REVIEW

A role for sphingolipids in the pathophysiology of obesity-induced inflammation

Benjamin T. Bikman

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Abstract Following the initial discovery that adipose tissue actively synthesizes and secretes cytokines, obesityinduced inflammation has been implicated in the etiology of a host of disease states related to obesity, including cardiovascular disease and type II diabetes. Interestingly, a growing body of evidence similarly implicates sphingolipids as prime instigators in these same diseases. From the recent discovery that obesity-related inflammatory pathways modulate sphingolipid metabolism comes a novel perspective—sphingolipids may act as the dominant mediators of deleterious events stemming from obesity-induced inflammation. This paradigm may identify sphingolipids as an effective target for future therapeutics aimed at ameliorating diseases associated with chronic inflammation.

Keywords Sphingolipid · Obesity · Inflammation · TLR4

Introduction

Despite an explosion in interventions aimed at holding back the tide of obesity and improving public health, the rising trends in its prevalence show little sign of slowing. The economic and personal toll of obesity is enormous. Medical costs associated with treating obesity and its complications are roughly \$150 billion annually, and obese individuals can expect average annual medical expenses to be approximately \$1,500 more than normal-weight persons [1]. The problem is not merely one of money, however,

B. T. Bikman (🖂)

considering that obesity is estimated to result in up to 20 years of life lost when compared to the non-obese [2]. If interventions aimed at averting obesity continue to fail, trends can be expected to increase with an estimated 2.4 million more adults becoming obese annually, followed closely by an ever-greater financial burden. The observation by the ancient Greek physician Hippocrates that "sudden death is more common in those who are naturally fat than in the lean" astutely describes the increase in mortality associated with obesity.

Of course, the increased health-related expenses and risk of mortality are not simply due to the mechanical discomforts and limitations that accompany an expanding fat mass. Obesity not only adversely affects tissue function but also contributes to an elevated risk of developing several fatal diseases, such as hypertension, atherosclerosis, type 2 diabetes mellitus, and nonalcoholic fatty liver disease [3]. As such, it is not surprising that extensive efforts have been devoted to understanding the role of obesity in the etiology of these prominent diseases. In particular, two deleterious factors often associated with obesity are implicated in increasing disease risk-lipotoxicity and inflammation. Lipotoxicity refers to the ectopic deposition of fat in tissues not intended as lipid storage sites. Of the myriad of bioactive lipids in tissues, however, the sphingolipids warrant particular attention due not only due to being highly correlated with the degree of obesity, but also implicated in the etiology of various diseases [4–7]. In addition, inflammation, defined by elevated levels of proinflammatory cytokines and increased presence and activity of monocytes/macrophages, is also present in obesity [8–11]. Interestingly, recent efforts reveal that these two seemingly distinct characteristics of obesity, lipotoxicity and inflammation, share a degree of linearity that might alter our perspective on the mechanism of inflammation-induced complications in various diseases.

Department of Physiology and Developmental Biology, Brigham Young University, 593 WIDB, Provo, UT 84602, USA e-mail: benjamin_bikman@byu.edu

Thus, it is the intention of this review to explore the role of sphingolipids in the pathophysiology of some of the prominent diseases associated with obesity-induced inflammation.

Obesity, inflammation, and sphingolipids

Obesity-induced inflammation

Gone are the days when adipose tissue was considered a passive organ whose exclusive functions are to accumulate triglycerides in hypertrophied adipocytes during caloric excess and release these lipids during caloric restriction. The exploration of obesity and inflammation began in adipose tissue. From the initial observation that adipose is capable of synthesizing and releasing tumor necrosis factor α (TNF α) [12], the total number of adipose-derived molecules (termed "adipokines") has ballooned. Compared to adipose from lean individuals, adipose tissue from the obese has increased levels of several inflammatory proteins, such as interleukin-6 (IL-6), iNOS, C-reactive protein (CRP), monocyte chemotactic protein-1 (MCP-1), and plasminogen activator inhibitor type-1 (PAI-1), among others [13-16]. Transcriptional profiling reveals that inflammatory genes are among the most abundantly regulated in adipose tissue in obesity [17]. However, adipose tissue is not simply a homogenous collection of adipocytes. Expanding adipose tissue is accompanied by increased infiltration of activated macrophages, and, while adipocytes themselves appear to express the cellular machinery to enable cytokine synthesis and responsiveness [18], these resident macrophages play a dominant role in adiposederived proinflammatory gene expression [19].

