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Sphingosine kinase and sphingosine-1-phosphate in liver pathobiology

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Abstract

Over 20 years ago, sphingosine-1-phosphate (S1P) was discovered to be a bioactive signaling molecule. Subsequent studies later identified two related kinases, sphingosine kinase 1 and 2, which are responsible for the phosphorylation of sphingosine to S1P. Many stimuli increase sphingosine kinase activity and S1P production and secretion. Outside the cell, S1P can bind to and activate five S1P-specific G protein-coupled receptors (S1PR1–5) to regulate many important cellular and physiological processes in an autocrine or paracrine manner. S1P is found in high concentrations in the blood where it functions to control vascular integrity and trafficking of lymphocytes. Obesity increases blood S1P levels in humans and mice. With the world wide increase in obesity linked to consumption of high-fat, high-sugar diets, S1P is emerging as an accomplice in liver pathobiology, including acute liver failure, metabolic syndrome, control of blood lipid and glucose homeostasis, nonalcoholic fatty liver disease, and liver fibrosis. Here, we review recent research on the importance of sphingosine kinases, S1P, and S1PRs in liver pathobiology, with a focus on exciting insights for new therapeutic modalities that target S1P signaling axes for a variety of liver diseases.

Keywords

Metabolic syndrome; nonalcoholic fatty liver disease; sphingosine-1-phosphate; acute liver failure; steatosis

Introduction

Sphingosine, which is produced by hydrolysis of the sphingolipid core ceramide, is phosphorylated to the pluripotent bioactive signaling molecule sphingosine-1-phosphate (S1P) by sphingosine kinase 1 and 2 (SphK1 and SphK2). In addition to its role as a signaling molecule, production of S1P is required for the complete degradation of sphingoid bases and serves to reduce the level of ceramide, another signaling sphingolipid. Many cellular stimuli and hormones induce phosphorylation and activation of cytosolic SphK1 and its translocation to the plasma membrane where its substrate sphingosine resides. These

Disclosure statement

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topics have been reviewed extensively (Hannun and Obeid 2008, Maceyka *et al.* 2012, Maceyka and Spiegel 2014).

Newly produced S1P can be exported out of the cell by ATP-binding cassette (ABC) transporters or by major facilitator superfamily member spinster 2 (Spns2) (reviewed in Nishi *et al.* 2014, Takabe and Spiegel 2014) where it then can bind and activate a family of five S1P-specific G protein-coupled receptors (S1PR1–5), whose activation initiates a diverse range of cellular responses (Pyne and Pyne 2010, Maceyka and Spiegel 2014). Recently, several direct intracellular targets of S1P have been identified. For example, SphK1-produced S1P binds to and activates TNF receptor-associated factor 2 (TRAF2) that serves as a platform for recruitment and stimulation of I κ B kinase, important for the activation of the transcription factor NF- κ B (Alvarez *et al.* 2010). In contrast to SphK1, SphK2 is typically found in intracellular compartments, including the endoplasmic reticulum, mitochondria, and nucleus (Maceyka and Spiegel 2014). Nuclear produced S1P has an important role in regulation of gene expression by acting as an endogenous inhibitor of histone deacetylases (Hait *et al.* 2009).

S1P is present in high levels in the blood and has emerged as a key mediator of numerous physiological and pathophysiological responses, such as cell growth and survival, differentiation, migration, vascular integrity, lymphocyte trafficking, and immune responses, to name a few (Pyne and Pyne 2010, Orr Gandy and Obeid 2013, Maceyka and Spiegel 2014). The liver is a vital organ in the body that is critical for many functions, including detoxification of various metabolites, and in carbohydrate, protein, amino acid, and lipid metabolism. Because of its multidimensional functions, liver disease can endanger survival of the whole organism. There has been a marked increase in the rate of obesity and obesity-related diseases linked to the Western diet, and S1P is emerging as an important player in liver pathobiology. Here, we discuss the emerging role of the SphK/S1P/S1PR axis in the functions and pathobiology of the liver including acute liver failure (ALF), metabolic syndrome, control of blood lipid and glucose homeostasis, nonalcoholic fatty liver disease (NAFLD), and liver fibrosis (Figure 1 and Table 1).

