

Lipid Metabolism and Toxicity in the Heart

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The heart has both the greatest caloric needs and the most robust oxidation of fatty acids (FAs). Under pathological conditions such as obesity and type 2 diabetes, cardiac uptake and oxidation are not balanced and hearts accumulate lipid potentially leading to cardiac lipotoxicity. We will first review the pathways utilized by the heart to acquire FAs from the circulation and to store triglyceride intracellularly. Then we will describe mouse models in which excess lipid accumulation causes heart dysfunction and experiments performed to alleviate this toxicity. Finally, the known relationships between heart lipid metabolism and dysfunction in humans will be summarized.

Introduction

Although the heart is far and away the most energy-requiring organ of the body, studies of cardiac lipid metabolism, especially *in vivo*, are relatively scarce compared to investigations in adipose tissue or liver. In adult fasting mammals, 60%–80% of cardiac energy metabolism relies on the oxidation of fatty acids (FAs) with glucose, lactate, and ketones providing substrates for the remainder (Neely et al., 1972). The adult heart, however, has the ability to switch to different substrates for ATP generation depending on feeding, hormonal status, and overall nutritional supply as characterized by the Randle cycle (Hue and Taegtmeyer, 2009). Of note, there are major species differences with mice relying more on glucose, lactate, and ketone bodies, and less on FAs (30%–40% from fat) (Stanley et al., 2005; Stowe et al., 2006). The fetal heart operates under low oxygen pressure and primarily depends on glucose and lactate for ATP generation, whereas the adult heart utilizes FAs but conserves the ability to switch other substrates. Older animals and humans use relatively fewer FAs and more glucose.

The heart avidly acquires lipids both from circulating nonesterified (free) fatty acids (FFAs) and esterified FAs bound to lipoproteins (Figure 1A). Observations made studying arterial venous differences in substrate concentrations showed that esterified FAs were a major source of lipids for the human heart (Ballard et al., 1960). More recent methods to study heart lipid metabolism have relied on tracers of FFAs in, for example, isolated perfused hearts. These studies quantify conversion of FFAs to CO₂ and TCA cycle intermediates under a variety of experimental conditions. *In vivo* studies can assess the uptake and loss of tracers from the heart. Although the heart can synthesize lipoproteins as it expresses both apoB and microsomal triglyceride transfer protein (Bartels et al., 2009; Nielsen et al., 1998), under most circumstances, the heart probably does not re-secrete appreciable amounts of glucose or lipids, and the uptake should indicate oxidation plus a relatively small amount of substrate that is stored and a small amount of substrate used for structural requirements of the cell.

In some situations the heart adjusts to maintain lipid homeostasis. Increases in work load (Goodwin et al., 1998) and myocardial ischemia (Lopaschuk et al., 2010) cause a rapid switch from fat to glucose utilization for ATP generation. This finding has led to several animal studies showing that administration of

compounds that reduce FA oxidation protect the heart from the consequences of ischemia and ischemia-reperfusion injury (Goodwin et al., 1998; Lopaschuk et al., 2010). This is presumed to be due to reduced oxygen requirements for non-FA substrates. Deleterious effects of cardiac ischemia could be due in part to excess cardiac lipid accumulation via the VLDL receptor (Perman et al., 2011). Similarly, in another mouse model of cardiomyocyte death adiponectin-induced activation of a ceramidase and reduction of ceramide was beneficial (Holland et al., 2011). Therefore, abnormal regulation of lipid uptake or its intracellular metabolism might play an important role in heart diseases other than metabolic dilated cardiomyopathy.

An imbalance between FA uptake and oxidation leads to accumulation of long-chain FAs that are incorporated into triglyceride (TG) and phospholipids, as well as a multitude of other lipid subspecies. Although TG is the most easily detected, other lipids are more likely to be toxic. Diacylglycerols (DAGs) and ceramides are signaling lipids that are thought to be toxic when their intracellular concentrations are increased. Defective mitochondrial FA oxidation could lead to accumulation of medium-chain acyl carnitines (Koves et al., 2008), another possible toxin. Finally, saturated long-chain FAs, most notably palmitate, are associated with toxicity in cells either because of their direct actions or their incorporation into phospholipids (Borradaile et al., 2006).

