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Ceramides in Metabolism: Key Lipotoxic Players

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Abstract

The global prevalence of metabolic diseases such as type 2 diabetes mellitus, steatohepatitis, myocardial infarction, and stroke has increased dramatically over the past two decades. These obesity-fueled disorders result, in part, from the aberrant accumulation of harmful lipid metabolites in tissues not suited for lipid storage (e.g., the liver, vasculature, heart, and pancreatic beta-cells). Among the numerous lipid subtypes that accumulate, sphingolipids such as ceramides are particularly impactful, as they elicit the selective insulin resistance, dyslipidemia, and ultimately cell death that underlie nearly all metabolic disorders. This review summarizes recent findings on the regulatory pathways controlling ceramide production, the molecular mechanisms linking the lipids to these discrete pathogenic events, and exciting attempts to develop therapeutics to reduce ceramide levels to combat metabolic disease.

Keywords

obesity; insulin resistance; diabetes; ceramides; dyslipidemia; atherosclerosis

INTRODUCTION

The epidemic of obesity and its comorbidities is creating an enormous worldwide health burden, threatening the quality of life of families and the economic stability of countries around the globe. The number of individuals afflicted with obesity-related metabolic disorders—including diabetes, nonalcoholic fatty liver disease/steatohepatitis (NAFLD/NASH), and cardiovascular diseases (CVDs)—is astonishing. More than 450 million adults are living with diabetes (1); ~25% of the adult population has NAFLD/NASH (2); and ~30% of the population will die from cardiovascular disease (3). Studies published over the last two decades indicate that ceramides, which are intermediates in the biosynthetic pathway that produces complex sphingolipids, cause many of the cellular defects that drive these debilitating diseases. These discoveries about ceramides, described at length herein, present

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opportunities to understand the evolutionary pressures that lead to metabolic pathologies and to identify new therapeutic options for combating them.

Many of the early mechanistic studies probing the link between obesity, dyslipidemia, and cardiometabolic disease focused on insulin resistance, a condition where insulin is unable to effectively clear glucose from the bloodstream and a risk factor for all of the diseases listed above. Pioneering studies of Sir Philip Randle revealed that glucose and fat served as competitive substrates, with elevated fat effectively blocking utilization of carbohydrates (4). Randle and colleagues postulated a mechanism involving intermediates produced during lipid metabolism, which inhibit steps of glycolysis and thus might slow rates of glucose utilization. Subsequently, researchers found that complex lipids derived from excess fatty acids inhibited insulin signaling, thus blocking the postprandial translocation of the GLUT4 glucose transporters to the plasma membrane and concomitant glucose entry into muscle and fat tissue (5). Initially, the focus was on the glycerol-containing lipids (e.g., diacylglycerols and triacylglycerols), as their intramyocellular levels generally correlate with the severity of insulin resistance. However, sphingolipids such as ceramides were also found to accrue under conditions of lipid oversupply (6). The enigmatic ceramides were found to be potent inhibitors of insulin-stimulated GLUT4 redistribution, owing to their ability to block activation of Akt/PKB, a serine/threonine kinase that is a master inducer of many anabolic processes (7). One could restore insulin signaling and glucose uptake in lipid-treated myotubes and muscle fibers by using interventions that lowered ceramides, but not levels of the more abundant glycerol-containing lipids, such as the diacylglycerols (8-11).

Building upon these in vitro studies, researchers turned to rodent models to gauge the importance of ceramides as modulators of insulin sensitivity. In rat or mouse models of obesity or dyslipidemia, the administration of pharmacological reagents and/or genetic engineering strategies to lower ceramide levels restored insulin sensitivity owing to improvements in glucose utilization in skeletal muscle and adipose tissue, as well as suppression of gluconeogenesis in the liver (11). Remarkably, inhibition of ceramide production also reversed all of the other complications of obesity and insulin resistance including type 2 diabetes, NAFLD/NASH, atherosclerosis, hypertension, and cardiomyopathy (11-26). These findings indicated that ceramides likely had a much broader array of tissue actions, beyond those involving Akt/PKB and the inhibition of glucose disposal, that contribute to metabolic disorders.

Subsequent research identified numerous new ceramide actions that alter metabolism and compromise tissue function. They also revealed ceramides as strong biomarkers of insulin resistance, diabetes, and cardiovascular disease in large human cohorts, to the extent that they are now being measured clinically as an indicator of disease risk (27). Herein, we summarize these studies and speculate about the evolutionary pressures that gave ceramides their deleterious attributes. Moreover, we discuss possible therapeutic strategies to modulate ceramide levels and combat the spectrum of cardiometabolic diseases associated with obesity.

DISCOVERY OF SPHINGOLIPIDS

In 1884, German-born physician Johann L.W. Thudichum published a treatise describing the chemical constituents of the brain. Armed with only the rudimentary procedures available during his era, he determined that most lipids within the central nervous system comprised a glycerophosphate, fatty acids, and a base. Curiously, he also found a distinct, less-abundant lipid that contained no glycerol moiety. He named the unknown backbone sphingosine, a reference to the mythological Sphinx, which has a woman's head attached to a lion's body. Like their namesake, these enigmatic, sphingosine-containing lipids (i.e., sphingolipids) had a dichotomous structure, which included a polar head attached to an unknown hydrophobic body.

Ensuing technological developments enabled a precise determination of the structure of sphingosine, which distinguishes sphingolipids from other lipid classes (28, 29). Thanks to the work of the LIPID MAPS Lipidomics Gateway consortium (<http://www.lipidmaps.org/>), scientists now know the composition of thousands of distinct sphingolipid species. Researchers also characterized the enzymatic steps that control ceramide production and metabolism, cloning the majority of enzymes driving these reactions (30). The advent of these technologies provided new opportunities to probe the role of these molecules in biology, revealing their important roles in the development of disease.

Sphingolipids [e.g., ceramides, sphingomyelins, sphingosine, sphingosine-1-phosphate (S1P) and gangliosides] are far less abundant than glycerolipids, typically representing 2–15% of the total cellular lipidome. Despite being relatively minor components of membranes, they have potent biological activities, altering the physiochemical properties of lipid bilayers and regulating the activity of receptors and intracellular proteins (28). Ceramides, which are particularly low in abundance, have emerged as signals of lipid excess that initiate a host of cellular stress responses, leading ultimately to the induction of apoptosis (31). These actions have enormous relevance to cardiometabolic disease.

PATHWAYS CONTROLLING SPHINGOLIPID SYNTHESIS AND METABOLISM

Enzymatic pathways controlling the biosynthesis, degradation, and regeneration of ceramides dictate the composition of the cellular sphingolipidome.

De Novo Biosynthesis

De novo biosynthesis of ceramides, the precursor of most complex sphingolipids, occurs on the cytosolic surface of the endoplasmic reticulum. The multistep enzymatic cascade starts with the condensation of palmitoyl-CoA (CoA) and serine to produce 3-ketosphinganine, a reaction that is catalyzed by serine palmitoyltransferase (SPT) (Figure 1). SPT comprises essential subunits encoded by the *Sptlc1*, *2*, and *3* gene products, which form heterodimers with varying CoA preferences. For example, complexes containing SPTLC1 and 2 display a preference for C₁₆-CoA, while complexes with SPTLC1 and 3 prefer C₁₄-CoA (28, 29, 32). Additional, small subunits termed SSTPSA and SSTPSB enhance activity of the SPT heterodimers and confer additional specificity regarding fatty acid preference (33).

Moreover, a family of small membrane-bound ORMDL proteins provide an addition level of regulation to SPT activity (34).

The SPT complex can occasionally use alternative amino acids (i.e., alanine and glycine), instead of serine, to produce deoxysphingolipids. These rare sphingolipids were initially found in studies of individuals with hereditary sensory neuropathy, where mutations in SPT subunits alter its substrate selectivity (35). This finding prompted speculation that deoxysphingolipids elicited cellular damage, launching new studies of their relevance to disease processes, including many of the disorders that are the focus of this review (36-41).