The adipokine profile secreted from adipose tissue is dynamic and is affected by fat mass status (Fig. 1). While proinflammatory cytokine expression (e.g., TNF α , MCP-1, IL-1 β) is increased with weight gain by fat mass expansion



Fig. 1 Adipocyte expansion, leading to obesity, is associated with reduced adiponectin levels and elevated $TNF\alpha$ and ceramide

[12], anti-inflammatory protein expression (e.g., adiponectin, IL-10) is elevated in adipose from lean or increased with weight loss in the obese [20, 21]. Additionally, circulating and tissue ceramide levels follow a trend similar to proinflammatory cytokines. Though once considered to be nothing more than correlative companions [22], we now know that sphingolipid biosynthesis requires activation of immune receptors and pathways.

TLR4 and sphingolipids

Recent findings have refuted the old notion that production of ceramide, the backbone of all higher-order sphingolipids, is exclusively controlled by substrate availability. Rather, it appears hormonal signals have a powerful influence to alter ceramide metabolism, resulting in ceramide accumulation or degradation. Similar to inflammatory status in lean and obese, ceramide metabolism and accumulation is altered in the tissues of lean and obese. Both rodent and human models of obesity have shown an increase in ceramide levels in a variety of tissues, such as skeletal muscle, liver, and hypothalamus [7, 23–26]. Also, paralleling the shift in inflammatory status with weight loss [11], a reduction in fat mass is associated with a reduction in tissue and circulating ceramide levels [7, 27, 28]. For example, Huang et al. [27] found that substantial surgeryinduced weight loss ($\sim 25\%$ of body weight) in morbidly obese humans correlated with a reduction in plasma ceramides. However, Dube et al. [29] found that even modest weight loss (2% body weight) is associated with a significant drop in ceramide levels.

The parallel changes in inflammatory status and ceramide metabolism are not coincidental. We recently found that the activation of inflammatory pathways is a necessary event in ceramide biosynthesis, with both saturated fatty acids and bacterial endotoxins converging on the Toll-like receptor 4 (TLR4) pathway to induce ceramide accumulation (Fig. 2)[23]. Using mutant mice lacking a functional TLR4 (C.C2- $Tlr4^{Lps-d}$), we found that TLR4 is required for saturated fatty acid- and endotoxin-induced ceramide biosynthesis in skeletal muscle and liver. Interestingly, while saturated fatty acids are known to induce both an inflammatory response and de novo ceramide biosynthesis, unsaturated fatty acids, which do not activate TLR4, fail to do both [23]. In addition to saturated fats, lipopolysaccharide (LPS), a major bacterial membrane lipid and TLR4 ligand, similarly induced ceramide biosynthesis in a TLR4dependent manner [23].

TLR4 initiates signaling through the canonical IKK β -NF- κ B pathway. Briefly, upon activation, IKK β phosphorylates and marks the NF- κ B inhibitor, I κ B α , for degradation, after which NF- κ B is liberated and free to migrate into the nucleus to initiate transcription of various



Fig. 2 The TLR4-induced activation of ceramide biosynthesis represents a common pathway between lipotoxicity- and inflammationinduced insulin resistance. Saturated fatty acids (SFA); lipopolysaccharide (LPS); tumor necrosis factor (TNF) α ; inhibitor of κ B kinase (IKK) β ; sphingomyelinase (SMAase)

cytokines. IKK β has been previously implicated in acting as a prime mediator of inflammation-induced metabolic disorders via its inhibitory effects on proximal insulin signaling [30, 31]. We found that IKK β was a necessary downstream factor in TLR4-induced ceramide biosynthesis. Not only was IKK β ablation associated with dramatically reduced ceramides in spite of TLR4 activation, it was necessary for the fatty acid- and inflammationinduced increase in transcription of several of the enzymes involved in de novo ceramide synthesis [23]. Additionally, diet-induced obese mice treated with the IKK β inhibitor sodium salicylate fail to accrue ceramides [23].