Sources of Heart Lipids

All tissues obtain lipids from FFAs associated with albumin, lipoproteins, and *de novo* synthesis (Figure 1A). Although *de novo* synthesis is thought to play a minor role in heart lipid metabolism, a recent study of deletion of FA synthetase in heart showed that *de novo* synthesis is important to maintain cardiac function during aortic constriction and aging (Razani et al., 2011). Loss of lipoprotein lipase (LpL)-derived lipids leads to increased glucose uptake in mouse hearts (Augustus et al., 2004). In humans, deficiency in CD36 is associated with increased glucose uptake (Fukuchi et al., 1999). CD36 appears most important in the setting of lower concentrations of FFAs (Coburn et al., 2000). Therefore, it is not surprising that when large amounts of FFA are generated during hydrolysis of large TG-rich lipoproteins like chylomicrons, heart uptake of lipids appears to be exclusive of this receptor (Bharadwaj et al., 2010). Lipolysis of

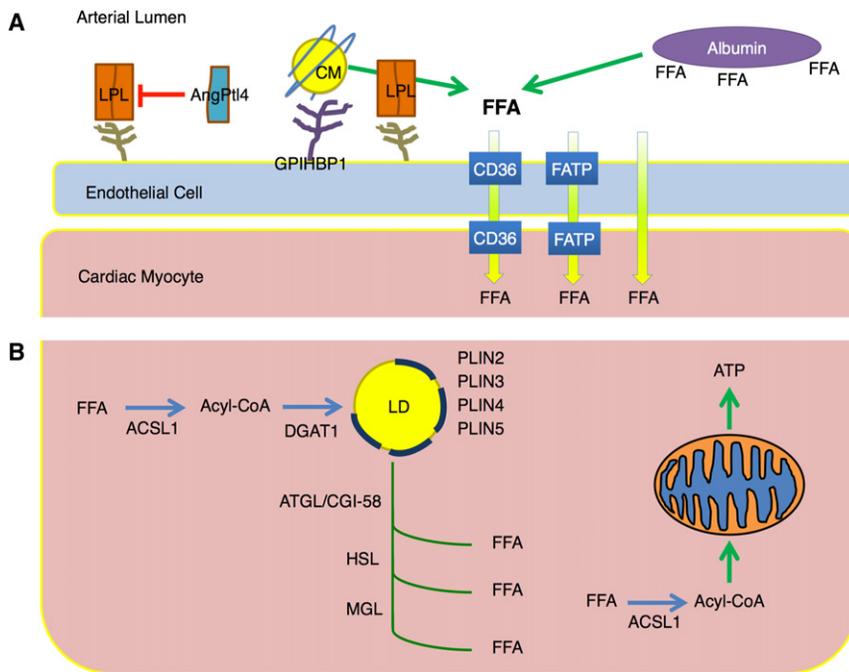


Figure 1. Regulation of Cardiomyocyte Lipid Storage

(A) Fatty acids esterified as triacylglycerol (TG) within lipoproteins require hydrolysis by lipoprotein lipase (LpL) associated with proteoglycans and GPIHBP1 on the luminal surface of endothelial cells. Angiopoietin-like protein 4 (ANGPTL4) is an LpL inhibitor. Nonesterified fatty acids (FFA) associated with albumin likely are internalized by membrane transporters such as CD36. These lipids must cross the endothelial barrier; how this occurs is unclear.

(B) Within the cardiomyocytes the FAs are esterified to CoA and either stored in the lipid droplet (LD) or used for energy. At least four lipid-droplet proteins (perilipins; PLINs) are expressed in the heart. The lipid droplet supplies some oxidized FAs via the actions of adipose triglyceride lipase (ATGL)/desnutrin and hormone-sensitive lipase (HSL). CGI58 is the ATGL coactivator. ATGL and LpL actions both provide ligands for PPAR activation.

lipoproteins is also a pathway for delivery of esterified core lipids such as cholesteryl esters and retinyl esters into the heart (Bharadwaj et al., 2010).