In the canonical SPT reaction, the product 3-ketosphinganine is rapidly converted to sphinganine by 3-ketosphinganine reductase. This scaffold quickly acquires additional fatty acid side chains through *n*-acylation reactions catalyzed by a family of six (dihydro)ceramide synthases (CERS1–6). The CERS enzymes show variable tissue distribution patterns and distinct specificities for fatty acyl-CoA substrates. They produce a large number of dihydroceramides that include variable acyl chain lengths ranging from 14- to 34-carbon atoms. The complexity of the CERS family suggests that the acyl composition of sphingolipids influences cellular physiology in distinct, but still somewhat unresolved ways (42). Recent studies suggest that these CERS enzymes may also respond to fatty acid engagement by translocating to the nucleus, where they serve as transcriptional repressors of lipase genes (43).

The next reaction introduces a 4,5-*trans*-double bond into the sphingoid backbone of the dihydroceramides, producing the ceramides that are the major scaffold for most complex sphingolipids. This double bond, which is inserted by dihydroceramide desaturases (*DEGS1* and 2 genes; DES1 and 2 proteins), bestows the unique biophysical properties that are instrumental for initiation of the stress responses that are the focus of this review (44). DES1 is a ubiquitous enzyme, while DES2 is an enzyme that can also produce phytoceramides and is enriched in the skin and intestinal epithelium (45, 46).

Once synthesized, ceramides are transported from the endoplasmic reticulum to the Golgi apparatus, where head groups are added to the oxygen residue at the first position to produce the complex sphingolipids (e.g., sphingomyelin, gangliosides, or ceramide 1-phosphate) (28, 32). At least one ceramide transporter protein (CERT1) has been identified, and this protein facilitates the passage of the ceramides to the Golgi apparatus for production of sphingomyelins (but not gangliosides). Ceramides can also acquire another acyl chain at this first position. Like their triglyceride counterparts in the parallel glycerolipid biosynthesis pathway, these tri-acylated ceramides can be stored in the lipid droplet (47). The reaction that produces 1-*O*-acylceramides uses fatty acyl-CoA esters generated by ASCL5, an acyl-CoA synthase, and ceramides as substrates. Diacylglycerol acyltransferase (DGAT)-2, an enzyme involved in the triglyceride synthesis (47), adds this acyl chain to ceramides, and thus plays roles in both the glycerolipid and sphingolipid synthesis pathways. In hepatocytes, channeling ceramide into 1-*O*-acylceramides inhibits apoptosis (47).

Ceramide Degradation

Ceramides are deacylated by ceramidases, which break them into their components: a variable free fatty acid and sphingosine (48). Until recently, the family of mammalian ceramidases included only five enzymes, classified by their pH optima [i.e., acid (ASAH1), neutral (ASAH2), and alkaline (ACER1–3) ceramidases]. These enzymes display different cellular locations and substrate preference (48). Interestingly, Scherer and colleagues found that adiponectin receptors (ADIPOR1 and 2) also have ceramidase activity (49, 50). Adiponectin is a cardioprotective and antidiabetic protein that is released from healthy adipose tissue and circulates at high concentrations. Upon ligand binding, the receptors undergo a conformational shift that enhances ceramidase activity. Substitution of inactivating residues in the ceramidase motif renders the receptor ineffectual (49). Granier and colleagues (51) purified the receptor in the presence of ceramide, determining that the resolved crystal structure contained a bound fatty acid located in a hydrophobic pocket; this region resembles similar structural domains found in the other ceramidases. This group's studies with the purified protein confirmed that it had ceramidase activity (51).

Sphingosine can be phosphorylated to produce S1P, a bioactive lipid that gets secreted from the cell and activates a family of G protein–coupled receptors. S1P elicits biological responses that often oppose those of ceramide. S1P can be degraded by either sphingosine phosphate phosphatase (SPP) or S1P lyase. SPP dephosphorylates S1P to sphingosine, allowing for the reformation of ceramide through the salvage pathway discussed below. S1P lyase irreversibly converts S1P to ethanolamine phosphate and hexadecanal, thus breaking the sphingoid backbone and enabling an exit from the sphingolipid pathway (52).

Ceramide Regeneration

Ceramides can also be reproduced by the catabolism of sphingomyelin (e.g., through the action of sphingomyelinases), which generates ceramides while liberating the choline head group (53, 54). This reaction is catalyzed by a family of sphingomyelinases that differ by subcellular location and pH optimum. Alternatively, the sphingosine generated by ceramidases can also be reacylated by CERS enzymes to regenerate ceramides (28, 32, 52). This reacylation process is termed the salvage pathway. Glycosphingolipids can also be hydrolyzed by glycosidases, but this step is less frequent and accounts for a smaller fraction of ceramides than the aforementioned sphingomyelinase and ceramidase reactions (55, 56).

CLINICAL RELATIONSHIPS BETWEEN SERUM OR TISSUE SPHINGOLIPIDS AND CARDIOMETABOLIC DISEASES

Advances in mass spectrometry and lipidomics have enabled sensitive and accurate characterization of the sphingolipidome in human serum and tissue samples. These studies have revealed strong relationships between circulating sphingolipids and insulin resistance (57, 58), diabetes (59, 60), CVD incidence (61), secondary CVD events (62), secondary CVD mortality (63–65), and coronary artery disease (66). Because of the strength of these relationships, which often surpass those of other lipid biomarkers (e.g., LDL cholesterol), clinics in the United States and Europe now offer diagnostic tests to measure circulating

ceramides as measures of CVD risk (27). Herein, we detail these clinical studies and discuss the potential of using serum sphingolipids for disease diagnosis and patient stratification.

Ceramides as Markers of Insulin Resistance and Diabetes

Large studies in Australians, Chinese Singaporeans, and Native Americans, some involving sample numbers in the thousands, have revealed strong associations between circulating ceramides, insulin resistance, and type 2 diabetes. Meikle and colleagues (67) profiled 640 individuals enrolled in the Australian Diabetes, Obesity and Lifestyle Study, finding that ceramides positively correlated with fasting blood glucose. Herr and colleagues (57, 58) profiled 2,302 ethnically Chinese Singaporeans, finding a positive relationship between most ceramide species and homeostatic model of insulin resistance (HOMA-IR) scores. In addition, Lemaitre et al. (58) profiled 2,086 Native Americans, finding that several ceramide species (e.g., C16:0 and C18:0) correlated with HOMA-IR scores. In a nutritional interventional study, Yki-Jävinen and colleagues (68) also found that overfeeding saturated fats increased serum ceramides in plasma while producing a concomitant decline in insulin sensitivity.

Longitudinal studies indicate that serum sphingolipid levels increase early in disease progression, and thus might serve as predictive biomarkers of future disease. For example, Thorens and colleagues (60) found that dihydroceramides, which are the less-abundant precursors of ceramides, were elevated in individuals up to 9 years prior to occurrence of type 2 diabetes. These relationships held true in two independent clinical cohorts. In general, we have found that the dihydroceramides, owing to their low abundance, change more dynamically than ceramides under conditions of increased flux both in vitro and in vivo (S.A. Summers, unpublished observations).

Studies evaluating tissue levels of ceramides also reveal positive associations between ceramides and insulin resistance. In liver, Yki-Jävinen and colleagues (69) analyzed 125 liver biopsies obtained from individuals of Finnish descent, finding that C16:0, C18:0, C20:0, C22:0, and C24:1 ceramide species correlated with insulin resistance. These relationships were independent of hepatic triglycerides and body weight.

In adipose tissue, Yki-Jävinen's team (70) similarly found that ceramides correlated with insulin resistance, independent of obesity. Moreover, Brüning and colleagues (26) profiled 439 individuals of European descent, finding numerous ceramide species that were elevated in the adipose tissues of obese individuals. His group also demonstrated that adipose *CERS6* transcripts correlated positively with body mass index (BMI), body fat content, and hyperglycemia (26). By comparison, *CERS6* negatively correlated with glucose infusion rates during euglycemic-hyperinsulinemic clamps (26). These data support rodent studies (discussed below) revealing that C16:0 ceramides derived from *CERS6* antagonize insulin action.