As mentioned, both saturated fatty acids (SFA) and bacterial endotoxins appear to activate TLR4, though whether they activate the receptor via similar or disparate mechanisms is not clear. The findings that SFA, but not unsaturated FA, activate TLR4 are widely reported [23, 32–34]. In contrast, Erridge et al. [35] found conflicting results, reporting that SFA failed to activate TLR4 in multiple cell types and implicating LPS contamination in fatty acid preparations as the source of TLR4 activation. However, this fails to explain the contrasting effects of SFA compared to unsaturated counterparts, which should be similarly contaminated [23]. Another argument against SFA and endotoxins affecting TLR4 via a single mechanism is the different kinetics of TLR4 activation between the two purported ligands. In exploring the inflammatory effects of SFA and LPS, Shi et al. [36] observed that orders of magnitude more SFA are necessary to get a comparable response seen with LPS, indicating LPS is a far more potent TLR4 activator, regardless of the mechanism. Similarly, LPS elicited a significantly greater increase in NF- κ B activation compared to SFA [36]. Additionally, non-TLR4 mechanisms may explain the ability of SFA to activate inflammatory pathways [37].

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Cytokines and sphingolipids

However, the absence or presence of TLR4 signaling does not explain every alteration in sphingolipid metabolism in response to the myriad of proinflammatory or anti-inflammatory stimuli. While TNF α and other proinflammatory cytokines (e.g., IL-1 β) can activate de novo ceramide synthesis [38], likely via increased IKK β -NF- κ B action, they can also exploit a 'backdoor'. Several proinflammatory cytokines result in ceramide accrual by activating sphingomyelinase, which mediates the conversion of sphingomyelin to ceramide. Specifically, TNF α and IL-1 β signaling activates acid and neutral sphingomyelinase activity, resulting in significant ceramide pooling [39–41]. Considering that sphingomyelin is the dominant membrane lipid, its role as a source of ceramide is potentially enormous.

In contrast, adiponectin has long been known to oppose proinflammatory cytokines. Whereas $TNF\alpha$ activates proinflammatory pathways and induces cell death, adiponectin inhibits proinflammatory pathways and promotes cell survival [42, 43]. Importantly, evidence suggests that the contrasting effects of the quintessential pro- and antiinflammatory cytokines TNFa and adiponectin, respectively, can be explained by how they modulate ceramide metabolism (Fig. 3). Holland et al. [44] recently demonstrated that adiponectin receptors contain inherent ceramidase activity [45], resulting in the degradation of ceramide to form sphingosine [46]. Sphingosine, in turn, is phosphorylated by sphingosine kinase, forming sphingosine 1-phosphate (S1P). Of the several metabolic fates available to ceramide, its deacylation and phosphorylation (via ceramidase and sphingosine kinase, respectively) to



Fig. 3 TNF α induces ceramide biosynthesis and accrual through multiple mechanisms, whereas adiponectin activates ceramide degradation and eventual formation of S1P. Ceramide (Cer); sphingosine 1-phosphoate (S1P)

form S1P is intriguing, inasmuch as ceramide and S1P exert completely different and even antagonistic actions. In stark contrast to ceramide, S1P has been repeatedly shown to activate Akt and promote cell survival and growth, even directly antagonizing ceramide [46–50]. Indeed, S1P has been shown to reduce ceramide synthesis by inhibiting de novo enzymatic activity [51], which gives rise to the theory that ceramide and S1P constitute a rheostat system [52]. Interestingly, S1P may be a critical mediator of adiponectin's anti-inflammatory profile. S1P has been shown to inhibit actions of pro-inflammatory cytokines [53] and regulate inflammation-related gene expression [54].