Cardiac Storage of Lipids

Excess lipid, especially TG, beyond that needed for cellular structures and ATP generation, is stored in lipid droplets (Figure 1B). Within the heart, there normally is little droplet accumulation, suggesting that uptake and oxidation are finely regulated. Lipid droplets are found in hearts of patients with diabetes and metabolic syndrome (Marfella et al., 2009; McGavock et al., 2007; Sharma et al., 2004) and in those of high-fat diet-fed rodents and genetically altered mice (see below and Table 1). In addition, after an overnight fast, lipid droplets appear in the hearts of wild-type mice (Suzuki et al., 2002).

Lipid-droplet protein makeup in the heart is different from that of adipocytes. In the heart, there is minimal expression of perilipin (Plin1). However, the other major lipid droplet proteins, adipophilin/adipose differentiation-related protein (ADRP/Plin2), tail-interacting protein 47 (Tip47/Plin3), S3-12 (Plin4), and OXPAT/myocardial LD protein/lipid-storage droplet protein 5 (Plin5) are all expressed in the heart (Paul et al., 2008). Plin2 expression might be most upregulated in some forms of lipotoxicity and be important for nontoxic lipid storage (Son et al., 2007). Plin5 appears to regulate TG oxidation by approximating lipid droplets and mitochondria (Bosma et al., 2011; Wang et al., 2011). Of these droplet proteins, only Plin2 has been deleted, and chow-fed *Plin2*^{-/-} mice do not have an obvious cardiac phenotype (Chang et al., 2006). Thus, knowledge of how and whether these proteins, and probably others, modulate heart-lipid accumulation and TG oxidation is likely to be forthcoming.

Lipid-droplet turnover is regulated by lipid-droplet-associated proteins, intracellular lipases, and acyltransferases (Meex et al., 2009). Cardiac myocytes and skeletal myocytes have similar

regulatory pathways that govern lipid metabolism. Lipid-droplet TG can be hydrolyzed by adipose triglyceride lipase/desnutrin (ATGL) and hormone-sensitive lipase (HSL), both of which are expressed in the heart. In the adipose tissue, insulin inhibits lipolysis, whereas catecholamines, thyroid hormone, and glucagon stimulate lipolysis. Whether similar regulation occurs in the heart is not known at present.

The roles of lipolytic enzymes in the heart have been studied using genetically modified mice (Table 1). Lipid-droplet accumulation in overnight fasting mice is prevented by overexpression of HSL (Suzuki et al., 2001). In the total HSL knockout mouse, cardiac TG lipase activity was decreased, but cardiac TG was not dramatically changed and there was no overt cardiac phenotype (Osuga et al., 2000). In contrast, *Atgl*^{-/-} mice have markedly reduced cardiac TG lipase activity, massive lipid accumulation, and severe cardiomyopathy (Haemmerle et al., 2006). In part, this is likely due to a defect in the hydrolysis of TG that the heart stores for potential energy (Banke et al., 2010). Treatment of *Atgl*^{-/-} mice with a PPAR α agonist corrected the cardiac phenotype (Haemmerle et al., 2011). Therefore, the excess accumulation of TG in the *Atgl*^{-/-} hearts was at least partially due to increased lipid storage secondary to defective FA oxidation. A less dramatic phenotype but one also associated with decreased FA oxidation activation occurred with cardiac deletion of acyl CoA synthetase 1 (Ellis et al., 2011). This study and another from this group (Ellis et al., 2010) suggest that PPAR activation is via a product of the CoA synthetase reaction.

Recent studies have elucidated the role of autophagy in hepatic TG lipolysis. In the rodent heart, autophagy has been investigated as a stress-response mechanism in myocardial infarction and pressure-induced cardiac hypertrophy. Fasting induces autophagy in the heart (Ogata et al., 2010), yet characterization of autophagy in the heart has not focused on lipid metabolic derangements. Inducible heart-specific autophagy knockouts develop cardiomyopathy (Nakai et al., 2007); whether