Hepatic production of apoM regulates S1P plasma levels

The liver is an important organ for regulating S1P levels in the blood, which in turn play a significant role in maintaining vascular and epithelial barriers, vascular tone and inflammation. Plasma S1P is mainly contributed by erythrocytes (Pappu *et al.* 2007), platelets, endothelial cells, and liver (Yatomi *et al.* 1997, Hanel *et al.* 2007, Bode *et al.* 2010, Jonnalagadda *et al.* 2014). Erythrocytes, which are an important source of plasma S1P, can phosphorylate sphingosine to generate as well as store S1P but the mechanism of their release of S1P is not well understood. Platelets also store S1P and release it upon activation (Jonnalagadda *et al.* 2014). In human plasma, the concentrations of S1P are around 1 lM and approximately 60% of S1P is associated with HDL and 40% with albumin (Aoki *et al.* 2005, Yatomi 2008, Christoffersen *et al.* 2017). Indeed, circulating S1P levels were shown to directly correlate with apoM levels in mice and humans (Christoffersen *et al.* 2011, Kurano *et al.*

2013, Liu *et al.* 2014, Frej *et al.* 2015). Crystal structure of the apoM-S1P complex and molecular dynamics studies indicate that apoM binds S1P in an amphipathic pocket of its lipocalin domain with high affinity and suggest that S1P is not spontaneously released from apoM but requires binding to other molecules such as the S1PRs (Christoffersen *et al.* 2011, Zhang *et al.* 2016). Binding of S1P to apoM was linked to the rate of secretion of apoM from hepatocytes as mutations which resulted in its signal sequence cleavage reduced the amount of S1P associated with HDL, decreased HDL particle size, and reduced circulating S1P *in vivo* (Liu *et al.* 2015). Intriguingly, LDL receptor overexpression also reduced circulating S1P and apoM levels and the rate of clearance of exogenous S1P–apoM complexes in an apoE-dependent manner (Kurano *et al.* 2015), suggesting multiple levels of regulation of circulating S1P levels by the liver.

Many lines of evidence have linked circulating S1P to the maintenance of endothelial barrier function (Christensen et al. 2016, Yanagida and Hla 2017). Current evidence suggests the ApoM-S1P-HDL complex secreted by the liver is the endogenous mediator of endothelial barrier integrity. HDL-delivered S1P activated S1PR1 on endothelial cells in vitro while HDL from apo $M^{-/-}$ mice, which have significantly reduced circulating S1P, did not (Christoffersen et al. 2011). ApoM/HDL induced S1PR1 internalization, activation of MAPK and Akt, and increased endothelial adherens junction formation. Other studies using selective agonists and antagonists have also linked vascular barrier integrity to S1PR1 but not to other S1PRs (Ruiz *et al.* 2017). Subsequently, it was shown that apo $M^{-/-}$ mice have increased basal lung vasculature leakage that was reversed with the S1PR1 agonist SEW2871. ApoM-delivered S1P also reduced edema in a subcutaneous inflammation model in a S1PR1-dependent fashion (Christensen et al. 2016). Plasma concentrations of apoM decrease during sepsis and in systemic inflammatory response syndrome in both mice and humans, and this reduction has been linked to cytokine-induced decreased hepatocyte production of apoM mRNA and protein (Feingold et al. 2008, Kumaraswamy et al. 2012). However, in many such studies, S1P levels were not directly measured. Similarly, it was shown that levels of S1P and apoM inversely correlated with vascular leakage in humans infected with dengue (Michels et al. 2015). Moreover, the ability of ApoM/HDL to act as a biased agonist on S1PR1 inhibits vascular inflammation, which may be related to the cardiovascular protective functions of HDL (Galvani et al. 2015). Together, these results indicate that circulating apoM-S1P produced by the liver is important for the maintenance of endothelial barrier integrity.

Role of S1P in acute liver failure

Though more rare than chronic liver diseases, ALF has high mortality rates. ALF has a variety of causes, including viral infection, toxins, and ischemia, and is mediated by hepatocyte dysfunction and/or apoptosis. Recent studies have implicated S1P in certain ALF models. Curiously, although in most studies S1P seems to contribute to liver damage, in others, it seems to have a protective role. For example, S1P is important for recruitment and activation of liver-damaging immune cells, and yet several *in vitro* studies indicate that S1P protects hepatocytes from apoptotic insults (Osawa *et al.* 2005, Liu *et al.* 2013, Nowatari *et al.* 2015).

