



Review

# Sphingolipids in Atherosclerosis: Chimeras in Structure and Function

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**Abstract:** Atherosclerosis—a systemic inflammatory disease—is the number one cause of mortality and morbidity worldwide. As such, the prevention of disease progression is of global interest in order to reduce annual deaths at a significant scale. Atherosclerosis is characterized by plaque formation in the arteries, resulting in vascular events such as ischemic stroke or myocardial infarction. A better understanding of the underlying pathophysiological processes at the cellular and molecular level is indispensable to identify novel therapeutic targets that may alleviate disease initiation or progression. Sphingolipids—a lipid class named after the chimeric creature sphinx—are considered to play a critical and, metaphorically, equally chimeric regulatory role in atherogenesis. Previous studies identified six common sphingolipids, namely dihydroceramide (DhCer), ceramide (Cer), sphingosine-1-phosphate (S1P), sphingomyelin (SM), lactosylceramide (LacCer), and glucosylceramide (GluCer) in carotid plaques, and demonstrated their potential as inducers of plaque inflammation. In this review, we point out their specific roles in atherosclerosis by focusing on different cell types, carrier molecules, enzymes, and receptors involved in atherogenesis. Whereas we assume mainly atheroprotective effects for GluCer and LacCer, the sphingolipids DhCer, Cer, SM and S1P mediate chimeric functions. Initial studies demonstrate the successful use of interventions in the sphingolipid pathway to prevent atherosclerosis. However, as atherosclerosis is a multifactorial disease with a variety of underlying cellular processes, it is imperative for future research to emphasize the circumstances in which sphingolipids exert protective or progressive functions and to evaluate their therapeutic benefits in a spatiotemporal manner.

**Keywords:** cardiovascular disease; atherosclerosis; sphingolipids; ceramide; sphingosine-1-phosphate; dihydroceramide; lactosylceramide; glucosylceramide; sphingomyelin



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## 1. Introduction

The enigmatic character of sphingolipids has been first highlighted by assigning their name to a new class of lipids first described in 1884 by the German physician and biochemist J. L. W. Thudichum [1]. In the 1880s, he found “sphingosine” with unique chemical characteristics, which directed him to name this brain-derived lipid after the Sphinx, a mythical creature with a human head and a lion's body. This iconic name became formative for the substance class of sphingolipids, but also adequately reflects the chimeric role of sphingolipids in the etiology of atherosclerosis.

Cardiovascular diseases (CVDs) are the leading cause of mortality, accounting for 17.9 million deaths per year worldwide [2,3]. Atherosclerotic cardiovascular disease is a progressive and lifestyle-dependent condition characterized by arterial lesions characterized by local oxidative stress and inflammation that initiate vasoconstriction, reduced

and/or turbulent flow, plaque formation, and/or hemostasis. These responses in combination with excessive plasma low-density lipoprotein (LDL) cholesterol levels, e.g., caused by poor dietary quality or sedentarism, lead to lipid deposition and atheromatous plaque formation resulting in functional and, ultimately, structural disintegrity of the arterial vessel wall [4–6]. This in turn triggers primary and secondary hemostasis that in combination with locally altered fluid mechanics is causally responsible for embolic complications, resulting in, e.g., myocardial infarction or ischemic strokes [7–9]. Therefore, a comprehensive understanding of the molecular mechanisms of disease initiation and progression is indispensable for the identification of possible therapeutic targets providing the spotlight for current atherosclerosis research.

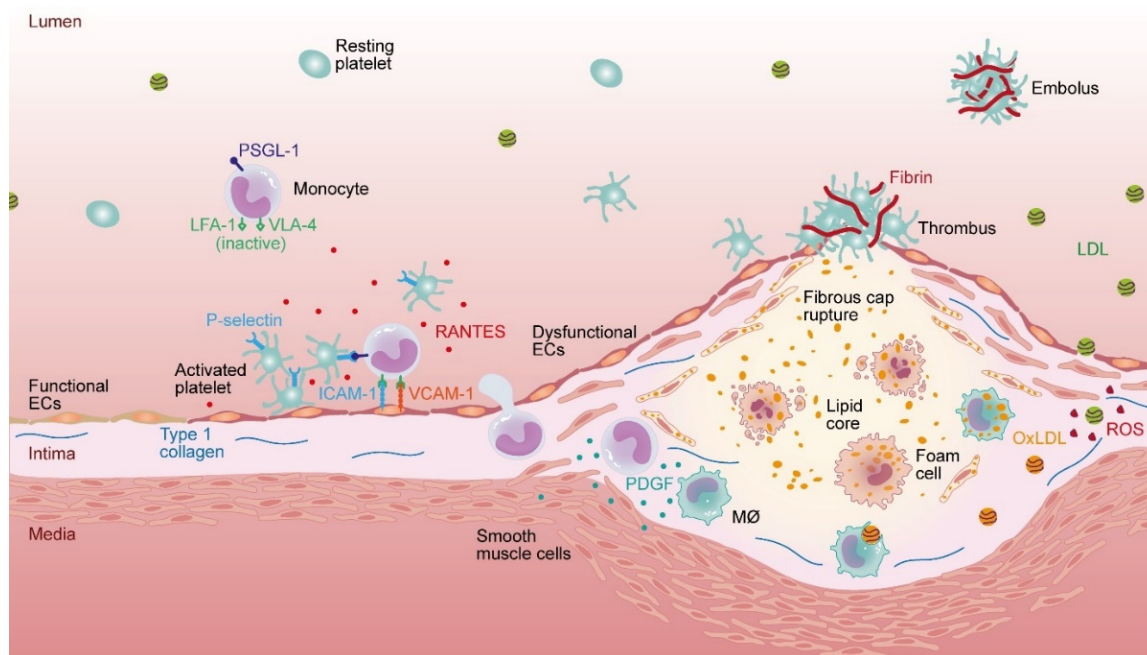
Atherosclerosis represents a subtype of arteriosclerosis. Arteriosclerosis is the most common form of adverse vascular remodeling and is usually limited to small arteries and arterioles. This vascular remodeling comprises thickening and hardening of the arterial vessel wall, resulting in an increase in vascular stiffness and a reduction of blood flow to the tissues. All subtypes of arteriosclerosis have these processes in common, and sub-classification is solely based on the cause and localization of the vascular remodeling. Atherosclerosis—the focus of this review—describes adverse vascular remodeling stemming specifically from the formation of atherosclerotic plaques in the arteries. This plaque formation leads to thickening of the vessel wall, subsequent calcification further reduces wall compliance, and both processes together increase arterial stiffness [10,11].