Table 1. Models of Cardiac Lipotoxicity

Cardiac Lipotoxicity	Reference	Corrections	Reference
MHC-ACS1	(Chiu et al., 2001)	Leptin treatment X MHC-DGAT1 α -Lipoic acid	(Lee et al., 2004, 2006) (Liu et al., 2009)
MHC-FATP1	(Chiu et al., 2005)		
Heart-specific PPAR δ knockout	(Cheng et al., 2004)		
MHC-PPAR α	(Finck et al., 2002)	X heart-specific LpL ko X CD36 $^{-/-}$ Medium-chain triglyceride diet	(Duncan et al., 2010) (Yang et al., 2007) (Finck et al., 2003)
MHC-PPAR γ	(Son et al., 2007)	X PPAR α knockout	(Son et al., 2010)
Leptin deficiency (<i>ob/ob</i>)	(Christoffersen et al., 2003) (Barouch et al., 2003)	Leptin infusion	(Barouch et al., 2003)
ATGL knockout	(Schweiger et al., 2006)	PPAR α agonists	(Haemmerle et al., 2011)
MHC-LpL ^{GPI}	(Yagyu et al., 2003)	Ceramide synthesis inhibition	(Park et al., 2008)
MHC-LPL X PPAR α $^{-/-}$	(Nöhammer et al., 2003)		
MHC-GLUT1 on HFD	(Yan et al., 2009)		

this is associated with increased lipid accumulation, as has been found in other organs (Singh et al., 2009), is unknown.

Creation of Cardiac Lipotoxicity in Mice

A number of reasons for the association between diabetes and heart dysfunction in the absence of underlying vascular disease have been proposed; one of these is excess accumulation of lipids in cardiomyocytes (Boudina and Abel, 2007). This possible cause of cardiomyopathy has been modeled by creating genetically modified animals in which lipid accumulation without generalized metabolic derangements leads to contractile impairment (Figure 2, Table 1). These animals have an imbalance between lipid uptake and oxidation due to either increased lipid uptake or decreased oxidation. Increased uptake of circulating FFAs or lipoprotein-derived lipids as occurs with transgenic expression of LpL (Nöhammer et al., 2003; Yagyu et al., 2003) leads to reduced heart function. Transgenic mice with cardiomyocyte-specific expression of FA transport protein 1 (Chiu et al., 2005) and acyl CoA synthetase 1 (Chiu et al., 2001) are thought to have increased FFA uptake or trapping in the heart leading to heart failure. PPAR transcription factors drive FA oxidation; however, the increased lipoprotein-lipid uptake in PPAR α transgenic mice (Duncan et al., 2010) must exceed the increased FA oxidation found in this model because the hearts have excess stored lipids. Cardiomyocyte PPAR γ overexpression leads to a similar phenotype (Son et al., 2007). Surprisingly, PPAR δ expression does not lead to cardiac dysfunction or toxicity (Burkart et al., 2007), presumably because upregulation of the LpL inhibitor angiopoietin-like protein 4 (Angptl4) prevents excess lipid uptake (Georgiadi et al., 2010).

LpL is the key enzyme for distribution of circulating lipids between organs. Perhaps the most clinically relevant model of cardiac lipotoxicity is one created by accident. Wang et al. deleted LpL using a skeletal muscle-specific promoter (Wang et al., 2009). The mice were meant to model physically inactive humans who also have reduced muscle LpL and FA oxidation. These mice have increased insulin sensitivity in skeletal muscle, as would be expected with reduced FA uptake, but develop

insulin resistance in the heart, often a precursor of eventual heart dysfunction.

Reduced lipid oxidation can also lead to lipid accumulation and cardiomyopathy. This occurs with cardiac-specific knockout of PPAR δ (Cheng et al., 2004) and a cardiac LpL transgene crossed onto the *Ppara* $^{-/-}$ background (Nöhammer et al., 2003). Similarly heart dysfunction with excess lipid accumulation is found when heart-specific GLUT1 overexpressing mice are placed on a high-fat diet (Yan et al., 2009). In contrast, when these animals eat chow, their function is improved in the presence of hypertension (Liao et al., 2002). So, in some situations, dietary-driven lipid uptake—a likely accompaniment of our western diet—needs to be added to create a lipotoxic environment.