Inflammation-induced liver injury

Hyperactivation of the immune system is a common pathological inducer of acute organ damage, including the liver, and mouse models have indicated that various liver-resident immune cells such as macrophages, natural killer T cells, and dendritic cells play important roles. Given the importance of S1P in regulation of trafficking and activation of a variety of immune cells, it is not surprising that S1P signaling is involved in induction of inflammation-mediated liver damage. In viral-induced ALF in rabbits, expression of SphK1, S1P levels, and S1PR1 are all increased in the liver, and correlated with TNF-α and IL-6 levels and disease severity (Crespo et al. 2016). In agreement, treatment with the S1PR1 functional antagonists FTY720 or KRP203, both of which induce lymphopenia, was protective in a concanavalin-A-mediated, T cell-dependent liver damage model (Kaneko et al. 2006). Similarly, in lipopolysaccharide (LPS)-induced liver failure, S1P accumulated and S1PR1 expression was upregulated in the liver (Rosenberg et al. 2016). In a different ALF model, treatment with LPS plus D-galactosamine (D-Gal) induced macrophage-dependent hepatocyte cell death and liver failure. LPS/D-Gal increased serum S1P levels and upregulated SphK1 expression in peripheral blood monocytes as well as in Kupffer cells, liver resident macrophages (Lei et al. 2015b). Treatment with a pan SphK inhibitor, N,Ndimethylsphingosine, reduced mortality and was associated with reduced liver inflammation and cell death, reduced serum markers of liver failure, AST and ALT, and serum levels of the cytokines TNF-α, IL-1, and IL-6. Further work implicated PKCδ in mediating the effects of SphK, as inhibition of SphK reduced the activation of PKC δ , and inhibition of PKC δ with Rottlerin also protected from LPS/D-Gal-induced ALF (Lei et al. 2015a). Another group extended these findings, demonstrating that LPS/D-Gal-induced ALF could be ameliorated with a more specific SphK1 inhibitor (Tian et al. 2016). Intriguingly, they showed that hepatic SphK1, S1PR1, and S1PR3 levels were all elevated in ALF, and that inhibitors of any one of these proteins reduced serum levels of the pro-inflammatory cytokines IL-6 and TNF-a. However, only inhibition of SphK1, but not S1PR1 or S1PR3, protected the liver by significantly reducing hepatic hemorrhage and caspase-3 activation as well as serum AST and ALT (Tian et al. 2016). An explanation for the discrepancy is that S1P can signal either through both S1PR1 and S1PR3 to induce ALF, or that SphK1 may promote ALF directly in the liver in a cytokine-independent manner. Another study also reported that LPS/D-Gal upregulated hepatic and PBMC SphK1 and that inhibition of the complement factor C5a receptor (C5aR), which is upregulated in liver and PBMCs during ALF, reduced SphK1 expression, serum TNF-a, IL-6, and IL-1β, liver damage, and mortality (Lei et al. 2016).

Ischemia/reperfusion-induced liver injury

Acute liver damage can also be induced by ischemia and reperfusion (I/R), as is often seen after liver transplantation. One group showed that activation of S1PR1, but not S1PR2 or S1PR3, protected the liver from I/R damage induced by restriction of blood flow *in situ*, reducing apoptosis and cytokine secretion and increasing vascular integrity in a MAPK- and Akt-dependent manner (Park *et al.* 2010). However, others demonstrated that S1P was increased in both I/R and transplanted livers. Treatment with a weak SphK2 inhibitor reduced S1P levels and improved survival and liver function (Shi *et al.* 2012). The SphK2 inhibitor also reduced activation of NF- κ B and production of TNF- α and IL-1, which was accompanied by a marked reduction in the infiltration of monocytes, neutrophils, and CD4 +

T cells. These results suggest that the ischemic liver produces S1P that initiates secretion of cytokines and recruitment of inflammatory cells leading to organ damage and failure. Different effects of S1P noted above may be related to whether S1P is increased in the circulation (Park *et al.* 2010) or generated in the liver itself (Shi *et al.* 2012). It also cannot be excluded that the SphK2 inhibitor has additional targets. These studies highlight the need for analysis of S1P levels both in circulation and in the organ itself during I/R as well as thorough determination of effects of inhibitors and specificity of S1PR agonists/antagonists used.