In skeletal muscle, Goodpaster and colleagues (71-74) identified positive correlations between muscle ceramides and insulin resistance in a large number of separate studies. In these studies, insulin-sensitizing interventions (e.g., exercise, metformin, pioglitazone, and

bariatric surgery) decreased muscle ceramide levels (71-74). Mandarino and colleagues (75) reported similar findings in muscle biopsies.

Ceramides as Biomarkers of Cardiovascular Disease

A particularly large number of studies, many conducted by Laaksonen and colleagues at Zora Biosciences, have identified serum ceramide species that are potent, independent, and predictive markers of coronary artery disease, major adverse cardiac events, and/or fatality (58, 63, 65, 76-84). They also determined that plasma ceramides predict recurrent heart attacks and cardiovascular-related deaths in patients that had undergone a prior cardiac event (62). Moreover, they found that statins decrease plasma ceramides (84, 85). In each of these studies, three ceramide species (C16:0, C18:0, and C24:1) reliably predicted the severity of cardiovascular disease. Using these data, the authors generated a ceramide-based score (i.e., CERT1) that included these species in proportion to C24:0, a benign ceramide that shows no relationship with disease and could thus be used for normalization (86). CERT1 predicted coronary death more than three times better than LDL cholesterol. Clinics are now offering CERT1 diagnostic tests (27). More recently, this group improved upon CERT1, producing a refined CERT2 score that included additional markers and showed enhanced prognostic utility (78).

Our group also determined that sphingolipids were strong, cholesterol-independent biomarkers of coronary artery disease (66). In this study, 30 different sphingolipids demarcated individuals with the disease. Using machine learning, we determined that several low-abundance sphingolipids, which are likely good markers of flux through the biosynthesis pathway, were particularly strong markers. Using this information, we created a Sphingolipid Inclusive Score (SIC), which proved superior to CERT1 and LDL cholesterol for that patient population.

In tissues, Schulze and colleagues found elevated myocardial ceramides in patients with advanced heart failure (18, 87).

Efficacy of Ceramide-Reduction Interventions in Rodents

Many of the interventional studies exploring the relevance of sphingolipids to cardiometabolic disorders were conducted with myriocin, an irreversible and high-affinity inhibitor of SPT. In rodents, chronic treatment with myriocin has been shown to prevent and/or reverse atherosclerosis (15-17, 22-24), insulin resistance (11, 88-91), diabetes (11), NAFLD/NASH (92), hypertension (90), and cardiomyopathy (18, 21). The astonishing utility of this single compound in such a large array of disease models has generated considerable excitement about the potential of therapeutics designed to lower ceramides. Subsequent studies, discussed below, confirmed the importance of ceramides in cardiometabolic disorders using genetic engineering strategies to inhibit rates of ceramide synthesis or accelerate rates of ceramide degradation. This research, in addition to revealing the involvement of sphingolipids in pathology, has enabled determination of tissue-specific roles of ceramides as drivers of tissue dysfunction.

Ceramide-lowering interventions resolve insulin resistance and prevent diabetes.—

Myriocin was shown to prevent and/or reverse insulin resistance in high-fat diet-fed mice (11, 25, 89, 91), lipid-infused rats (11), fructose-fed hamsters (93), and leptin-deficient mice and rats (i.e., Zucker *fa/fa* rats and *ob/ob* mice) (11). Moreover, myriocin showed efficacy in models where the insulin resistance progresses to frank diabetes. Specifically, Zucker diabetic fatty rats transform from a prediabetic state characterized by insulin resistance and hyperinsulinemia to frank diabetes that includes fasting hyperglycemia, decreased insulin, and declining beta-cell mass. Myriocin protects these animals from the progression of beta-cell failure and the concomitant development of diabetes (11). Unger and colleagues (94) also found that cycloserine, a broad-spectrum antibiotic that also inhibits SPT, prevents beta-cell failure in Zucker diabetic fatty rats.

In mice fed a high-fat diet, the insulin-sensitizing effects of myriocin could be recapitulated with fenretinide (95, 96), a synthetic retinoid that inhibits DES1 (96). The beneficial actions of fenretinide were initially attributed to its effects on retinol-binding protein 4 (RBP4), which has also been implicated in the development of insulin resistance. However, fenretinide shows efficacy in RBP4-knockout mice (97), suggesting that the compound has other insulin-sensitizing mechanisms. Indeed, it dramatically lowered ceramide levels (and increased dihydroceramides) in concert with its inhibition of DES1 and improvement in metabolic homeostasis (96).

Additional studies reveal that removal of genes required for ceramide biosynthesis or overexpression of those that catalyze ceramide degradation is also insulin sensitizing. For example, we found that inducible depletion of *Degs1* in the entire periphery, as well as tissue-specific removal from the liver, adipose tissue, or the liver plus adipose tissue, was sufficient to restore insulin sensitivity in high-fat diet-fed and/or leptin-deficient *ob/ob* mice (12). The following interventions also improved insulin sensitivity in mice fed a high-fat diet: (a) adipose-specific depletion of a gene encoding a key subunit of SPT (i.e., *Sptlc2*) (88), (b) whole-body or brown adipose tissue-specific depletion of *Cers6* (26, 98), or (c) whole-body or muscle-specific depletion of *Cers1* (99). Scherer and colleagues (100) also found that overexpression of acid ceramidase (*Asah1*), which lowers ceramides by inducing their conversion to sphingosine, in either adipose tissue or the liver, improved insulin sensitivity in high-fat diet-fed mice.

Ceramide-lowering interventions resolve dyslipidemia and NAFLD/NASH.—

Most of the models described in the preceding section also evaluated liver and serum lipid levels, revealing that ceramide-lowering interventions led to improvements in lipid homeostasis. For example, treatment with myriocin (11, 20, 25) or fenretinide (96), as well as depletion of *Sptlc2* (adipose) (88), *Cers6* (whole body or brown adipose tissue) (26, 98), or *Degs1* (whole body, adipose tissue, or liver) (12), resolved hepatic steatosis and/or reduced circulating triglycerides. Liver- or adipose-specific overexpression of *Asah1* also resolved steatosis and dyslipidemia. Recently, Jiang et al. (92) tested myriocin in a rat model that progresses from hepatic steatosis (NAFLD) to NASH, characterized by pronounced liver fibrosis. Myriocin prevented the progression of fibrosis that defines NASH.

Ceramide-lowering interventions resolve atherosclerosis.—Myriocin also prevents development of atherosclerotic lesion formation in apolipoprotein E (ApoE) knockout mice (17, 22, 24). In some instances, the intervention was also shown to enable regression of preformed plaques. Genetically, depletion of *Sptlc2* from myeloid cells was sufficient to reduce the atherosclerotic plaque area in LDL-receptor knockout mice (101).

Ceramide-lowering interventions ameliorate vascular dysfunction.—In mice fed a high-fat diet, myriocin and haploinsufficiency for *Degs1* resolved hypertension (90, 102). The effects could be recapitulated in isolated blood vessels exposed to high levels of saturated fat, with the studies indicating that these ceramide actions likely occurred in the endothelial cell.

Ceramide-lowering interventions ameliorate cardiomyopathy and heart failure.—Ceramide-reducing strategies, including either myriocin or haploinsufficiency for *Sptlc2*, have also shown to correct hypertrophy and improve cardiac function in the following mouse models of heart disease: (a) lipotoxic cardiomyopathy (i.e., overexpression of lipoprotein lipase in cardiomyocytes) (21) and (b) coronary occlusion (18, 103). Russo et al. (104) identified lipids produced by CERS5 as possible mediators of cardiac hypertrophy.

Fundamental Cellular Mechanisms Linking Ceramides to Cardiometabolic Disease

Studies conducted in mice that involve whole-body or tissue-specific manipulations of SPT (e.g., the SPTLC2 subunit), DES1, ASAH1, or the various CERS isoforms have revealed important information about the mechanisms linking ceramides to the tissue dysfunction that drives cardiometabolic disease. Complementary studies in cell culture systems have further illuminated downstream pathways and intracellular mediators of these ceramide actions. In this section, we discuss the data (a) resolving which ceramides are lipotoxic and (b) elucidating fundamental ceramide actions that are relevant to disease states.