Perhaps of most interest for understanding heart-lipid metabolism and toxicity are the situations where reduced FA oxidation does not lead to lipid accumulation. In many cases the reason for this has not been investigated, but we would presume that there is a compensation, such as a reduction in lipid uptake. This occurs with genetic or pharmacologic deficiency of DGAT1, which markedly reduces CD36 expression (Liu et al., 2011). Pharmacologic FA oxidation inhibitors proposed for reduction of ischemia (Lopaschuk et al., 2010) and genetically engineered defects in FA oxidation (Dyck et al., 2006) would be expected to also create cardiac lipid accumulation and toxicity. That this does not occur indicates the existence of some processes that balance reduced oxidation or that lead to nontoxic lipid storage.

Lipid Stores and the Causes of Toxicity

Although the most obvious and easiest to measure accumulated lipid is TG, TG itself might not be toxic. Consistent with cellular studies (Listenberger et al., 2003), several experimental situations have dissociated TG accumulation from toxicity. Total body knockout of HSL was associated with more refeeding TG accumulation, but no toxicity (Suzuki et al., 2009). A cross of the PPAR γ transgene onto the *Ppara* $^{-/-}$ background corrected toxicity without reducing heart TG, ceramide, or DAGs, but redistributed the lipids into larger droplets (Son et al., 2010).

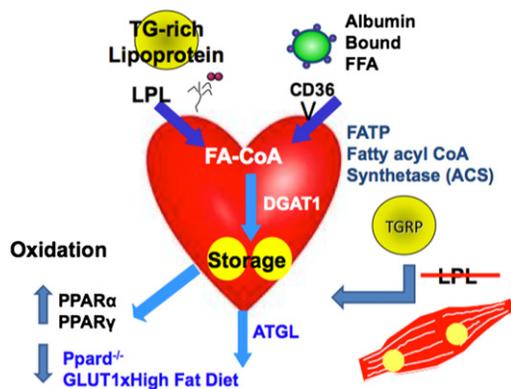


Figure 2. Lipotoxicity Is Created by an Imbalance of Lipid Uptake and Oxidation

Genetically modified mice have been created that have either increased lipid uptake or decreased oxidation. Uptake occurs via the cell-surface molecules lipoprotein lipase (LpL) and perhaps the FA transporter CD36 and/or FATPs. More FAs are “trapped” by complexing to CoA. Stored triglyceride accumulates with defective hydrolysis due to deletion of ATGL. Two interventions, loss of PPAR δ and high-fat feeding in mice overexpressing Glut1, lead to reduced lipid oxidation. Transgenic expression of PPAR α and PPAR γ , which induce lipid oxidation genes, also cause lipid accumulation, presumably because uptake exceeds oxidation.

Although the preferred substrate for the heart is FA, a case has been made that excess reliance on FA oxidation is harmful, even under nonischemic conditions. Some experimental data suggest the opposite. High-fat diets, which usually increase reliance on FA oxidation, may be beneficial in the setting of nonischemic heart failure (Okere et al., 2006b). Increased FA oxidation has been found with a *Ppar γ* transgene crossed onto the *Ppara*^{-/-} background (Son et al., 2010), *Dgat1* transgenic expression (Liu et al., 2009), and most recently with PPAR α agonist treatment of *Atgl*^{-/-} mice (Haemmerle et al., 2011), all of which improve heart failure.

Heart-lipid content can also be increased by nongenetic means. While in some ways these models may more closely reflect human pathology, they suffer from the many other systemic effects of overnutrition or diabetes. In addition, the heart phenotypes are relatively mild compared to those found with genetic modifications. Mice fed a high-fat diet rapidly develop cardiac insulin resistance, suggesting that lipid accumulation rapidly causes changes in heart metabolism allowing it to rely more on FAs as its fuel (Park et al., 2005). But the effects of high fat on cardiac function are mixed. Hypertensive rats fed a 60% fat diet had less left ventricular hypertrophy and systolic dysfunction than animals fed 10% fat (Okere et al., 2006b). Similarly, rats fed a high-fat diet appeared to compensate by increasing FA oxidation, whereas those eating a lower fat but higher carbohydrate western diet developed dysfunction (Wilson et al., 2007). Not all fats are equal. Saturated FA-rich diets alone increased cardiomyocyte apoptosis, perhaps due to accumulation of ceramide (Okere et al., 2006a). In contrast, medium-chain FAs are protective against cardiac dysfunction in some situations (Irie et al., 2003; Labarthe et al., 2005).