Sphingolipids in the pathophysiology of obesity-induced inflammation

The inflammatory state associated with obesity is implicated in several clinically important complications. Similarly, a growing body of literature suggests that ceramide and other sphingolipids are also present in these disease states and may play a prominent role in the etiology. In light of findings suggesting that sphingolipid metabolism is affected by inflammatory profile [23, 44], it seems likely that sphingolipids are downstream effectors of obesity-induced inflammation and are critical mediators of inflammation-associated diseases. This section will highlight the role of sphingolipids in the etiology of several diseases associated with obesity-induced inflammation (Fig. 4).

Cardiovascular disease

A role for inflammation in the etiology of cardiovascular disease, a term used to collectively describe diseases that involve the heart or blood vessels, has become so well



Fig. 4 Ceramide has been shown to cause virtually all of the pathological states elicited by obesity-induced inflammation

established over the past two decades that a number of inflammatory markers are now measured as potential predictors of prevalent or incident cardiovascular disease [55]. Additionally, anti-inflammatory medications are now used to reduce the risk of cardiovascular disease [55, 56]. Similarly, sphingolipids have been shown to play a role in the regulation of vascular growth and tone, thus impacting cardiovascular function. We will explore the impact of sphingolipids in three prominent cardiovascular disorders commonly associated with obesity-induced inflammation, namely hypertension, atherosclerosis, and cardiomyopathy.

Hypertension

Hypertension, identified as clinically elevated blood pressure, is a major risk factor for cardiac and cerebrovascular disease. Noting the vascular effects of sphingolipids, Spijkers et al. [57] sought to determine the role of sphingolipids in essential hypertension. They found that shifting the ceramide/S1P ratio towards ceramide dominance by administering a sphingosine kinase inhibitor or by exogenous sphingomyelinase induced pronounced endotheliumdependent contractions in isolated carotid arteries. Additionally, in vivo administration of a sphingosine kinase inhibitor resulted in a marked rise in blood pressure in spontaneous hypertensive rats, and, further implicating ceramide, hypertensive rats have significantly increased levels of total ceramides in arterial tissues. Moreover, both hypertensive rats and humans have elevated plasma ceramide levels [57]. Similarly, in an effort to decipher the genetics of hypertension, Fenger et al. [58] conducted a genomic analysis focusing on components of ceramide metabolism and concluded that the ceramide/S1P rheostat has a substantial influence on blood pressure regulation. Interestingly, they found that genes involved in de novo ceramide synthesis, rather than ceramide formation via sphingomyelinase, were the most important sources of ceramide in a hypertensive population.

Atherosclerosis

Several early reports have cited a consistent positive correlation between circulating sphingolipids and occurrence of cardiovascular complications [59–62]. In fact, the correlation is so consistently observed that sphingomyelin levels are considered an independent risk factor for coronary artery disease [63]. In support of these observations, recent findings have established that sphingolipids indeed play a causative role in the etiology of atherosclerosis. Park et al. [64] found that treating apoE KO mice, which develop advanced atherosclerosis and have elevated circulating sphingolipids [59], with an inhibitor of de novo ceramide synthesis, myriocin, not only reduced plasma ceramide and sphingomyelin, but also total plasma cholesterol and triglycerides. Further, inhibition of ceramide synthesis in the KO mice resulted in a substantial reduction in atherosclerotic lesion area in the aortic root and slowed the progression of atherosclerosis in the brachiocephalic artery. Macrophage content in the aortic root was similarly reduced in myriocin-treated mice [59]. Interestingly, inhibition of de novo ceramide synthesis also results in increased hepatic apoA-I synthesis and elevated circulating levels of favorable HDL cholesterol [65]. Similar to treatments using inhibitors of ceramide synthesis, the use of a myriocin-based drug FTY720, an S1P homologue, also inhibits atherosclerosis [66].

