medicine, New Haven, CT, USA, personal communication).

Hispanic adults and children are also a large ethnic group at risk of developing non-alcoholic fatty liver disease and insulin resistance.^{94,95} Use of genetic screening has identified a polymorphism rs738409 within patatin-like phospholipase domain containing 3 (PNPLA3 population or adiponutrin) that is prevalent in the Hispanic population, and is highly associated with non-alcoholic fatty liver disease.⁹⁶ This polymorphism results in a missense mutation I148M within PNPLA3 that renders the protein incapable of triglyceride hydrolysis.⁹⁷ Though the association between this polymorphism and liver triglyceride content has been noted in other populations, there is no association with worsening insulin resistance.^{98,99} When Kantartzis and colleagues⁹⁸ analysed only patients with non-alcoholic fatty liver disease, those with the polymorphism actually seemed to have increased insulin sensitivity. The results of these studies^{98,99} contrast with the findings obtained with *ApoC3* gene variants. Whereas individuals with *ApoC3* gene variants and non-alcoholic fatty liver disease are insulin resistant, those with non-alcoholic fatty liver disease and PNPLA3 variants do not seem to have worsening insulin resistance. Further studies are needed to better define tissue-specific insulin resistance in liver and muscle, how these different polymorphisms might affect the composition and cellular localisation of intracellular lipids, and identify additional genetic factors that might affect the development of insulin resistance.

PKC ϵ , hepatic steatosis, insulin resistance

Wild-type mice and rats develop hepatic steatosis after a few days of high-fat feeding that is associated with hepatic insulin resistance, without much change in muscle lipid content or peripheral insulin action.¹⁰⁰ Moreover, by promotion of mitochondrial fatty acid oxidation with low doses of the mitochondrial uncoupler 2,4-dinitrophenol, rats were protected from fat-induced hepatic steatosis and hepatic insulin resistance.¹⁰⁰ In this model, hepatic steatosis was associated with proximal defects in insulin signalling, with decreased tyrosine phosphorylation of IRS1 and IRS2 by the insulin receptor, ultimately impairing the ability of insulin to activate hepatic glycogen synthesis and suppress hepatic glucose production. This

inability of insulin to regulate hepatic glycogen synthesis and glucose production has been shown in patients with type 2 diabetes mellitus.^{101,102} PKCs again were the logical link between hepatic steatosis and hepatic insulin resistance. Though PKC ϵ is poorly expressed in the liver, PKC ϵ , another novel PKC (figure 2), is expressed in high concentrations and activated in the fatty liver. If hepatic steatosis was prevented with the use of 2,4-dinitrophenol, PKC ϵ activation was also prevented. The association between PKC ϵ and hepatic insulin resistance has now been shown in other rodent models.^{53,103–106}

The specific role of PKC ϵ in the pathogenesis of hepatic insulin resistance was assessed by use of antisense oligonucleotides containing a modified 2'-O-(2-methoxy) ethyl and phosphorothioate bond to enhance potency, stability, and cell permeability.¹⁰⁷ They are taken up preferentially in the liver, adipose tissue, and kidney, though not in other key tissues such as muscle, brain, or β cells. The effect of a specific PKC ϵ antisense oligonucleotide was assessed in rats fed a high-fat diet for 3 days. Though fat accumulation, and specifically diacylglycerol accumulation, was equal in all groups, the PKC ϵ antisense oligonucleotide improved hepatic insulin sensitivity and insulin signalling. Specifically, PKC ϵ antisense oligonucleotide prevented the impairment in insulin receptor kinase activity noted with high-fat feeding. Thus, by blockage of PKC ϵ , hepatic insulin action was preserved despite the development of fatty liver.

Insulin resistance and lipodystrophy

One challenge in the assessment of the specific role of non-alcoholic fatty liver disease in the development of hepatic insulin resistance is the close association between obesity and non-alcoholic fatty liver disease. Thus, the changes in liver insulin action due to steatosis and those attributable to adiposity and associated changes, such as inflammation, are difficult to ascertain.^{108–110} The lipodystrophies offer an opportunity to assess the role of ectopic lipid deposition without any contribution from an expansion in peripheral or visceral adipose tissue mass. Lipo dystrophy and lipoatrophy are discrete genetic and acquired disorders with a lack of adipocytes, either through impaired fat-cell formation (eg, lipoatrophy) or acquired destruction of fat cells often noted in patients treated with antiretroviral drugs.^{111,112} Individuals with severe, generalised lipodystrophy have a substantial reduction in fat cells, are hypoleptinaemic, and consequentially many are hyperphagic. The lack of subcutaneous fat leads to hypertriglyceridaemia, insulin resistance, and ectopic fat deposition, including substantial hepatic steatosis.