Mechanistically, the pathogenesis of atherosclerosis comprises a diversity of cell types and molecules (Figure 1). Atherogenesis is exacerbated by various environmental risk factors such as cigarette smoking and hypercaloric diet or by preexisting conditions such as hypercholesterolemia, hyperglycemia, or hypertension [12–14]. Underlying these risk factors is the presence of oxidative stress and, consequently, endothelial dysfunction. NO is continuously produced and released by endothelial NO synthase (eNOS) in endothelial cells at baseline. NO primarily exhibits anti-inflammatory and antithrombotic functions such as attenuation of platelet adhesion, aggregation and leukocyte adhesion [15,16]. In a stable endothelium, protective NO and harmful ROS are in balance since ROS and NO react to peroxynitrate [17]. In the progression of atherosclerosis or diabetes, eNOS produces ROS, e.g., superoxide anion instead of NO; a process also known as “eNOS uncoupling” [18]. The eNOS uncoupling further enhances superoxide anion production [19] and activation of NAD(P)H oxidase [20], which, in turn, represents a major source of the superoxide anion [20–22]. The subsequent imbalance toward ROS results in endothelial dysfunction, which culminates in increased vascular permeability [23–25] and extravasation of LDL into the intima [26–28]. Furthermore, endothelial dysfunction fosters platelet adhesion to von Willebrand factor (vWF) and consequential platelet activation by the release of paracrine mediators, such as adenosine diphosphate (ADP) and thromboxane (TxA<sub>2</sub>) [29]. Activated platelets also secrete the chemokine RANTES (CCL5) that immobilizes on the surface of inflamed microvascular or aortic endothelium and allows for shear-resistant monocyte arrest under flow conditions [30].

Parallel endothelial dysfunction is associated with the expression of cell adhesion molecules (CAMs), i.e., ICAM-1 and VCAM-1, on vascular endothelial cells, and subsequent tethering, rolling and adhesion of monocytes on the endothelium—a hallmark of atherogenesis [31–33]. The accompanying morphological change allows monocytes to transmigrate across the endothelium into the intima in a process called diapedesis. Upon activation, monocytes become synthesizers of ROS, i.e., superoxide, hydroxyl radicals, and peroxy radicals that support protein degradation and DNA oxidation, but, most importantly, lipid peroxidation, which is a hallmark of chronic inflammatory diseases including atherosclerosis [34–37]. In this milieu, ROS can oxidize native non-atherogenic LDL to oxidized low-density lipoprotein (oxLDL). These aid in the activation of monocytes through scavenger receptor pathways, which in turn mature to macrophages and, subsequently, cholesterol rich foam cells. The differentiation of monocytes into macrophages is a multi-step process initiated by the recruitment of monocytes to the lesion site accompanied by the

secretion of granulocyte–macrophage colony-stimulating factor (GM-CSF) and macrophage colony-stimulating factor (M-CSF). These factors, in turn, drive the proliferation of intimal cells in the early phase of atherosclerosis [38] and promote advanced plaque progression by increasing macrophage apoptosis susceptibility [39]. The altered transcriptional program in the activated monocytes promotes macrophage maturation [40,41]. Specifically, by expression of atherogenic scavenger receptors including CD36, macrophages become enabled to internalize oxidatively modified proteins such as oxLDL. This oxLDL uptake by CD36 promotes macrophage differentiation and foam cell formation as illustrated by the fact that apolipoprotein E (ApoE)-deficient animals that lack expression of CD36 show a marked reduction in atherosclerotic lesions as compared to ApoE-deficient mice expressing CD36 [42,43]. Due to the prevailing oxidative stress by oxLDL, smooth muscle cells (SMC) express scavenger receptors and take up oxLDL, resulting in the formation of foam cells [44–48]. Concomitantly, macrophages proliferate in the intima and amplify the maladaptive inflammatory process through the release of cytokines and matrix metalloproteinases (MMPs), which can degrade the arterial extracellular matrix and promote further differentiation of macrophages into foam cells following uptake of oxLDL [42]. Cytokines from activated macrophages and endothelial cells result in the release of platelet-derived growth factor (PDGF), which, in turn, stimulates the migration of vascular SMCs from the media into the intima and support their proliferation [49,50]. Ultimately, foam cells derived from SMCs and macrophages die through both necrotic and apoptotic processes, thereby releasing their contents [51], and in this way, attract further macrophages. Secreted oxLDL molecules and dying foam cells accumulate in a necrotic core, a condensation site for further cellular debris of apoptotic macrophages and SMCs, which is surrounded by an endothelial layer and migrated SMCs. As the necrotic core progresses, calcium deposits further establish the atherosclerotic plaque that thins its fibrous cap along maturation and eventually becomes vulnerable to rupture [52]. When this luminal surface of the plaque is disrupted, the highly thrombogenic core is exposed, which ultimately leads to primary hemostasis. Locally impaired release of, e.g., the tissue factor pathway inhibitor, thrombomodulin or reduced expression of the endothelial protein C receptor on dysfunctional endothelial cells at the site of plaque rupture, further supports thrombus formation and prompts vessel stenosis, complete occlusion, and/or embolism [53–57].