Identification of Roles for C16:0 and C18:0 Ceramides as Key Lipotoxic Species in Adipose Tissue, Skeletal Muscle, and/or the Liver

Though numerous different sphingolipid species likely contribute to the decline in tissue dysfunction in obesity and cardiometabolic disease, ceramides containing the C16 and C18 acyl chains (C16:0 and C18:0) have proven to be particularly damaging. By contrast, those containing much longer chains (e.g., C24 or C24:1) appear to be either protective or benign.

In 2014, Brüning and colleagues (26) provided the strongest evidence to date that C16:0 ceramides drove the decline in adipose and liver function in obesity and the concomitant disruption in glucose and lipid homeostasis. They generated knockout mice allowing for whole-body or tissue-specific (i.e., liver or brown adipose tissue) ablation of the gene (*Cers6*) encoding CERS6, the enzyme that produces C16:0 ceramides. Depleting *Cers6* from any of these locales resolved high-fat diet–induced obesity, glucose tolerance, and insulin resistance in mice (26). They later determined that these improvements were due, in part, to a reduction in mitochondrial fission (discussed below) (98). In parallel, our laboratory found that mice that were haploinsufficient for the *Cers2* gene that encodes CERS2, which produces the C20–C24 ceramide species, demonstrated a worsening of glucose tolerance

and hepatic steatosis (25). This surprising result became understandable when we observed a large, compensatory induction of the CERS6 isoform and a concomitant increase in C16:0 ceramides. Moreover, we found that overexpression of CERS6, but not CERS2, induced fat accumulation and antagonized insulin action in primary hepatocytes. These complementary papers, published by the Brüning (26) and Summers (25) laboratories in the same issue of *Cell Metabolism*, revealed remarkably similar mechanisms involving C16:0 ceramide actions on mitochondrial efficiency and insulin signaling. Shortly thereafter, researchers at Sanofi used an antisense oligonucleotide approach to deplete CERS6 from mice. In this study, Tennagels and colleagues (105) demonstrated that inhibition of CERS6 reduced C16:0 ceramides and fat content and improved glucose tolerance and insulin sensitivity. Hoch and colleagues (106) obtained similar results in mice lacking CERS5, another enzyme that produces C16:0 ceramides.

Brüning and colleagues (99) also studied mice lacking CERS1, the enzyme that produces C18:0 ceramides and the major ceramide synthase isoform in skeletal muscle. They determined that depletion of the gene globally (i.e., in all tissues) or selectively in skeletal muscle conferred protection from obesity-induced insulin resistance. Don and colleagues (107) also synthesized a selective inhibitor of CERS1, termed P053. Chronic administration of P053 to obese mice impeded fat deposition, though it had unimpressive effects on insulin resistance (107). Mechanistically, these actions of CERS1 were attributed to effects on fatty acid metabolism in the skeletal muscle and increased secretion of fibroblast growth factor 21 (99, 107). Additional studies are necessary to better clarify the importance of C18:0 ceramides in cardiometabolic disease.

Several studies conducted either in vivo or in vitro evaluated the relevance of the ceramide headgroups in insulin resistance and metabolic dysfunction. Deletion of sphingomyelin synthase 1 from mice was shown to be insulin sensitizing, but the animals were resistant to weight gain (108); the disparity in body mass thus made it difficult to discern whether sphingomyelin is a direct regulator of insulin sensitivity. Studies conducted in vitro suggest that ceramides, rather than the more abundant sphingomyelins, were the key modulators of metabolism. Inhibition of sphingomyelin synthase isoforms induced ceramide accumulation by blocking its conversion into sphingomyelin (109). This intervention elicited many of the downstream mechanisms described below (i.e., inhibition of insulin signaling and impaired mitochondrial respiration) (109).

Additional studies evaluated the role of glucosylated ceramides, including the gangliosides, as drivers of metabolic dysfunction. Inhibitors of glucosylceramide synthase have shown efficacy in models of steatosis and insulin resistance (110-112), with mixed results seen in models of atherosclerosis (113-115). Moreover, knockout of GM3 synthase, which produces higher-order gangliosides, is also insulin sensitizing in vitro and in vivo (116-118). These data indicate that glucosylated ceramides also contribute to cardiometabolic disease, in part by inhibiting insulin signaling. Our own studies revealed that ceramides and glucosylceramides have separable and independent actions on insulin signaling, depending on the tissue type being studied (119).

Ceramide Regulation of Mitochondrial Biology (i.e., Energetics and Apoptosis)

Ceramides exhibit numerous actions on mitochondria, leading to diminished electron transport chain activity and increased mitochondrial outer membrane permeability. These actions lead to decreased mitochondrial respiratory capacity and increased tissue apoptosis, respectively. Both of these are hallmarks of diabetes and cardiovascular disease.

Ceramides have been shown to decrease mitochondrial respiration by inhibiting oxidative phosphorylation and promoting mitochondrial fragmentation, leading to an induction of reactive oxygen species (25, 26, 98) (Figure 2). Treating cells or isolated mitochondria with ceramide analogs acutely disrupts components of the electron transport chain (120-122). Similar findings were obtained after overexpressing CERS6, which produced mitochondrial C16:0 ceramides and inhibited complex II (25). Moreover, depletion of sphingomyelinase synthase 2 produced impairments in mitochondrial respiration in association with an elevation in intracellular ceramides (109). Lastly, knockout of the genes encoding SPTLC2, CERT1, CERS6, CERS2, and DEGS1 leads to changes in mitochondrial energetics that inversely mirror tissue ceramide levels (12, 25, 26, 88, 98, 99, 107).

Hammerschmidt et al. (98) determined that ceramides were potent regulators of mitochondrial fission. Using a proteomics approach to detect proteins that interacted with sphingolipids produced by CERS6, they determined that mitochondrial fission factor (MFF) was a ceramide target. The protein linked ceramides to an acceleration of mitochondrial fission, which resulted in a change in mitochondrial morphology and a decrease in respiratory capacity (98). Moreover, they found that CERS6-knockout mice contained numerous large, highly efficient mitochondria (98). While these actions do not explain the acute effects of ceramides on mitochondrial respiration, they do provide one mechanism by which chronic increases in ceramides disrupt mitochondrial respiration.

As ceramide levels continue to rise, they increase permeability of the mitochondrial outer membrane. This leads to the release of cytochrome *c* and initiation of apoptosis (123). Ceramides promote translocation of the proapoptotic Bcl2 family proteins (e.g., BAX) to mitochondrial membranes (124). They also bind voltage-dependent anion channel 2 (VDAC2), which further increases mitochondrial membrane permeability (125). Lastly, C16:0 ceramides form channels in mitochondrial membranes that can allow passage of cytochrome *c* (126-129).

Ceramides Inhibit Insulin Signaling to Block Glucose Uptake and Storage

Ceramides inhibit insulin and growth factor action by preventing activation of Akt/PKB, a serine/threonine kinase that stimulates glucose utilization, promotes cell survival, and activates enzymes driving fatty acid, protein, and triglyceride synthesis. Ceramides influence Akt/PKB via two distinguishable actions on distinct kinase domains. First, they accelerate the dephosphorylation of two important Akt/PKB residues on Akt/PKB via protein phosphatase 2A (PP2A) (9, 130-132). The action of ceramides on PP2A may be direct (133), but they also regulate the enzyme by displacing the PP2A inhibitory protein I2PP2A (102). In certain cell types, the inhibitory actions of ceramides on Akt/PKB could be negated either by treating with PP2A inhibitors (e.g., okadaic acid) or by overexpressing the SV40 small T

antigen (which blocks access of PP2A to its substrates) (8, 9, 130-132). Second, ceramides have also been shown to block Akt/PKB translocation to the membrane by stimulating its phosphorylation on a separable, inhibitory residue in the pleckstrin homology domain (134, 135). This mechanism requires a ceramide-driven association of protein kinase C ζ (PKC ζ) with caveolar domains (136). Both of these pathways can exist within the same cell type (137). In insulin-responsive tissues, the ceramide actions on Akt/PKB account for the impairments in insulin-stimulated GLUT4 glucose transporter translocation to the plasma membrane (7).