Underlining the connection between inflammation and sphingolipids, LDL receptor KO mice that lack the ability to generate sphingomyelin in macrophages, the archetypal immune cell, have decreased atherosclerotic lesions in the entire aorta. Moreover, plaque morphology analysis from brachiocephalic arteries of LDL receptor KO mice reveal reduced necrotic core area [67].

Cardiomyopathy

Like most cardiovascular complications, cardiomyopathy, or weakening of the heart, has strong ties to inflammation [68, 69]. However, inflammation per se may be a sufficient, but unnecessary factor in cardiomyopathy etiology. To tease out the role of inflammation versus the inflammationrelated factor NF- κ B, Kawamura et al. [70] blocked the activation of cardiac NF- κ B by crossing transgenic mice harboring cardiac-specific $TNF\alpha$ overexpression with mutant mice carrying a disrupted NF-kB subunit. Interestingly, they found that while NF- κ B blockade did not ameliorate myocardial inflammation as determined by inflammatory cell infiltration, it significantly improved cardiac function and survival, hinting that the classic inflammation-immune response is less important than the activation of NF- κ B in cardiomyopathy. Combined with the separate observation that TLR4 deficiency protects against cardiomyopathy [63], and the fact that TLR4 and NF- κ B are regulators of de novo sphingolipid synthesis [23], these findings may suggest that a product(s) of inflammation, rather than inflammation per se, is responsible for the inflammation-induced cardiomyopathy.

An oft-used rodent model of cardiomyopathy is the cardiomyocyte overexpression of a glycosylphatidylinositol membrane-anchored form of lipoprotein lipase (LpL^{GPI}). Park et al. [63] explored the role of ceramide in this model by (a) crossing the LpL^{GPI} mouse with a heterozygous deletion of LCB1, a SPT subunit, and (b) treating LpL^{GPI} mice with myriocin. In addition to finding that the hearts from LpL^{GPI} mice contained 45% more ceramide than control levels, they found that both genetic (LpL^{GPI}-LCB1

cross) and pharmacological (myriocin) inhibition of ceramide synthesis improves cardiac function and reduces expression of heart failure markers [63].

Non-alcoholic fatty liver disease

Non-alcoholic fatty liver disease (NAFLD) is defined as fat accumulation in excess of 5 to 10% of liver weight [71]. While seemingly benign at onset, NAFLD, or hepatic steatosis, can develop into considerably less benign conditions, such as steatohepatitis, cirrhosis, and even hepatocellular carcinoma [72, 73]. In fact, NALFD is the most common cause of liver dysfunction in the United States [72].

A large part of NALFD's prevalence is its association with obesity. The hyperlipidemia and chronic inflammation associated with obesity impact NAFLD risk. Although lipid accumulation is the hallmark of early NAFLD, latter stages are marked by elevated inflammation [74]. Patients with NALFD have elevated $TNF\alpha$ levels compared to obese and non-obese controls [75], and TNF α receptor level and TNF α gene expression are increased in the livers of those with NAFLD when compared to healthy livers [76]. In contrast, adiponectin levels are reduced in patients with NAFLD [75]. Moreover, adiponectin transcript expression in higher-fat livers is negatively associated with hepatic ceramides and sphingomyelinase transcript levels [22]. By investigating adiponectin and its receptors, Peng et al. [77] found that circulating adiponectin was decreased in diet-induced obese (DIO) mice compared to lean controls, and that DIO mice have dramatically reduced hepatic expression of adipoR2, the predominant adiponectin receptor in the liver. Similar examinations in humans have revealed comparable results-a reduction in hepatic adiponectin and adipoR2 expression with advancing steatosis [78, 79].