A potential relationship between sphingolipids and atherosclerosis was first described by Smith in 1960 [58]. She reported that in the area of advanced lesions, human aortas present a higher proportion of lipids in the intima and media of the vessel wall. Specifically, sphingomyelin (SM) is increased in the intima of lesions sites compared to areas with less advanced lesions [58]. Sphingomyelin was found to account for 70–80% of all phospholipids in the necrotic core, indicating a potential pathophysiological role of sphingolipids in atherosclerosis—an observation that has been confirmed since then on several occasions [59–62]. Beyond SM, the presence of dihydroceramides (DhCer), ceramides (Cer), lactosylceramides (LacCer), glucosylceramides (GluCer), and sphingosine-1-phosphates (S1P) was subsequently identified as a common sphingolipid signature of carotid plaques [59]. In this review, we provide an overview of the disease modulating anti- and pro-atherogenic functions of each of these sphingolipids and discuss open aspects of the mechanistic pathophysiological relationship of these sphingolipids in the onset and progression of atherosclerosis.

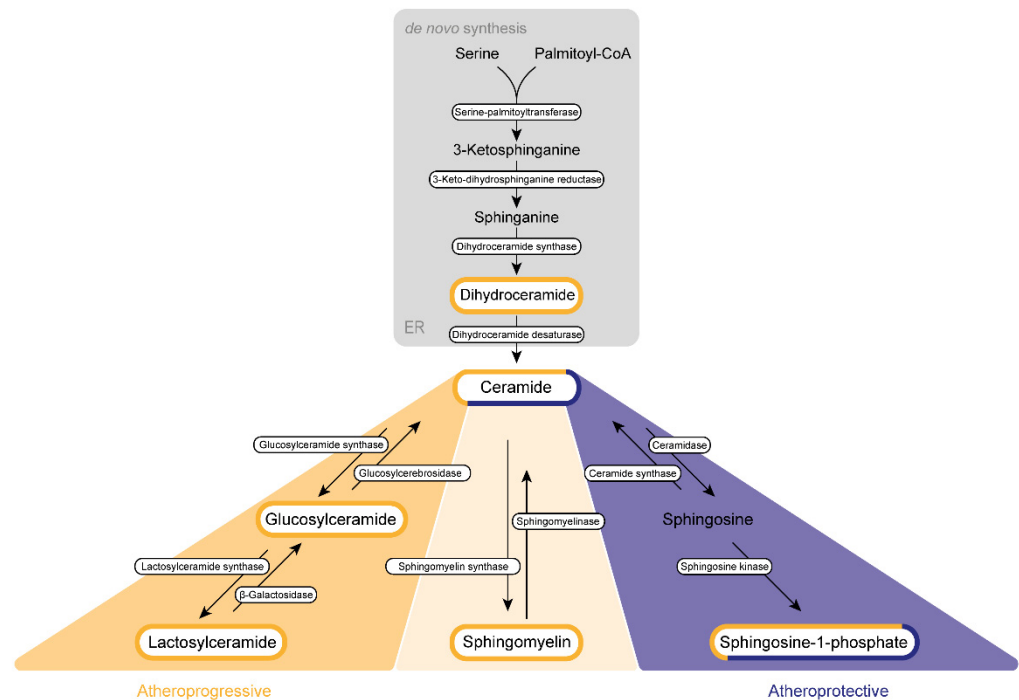


**Figure 1.** Cellular pathomechanisms of atherosclerosis and progression. Environmental risk factors such as cigarette smoking and hypercaloric diet or preexisting conditions such as hypercholesterolemia, hyperglycemia or hypertension promote endothelial dysfunction and increase vascular permeability and retention of LDL in the vascular intima. Endothelial dysfunction further promotes platelet adhesion through the release of von Willebrand factor (vWF) and platelet activation by mediators such as adenosine diphosphate (ADP) and thromboxane (TxA<sub>2</sub>). Activated platelets secrete the chemokine RANTES (CCL5), which enables monocytes to adhere under flow conditions. The adhesion is further promoted by cellular adhesion molecules (CAM) expressed by activated endothelial cells. The lymphocyte function-associated antigen 1 (LFA-1) on the surface of monocytes enables their binding to intercellular adhesion molecule 1 (ICAM-1) expressed by endothelial cells. This cellular interaction is strengthened by monocytic integrin  $\alpha 4\beta 1$  (VLA-4) binding to vascular cell adhesion molecule 1 (VCAM-1), further mediating lateral migration and transendothelial diapedesis of monocytes into the intima. Intimal LDL is oxidized by ROS to oxidized LDL (oxLDL), which aids in the recruitment of monocytes and initiates differentiation into macrophages by scavenger receptor mediated uptake of oxLDL. Activated macrophages secrete platelet-derived growth factor (PDGF), which stimulates smooth muscle cells (SMCs) to migrate into the intima where they proliferate and produce extracellular matrix and again incorporate oxLDL. Uptake of oxLDL by SMC and macrophages leads to their differentiation into foam cells, which degrade and, in turn, release oxLDL. This self-amplifying process further attracts macrophages and SMCs that accumulate oxLDL and dying cells—the necrotic core of the atheromatous plaque. This process is accompanied by thickening of the intima limiting blood flow through the lumen and results in weakening of the fibrous cap of the vulnerable plaque. As the disease progresses, the vascular lumen becomes gradually occluded, leading to turbulent blood flow, which supports endothelial dysfunction, the expression of CAMs, and the formation of vascular lesions. Increasing instability culminates in plaque rupture and subsequent thrombus formation.