S1P is an important mediator of liver fibrosis

Chronic liver injury is a serious problem due to the associated apoptosis, inflammation and fibrosis. The latter is the most concerning aspect as it leads to cirrhosis, liver failure and ultimately requires liver transplantation (Bataller and Brenner 2005). Fibrosis is induced when quiescent hepatic stellate cells are induced to transdifferentiate to myofibroblasts, the cells responsible for laying down the fibrotic extracellular matrix (Pellicoro *et al.* 2014). It was suggested that the myofibroblasts, which express a-smooth muscle actin and secrete collagen I to form the fibrotic extracellular matrix, are dependent on the S1P axis (Sato *et al.* 2016). However, liver samples taken during resection for hepatocellular carcinoma showed no detectable differences in S1P levels between normal and fibrotic liver (Sato *et al.* 2016). Because SPNS2, a transporter of S1P, seemed to be elevated, it was suggested that S1P is transported out of hepatocytes and signals through S1PR2 in an autocrine loop that promotes fibrosis (Sato *et al.* 2016).

In contrast, another study found that S1P levels measured by HPLC were elevated in fibrotic human livers, as well as myofibroblast expression of S1PR1 and S1PR3, but not S1PR2 (Li et al. 2011). Migration of human myofibroblasts in vitro was S1PR1/3-dependent but independent of S1PR2 (Li et al. 2011). In subsequent studies, this group used a carbon tetrachloride-induced model of liver fibrosis in mice, and demonstrated that S1PR3 promoted recruitment of bone marrow mesenchymal stem cells to the liver that then differentiated into myofibroblasts. Moreover, prevention of the upregulation of S1PR3 via the RNA binding protein HuR reduced injury-induced migration of bone marrow mesenchymal stem cells to the liver (Chang et al. 2017). Furthermore, it has been suggested that S1P is synthesized in hepatocytes in response to palmitate and released into the extracellular environment leading to activation of hepatic stellate cells by binding to S1PR3 (Al Fadel et al. 2016). In contrast, others reported that the S1PR2 antagonist JTE-013 blocked liver fibrosis (Wang et al. 2015). Similar results were observed in a bile duct ligation-induced model of liver fibrosis in which knockout of S1PR2 was protective (Wang et al. 2017). Disruption of the hepatocyte-endothelium interaction in the injured liver frequently results in impaired regeneration and fibrosis. A recent study has shown that activation of endothelial S1PR1 by S1P bound to HDL enhances regeneration and inhibits fibrosis in the liver (Ding et al. 2016). Together, these studies suggest that there are multiple roles for S1P signaling through different S1PRs at various stages of liver fibrosis, which at times may be opposing. For example, while S1P signaling may increase fibrosis through recruitment and activation of hepatic stellate cells, S1PR1 signaling in the endothelium promotes proper neo-vascularization of damaged liver and prevention of further liver

damage (Ding *et al.* 2016). Further studies are needed to determine at which stages S1P plays a role in fibrosis or inflammation and whether altered S1P metabolism in fibrotic liver could be a therapeutic target.

S1P signaling in viral hepatitis

Hepatitis B virus (HBV) and hepatitis C virus (HCV) induce liver fibrosis in chronically infected patients and are two of the most common hepatic infections in the world (Rehermann and Nascimbeni 2005, Perz et al. 2006). Both viruses have an outer membrane derived from cellular membranes, and like most membrane viruses, sphingolipids are a crucial constituent of these viruses (Sakamoto et al. 2005, Grammatikos et al. 2015, Schneider-Schaulies and Schneider-Schaulies 2015). Indeed, myriocin, an inhibitor of SPT, the first and committed step in *de novo* sphingolipid biosynthesis, suppressed replication of HCV in cultured cells and in animals (Umehara et al. 2006). Yet, not much is known about the role of S1P or SphKs in HBV and HCV infections. Chronic infection with HBV contributes to development of HCC. The HBV X protein, a multifunctional regulatory protein that plays an important role in this process, upregulates SphK1 through the transcription factor AP2a, promoting proliferation of hepatoma cells (Lu et al. 2015). However, in contrast to SphK1 that promotes viral replication, SphK2 inhibits it (Yamane et al. 2014). These opposing effects were linked to differential effects on lipid peroxidation, with SphK2 promoted-lipid peroxidation reducing viral yields (Yamane et al. 2014). In HCV patients, serum S1P levels were reduced relative to HBV patients as measured by mass spectrometry (Grammatikos et al. 2015). Moreover, increased serum levels of the SphK substrates sphingosine and sphinganine correlated with liver fibrosis progression and poor treatment outcome in HCV but not in HBV infection (Grammatikos et al. 2015). This study suggests that sphingosine and sphinganine may be biomarkers for the progression of HCV infections and further studies are needed to evaluate their usefulness for noninvasive prediction of liver fibrosis.