Ceramides Stimulate Lipid Uptake and Storage

Within the last few years, ceramides were found to elicit a new spectrum of actions that altered lipid handling in adipocytes and hepatocytes. First, they facilitate the uptake and esterification of fatty acids by promoting the redistribution of fatty acid translocases like CD36 (12, 49). Second, they induce genes (e.g., *SREBP*) that promote the incorporation of free fatty acid into triglycerides to facilitate their storage in lipid droplets (12, 138). Third, they inhibit lipolysis by blocking activation of hormone-sensitive lipase (HSL) (12). These effects are also attributable to the aforementioned ceramide effects on PKC ζ and PP2A (12, 49, 138).

TISSUE-SPECIFIC ROLES OF CERAMIDES IN CARDIOMETABOLIC DISEASE

Tissue-specific interventions revealed how the fundamental ceramide actions described above manifest themselves in mammals. For a comprehensive list of transgenic mice used to discern the role of ceramide and ceramide metabolites in metabolic diseases, see Table 1.

Ceramides in the Adipocyte

In mice and rats, myriocin decreases adipocyte size, elevates numbers of brown/beige adipocytes, and increases recruitment of M2 macrophages into the subcutaneous adipose bed (88). These effects were most robust in the subcutaneous depots. Similar results were obtained following adipocyte-specific depletion of *Sptlc2*, which induced a comparable spectrum of benefits and was sufficient to improve insulin sensitivity and glucose tolerance, resolve hepatic steatosis, increase whole-body energy expenditure, and enhance adipocyte respiration (88).

We have conducted analogous studies by depleting *Degs1* from adipocytes (12). The absence of the DES1 protein reduced adipocyte size, increased glucose uptake into the adipose tissue, and enhanced mitochondrial respiration. These effects were strong enough to improve glucose tolerance and insulin sensitivity and resolve hepatic steatosis. Similar findings were obtained following brown adipose tissue-specific *CerS6* depletion (26) or adipose-specific *Asah1* overexpression (100). Because of the comparable phenotypes observed in all of these studies, we conclude that ceramides act as nutrient signals that direct the adipocyte toward a hypometabolic phenotype.

We note one important distinction between the studies involving SPT inhibition or *Sptlc2* depletion and those involving manipulation of other enzymes in the pathway. Unlike inhibition of SPT, neither removal of *DeGs1* nor overexpression of *Asah1* induced browning/beiging of the adipose depots. We speculate that the discordance may be attributable to intermediates in the biosynthetic pathway, which are downstream of SPT and upstream of DES1. These intermediates (e.g., the CERS enzymes, which can act as transcriptional repressors) may influence the browning program.

Notably, studies by the Proia (139) and Park (140) laboratories found that ablation of either *Sptlc1* or *2*, respectively, in adipose tissue impaired adipose differentiation and elicited lipodystrophy. In these studies, the authors used an adiponectin-Cre-recombinase mouse (141) that expressed the transgene early in development (142). By comparison, the adiponectin-Cre mouse we used, from the Scherer laboratory, used a different promoter fragment that expresses the recombinase weaker and later (88). In concordance with their work, our studies in primary cells showed that inhibition of SPT inhibits adipocyte differentiation in vitro (88).

Ceramides in the Hepatocyte

Treating mice with either myriocin or fenretinide reduces hepatic lipid accumulation and represses hepatic gluconeogenesis (11, 96). Tissue-specific ablation of DES1, either by genetic knockout or knockdown using short hairpins, from the liver of mice conferred comparable metabolic benefits (12). These findings, and complementary data from researchers studying cultured hepatocytes, confirm that ceramides have cell-autonomous actions within the hepatocyte. Mechanistically, these effects result from ceramide's ability to (a) induce triglyceride synthesis by modulating *Srebf1* expression, (b) promote fatty acid uptake by altering CD36's subcellular distribution, and (c) enhance gluconeogenesis by blocking AKT/PKB phosphorylation (12). These actions of ceramides lead to increased fat deposition and glucose output, thus producing the "selective insulin resistance" (143) that defines the prediabetic condition.

The Scherer group (100) previously conducted an elegant study that elucidated many of these mechanistic details. Their group found that liver-specific overexpression of acid ceramidase reduced hepatic ceramide content, improved insulin sensitivity, and resolved hepatic steatosis (100). This study was the first to show that ceramide induced translocation of CD36 via PKC ζ (100).

Brüning and colleagues (26, 98) also performed liver-specific interventions with CERS6, finding that its removal from the liver protected mice from steatosis and insulin resistance. A major mechanism was related to the aforementioned improvements in mitochondrial function. We similarly found that overexpressing CERS6 in hepatocytes induced triglyceride production, inhibited insulin signaling, and induced mitochondrial dysfunction (25).

Ceramide Actions in the Myocyte

The question of whether ceramides modulate metabolism in skeletal muscle has been an area of debate. The disagreement is largely due to a discordance in lipidomic profiling studies. Though many researchers found correlations between muscle ceramides and insulin

sensitivity (described above), a small subset did not (144, 145). Despite this contention, interventional studies unequivocally support roles for ceramides as modulators of insulin signaling and mitochondrial function in rodents and cultured myotubes. In cellular models, ceramides invariably and potently inhibit insulin signaling and decrease mitochondrial respiration (9, 109, 146, 147). Moreover, in myotubes or rodent muscle fibers, blocking ceramide synthesis or accelerating ceramide degradation attenuates palmitate- or TLR4-induced insulin resistance (i.e., inhibition of insulin signaling and/or glucose uptake) (8, 9, 11, 119, 135, 147). In rodents, pharmacological inhibition of SPT or DES1, as well as depletion of *Degs1*, improved muscle insulin sensitivity in mice (11). Such experiments have been conducted in mice or rats fed a high-fat diet, treated with dexamethasone, intravenously infused with lipid cocktails, or lacking leptin (11).

These data indicate that ceramides produced within myocytes can operate as cell-autonomous regulators of metabolism. Nonetheless, Watt and colleagues (148) have presented the provocative idea that circulating ceramides may be even more important as determinants of muscle insulin sensitivity. His group found that ceramides present in LDLs were sufficient to induce insulin resistance (148). Lipoprotein particles lacking ceramides were incapable of antagonizing insulin action in muscle, but those containing the sphingolipid had potent effects. This finding suggests that lipoprotein-bound ceramides may be important drivers of muscle insulin resistance.

Ceramide Action in the Pancreatic Beta-Cell

Myriocin preserves pancreatic beta-cell mass in Zucker diabetic fatty rats (11). While this could simply be secondary to improved insulin sensitivity, which would alleviate the burden on the beta-cell, several *in vitro* studies suggest that ceramides might have cell-autonomous actions within the islet. Indeed, the earliest studies relating ceramides to diabetes were conducted by Roger Unger's laboratory, which found that ceramides were responsible for palmitate-induced beta-cell apoptosis (94, 149). Subsequent studies also implicated ceramides in the impairment of insulin secretion and insulin gene transcription (94, 150-157). Glycosylated derivatives of ceramide (i.e., gangliosides) have also been identified as putative antigens that contribute to the auto-immune response (158-161). These *in vitro* studies suggest that ceramides could contribute to the decline in beta-cell function that underlies diabetes. Nonetheless, the critical animal experiments allowing for beta-cell-specific modulation of ceramides *in vivo* have not been conducted.