## 2. Dihydroceramide in Atherosclerosis Progression

### 2.1. Synthesis and Metabolism

The *de novo* synthesis of sphingolipid is initiated by a highly coordinated sequence of actions involving serine palmitoyltransferase, 3-keto-dihydrosphingosine reductase, and dihydroceramide synthase, which convert cytosolic serine and palmitoyl CoA molecules via sphinganine into DhCer (Figure 2).



**Figure 2.** Sphingolipid biogenesis in atherosclerosis. Sphingolipids are synthesized de novo in the endoplasmic reticulum (ER) and the Golgi apparatus. Subsequently, they are transported via vesicles to the plasma membrane and the endosomes. The amino acid serine and palmitoyl-CoA provide the basis for the synthesis of 3-keto-sphinganine, which is reduced to sphinganine via 3-keto-dihydrosphinganine reductase. The dihydroceramide synthases form dihydroceramide, which can be catalyzed to ceramide, the backbone of all sphingolipids, by dihydroceramide desaturase. Ceramide itself can be converted into three further sphingolipid species. Glucosylceramide synthase mediates the production of glucosylceramide, which can be further modified to lactosylceramide through the enzyme lactosylceramide synthase. This modification can be reversed by  $\beta$ -galactosidase and glucosylcerebrosidase, respectively. Ceramide also provides the backbone for the generation of sphingomyelin via the activity of sphingomyelin synthase. Sphingosine-1-phosphate can be synthesized by ceramidase and sphingosine kinase. Several sphingolipids shown are assumed to exert influence on the progression of atherosclerosis. This impact can be categorized either as atherogenic (yellow) or as protective (purple) or can display characteristics of both categories (mixed).

DhCer is further processed at the endoplasmic reticulum (ER) membrane. Here, DhCer serves as a substrate for dihydroceramide desaturase that introduces a 4,5-trans-double bond to the sphingolipid backbone, thus generating Cer, which is further catalyzed by ceramidase and sphingosine kinases to first sphingosine and then S1P in the Golgi apparatus. Similar to most sphingolipids, DhCer is elevated in atherosclerotic plaques and is associated with inflammation and plaque instability [59].

## 2.2. Regulation of Inflammation

For a long time, no specific cellular function was attributed to DhCer, yet this notion has changed over the past 15 years, as DhCer was shown to impact autophagy, cell proliferation, cell survival and cell death in cancer and metabolic diseases [63–67]. In atheromatous plaques, DhCer levels positively correlate with proinflammatory cytokines such as monocyte chemoattractant protein-1, interleukin 6 (IL-6), and macrophage inflammatory protein-1  $\beta$ . Over and above that, DhCer is able to induce the release of IL-6 in human coronary smooth muscle cells without inducing apoptosis [59]. However, caution is warranted in the interpretation of experimental results focusing on the specific function of DhCer, as pharmacological or genetic inhibition of enzymes involved in the de novo pathway will not only affect DhCer levels but also Cer concentration [68].

### 2.3. Regulation of Autophagy

In line with a potential functional role of DhCer in inflammatory processes *per se*, DhCer has been found to promote autophagy as demonstrated by the formation of autophagosomes in prostate cancer cells after stimulation with a DhCer desaturase inhibitor [69]. Of note, similar results were obtained by exogenous addition of short-chain DhCer [69]. Similarly, exogenous addition of DhCer analogues or treatment with DhCer desaturase inhibitors led to the accumulation of DhCer and promoted autophagy in cancer cells without causing cell death [64,65]. While a mechanistic link between DhCer and autophagy has thus been established, it remains a matter of controversy whether autophagy has a protective or a progressive effect on atherosclerosis. Normal autophagy flux is involved in vascular homeostasis, yet abnormal activity results in mechanisms aggravating atherosclerosis such as inducing thrombosis in endothelial cells, the secretion of pro-inflammatory cytokines by macrophages and abnormal remodeling of SMC in the intima. These characteristics can finally cause cell death and plaque instability [70]. Since short-chain DhCer can favor the formation of autophagosomes, it is appealing to hypothesize that short-chain DhCer also promotes autophagy in a pathophysiological context that may drive the progression of atherosclerosis. Moreover, the influence of DhCer on atherosclerosis promoting as well as atheroprotective mechanisms appears not to be restricted to autophagy only. DhCer has also been proposed to diminish apoptosis by inhibiting the formation of pores on the outer mitochondrial membrane, thereby impeding an essential step of the apoptotic cascade [71]. It remains to be evaluated whether and how this effect of DhCer on apoptosis influences atherosclerosis progression. In addition, DhCer affects oxidative stress by inducing ER stress. In contrast, DhCer levels are also elevated in the presence of oxidative stress, which can be explained by the inhibition of DhCer desaturase [72,73]. To investigate which effect provides the initiator for the other, further research is needed.

## 3. Ceramide

The hydrophobic properties of ceramides restrict their solubility in an aqueous environment. Ceramides in plasma are therefore either bound to carrier proteins such as lipid transfer proteins or are associated with lipoproteins such as LDL and high-density lipoprotein (HDL). Cer provides the acyl-backbone for other sphingolipids such as S1P, GluCer, LacCer and SM. Besides the *de novo* pathway, the most physiologically relevant means of Cer synthesis is the acyl-CoA-dependent conversion of sphingosine and non-esterified fatty acids by the activity of a family of six ceramide synthases (CerS1-6) [74,75] into ceramides with distinct acyl chain lengths. Alternatively, ceramides can be metabolized by sphingomyelinases (SMases)-induced hydrolysis of sphingomyelin to Cer.