Role of S1P signaling in fatty liver disease

NAFLD is a prevailing health concern in the USA and is a spectrum disorder affecting the liver due primarily to overconsumption of fat-enriched food. Initially, accumulation of lipids leads to a relatively benign state of steatosis. However, over time, accumulation of lipids induces liver inflammation, termed steatohepatitis (NASH), characterized by increased immune cell infiltration followed by fibrosis. Further disease exacerbation leads to liver cirrhosis and ultimately hepatocellular carcinoma. Moreover, during disease progression, other metabolic issues typically arise, including insulin resistance and elevated triglycerides (Cohen and Chun 2011, Satapathy and Sanyal 2015).

High-fat diets (HFDs) are typically characterized by elevated levels of palmitate, the primary precursor of ceramide and other sphingolipids. Indeed, palmitate has been shown to induce sphingolipid synthesis in hepatocytes (Watt *et al.* 2012) and numerous reports have documented the relationship between a HFD, increased ceramide levels, and NAFLD. Elegant reviews have highlighted the numerous metabolic complications mediated by elevated ceramide that are associated with NAFLD (Chavez and Summers 2012, Pagadala *et*

al. 2012). It has been suggested that insulin resistance in obesity is associated with increases of pro-inflammatory cytokines TNF-a and IL-6 that can induce ceramide production in the liver. Ceramide, in turn, activates PP2A and attenuates Akt signaling, inducing apoptosis, causing hepatic insulin resistance (Chavez and Summers 2012). Ceramide also contributes to the development of NAFLD and progression to NASH (Pagadala *et al.* 2012). However, much less is known of the role of the ceramide metabolites sphingosine and S1P in NAFLD and metabolic syndrome.

Lipidomic analyzes of tissues from obese mice that were fed for 16-52 weeks on a HFD (42% kcal fat) provided some insights into how HFD affects sphingosine and S1P levels. As expected, these mice had elevated liver levels of palmitate-containing sphingolipids as well as sphingosine and S1P compared to normal chow fed mice. Moreover, these mice also had elevated levels of sphinganine and sphinganine-1-phosphate, suggesting that the increased dietary fat was inducing *de novo* sphingolipid synthesis (Sanyal and Pacana 2015). Consistent with these results, adding palmitic acid to primary mouse hepatocytes increased SphK1 expression, raised S1P levels, and increased lipid accumulation. Consistent with this, it was observed that SphK1 expression is increased in livers from humans with NAFLD and also in mice with high saturated fat feeding, which recapitulated the human disease (Geng et al. 2015). From these initial findings, it seems likely that SphK1 and formation of S1P play a role in regulation of lipid storage. Additional in vitro studies support the notion that SphK1 is a pro-steatosis gene. For example, SphK1 expression is closely linked to expression of the lipid storage gene PPARc (Ross et al. 2013, Geng et al. 2015). Further support came from the demonstration that weight gain and triglyceride levels in mice fed a 21% kcal fat diet enriched with fructose for 24 weeks were lower in SphK1 knockout mice compared to wild type mice. Importantly, hepatic pathology was less severe in the SphK1^{-/-} mice as determined by the NAFLD Activity Score (Chen et al. 2016a). However, the role of SphK1 in NAFLD is not limited to steatosis (Figure 2). Mouse and rat models of NAFLD have shown that SphK1 is important in hepatic inflammation and progression to steatohepatitis (Wang et al. 2013, Geng et al. 2015). After 18 weeks on a high saturated fat diet, there were elevated numbers of macrophages recruited to the liver that correlated with increased inflammatory cytokines such as TNF-a and MCP1 (Geng et al. 2015). Moreover, although SphK1 knockout mice gained weight to a similar extent as their wild type littermates, they had reduced liver lipid accumulation, cytokine signaling, and macrophage accumulation. In *vitro*, the saturated fat palmitate induced hepatocyte expression of SphK1, TNF- α , and MCP1, and this effect could be blocked by knockdown of either SphK1 or S1PR1. These data suggest that saturated fats upregulate SphK1 and S1P that signals through S1PRs to promote inflammatory signaling in the liver (Geng et al. 2015). This may have clinical relevance as patients with nonalcoholic steatohepatitis have elevated liver levels of SphK1. Elevation of liver SphK1 and S1PRs was also observed in a rat model of steatohepatitis induced by a high fructose diet (Wang et al. 2013). Similarly, in a rat model of high fatinduced NAFLD, inhibition of sphingolipid synthesis with myriocin decreased ceramide and sphinganine and concomitantly reduced signs of NAFLD, including reduced liver enzymes (Kurek et al. 2014). Myriocin-treated animals on the high fat diet also had reduced liver fat accumulation and reduced immune cell infiltration (Chavez and Summers 2012).