Considering the contrasting presence of TNF α and adiponectin in NAFLD, and given that TLR4 has been implicated in the etiology of NAFLD complications [80], it is not surprising that modulation of sphingolipid metabolism impacts NAFLD. Memon et al. [81] observed that LPS treatments, a TLR4 activator, resulted in a twofold increase in hepatic SPT gene expression and enzyme activity, which was supported by a 75 and 200% increase in hepatic sphingomyelin and ceramide, respectively. Further, they found that IL-1 β injection increased hepatic SPT transcription and activity in vivo and both IL-1 β and TNF α induced SPT transcript levels in cultured hepatocytes [81]. Additionally, SPT inhibition with myriocin results in a significant reduction in hepatic triglycerides in DIO mice [82].

In addition to implicating the de novo ceramide synthesis pathway, several groups have explored the role of the salvage pathway in NAFLD, where ceramide is synthesized from sphingomyelin (SM) via sphingomyelinase (SMase). Deevska et al. [83] demonstrated that mice lacking acid SMase and functional LDL receptors are protected from diet-induced hepatic steatosis when compared to littermates with functional acid SMase. Moreover, inhibition of SMase in palmitic acid-treated hepatocytes exhibited significantly reduced triglyceride levels compared to untreated controls, suggesting a role for SMase in mediating steatosis in response to elevated fatty acids in vivo [83]. SMase is mediated by numerous stimuli, including TNF α [84–90], which induces SMase by binding the p55 TNF α receptor (also known as TNF type 1 receptor)[91]. This is noteworthy, given that p55 TNFR KO mice are resistant to diet-induced steatosis and liver injury [92].

Insulin resistance

The development of the concept that obesity-induced inflammation mediates disease onset started with a pivotal publication by Hotamisligil et al. [9], demonstrating that adipose-released cytokines inhibit insulin signaling. Insulin resistance mediates a surplus of diseases, including cardiovascular disease [93], type II diabetes, and even some cancers [94, 95]. Hence, understanding the etiology of insulin resistance with regards to inflammatory pathways is intensely pursued.

ΙΚΚβ

Following their novel observation that adipose-secreted TNF α inhibits insulin signaling [9], Hotamisligil et al. [96] found that the molecular mechanisms linking TNF α to insulin resistance involved serine phosphorylation of insulin receptor substrate (IRS)-1, inhibiting normal tyrosine phosphorylation and reducing IRS-1 action. This was later revealed to be through the actions of the serine kinase inhibitor of κ B kinase- β (IKK- β) that phosphorylates IRS-1 and impairs the ability of IRS-1 to associate with the insulin receptor, which subsequently inhibits insulin-stimulated tyrosine phosphorylation and activation of IRS-1-associated PI3-kinase [11, 97].

The realization that IKK β mediates insulin resistance elicited a rediscovery of insulin-sensitizing therapies. Over a century ago, Williamson et al. [98] showed that salicylate treatment reduced symptoms associated with diabetes. This was later further confirmed when Reid et al. [99] demonstrated that a regimen of aspirin improved glucose tolerance in diabetic patients. More recently, salicylate was found to inhibit IKK β activity [100] and the Shoelson lab was instrumental in establishing that salicylates, via IKK β disruption, were effective in preventing inflammationinduced insulin resistance [101].

Research has continued to support a role for IKK β in mediating inflammation-induced insulin resistance, though the originally proposed mechanism, namely serine phosphorylation of IRS-1, is debated [102]. A novel perspective to IKK β -induced insulin resistance is the observation that IKK β regulates ceramide synthesis. In murine myotubes overexpressing a kinase dead IKK β , ceramide levels are dramatically reduced compared to wild-type cells, and the lack of functional IKK β prevents ceramide biosynthesis in response to common insults, like saturated fatty acids [23]. This was found to be a result of IKK β -NF- κ B-mediated transcription of enzymes involved in de novo ceramide biosynthesis, including SPT2, various ceramide synthase isoforms, and dihydroceramide desaturase 1. Additionally, further evidence for ceramide as a mediator of IKK β induced insulin resistance comes from treatment of DIO mice with sodium salicylate, an IKK β inhibitor. Sodium salicylate-treated mice fail to accrue ceramide in skeletal muscle and liver and remain insulin sensitivity despite DIO [23].