Importantly, Cer concentrations correlate with the risk for cardiovascular disease (CVD) in general and atherosclerosis specifically; as such, Cer qualifies as a prognostic marker for CVD as well as for sphingomyelin (SM) [76–78]. Since Cer is present in significantly enriched amounts in atherosclerotic plaques and has been shown to be correlated with aggregated [79] and circulating LDL [80], a causal relationship between Cer and atherosclerotic plaque progression has been assumed.

### 3.1. Sphingomyelinases (SMases)

It seems that an athero-promoting effect of Cer is mediated by specific types of SMases, e.g., Cer can be hydrolyzed from multiple SMases such as secreted lysosomal (L-SMase), acidic sphingomyelinase (A-SMase) and membrane neutral SMase (N-SMase). L-SMase and A-SMase are located in the endosome but can be translocated to the outer plasma membrane under certain conditions [81,82]. N-SMase, however, is synthesized predominantly in the ER and Golgi apparatus, but also in the inner leaflet of the plasma membrane. All three forms of SMase have been implicated in atheroprotection in distinct manners. High density lipoprotein (HDL) is one out of five major lipoproteins that transports lipid molecules within the body. HDL is usually referred to as “good cholesterol”, as it captures

lipid molecules in the artery walls and thereby prevents atheroprogession [83,84]. HDL molecules mainly consist of apolipoprotein A (ApoA) and further apolipoprotein C (ApoC). The main function of ApoC-1 protein is the inhibition of cholesterol ester transfer protein (CETP) and inhibiting the lipoprotein binding to the “bad cholesterol” high density lipoprotein (HDL) and very low density lipoprotein (VLDL). Mutations reducing the function of CETP have thereby been associated with elevated atherosclerosis progression [85]. This pathophysiological mechanism seems to be of crucial role in terms of the involvement of Cer in atherogenesis, since ApoC-1-enriched HDL induces apoptosis and cell death of vascular smooth muscle cells (VSMC) via N-SMase activation [86]. Furthermore, oxLDL induces proliferation of VSMC via N-SMase [87,88]. Since both apoptosis and proliferation of VSMC are mechanisms associated with atherogenesis, these findings may suggest an atheroprogessive effect of N-SMase activation. Similar pro-atherogenic effects have been described for A-SMase. Endothelial cells secrete A-SMase, which hydrolyses SM on the surface of atherogenic lipoproteins to Cer and thus mediates the fusion, aggregation and affinity of lipoprotein particles with/at/toward the endothelium of arteries [89]. Analyses of *ApoE<sup>-/-</sup>/Ldlr<sup>-/-</sup>/Smpd1<sup>-/-</sup>* triple knockout mice highlighted the impact of A-SMase on atherogenesis, since the absence of A-SMase reduced the formation of atherosclerotic lesions and arterial trapping of atherogenic lipoproteins in the otherwise atheroprone *ApoE<sup>-/-</sup>/Ldlr<sup>-/-</sup>* mice [90]. Similar to A-SMase, L-SMase has also been found to promote the pathogenesis of atherosclerosis. As a result of ligand binding to TNF receptors, activation and translocation of L-SMase proceeds. Grassme et al. identified a mechanism by which L-SMase seems to enhance atherosclerosis [91] through the formation of Cer-enriched domains. These domains are formed by receptor-mediated translocation of L-SMase. L-SMase is primarily localized in the endolysosomal compartment and can be relocated to the outer leaflet of the plasma membrane upon stimulation via CD95 receptor [92–94]. Due to this translocation, sphingolipid-rich domains accumulate and release extracellularly orientated Cer. Accumulation of Cer leads to the formation of Cer-enriched platforms on the surface, which, in turn, efficiently initiate apoptosis signaling by trapping and clustering the receptors. The aggravating effect on atherosclerosis is postulated since the Cer-enriched membrane domains in VSMC and EC impair the vasodilatory properties in ECs and VSMC [95,96] and enhance muscarinic-1 receptor-mediated constriction of coronary arteries [97].

Overall, the three types of SMases have been implicated at several levels in the progression of atherosclerosis. However, as these studies have been typically performed in different models without back-to-back comparisons of the role of different SMases, it remains to be shown whether the individual roles of L-SMase vs. A-SMase or N-SMase in atherogenesis are specific or redundant.

### 3.2. Regulation by Matrix Metalloproteinases (MMPs)

Activation of the oxLDL-induced SM/Cer pathway and subsequent activation of ERK1/2 is regulated by MMPs, a large family of zinc proteases [98]. In principle, MMP content is increased in atheromatous plaques and has been associated with plaque instability and the formation of stenotic lesions that recur after treatment [99]. The expression of these MMPs is regulated and activated by major triggers of vascular remodeling such as inflammation or oxidative stress [100]. In SMC, the connection between MMPs and atherogenesis is considered to be mediated by oxLDL-induced activation of N-SMase, in that inhibition of MMP-2 inhibits N-SMase and as such, Cer production. Conversely, exogenous MMP-2 activates the SM/Cer pathway, supporting the notion of an oxLDL-induced activation of the Cer pathway via activation of N-SMase [98]. However, the exact mechanism by which oxLDL activates SMases via MMPs is currently unclear and remains the scope for future research. These findings highlight, thus far, the atheroprogessive functions of Cer and related mediators as SMases and MMPs. As we will discuss in the next paragraph, inflammatory mediators may exert an additional influence on the effects of sphingolipids on cellular mechanisms such as apoptosis or vasodilation.