Deposition of fat and free fatty acids in the liver can cause lipotoxicity involved in the development and progression of fatty liver diseases. SphK1 expression suppressed ER stress-mediated pro-apoptotic pathways and prevented IRE1a activation and JNK phosphorylation (Qi et al. 2015). Thus, this study provides further evidence for a role of SphK1 in hepatocyte survival and uncovers a novel mechanism of protection against ER stress-mediated cell death (Qi et al. 2015). SphK2 has been less well studied but also appears to have a role in NAFLD. SphK2 was upregulated in murine liver after administration of the ER-stress inducing agent tunicamycin (Lee et al. 2015). Lentiviral overexpression of SphK2 in liver of mice led to increased phosphorylation of Akt in liver, muscle and adipose tissue. In addition, mice overexpressing SphK2 demonstrated improved glucose tolerance while on a 60% kcal fat diet compared to wild type mice. Moreover, fatty acid oxidation was increased in these mice that were accompanied by elevated S1P levels (Lee et al. 2015). Similarly, it was found that in contrast to SphK1 knockout mice, SphK2 knockout mice rapidly developed steatosis on a HFD (Nagahashi et al. 2015). The increased liver fat accumulation was linked in this study to the role of SphK2 in promoting transcription by producing S1P in the nucleus that inhibited HDAC activity, by showing that SphK2 knockout mice had reduced expression of a number of genes involved in hepatic lipid metabolism. Intriguingly, it was also found that S1PR2 knockout mice had a similar phenotype as the SphK2 knockouts that was suggested to be due to the inability of the S1PR2 mice to upregulate SphK2 in response to a HFD (Nagahashi et al. 2015).

Additional insight into the roles SphKs and S1P in NAFLD has come from studies using FTY720, a prodrug that is phosphorylated mainly by SphK2 to produce FTY720-phosphate that is a S1P mimetic that is a ligand of all of the S1PRs except S1PR2 (Brinkmann *et al.* 2010). FTY720-phosphate also acts as a functional antagonist for S1PR1 as it induces its degradation. Many studies have demonstrated that this effect on S1PR1 interferes with and ultimately mitigates immune responses (Brinkmann *et al.* 2010). In a murine model of NAFLD, administration of FTY720 reduced liver/body weight ratio along with decreased hepatic inflammation, fibrosis and hepatocyte ballooning (Mauer *et al.* 2017). Perhaps most importantly, FTY720 treatment was able to reverse NAFLD symptoms even after they had already been initiated, suggesting that FTY720 may be a useful therapeutic for these liver disorders.

Role of liver S1P signaling in metabolic syndrome

Though it has many causes, NAFLD is commonly associated with metabolic syndrome, the name given to a collection of symptoms, including obesity, insulin resistance, dyslipidemia, and hypertension, that predispose individuals to type 2 diabetes, cancer, and cardiovascular diseases. Metabolic syndrome is a serious health issue, affecting nearly 1 in 4 US adults (Beltran-Sanchez *et al.* 2013), and involves multiple organs, including the pancreas, skeletal muscle, adipocytes, and the immune system. Because the role of S1P in these various organs and metabolic diseases has been recently reviewed (Brice and Cowart 2011, Chen *et al.* 2016b), this section will focus specifically on the role of S1P signaling in the liver in metabolic syndrome. The liver plays a very important function in maintaining systemic glucose levels by producing glucose during periods of glucose deprivation (Saltiel and Kahn 2001). Adenoviral delivery of SphK1 to the liver greatly reduced blood glucose levels and