Ceramides in Cardiomyocytes and the Vascular Endothelium

Myriocin prevents or reverses numerous cardiovascular conditions (e.g., atherosclerosis, hypertension, cardiomyopathy, and heart failure) in rodents (22-24, 102, 162). Though these effects could be a consequence of improvements in insulin sensitivity and dyslipidemia, additional studies suggest that ceramides have deleterious actions in cardiac and vascular tissues. For example, the atherogenic effects of ceramide may result from direct actions in blood vessel walls, as they induce transcytosis of oxidized low-density lipoproteins. This action leads to lipid retention (163) and monocyte adhesion (164), thus driving plaque formation. In addition, ceramides may alter vascular reactivity. Studies in isolated blood vessels indicate that ceramides induced colocalization of PP2A with eNOS, leading to the

inhibition of eNOS phosphorylation and its dissociation from Akt/PKB (90). Lastly, ceramides may directly compromise cardiomyocyte function, as pharmacological or genetic inhibition of ceramide synthesis in a model of lipotoxic cardiomyopathy (i.e., cardiac-specific overexpression of human lipoprotein lipase) abrogates apoptosis and cardiac contraction (24). Unfortunately, heart-specific deletion of *Sptlc2* impairs cardiac function (140), making it difficult to ascertain whether the ceramides that regulate cardio-lipototoxicity are generated within the cardiomyocyte.

Ceramides in the Hypothalamus

Ceramides accumulate in the hypothalamus of mice following either high-fat diet feeding or acute lipid infusion (89). They are presumed to serve as nutritional signals that modulate appetite or metabolic rate. Experimental delivery of ceramides into the hypothalamus promotes weight gain and reduces brown adipose tissue thermogenesis (165). By contrast, intracerebroventricular infusion of myriocin reduces ceramide accumulation and improves hypothalamic insulin sensitivity, improving glucose tolerance and restoring beta-cell mass (166). Additional studies suggest that glycosylated ceramides may serve as important modulators of central nervous system function. Mice deficient in glycosphingolipids in the hypothalamus develop progressive obesity due to decreased thermogenesis (167). Moreover, adeno-associated virus (AAV)-mediated *Ugcg* (encoding for GCS) delivery to the hypothalamic arcuate nucleus led to ensuing elevations in nuclear glucosylceramides and subsequent weight loss. Mechanistically, GCS-depleted neurons displayed inadequate leptin receptor activation, requiring neuronal gangliosides GM1 and GD1a to be recruited to the leptin receptor upon ligand stimulation (167). In a follow-up study, Nordström and colleagues (168) ablated *Ugcg* in the neurons that target the mediobasal hypothalamus. They found that glucosylceramides regulate fasting-induced lipolysis by regulating norepinephrine content in the adipose depots. These studies raise the possibility that products of the sphingolipid pathway serve as important gauges of fuel load in the central nervous system.

REGULATION OF CERAMIDE SYNTHESIS AND METABOLISM

The ceramide biosynthesis pathway serves as a gauge of nutritional excess that reflects the supply of precursor substrates such as palmitoyl-CoA and serine (169). In human studies, diets enriched in saturated fat, but not unsaturated fat, increase ceramide accumulation (68). Similar findings are reported in rodents, with C16:0 ceramides emerging in unbiased metabolomic studies across multiple studies and strains as good markers of diet-induced metabolic pathologies (170). Beyond nutritional load, however, researchers have determined that several regulatory factors control rates of ceramide synthesis and degradation (Figure 3).

Inflammation

Obesity is associated with increased systemic inflammation, including a particularly robust recruitment and activation of macrophages within the expanded adipose depots (171). The resulting increase in inflammatory cytokines in serum and tissues enhances rates of ceramide synthesis. Indeed, tumor necrosis factor (TNF)- α , interleukins, and chemokines selectively increase levels of sphingolipids, typically without affecting glycerolipids (172). The combination of increased nutrient supply and inflammation in obesity thus leads to

synergistic enhancement of ceramide production (173). Indeed, inflammatory cytokines show a particularly tight association with ceramides in subjects with insulin resistance (172, 174).

One of the key modulators of ceramide biosynthesis is Toll-like receptor-4 (TLR4), a pattern recognition receptor that activates NF- κ B and drives innate immunity. A particularly well-known receptor agonist is lipopolysaccharide, but the receptor also binds to low-density lipoprotein, viral proteins, polysaccharide, and several other ligands. One of the more intriguing findings relates to saturated fatty acids, which were initially identified as putative ligands (175) but were later found to amplify receptor signaling (176). The presence of an intact TLR4 signaling network is essential for saturated fat-induced insulin resistance and for the induction of ceramide synthesis (89, 175). TLR4 agonists induce transcription of several enzymes in the de novo ceramide biosynthesis pathway (89, 173, 177).

TNF- α , another cytokine that is elevated in obesity, also increases ceramide levels. Studies in cultured cell reveal that it accomplishes this via coordinated activation and enhanced expression of the ceramide generating (e.g., SPT) and metabolizing enzymes (e.g., sphingomyelinase) (30, 123, 178). In cultured myotubes, we determined that TNF- α effects on ceramides are biphasic, with an acute increase resulting from activation of sphingomyelinase followed by a sustained upregulation resulting from enhanced de novo biosynthesis (S.A. Summers, unpublished observation).

Some data suggest that ceramides may themselves stimulate inflammation, which would enable a positive feedback mechanism that would exacerbate tissue damage. For example, they may activate the Nod-like receptor (Nlrp3) inflammasome, which induces caspase 1 cleavage in macrophages and adipose tissue and enhances cytokine release (179).

Endocannabinoids

Cannabinoids, the active components of marijuana (*Cannabis sativa*), exert their psychoactive and peripheral effects by activating G_{i/o} protein-coupled receptors (cannabinoid receptor CB₁R) that modulate ion channels, adenylyl cyclase, and extracellular signal-regulated kinases (180). Scattered reports indicate that cannabinoid receptors may induce ceramides by enhancing sphingomyelin hydrolysis and by stimulating de novo biosynthesis (180, 181). While the actions of ceramides in neurotransmitter release are not clearly elucidated, one study has implicated this cannabinoid action in obesity-driven disease (182). In obese mice, inhibition of CB₁R using the inverse agonist JD5037 or genetic deletion of hepatic CB₁R reversed steatosis and improved glucose tolerance and insulin sensitivity (183). These interventions lowered ceramide levels in the liver by reducing SPT activity, decreasing transcripts encoding biosynthetic genes (i.e., *Sptlc3*, *CerS1*, and *CerS6*), and increasing transcripts encoding ceramidases (i.e., *Asah1* and *Asah2*) (183). Moreover, the SPT inhibitor myriocin abrogated endocannabinoid-induced hepatic insulin resistance (183).

β -Adrenergic Agonists

In adipocytes, β -adrenergic receptors stimulate heat production by increasing mitochondrial activity and uncoupling (184). As noted above, ceramides are potent inhibitors of this

thermogenic response, owing in part to their ability to inhibit mitochondrial activity and block glucose utilization (88, 185). Using a newly developed flux assay to monitor rates of ceramide production, we found that β -adrenergic agonists rapidly and completely shut down ceramide biosynthesis (12), even under conditions where exogenous fatty acids are plentiful. This work reveals a novel regulatory mechanism that controls rates of ceramide biosynthesis to control metabolic activity of adipose depots.

Adiponectin

Adiponectin, a protein hormone secreted from metabolically healthy adipocytes, elicits a broad spectrum of anti-inflammatory, antiatherogenic, and antidiabetic actions (50). For example, the adipokine improves glucose homeostasis by enhancing insulin sensitivity in adipose tissue and the liver and promoting survival of insulin-secreting beta-cells. It improves lipid handling by modulating rates of triglyceride synthesis, oxidation, and lipolysis in adipocytes and the liver. It also reduces inflammation through direct actions on immune cells. These beneficial actions were initially thought to be mediated by AMPK, a serine/threonine kinase activated during starvation (186). However, although AMPK can be activated by adiponectin, it is dispensable for most of the aforementioned physiological responses (49).