Toll-like receptor 4

Obesity is the most important known risk factor contributing to insulin resistance [103]. As mentioned earlier, efforts to identify the molecular mechanisms to explain obesity-induced insulin resistance have revealed lipotoxicity and inflammation as two prominent explanations linking obesity to insulin resistance. A milestone was reached when the Flier laboratory [36] revealed that TLR4, a pattern-recognition receptor that plays a critical role in innate immunity by activating the canonical NF- κ B inflammatory pathway, is a carrefour of lipid- and inflammation-induced insulin resistance. They found that saturated fatty acids, which are elevated in obesity, activate TLR4 and evoke a TLR4-dependent inflammatory response. Additionally, mice lacking TLR4 were protected from acute lipid- and chronic HFD-induced insulin resistance, despite weight gain [36]. Similar to earlier reports, IRS-1 serine phosphorylation was increased in insulinresistant WT mice, but not in insulin-sensitive TLR4deficient mice. However, as with IKK β , accumulation of ceramide might provide an alternate explanation to TLR4incuced insulin resistance.

An association between TLR4 and ceramides has long been known, showing that TLR4 agonists activate ceramide-generating pathways, including de novo synthesis and sphingomyelin salvage [104–106]. Similar to Flier's observations [36], we found that mice lacking functional TLR4 were protected from a variety of insulin-desensitizing interventions, such as acute lipid infusions, diet-induced obesity, and LPS infusions. Additionally, we found that TLR4 mutant mice, in contrast to controls, failed to accrue ceramides in response to HFD in insulin-responsive tissues, like the muscle, liver, and hypothalamus. Therefore, we posit that TLR4 stimulates ceramide synthesis, which represents the common molecular mediator behind lipid- and inflammation-induced insulin resistance.

Adiponectin

In contrast to many of its fellow adipose-secreted hormones, adiponectin is known for its anti-inflammatory and insulinsensitizing functions. Similarly, rather than increasing with obesity, like many adipokines, adiponectin secretion is reduced with fat mass expansion [107] and increased with weight loss [21]. It is therefore little surprise that adiponectin reverses insulin resistance in obesity. Yamauchi et al. [108] found that obese insulin-resistant mice experienced significant improvements in insulin sensitivity and glucose tolerance following adiponectin treatment.

For some time, the positive effects of adiponectin treatment were thought to be mediated almost entirely by its activation of AMP-activated protein kinase (AMPK)[109], though this is not necessarily the case [110]. Adiponectin research took a significant step forward when it was found that adiponectin receptors contain inherent ceramidase activity, degrading ceramide and forming sphingosine and S1P [45, 111]. Indeed, it appears that many of adiponectin's beneficial effects are mediated by accumulation of S1P, including the activation of AMPK [44]. Unsurprisingly, where its proinflammatory counterparts inhibit insulin sensitivity, the anti-inflammatory adiponectin improves insulin sensitivity. In treating two models of obesity-induced insulin resistance with adiponectin, Holland et al. [44] found that the usual accumulation of tissue ceramides evident in $Lep^{ob/ob}$ and HFD-fed mice was conspicuously absent, and the reduction in ceramides was associated with a significant improvement in whole-body insulin sensitivity in both models. Moreover, tissue ceramide levels in disparate adiponectin models (transgenic overexpression or adiponectin-null mice) reveal that adiponectin function is inversely correlated with ceramide levels, and reduced ceramides convey protection against insulin resistance [44].

Given the widely acknowledged benefits of AMPK activation, pharmacological activators of AMPK are actively studied. In particular, metformin, the most widely prescribed anti-diabetic drug, activates AMPK, which has been shown to potently inhibit inflammatory mediators [30, 112–114]. Moreover, AMPK inhibits de novo ceramide synthesis in astrocytes [115] and myotubes (Bikman BT, unpublished observation) via SPT inhibition. However, as is the case with adiponectin, it is yet unknown whether the beneficial effects of metformin-induced AMPK activation require the subsequent actions of S1P [44].