dyslipidemia in the KK/Ay mouse model of type 2 diabetes (Ma et al. 2007). The protective effect of SphK1 was linked to its enhancement of insulin-stimulated kinases Akt and GSK3β. Although hepatic SphK1 was downregulated in mice fed a high-fat high-sucrose diet, overexpression of SphK1 in liver reduced triglyceride content in mice fed a low but not HFD and had no effect on glucose tolerance (Kowalski et al. 2015). Conversely, in hepatocytes, palmitate was shown to induce S1P production that opposed insulin signaling (Fayyaz et al. 2014). In addition, S1P was elevated in the livers of New Zealand obese mice, and S1PR2 inhibition reversed the insulin resistance in these mice. Interestingly, glucose stimulated SphK2 and increased S1P in pancreatic islets that correlated with increased glucose-stimulated insulin secretion (Cantrell Stanford et al. 2012). Moreover, treatment of mice with a SphK inhibitor impaired glucose disposal due to decreased plasma insulin levels, suggesting that S1P may be involved in glucose stimulated insulin secretion from beta cells (Cantrell Stanford et al. 2012). Hepatic overexpression of SphK2 in mice fed a HFD also increased S1P, and also decreased hepatic accumulation of lipid droplets by a SphK2-dependent increase in fatty acid oxidation in liver. In addition, glucose intolerance and insulin resistance were markedly reduced by improved hepatic insulin signaling through SphK2 upregulation (Lee et al. 2015). Because the ER stress-mediated UPR pathways upregulate SphK2 expression, it was suggested that this ameliorates hepatic steatosis and insulin resistance in mice (Lee et al. 2015).

Another area of research that has recently received much attention is related to the effects of adiponectin, which controls glucose and lipid homeostasis, on sphingolipid metabolism. A series of studies have shown that binding of adiponectin to its receptors AdipoR1 and AdipoR2, led to enhanced deacylation of ceramide to sphingosine and increased conversion to S1P. This conversion promoted survival of functional beta cells, allowing for sufficient insulin production to meet insulin demands. Alleviation of ceramide accumulation in the liver also led to improvements in hepatic insulin action (Holland *et al.* 2011, 2013).

Concluding remarks

S1P, the enzymes that control its levels through synthesis and degradation, and the multiple receptors that respond to it have important roles in numerous physiological hepatic functions and are critically involved in hepatic pathologies, particularly NAFLD and HCC. Thus, to mitigate their pathological contributions, development of targeted pharmacological interventions is being actively pursued. However, utilization of these approaches will require significant preliminary studies as S1P has pleiotropic effects depending on the receptor targeted or even the cell types in which S1PRs are expressed. Given the critical roles for S1P reviewed here, it is hoped that further pharmacological studies will not only provide useful therapeutics but will also aid in elucidating the mechanisms for the actions of S1P in different hepatic pathologies.

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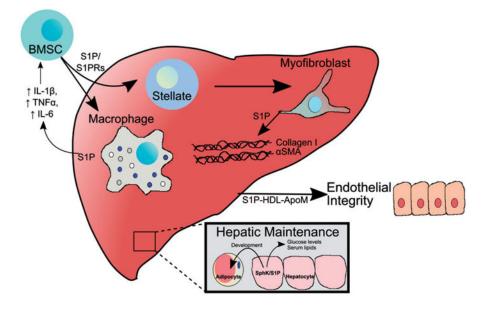


Figure 1.

S1P signaling in the liver is involved in multiple physiological and pathophysiological processes. This figure depicts several processes regulated by S1P described in the text. S1P promotes recruitment of bone marrow mesenchymal stem cells (BMSC) through activation of S1PRs. In the liver, these cells differentiate into hepatic stellate cells. When quiescent stellate cells are activated due to liver injury, they transdifferentiate to myofibroblasts which express α-smooth muscle actin and secrete collagen I to form the fibrotic extracellular matrix in a S1P-dependent manner. The liver also secretes S1P-containing apoM and S1P/ apoM in HDL promotes endothelial barrier integrity. S1P also regulates hepatic maintenance of serum lipids, glucose levels, and adipocyte development. S1P released from the liver in response to injury promotes recruitment of a variety of immune cells, including liver-resident macrophages known as Kupffer cells leading to hepatic inflammation (see color version of this figure at www.tandfonline.com/ibmg).