Ceramidases bear sequence homology to the receptors within the progestin and adipoQ (PAQR) superfamily, including the adiponectin receptors ADIPOR1 and ADIPOR2. Villa and colleagues (187) determined that these PAQRs promoted accumulation of the free sphingoid base in yeast. Moreover, receptor signaling can be abrogated by the inclusion of ceramidase inhibitors (188). Holland et al. (49) thus hypothesized that adiponectin might elicit its broad spectrum of metabolic improvements by degrading ceramides. These researchers demonstrated that adiponectin, via ADIPOR1 and ADIPOR2, deacylated ceramide and induced sphingosine and S1P (49, 189). They also found that mutation of critical residues in the receptor's ceramidase motif abrogated adiponectin action (49). Proof of the bioactivity of this domain came from studies on the crystal structure of the ADIPOR isoforms. In an initial structural assessment, Tanabe et al. (190) found that human adiponectin receptors possess a hydrophobic binding pocket that resembled a comparable structure in other known ceramidases. Subsequently, Vasiliauskaite-Brooks et al. (51) elucidated the crystal structure of the receptor in the presence of ceramide, determining that it had a bound fatty acid and that the purified protein contained intrinsic ceramidase activity. These data unveiled a new paradigm in receptor signaling, revealing that PAQR ligands could alter tissue function by acutely degrading ceramides.

To test this explicitly, Scherer and colleagues (100) generated mice overexpressing AdipoR selectively in liver. The increase in hepatic AdipoR reduced hepatic ceramide content and increased AdipoR-induced ceramidase activity, leading to improved hepatic insulin resistance and resolved hepatic steatosis.

Intestinal Farnesoid X Receptors

The farnesoid X receptor (FXR) is a nuclear hormone receptor present in the liver and intestines. Bile acids, which are its primary ligands, cause FXR to translocate to the nucleus,

where it suppresses expression of genes (e.g., cholesterol 7 alpha-hydroxylase) required to synthesize bile acids from cholesterol. This negative feedback pathway limits production of new bile acids when enterohepatic levels are high.

The receptor also regulates genes involved in lipid and glucose metabolism and displays a complex and multifaceted role in the development of cardiometabolic pathologies. FXR agonists are in clinical development as a means of reducing hepatic steatosis (191). However, Gonzalez and colleagues found that intestinal FXR enhanced metabolic disorders, owing to its ability to act as a sensor of the gut lumen environment. They found that intestinal FXR enhances hepatic steatosis and gluconeogenesis by stimulating intestinal ceramide biosynthesis (138, 185, 192). Specifically, FXR stimulated transcription of genes relevant to de novo sphingolipid biosynthesis, which increases circulating levels of ceramides and produces a commensurate ceramide-driven induction of genes driving fat accumulation and gluconeogenesis in the liver. These actions limit the therapeutic potential of the FXR agonists and suggest that a combination of tissue-specific pharmaceuticals may be needed.

A UNIFYING HYPOTHESIS ABOUT THE EVOLUTION OF CERAMIDES AS NUTRITIONAL GAUGES

Since fatty acids are detergent-like molecules that can perturb membrane structures, cells go to great lengths to keep free fatty acid levels low. They accomplish this by storing excess fatty acids in macromolecules, with glycerolipids, including the triglycerides, serving as the major storage forms. When the energetic needs of the tissue have been met and the storage capacity starts to fill, ceramides initiate cellular responses that change fuel preference and induce lipid storage. As described herein, they inhibit glucose and amino acid utilization and induce the uptake of fatty acids and their conversion into triglycerides. They also alter mitochondrial efficiency, which may allow cells to metabolize more fatty acids by limiting their impact on mitochondrial membrane potential. When ceramides rise to critical levels, they induce apoptosis to control the death of the compromised and unstable cell. We hypothesize that these responses are part of an evolutionary conserved adaptation to excessive fatty acids.

Though we acknowledge this as speculation, the following ceramide actions support the provocative idea that cells produce ceramides to protect themselves from the deluge of excess fat:

1. Ceramides limit the damage free fatty acids can do to cellular membranes by enhancing their safe passage through bilayers and enabling their esterification (e.g., via CD36) (12, 100).
2. Ceramides facilitate the safe storage of fatty acids by inducing expression of *Srebf1*, a master regulator of genes required for triglyceride synthesis and lipid droplet assembly (12, 138).
3. Ceramides slow release of fatty acids from lipid droplets (via HSL) (12).

4. Ceramides alter fuel preference by limiting the entry of glucose and amino acids into cells (7, 193-195).
5. Ceramides reduce mitochondrial efficiency (12, 25, 26, 98, 122, 196), which allows cells to minimize the effect of the excessive fatty acids on mitochondrial membrane potential.
6. Ceramides mediate fatty acid-induced apoptosis (123).
7. The CERS enzymes themselves, beyond producing ceramides, respond to fatty acids by repressing transcription of lipases (43, 197).

These data clearly identify ceramides as major regulators of cellular metabolism during times of fuel surplus.

OUTLOOK AND FUTURE DIRECTIONS

The astounding number of studies in numerous complementary and redundant rodent models make one thing clear: Ceramides are important nutrient signals that contribute to cardiometabolic disease. Clinical profiling studies further indicate that they are relevant to human disease. These transformative discoveries raise the exciting possibility that ceramide-lowering strategies could have therapeutic value in a wide range of cardiometabolic disease processes. Nonetheless, several important topics remain underexplored.

1. The precise, tissue-specific roles of ceramides are not fully defined. The availability of new tools to enable tissue-specific control of ceramide synthesis, including gain-of-function models, will likely reveal which tissues are most sensitive to ceramides.
2. Although several key mechanisms have been identified (e.g., ceramide regulation of AKT), we do not yet have a very satisfying explanation for how ceramides are detected in cells. For example, the reasons that the double bond in the sphingoid base and the length of the acyl chain are such critical determinants of ceramide action are largely unknown.
3. Little information exists about the genetic determinants of ceramide accumulation in the population. Several single nucleotide polymorphisms have been identified in ceramide-modifying enzymes. For example, CERS2 polymorphisms are associated with increased HbA1c and risk of diabetic kidney disease (198-200). This common natural variant results in the replacement of a glutamate to an alanine in an important protein domain and is predicted to be inactivating. The impact of this and other rare and common variants on sphingolipid accumulation and disease risk merits additional attention.
4. The regulatory elements that influence rates of ceramide production, including the activity and expression of ceramide-synthesizing and -metabolizing enzymes, are still not fully resolved.

5. Therapeutics to safely lower ceramides in humans have not been developed. Alternatively, insufficient information exists regarding specific behavioral interventions that lower ceramides and improve patient health.

Despite these areas of uncertainty, ceramides undoubtedly serve as a nexus in nutrient signaling networks and have profound effects on a wide variety of metabolic disease processes. Research on these exciting lipids holds great promise as a means of combating the metabolic disease epidemic.

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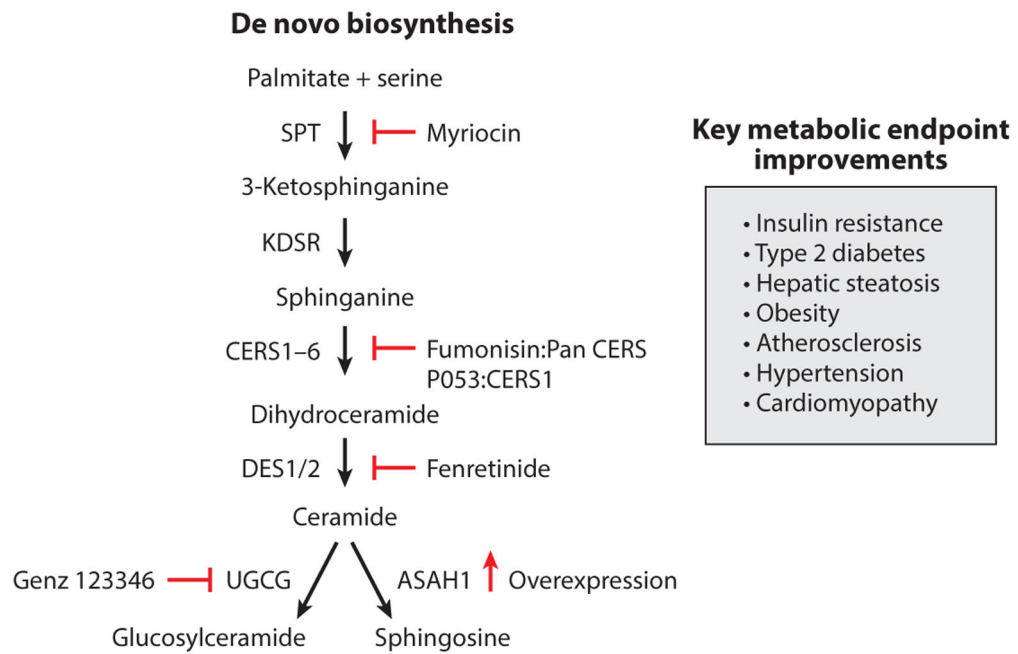


Figure 1.