### 3.3. Regulation by Tumor Necrosis Factor Alpha (TNF $\alpha$ )

Tumor necrosis factor alpha (TNF $\alpha$ ) is likely a central factor that further increases Cer concentrations in atherosclerotic lesions [101,102]. TNF $\alpha$  contributes to endothelial dysfunction by stimulating ROS production and induces the expression of various inflammatory cytokines and chemokines [103–106]. Acting on the vascular endothelium, TNF $\alpha$  thus emerges as a key driver for the progression of atherosclerosis. Linking TNF $\alpha$  to sphingolipids, Sawada et al. proposed a TNF $\alpha$ -induced increase in Cer levels in human glioma cells via two different pathways, both of which are initiated by activation of caspase-8: first, a p53 and ROS-dependent pathway that leads to N-SMase activation via GSH depletion and thus to increased production of Cer; a second pathway activates A-SMase directly via caspase-8, and, thus, causes a ROS-independent increase in Cer levels resulting in a TNF $\alpha$ -induced apoptosis of human glioma cells [107]. Analogously, clinical studies have shown that the ischemic myocardium is stimulated by inflammatory cytokines such as TNF $\alpha$ , interleukin 2 and endostatin, similarly resulting in an A-SMase- and N-SMase-dependent elevation of Cer levels [108,109]. However, the effect of TNF $\alpha$  on Cer production is not unidirectional. TNF $\alpha$  can also be induced by stimulating human umbilical vein endothelial cells with C2-Cer [110]. It may thus be inferred that TNF $\alpha$  not only stimulates Cer production, but conversely, Cer synthesis also stimulates TNF $\alpha$  release—thus establishing a pathological feedback loop. This notion is in line with studies showing that anti-TNF $\alpha$  therapy is able to improve endothelial function in humans with vascular inflammation [111,112]. Nevertheless, it remains to be shown whether anti-TNF $\alpha$  treatment may reduce vascular ceramide production and attenuate CVD and atherosclerosis. Of interest, changes in amino acid metabolism may also affect Cer de novo synthesis, as homocysteine leads to increased formation of superoxide anions by stimulation of the NADPH oxidase pathway [113]. In agreement with this hypothesis, ceramide levels increase in response to rising homocysteine concentrations via the de novo synthesis pathway rather than the SMase pathway, as treatment with myriocin (a highly selective serine palmitoyltransferase inhibitor) reduced homocysteine-induced ceramide production in rats [114]. In summary, ceramide is produced by two independent synthesis pathways: (i) SMase-dependent hydrolysis from sphingomyelin and (ii) de novo synthesis via ceramide synthase, both of which are assumed to be stimulated in atherosclerosis in general and by inflammatory cytokines such as TNF $\alpha$  specifically.

In conclusion, Cer has been demonstrated to be detrimental in atherosclerosis as (i) being enriched in atherosclerotic plaques, (ii) SMases being involved in formation of aortic lesions and processes involved in atherogenesis such as apoptosis or lipoprotein trapping and (iii) Cer levels being elevated in response to MMPs and TNF $\alpha$ —which are also elevated in atherosclerotic lesions—via SMase activation. The underlying mechanisms of action are probably diverse, only partially elucidated, and will be discussed in the following sections.

### 3.4. Regulation of NO Production

Under physiological conditions, vascular NO production is stimulated by shear stress, catalyzed by the endothelial NO synthase (eNOS), and constitutes an essential feature of endothelial cell function and vascular homeostasis. Reduced NO release or impaired NO bioavailability are key factors in the progression of endothelial dysfunction, manifested by loss of endothelium-dependent vasorelaxation. Cer is considered an important negative regulator of endothelial NO production, as it decreases the release of NO from human umbilical vein endothelial cells [115,116] and initiates the production of superoxide anions [117–119]. As such, Cer may promote endothelial dysfunction by decreasing NO and increasing ROS production, and thus promote the development of atherosclerosis.

### 3.5. Regulation of LDL Aggregation

Another essential role of ceramide in the development of atherosclerosis is the ceramide-induced aggregation of LDL. Increased levels of Cer correlate with the ability of LDL to form aggregates [120–123]. During atherogenesis, LDL is enriched at the vessel membrane where it is exposed to SMase. OxLDL activates SMase to convert LDL-SM to Cer



within atherosclerotic lesions [75,122]. Cer, in turn, enables a conformational change in apolipoprotein B100 (ApoB100), which provides the essential step for LDL molecules to aggregate [124–126]. This process is further accompanied by macrophage-mediated phagocytosis and foam cell formation, aggravating atherosclerotic lesion formation [92,127,128]. In line with this concept, the use of the sphingolipid synthesis inhibitor myriocin prevents aggregation of LDL and succeeds in a reduction of plaque formation [120,127].

In addition to its ability to promote oxidative stress and to enhance LDL aggregation, Cer causes apoptosis and necrosis in human coronary artery smooth muscle cells *in vitro* [59], which further accentuate its pro-atherosclerotic function.

More recent findings have also taken into account a more differentiated view on the distinct role of certain molecular species of ceramide. Long chain (C11–C20), very long chain (C21–C24) and ultra-long chain (>C24) ceramide species are formed in the sphingolipid synthesis pathway by six different Cer synthases (CerS1–6) with specific affinities for the chain length of the fatty acyl-CoA. Deletion or pharmacologic inhibition on N-SMase2 in the ApoE<sup>-/-</sup> mouse model reduced atherosclerotic lesions and decreased macrophage infiltration and lipid deposition via small interfering RNAs in the nuclear factor erythroid 2-related factor 2 pathway [129]. This species-dependent effect on the biological activities of Cer was underscored by overexpression of CerS4 and CerS6, which generate long chain Cer to inhibit cell proliferation while inducing apoptosis, respectively. CerS2, in turn, forms very long chain Cer that increases cell proliferation [128,130]. This highlights the importance of the activity of specific CerS and subsequent changes in Cer species composition in the initiation and progression of atherosclerosis and remains a point of consideration in the understanding of the pathophysiology of CVD.