Other pharmacologic and genetic alterations change heart TG content. Using oxfenicine to block CPT-1, mice fed a diet enriched in long-chain saturated FA accumulated TG but had no

changes in left ventricular dimensions or systolic function, while PPAR-regulated genes were upregulated (Okere et al., 2007). *Ob/ob* and *db/db* mice have increased heart FA oxidation that develops prior to hyperglycemia (Buchanan et al., 2005). Eventually these mice develop decreased contractile function (Boudina et al., 2005).

Correction of Lipotoxic Heart Disease

A role of animal models is to test interventions that might be beneficial in human disease. Genetic approaches define targets, which might be amenable to pharmacologic or dietary interventions. Although a common underlying theme of lipotoxic cardiomyopathy is that it is created via an alteration in lipid metabolism, a single underlying toxic lipid species might not be causative in all cases. Thus, interventions in one model might not prove to be beneficial in another. Similarly, the genetic and dietary variation amongst humans might also lead to multiple causes of cardiomyopathy associated with lipid accumulation.

Altering the amount or type of FAs acquired by the heart will prevent toxicity both in genetic and dietary models of lipid toxicity (Table 1). Deletion of either CD36 or LpL corrected the cardiac toxicity associated with cardiomyocyte overexpression of PPAR α (Duncan et al., 2010; Yang et al., 2007). CD36 deletion was also reported to improve heart function in aged mice that were eating a diet enriched in medium-chain FAs (Koonen et al., 2007). Medium-chain FA-rich diets were also beneficial in PPAR α transgenic mice (Finck et al., 2003), presumably because of a reduction in saturated long-chain fat-enriched lipid accumulation. Other methods to reduce heart lipid content may also correct or prevent toxicity. The approaches to do that have included transgenic expression of HSL to increase lipolysis of stored lipids (Ueno et al., 2008), reduced expression glycerol-3-phosphate acyltransferase-1 to decrease TG accumulation (Lewin et al., 2008), and overexpression of apolipoproteinB to increase cardiac lipid secretion (Yokoyama et al., 2004).

A targeted approach to modify one lipid species would seem to be an ideal way to both treat the disease and also define the toxic lipid species. Such an approach was taken by reducing ceramide levels in LpL-overexpressing mice using the serine palmitoyl transferase (SPT) inhibitor myriocin and SPT-deficient mice (Park et al., 2008). A similar attempt to specifically modify another toxic lipid, DAG, was attempted by overexpressing diacylglycerol acyl transferase 1 (DGAT1) in cardiomyocytes. Although this intervention reduced DAG levels and heart dysfunction, ceramide was also reduced (Liu et al., 2009). This study suggests that alteration of a single lipid species in the heart might be difficult due to the interrelationship of lipid-metabolism pathways.

Evidence that Human Cardiac Dysfunction Is Associated with Excess Lipid

Unger has speculated that heart dysfunction due to excess accumulation of lipid, termed lipotoxic cardiomyopathy or fatty heart, is an unappreciated clinical entity (Szczepaniak et al., 2007). While this claim is controversial due to the confounding effects of classical risk factors coexisting in obese people such as physical inactivity, hypertension, hyperlipidemia, and diabetes, cardiac lipotoxicity alone or in combination with other risk factors might be an additional pathophysiologic abnormality