De novo ceramide synthesis pathway and known inhibitors shown to improve metabolic diseases. Schematic of the de novo ceramide biosynthesis pathway highlighting the key enzymes involved in ceramide biosynthesis and their respective inhibitors. Abbreviations: ASAH1, acid ceramidase; CERS, ceramide synthase; DES, dihydroceramide desaturase; KDSR, 3-ketodihydrospingosine reductase; SPT, serine palmitoyltransferase; UGCG, UDP-glucose ceramide glucosyltransferase.

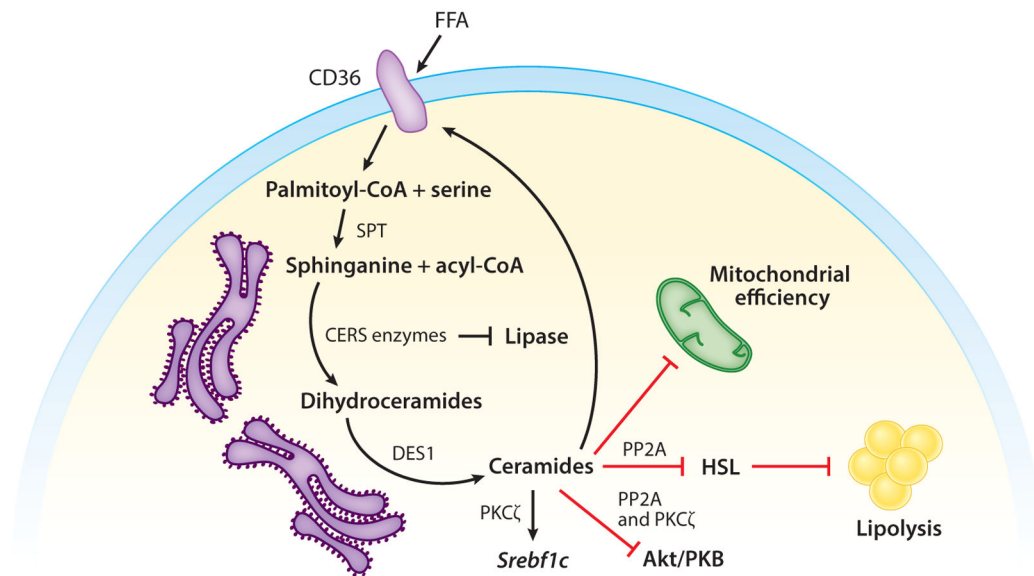


Figure 2.

Molecular mechanisms via which ceramides gauge intracellular lipids. Ceramides regulate metabolism through the following mechanisms: translocation of CD36 toward the plasma membrane, thus facilitating the safe uptake of FFA and enhancing the conversion of fatty acids into acyl-CoA, induction of *Srebf1c*, inhibition of Akt/PKB, inhibition of HSL, safe storage of ceramide as 1-*O*-acylceramide inhibition of mitochondrial metabolic efficiency, and inhibition of lipases. Abbreviations: Akt/PKB, protein kinase B; CD36, cluster of differentiation 36; CERS, ceramide synthase; DES1, dihydroceramide desaturase 1; FFA, free fatty acid; HSL, hormone-sensitive lipase; PKC ζ , protein kinase zeta; PP2A, protein phosphatase 2A; SPT, serine palmitoyl transferase; *Srebf1c*, sterol regulatory element-binding protein 1.

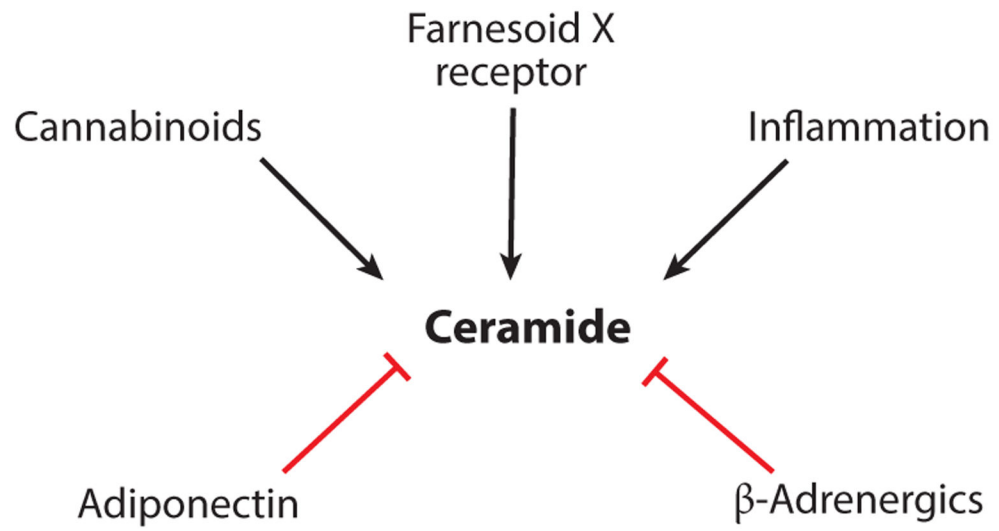


Figure 3. Key regulators of ceramide accumulation. Depicted are the key regulators of ceramide synthesis thus far identified and involved in metabolic diseases.

Table 1

Summary of mouse models used for manipulating de novo ceramide synthesis pathway in vivo to determine their role in metabolic diseases

Gene	Knockout/overexpression	Phenotype
<i>Sptlc1</i>	Whole body	Embryonic lethal (201)
	Adipose tissue	Lipodystrophy (139)
<i>Sptlc2</i>	Whole body	Embryonic lethal (201, 202)
	Haploinsufficiency	Improve insulin sensitivity and resolve hepatic steatosis (108)
	Liver	
	Adipose tissue	Increase energy expenditure, improve insulin sensitivity, and resolve hepatic steatosis (88), lipodystrophy (140)
	Myeloid cells	No phenotype (88)
	Intestinal cells	Lethal (203)
<i>CerS1</i>	Whole body	Increase energy expenditure, improve insulin sensitivity (99)
	Skeletal muscle	
<i>CerS2</i>	Haploinsufficiency	Impair insulin sensitivity and develop hepatic steatosis (25)
<i>CerS3</i>	Whole body	Impair skin barrier function, lethal (204)
<i>CerS4</i>	Whole body	Alopecia (205)
<i>CerS5</i>	Whole body	Two studies: one demonstrating improved insulin sensitivity (106) and another showing no phenotype (26)
	Skeletal muscle	No phenotype (99)
<i>CerS6</i>	Whole body, liver, brown adipose tissue	Increase energy expenditure, improve insulin sensitivity, and resolve hepatic steatosis (26)
	Myeloid cells	No phenotype (26)
	Skeletal muscle	No phenotype (99)
<i>Degs1</i>	Whole body, liver, adipose tissue	Improve insulin sensitivity and resolve hepatic steatosis (12)
	Myeloid cells	No phenotype (12)
	Intestinal cells	No phenotype (12)
<i>Asah1</i>	Liver (overexpression) Adipose tissue (overexpression)	Improve insulin sensitivity and resolve hepatic steatosis (100)
<i>Ugcg</i>	Intestinal cells	Regulate nutrient uptake (206)
	Central nervous system and hypothalamus	Regulate energy homeostasis and peripheral lipolysis (167, 168)