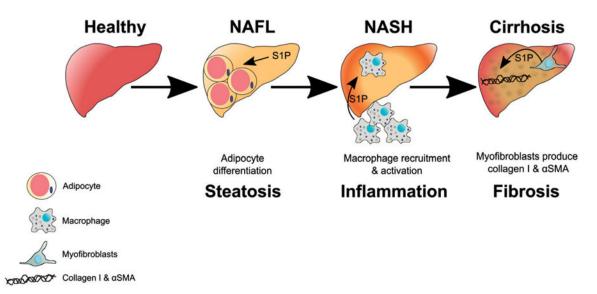


Figure 2.

S1P regulates progression of nonalcoholic fatty liver disease (NAFLD). This figure illustrates potential roles of S1P in the progression of NAFLD from nonalcoholic fatty liver (NAFL) to nonalcoholic steatohepatitis (NASH) and finally to cirrhosis. Initially S1P increases adipocyte differentiation and accumulation of lipids in the liver. S1P produced in the liver increases pro-inflammatory cytokines such as IL-1 β , TNFa, and IL-6, which recruit and activate macrophages to the liver. Lastly, S1P stimulates myofibroblasts to secrete collagen I and a smooth muscle actin (aSMA) (see color version of this figure at www.tandfonline.com/ibmg).

Table 1

Liver pathophysiology associated with the sphingosine-1-phosphate axis.

Pathophysiology	Effects of sphingosine-1-phosphate
Glucose tolerance	Systemic glucose •↑ SIP levels associated with ↑ glucose (Fox et al. 2011, Kowalski et al. 2013) Liver •↑ Ceramide leads to ↑ Akt signaling and ↑ glucose tolerance (Osawa et al. 2011) •↑ SphK2 leads to ↑ hepatic glucose tolerance (Lee et al. 2015) •↑ SIP associates with ↑ insulin resistance (Fayyaz et al. 2014) Pancreas •↑ glucose associated with ↑ SIP and ↑ insulin secretion (Cantrell Stanford et al. 2012)
Nonalcoholic fatty liver disease (NAFLD)	 Liver SIP caused ↑ in NF-κB, IL-1B, IL-6, and TNF-α induced inflammation (Wang <i>et al.</i> 2013) ↓ Loss of SphK1 associated with ↓ weight and ↓ lipid accumulation (Geng <i>et al.</i> 2015, Chen <i>et al.</i> 2016a) ↑ in palmitate causes ↑ lipid and triglyceride accumulation (Chen <i>et al.</i> 2016a) ↑ SphK2 expression improved weight, lipid levels, and ↑ glucose tolerance (Lee <i>et al.</i> 2015, Nagahashi <i>et al.</i> 2015) ↓ SphK2 correlated with ↑ hepatosteatosis along with systemic cholesterol and triglycerides (Nagahashi <i>et al.</i> 2015) ↑ ER stress leads to ↑ SphK2 and ↑ S1P with improved glucose tolerance and ↓ hepatosteatosis (Lee <i>et al.</i> 2015)
Obesity	Circulation •↑ S1P levels correlated with ↑ body fat percentage and BMI (Kowalski <i>et al.</i> 2013) Adipocytes •↓ S1PR2/S1P signaling associated with ↓ adipose tissue differentiation (Jeong <i>et al.</i> 2015) and ↓ lipid accumulation (Moon <i>et al.</i> 2014)
Hepatic fibrosis and liver injury	 Fibrogenesis ↑ SIP inhibits growth of human hepatic fibroblasts (Davaille <i>et al.</i> 2000). ↑ SIP induces apoptosis of human hepatic myofibroblasts in receptor independent manner (Davaille <i>et al.</i> 2002) ↑ SIP signals through Edg receptor to also activate Akt, ERK1/2 for survival in human hepatic fibroblasts (Davaille <i>et al.</i> 2002). • ↓ SIP plasma concentration associated with progression of CCl4 induced liver fibrosis (Ikeda <i>et al.</i> 2010) • ↑ SIP endas to Col a1 (I) and Col a1 (III) fibrosis development (Xiu <i>et al.</i> 2015) • ↑ SIP enhanced human hepatic myofibroblasts (Li <i>et al.</i> 2011). • SphK1 involved with hepatic fibrosis angiogenesis (Yang <i>et al.</i> 2013) <i>Migration</i> • SIPR2/3 activate Akt for bone marrow derived macrophage/monocyte migration (Yang <i>et al.</i> 2015) <i>Liver injury</i> • SIPR2 has inhibitory role in liver regeneration after acute liver injury (Ikeda <i>et al.</i> 2009)