Lipodystrophy can be modelled in mice. Moitra and colleagues¹¹³ created mice that were devoid of white adipose tissue by expressing a dominant negative protein A-ZIP/F under the control of the adipocyte specific AP-1 promoter. As in human generalised lipodystrophy, A-ZIP/F mice have no adipocytes and develop fat accumulation in the liver and skeletal muscle, and have profound peripheral and hepatic insulin resistance.¹¹⁴ Transplantation of fat pads into these mice from wild-type littermates rescued the phenotype and normalised the concentrations of tissue lipid, and hepatic and muscle insulin signalling and action. Shimomura and colleagues¹¹⁵ showed the potential to correct many of the metabolic defects associated with lipodystrophy by administering leptin to lipodystrophic mice.

Consistent with the results of the studies in mice, recombinant leptin restored plasma leptin to physiological concentrations, normalised plasma concentrations of glucose and lipids under fasting conditions, and corrected abnormalities in liver function tests in patients with severe congenital, generalised lipodystrophy.^{82,116} Before leptin replacement therapy, patients with lipodystrophy had higher basal rates of glucose production than did controls matched for age, weight, and sex; they also could not suppress hepatic glucose production and could not stimulate peripheral glucose uptake during hyperinsulinaemic-

normoglycaemic conditions.⁸² After leptin replacement therapy, hepatic triglyceride content decreased by about 90% with improvements in hepatic insulin responsiveness. Similarly, the roughly 30% reduction in muscle triglyceride content was associated with a near doubling in insulin-stimulated whole-body glucose disposal. The results of these studies in patients with lipodystrophy and mouse models of severe lipodystrophy show that ectopic accumulation of lipids can lead to insulin resistance, even in the absence of peripheral and visceral adiposity.

Other hypotheses

The data presented so far have supported a unifying theme—namely, that the accumulation of diacylglycerol within insulin-sensitive tissues activates novel PKCs that interfere with insulin signalling and cause insulin resistance. However, other mechanisms have been proposed to explain insulin resistance in obesity. These are only briefly discussed here, since other reviews are available.^{117,118}

Though we have endeavoured to show how accumulation of diacylglycerol leads to insulin resistance, not all lipid species are pathogenetic—eg, omega-3 fatty acids have many beneficial effects. They might promote hepatic fat oxidation in rodents through a PPAR α -dependent mechanism and increase plasma adiponectin through PPAR γ activation,¹¹⁹ thereby preventing ectopic diacylglycerol accumulation and preserving insulin action despite a high-fat diet.^{120–122} Palmitoleate (C16:1n7) has also been suggested as an endogenously produced fatty acid that is secreted from adipocytes (ie, a lipokine); it might improve insulin action in liver and muscle in mice,¹²³ and higher concentrations are associated with increased insulin sensitivity in people.¹²⁴

Clinically, insulin resistance and a proinflammatory state are both associated with the metabolic syndrome. Mechanistically, inflammatory signals affect cellular pathways that intersect with insulin action.¹¹⁷ Specifically, inflammatory signals such as tumour necrosis factor α and interleukin 6 activate serine and threonine kinases such as I κ kinase β and jun-N terminal kinase, which have been implicated in increased serine phosphorylation of IRS1. The development of hepatic insulin resistance after lipid infusions was associated with activation of the I κ kinase β -nuclear factor- κ B pathway.¹²⁵ Cai and colleagues¹²⁶ showed the potential for inflammation to cause insulin resistance by genetically activating this pathway specifically in the livers of mice. These mice developed both hepatic and peripheral insulin resistance. Moreover, the insulin resistance was ameliorated with salicylates, which also protected against insulin resistance in rodents after acute lipid infusion,¹²⁷ and reversed insulin resistance in people with type 2 diabetes mellitus.¹²⁸

Inflammatory pathways might also be activated in response to endoplasmic-reticulum stress.¹¹⁸ These pathways protect cells from producing aberrant proteins, hence referred to as the unfolded protein response. Their activation has been shown in rodent models of obesity¹²⁹ and in human obesity.¹³⁰ Modulation of endoplasmicreticulum stress by genetic or chemical means ameliorates jun-N terminal kinase activation and the development of insulin resistance.¹³¹ Substantial weight loss in patients after bariatric surgery has also been associated with both improvements in insulin sensitivity and reduction in markers of endoplasmic-reticulum stress.¹³²

Correction of hepatic steatosis

Thiazolidinediones, which are potent PPAR γ agonists, can effectively reduce hepatic steatosis.¹³³ Though PPAR γ is mainly expressed in adipocytes, it has effects on hepatic and muscle insulin sensitivity. On the basis of this discordance between the site of PPAR γ expression and the site of drug effects, the hypothesis was that thiazolidinediones redistribute fat from the liver and muscle into the adipocyte.⁴⁶ Mayerson and colleagues¹³³

tested this hypothesis, using rosiglitazone in patients with type 2 diabetes mellitus. Treatment for 3 months with rosiglitazone was associated with an almost 40% reduction in hepatic triglyceride content, and roughly 40% increase in extramyocellular triglyceride concentration, and improved suppression of adipocyte lipolysis.¹³³ Though no reductions in the concentration of intramyocellular triglyceride were noted in this study, rosiglitazone improved insulin-mediated whole-body glucose disposal, consistent with improvements in muscle insulin sensitivity. This disconnection between intramyocellular triglyceride and peripheral insulin action confirms that intra myocellular triglyceride is only a crude marker for the active metabolite (putatively diacylglycerol) that causes fat-induced insulin resistance.³⁸ Reductions in hepatic steatosis with both pioglitazone and rosiglitazone ameliorate hepatic insulin resistance.¹³⁴⁻¹³⁵ These data support the hypothesis that thiazolidinediones exert their beneficial effects through reversal of insulin resistance in patients with type 2 diabetes mellitus by shifting intracellular lipid from liver and muscle into adipose tissue.