that develops in obesity. It is argued that patients with type 2 diabetes, metabolic syndrome, and obesity accumulate excess intramyocardial lipid and exhibit decreased systolic or diastolic function (Ernande et al., 2010; Ng et al., 2009). Clinical data show that both obesity and diabetes markedly increase risk of heart failure even in the absence of ischemic vascular disease (Kannel et al., 1974; Regan et al., 1977). The underlying molecular mechanisms could be either increased lipid uptake or impaired mitochondrial oxidative function leading to accumulation of TGs and toxic lipid species such as ceramides, which cause myocyte loss through apoptosis, induction of iNOS, and prohypertrophic signaling (Unger and Orci, 2001). More recent data have, however, found conflicting results with TG accumulation being detectable both in patients with and without systolic and diastolic cardiac dysfunction, and it has been claimed that levels of TG are rather a phenomenon defined by the type of specific cardiomyopathy and not directly related to cardiac dysfunction itself (Ernande et al., 2010; Kankaanpää et al., 2006; Nakae et al., 2010; Ng et al., 2009). Therefore, the specific form of excess cardiac lipid compounds (TG, ceramides, DAG), their cellular compartmentalization and storage form (lipid droplets), and the specific cause of heart failure are likely to determine the importance of lipotoxicity in human disease.

Studies of human cardiac function and metabolism rely on imaging methods that are relatively noninvasive. PET scanning assesses the uptake of various tracers into the heart. This technique is well standardized for both glucose and FFA uptake. Myocardial PET imaging has consistently shown increased FFA uptake and oxidation, as well as impaired glucose uptake, in diabetic patients with normal systolic and mildly impaired diastolic function (Rijzewijk et al., 2009). More recently, magnetic resonance protocols have been developed to track TG metabolism such as ^1H magnetic resonance spectroscopy (MRS) (O'Connor et al., 2011).

Although hypertension and coronary artery disease are common in obese and diabetic patients, reduced heart function independent of these underlying disorders may relate to toxicities from excess metabolic substrates and defective insulin action. Some studies in patients with obesity and diabetes correlated TG accumulation with left ventricular hypertrophy (Herrero et al., 2006; Szczepaniak et al., 2003). More TG has also been found in failing hearts of patients with obesity or diabetes at the time of transplantation (Sharma et al., 2004). Similarly, a modest correlation exists between plasma FFA levels and reduced diastolic function (Leichman et al., 2006). Limited data suggest that weight loss leads to reduced cardiac TG levels and reduced FA uptake as well as improved diastolic function (Hammer et al., 2008; Viljanen et al., 2009).

Reducing plasma lipids to reduce lipid uptake and converting oxidation to more glucose and less FA might be a method to treat patients with lipotoxic and ischemic heart failure. Agents that inhibit FA oxidation have been used for angina. Perhexiline is a drug that blocks carnitine palmitoyl transferase-1 (CPT-1) and CPT-2 and mitochondrial FA uptake. Heart-failure patients with both ischemic and nonischemic heart failure treated with perhexiline had improved cardiac function and symptoms (Lee et al., 2005; Tuunanen et al., 2008). Trimetazidine, another drug thought to reduce FA oxidation, modestly improved heart function and also improved overall insulin sensitivity in patients

with idiopathic dilated ischemia (Tuunanen et al., 2008). In contrast, depletion of circulating FAs in order to reduce FA uptake and storage into TGs did not affect cardiac function in patients with heart failure (Halbirk et al., 2010)—and in one study actually was harmful (Tuunanen et al., 2006). This latter intervention study measuring myocardial function and metabolism before and after administration of acipimox, a nicotinic acid-like inhibitor of lipolysis, was an acute study in which in each subject served as his or her own control. Therefore, the overall benefit of reducing FA oxidation in heart failure is still unclear and might require studies in which the etiology of the heart failure is more precisely defined.

Conclusions

Although lipid-induced cardiac toxicity clearly can be created, a number of clinical and experimental issues await clarification. We do not know about the exact clinical implications of this pathophysiologic phenomenon. Is it really a primary cause of cardiac dysfunction in patients with type 2 diabetes and metabolic syndrome? If so, does this occur because of unregulated cardiac uptake of FAs or reduced FA oxidation? If the latter, one could then postulate that in the setting of ischemia and afterload-induced heart failure, reduced FA oxidation should also promote accumulation of toxic lipids. Or, as has been postulated in ischemia, does a disproportionate use of FAs create toxicity? Which lipids are really toxic and which pathways are induced that lead to cellular dysfunction? A marriage of human and animal experimentation should sort out the answers to many of these questions in the next decade.

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