#### 4. Sphingosine-1-Phosphate

The cleavage of fatty acids from the sphingolipid backbone of Cer by ceramidases releases sphingosine, which can be further phosphorylated by the activation of sphingosine kinase isoenzymes 1 and 2 (Sphk1, Sphk2) to sphingosine-1-phosphate [131]. Sphk1 and Sphk2 are highly conserved and present in most mammalian cells and tissues, including platelets [132], erythrocytes [133], and the endothelium itself [134] which secrete S1P by the specific S1P-transporters major facilitator superfamily domain containing 2B (MFSD2B, erythrocytes and platelets) and spinster-homologue-2 (SPNS2, endothelial cells) into plasma and lymph [135–138]. Here, S1P signals as a bioactive lipid mediator by targeting five different G protein-coupled S1P-receptors (S1PR1–5) on various hematopoietic and vascular cells, and thereby controls cellular proliferation, apoptosis and cell migration in the blood vasculature and interstitial spaces and regulates endothelial barrier function [139]. Therefore, S1P/S1PR signaling may infer a significant role in the pathogenesis of atherosclerotic cardiovascular disease. Serum S1P is a strong and robust predictor of the occurrence of obstructive coronary artery disease [140], suggesting a correlation with atherogenic effects. Furthermore, the S1PR modulator FTY720, which acts upon all S1PRs except S1PR2 [141], effectively attenuates atherogenesis in ApoE<sup>-</sup> and LDL-receptor (LDL-R) deficient mice, respectively [142,143], implicating an atheroprotective effect. Future research should further confirm these contradictory initial findings.

To realize signaling in health and disease, S1P has to bind to chaperone proteins including apolipoprotein M (ApoM) on HDL (~65% of all free plasma S1P) or albumin (~30% of all free plasma S1P) and LDL or VLDL (<5% of all plasma S1P), as its hydrophobic backbone and polar phosphate head group restrict the membrane permeability of S1P [144–146]. The plasma S1P levels also closely correlate to levels of total cholesterol, LDL cholesterol and HDL cholesterol in normolipidemic healthy subjects [147,148]. These associations may be of mutual functional relevance, e.g., the interaction of S1P and HDL has been proposed to reinforce their anti-thrombotic, anti-inflammatory and antioxidant properties [149]. The S1P/cholesterol interrelation has been experimentally validated by gain-of-function mutations of the LDL-R in livers of mice, which reduced S1P and ApoM levels in wildtype but not in ApoE-deficient mice. This finding suggests ApoE-dependent

clearance of ApoM-associated S1P [150]. In line with this notion, statin treatment reduced serum ApoM levels in type 2 diabetes mellitus patients [151]. Further, only the S1P/ApoM complex on HDL is able to activate endothelial S1PR<sub>1</sub> Gi-signaling and downstream ERK- and Akt-signaling, preserving endothelial adherent junctions [145] and decreasing TNF $\alpha$ -induced activation of nuclear factor kappa B (NF $\kappa$ B) and expression of ICAM-1 [152], while this endothelium-protective signaling cascade is only insufficiently activated by the S1P/albumin complex [153].

The majority of receptor-associated actions of S1PR-mediated intracellular processes are atheroprotective. Evidence from *in vivo* experiments shows that S1PR<sub>1</sub> and S1PR<sub>3</sub> are essential for both maintenance of endothelial barrier function, as the receptors' downstream signaling cascade stabilizes endothelial cell–cell junctions [154] and attenuates endothelial contraction [155], and vascular relaxation by phosphorylation of eNOS and subsequently increased endothelial NO release [156–158]. In addition, a protective function against the development of atherosclerotic lesions has been suggested, as expression of the adhesion molecules VCAM-1 and ICAM-1 can be inhibited by S1PR<sub>1</sub> signaling, thus reducing leukocyte adhesion and subsequent extravasation [157,158]. Analogously, S1P signaling via S1PR<sub>3</sub> can inhibit the recruitment of inflammatory neutrophils and suppress apoptosis of cardiomyocytes. S1PR<sub>3</sub>-deficient mice are accordingly more susceptible for infarction in a mouse model of myocardial ischemia/reperfusion as compared to their corresponding wild type [159]. In contrast to this anti-inflammatory role of S1PR<sub>3</sub> signaling, however, S1PR<sub>3</sub> deficiency in ApoE<sup>-/-</sup> mice was found to strongly reduce monocyte recruitment by decreasing monocyte chemoattractant protein-1 secretion without affecting the size of atherogenic lesions [160]. These pro-inflammatory and, hence, potentially atherogenic properties of S1P signaling are further supported by the finding that S1PR<sub>1</sub> enhances chemotaxis of lymphocytes and natural killer cells (NK) and, thus, has pro-inflammatory and pro-atherosclerotic properties [153]. S1P signaling through S1PR<sub>2</sub> is even more likely to be associated with atherogenic functions. Although S1PR<sub>2</sub> has been shown to inhibit SMC migration [161], it is centrally involved in the recruitment of inflammatory macrophages [162]. As such, S1PR<sub>2</sub><sup>-/-</sup>/ApoE<sup>-/-</sup>-double-deficient mice show reduced release of IL-18 and IL-1 $\beta$ , leading to impaired interstitial macrophage recruitment and, consequently, reduced formation of atherosclerotic plaques and necrotic cores in comparison to S1PR<sub>2</sub>-proficient mice [163]. Consistently, S1PR<sub>2</sub>-deficient macrophages express less CD36 and scavenger receptors *ex vivo* and increase cholesterol efflux while decreasing oxLDL uptake [163]. Atherogenic effects of S1PR<sub>2</sub> signaling have also been suggested based on the fact that S1P can impair endothelial barrier function via the S1PR<sub>2</sub>/Rho/ROCK pathway [164]. However, S1PR<sub>2</sub> deficiency in mice is associated with an increased risk of seizures and the development of B-cell lymphomas, arguing against the suitability of this receptor as a therapeutic target in atherosclerosis [165–167]. S1PR<sub>4</sub>, expressed on leukocytes, NK cells and airway SMC [168], and S1PR<sub>5</sub> expressed on NK cells and oligodendrocytes [169] have not been associated with atheroprotection to date, even though S1PR<sub>4</sub> stimulates IL-10 secretion from T-cells and simultaneously inhibits interleukin 4 and interferon- $\gamma$  production [170], while S1PR<sub>5</sub> mobilizes NK cells during infections. Although these findings may suggest an indirect involvement of S1PR<sub>4</sub> and S1PR<sub>5</sub> in atherosclerosis-associated inflammation, the latter receptors seem to have less of a direct impact on atherosclerosis pathology.