Small amounts of weight loss can return concentrations of plasma glucose to normal in patients with type 2 diabetes mellitus.¹³⁶⁻¹³⁷ In obese patients with type 2 diabetes mellitus, after an average weight loss of 8 kg over 7 weeks on a hypocaloric (1200 kcal) low-fat diet, intramyocellular lipid concentrations or muscle insulin sensitivity did not change; however, there were substantial reductions in hepatic triglyceride concentrations (from nearly 12% to 2%), with concordant improvements in hepatic insulin sensitivity and correction of fasting hyperglycaemia.⁸¹ Additionally, fitness affects the response to calorie reduction; obese individuals with a high baseline cardiorespiratory fitness might have a greater reduction in liver fat with diet-induced weight loss.¹³⁸ Thus improvement of fitness might improve the resolution of non-alcoholic fatty liver disease with weight loss.

Way forward

Achievement of sustainable weight loss, without bariatric surgery, is an enormously difficult task and, although small steps are being taken for the prevention of obesity, many hurdles remain. Until societal, political, and economic forces align to promote healthy lifestyles, the incidence of obesity, and consequently insulin resistance and type 2 diabetes mellitus, will probably increase. The development of new effective treatments for insulin resistance requires an elucidation of the underlying primary mechanism. Inflammation, endoplasmic-reticulum stress, adipokines, and lipokines have roles in the pathogenesis of insulin resistance in liver and skeletal muscle. However, because these do not change in lean insulin-resistant young and elderly individuals, they are probably secondary in nature, becoming manifest later in the course of disease, and are associated with the development of obesity. The unifying model of diacylglycerol-induced insulin resistance accounts for the insulin resistance seen in obesity, and in other disorders such as congenital and acquired lipodystrophy,⁸² in young lean offspring of patients with type 2 diabetes mellitus, and with increase in age.⁶⁶ Moreover, reductions in intracellular diacylglycerol content account for the improvements in insulin sensitivity after weight loss⁸¹ and therapy with thiazolidinediones.¹³³ Further studies are needed to improve our understanding of how tissue-specific diacylglycerol accumulation activates specific novel PKCs, whether the intracellular distribution of this pool differs in insulin-sensitive and insulin-resistant states, and to better understand how activation of novel PKCs impairs insulin signalling. These studies will hopefully serve the development of new treatments to correct insulin resistance and halt its progression to type 2 diabetes mellitus.

Search strategy and selection criteria

We searched PubMed with the search terms “insulin resistance” in combination with “skeletal muscle”, “liver”, “lipids”, or “diacylglycerol” from January, 1963, until February, 2010. We also searched with “protein kinase C” and “diacylglycerol”. Papers were restricted to those published in the English language. We gave preference to recent and relevant reports, and important papers that addressed the main themes reviewed in this Seminar. Relevant review articles were selected to provide more comprehensive reference lists than were provided in this Seminar.

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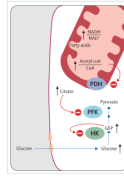


Figure 1. Glucose-fatty-acid cycle proposed by Randle and colleagues

CoA=coenzyme A. PDH=pyruvate dehydrogenase. PFK=phosphofructokinase.

G6P=glucose-6-phosphate. HK=hexokinase. Red circle with minus sign represents

inhibition. Black line with arrowhead represents increase or accumulation of substrate. Blue

dotted line with arrowhead indicates a pathway that is inhibited.



Figure 2. Mechanisms of insulin sensitivity and resistance in muscle and liver

(A) Insulin-sensitive muscle. (B) Insulin-resistant muscle. (C) Insulin-sensitive liver. (D) Insulin-resistant liver. IRS=insulin-receptor substrate. IR=insulin receptor. PI3K=1-phosphatidylinositol 3-kinase. GLUT4=glucose transporter 4. DAG=diacylglycerol. PKC=protein kinase C. Ser=serine. Thr=threonine. FOXO1=forkhead box O1. FOXA2=forkhead box A2. G6P=glucose-6-phosphate. GS=glycogen synthase. GSK=glycogen synthase kinase. Green circle with plus sign represents activation. Red circle with minus sign represents inactivation. Solid line with arrowhead represents increase or accumulation of substrate. Dotted line indicates inhibition of pathway.