Non-ceramide sphingolipids

While most research centered on sphingolipids and insulin resistance has focused on ceramides as the dominant sphingolipid responsible for insulin resistance, atherosclerosis, etc., caution is required. Because the majority of interventions used to explore ceramides as a mediator of insulin resistance have used inhibitors of de novo sphingolipid synthesis, it is likely that all sphingolipids downstream of ceramide are also reduced and, hence, may be important mediators of effects assigned exclusively to ceramide. Indeed, evidence exists to support a role for both glucosylceramides [116] and sphingomyelin [117] as necessary for a host of health complications, including insulin resistance and NAFLD. However, while other sphingolipids are clearly capable of inducing deleterious consequences, and indeed may be the prime sphingolipids mediating these effects, there is somewhat conflicted evidence concerning these lipids and their responsiveness to inflammatory signals (Fig. 5). The sphingomyelin-degrading actions of $TNF\alpha$ and other inflammatory cytokines via sphingomyelinase are well established [39-41] and covered above (See "Cytokines and sphingolipids"). The data surrounding glucosylceramides is less clear. While we and others have shown that TLR4 activation in macrophages elicits an increase in glucosylceramides [23, 105], unlike ceramides, this effect is not observed in other cell types or whole tissues (e.g., myocytes or whole muscle)[23]. However, given that macrophages reside in tissues that play a large role in metabolic function (e.g., muscle and liver), this may prove to be an important factor. Nevertheless, until the effects of macrophage-derived sphingolipids are more clearly understood, it is unknown whether inflammation-induced macrophage-derived glucosylceramides mediate deleterious metabolic outcomes.

Ceramides and inflammation

Although the majority of research, and the paradigm of this review, indicates that inflammatory signals activate ceramide biosynthesis, a conflicting perspective exists. After observing that the nucleotide-binding domain, leucine-richcontaining family, pyrin domain-containing-3 (Nlrp3) inflammasome, a Nod-like receptor, correlated with degree of obesity-induced insulin resistance, Vandanmagsar et al. [118] found that ablation of the Nlrp3 inflammasome prevents HFD-induced insulin resistance. Interestingly, they report that ceramide activates the Nlrp3 inflammasome, though it is noteworthy that ceramide is not used alone in treatments, but only in combination with LPS [118]. This combination of lipid and LPS requires the reported results to be viewed with an added measure of caution given the synergistic amplification of the inflammatory response in cells exposed to SFA and LPS together compared to either Fig. 5 Several sphingolipids elicit metabolically deleterious results, similar to ceramide. However, only ceramide formation (red pathways) has been shown to mediate inflammation-induced disturbances. Inhibition of the indicated enzymes invites the indicated results. Serine palmitoyltransferase (SPT); dihydroceramide desaturase 1 (Des1); sphingomyelinase synthase (SMS); sphingomyelinase (SMase); glucosylceramide synthase (GCS); sphingosine kinase (SK)



insult alone [119], which may be a result of a lipid-induced TLR4 dimerization [120].

Conclusions

Since its initial discovery, obesity-induced inflammation has lead to an eruption in research implicating inflammatory pathways in the etiology of several prominent obesityrelated diseases. The purpose of this review has been to highlight the role of inflammatory pathways in sphingolipid biosynthesis and, to a degree, establish a role for sphingolipids as critical mediators of many of the deleterious effects of obesity-induced inflammation. While other mechanisms may exist, it is clear that sphingolipids play important roles in the etiology and lethality of obesityrelated disease states commonly linked with inflammation, namely cardiovascular complications, NAFLD, and insulin resistance. Future efforts will not only further elucidate the role of sphingolipids in the pathophysiology of obesityinduced inflammation, but also likely find even greater obesity-related disease states mediated by sphingolipids.

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