## 5. Sphingomyelin (SM)

SM is the most abundant sphingolipid in mammalian tissues, where it serves as an important structural component of cell and plasma membranes [171]. Importantly, in the context of atherosclerosis, SM is also involved in maintaining cholesterol homeostasis, as addition of exogenous SM to cells increases cholesterol biosynthesis and affects LDL binding to cell surface receptors [172,173]. However, there is further evidence implicating SM in the pathogenesis of atherosclerosis. SM has been identified as one component of human atherosclerotic plaques, and its abundance correlates with histological markers of plaque instability and is associated with the expression of pro-inflammatory cytokines. In

accordance with this observation, stimulation of human coronary smooth muscle cells with SM *in vitro* induces a pro-inflammatory response reflected by IL-6 release [59]. SM plasma levels of atherosclerotic ApoE<sup>-/-</sup> mice are also elevated in comparison to WT mice [174]. Likewise, rabbits with hypercholesterolemia show elevated levels of SM compared with other lipids in atherosclerotic lesions [175]. Similar to S1P, SM in plasma is associated with VLDL/HDL cholesterol (63–75%) and LDL cholesterol (25–35%). The emerging notion that elevated SM levels in plasma are associated with pro-atherogenic properties is further supported by the fact that a decrease in HDL SM content is associated with smaller and more dense HDL. These complex lipoprotein particles favor cholesterol efflux, anti-oxidative activity toward LDL oxidation, antithrombotic activity in human platelets, as well as anti-inflammatory and anti-apoptotic activity [176]. In accordance, anti-apoptotic and anti-oxidative activities of small compact HDL cholesterol have been associated with SM degradation [177].

Unlike SMase, which hydrolyzes SM to Cer, the sphingomyelin synthase (SMS) catalyzes the synthesis of SM from Cer. SMS represents a family of different isoforms: SMS1 is primarily localized in the Golgi apparatus, whereas SMS2 primarily in plasma membranes [178,179].

Inhibition of SMS1 has been proposed as a potential therapeutic approach in atherosclerosis, as SMS1<sup>-/-</sup> mice show a decreased atherosclerotic phenotype characterized by reduced atherosclerotic lesions in the entire aortas as well as decreased macrophage content in these lesions [180]. Similar effects have been achieved in SMS2-deficient mice. These mice are marked by a reduction in secretion of pro-inflammatory cytokines, which is accompanied by the reduction of atherosclerotic lesions, necrotic core formation, macrophage content and collagen content compared to wild-type mice [181]. The pro-atherogenic capabilities of SM are further confirmed, as adenovirus-mediated insertion of SMS2 in ApoE<sup>-/-</sup> mice results in an increase in atherosclerotic lesions [182]. Similarly, SMS2 is shown to act as a modulator of NF-κB activation in HEK193 cells and macrophages from SMS2-deficient mice. This could provide one mechanistic explanation of the pro-atherogenic function of SM [183]. Consistent with this pro-atherogenic character of SM, overexpression of SMS1 and SMS2 increases the lipoprotein atherogenic potential in mice [184], whereas the simultaneous deficiency of SMS1 and SMS2 leads to a reduction in plasma SM and pro-inflammatory cytokine secretion [180]. In this context, it is remarkable that the inhibition of SMS1 alone leads to a decrease in the SM content in plasma, but simultaneously to an increase in DhCer and Cer in the plasma. Considering those two being associated with both atheroprotective and atherogenic effects, an explicit categorization of SM as an atheroprotective should only be made with caution. Further, it will be crucial to determine the mechanistic interplay between the inhibition of SMS1 and the increase in DhCer and Cer in order to identify a definite therapeutic signaling cascade. With regard to the identification of potential novel therapeutic targets, it is furthermore relevant to consider that loss-of-function by deletion of SMS1 (similar to S1PR2, *vide supra*) entails serious side effects such as low-frequency hearing loss [179,185], impaired insulin secretion [186], or CD4<sup>+</sup> cell dysfunction [187].

## 6. LacCer and GluCer—Sphingolipids with Non-Chimeric Functions?

Lactosylceramide synthase (LacCerS) generates LacCer by transferring galactose from uridine diphosphate-galactose to GluCer. LacCer and GluCer are classified as glycosphingolipids whose synthesis can be inhibited by D-threo-1-phenyl-2-decanoylamino-3-morpholino-1-propanol (D-PDMP). D-PDMP is an analog of glucosylceramide originally synthesized to inhibit glucosylceramide synthase in patients with Gaucher's disease [188,189]. However, D-PDMP has also been shown to be directly capable of LacCerS [188]. These inhibitory effects of D-PDMP of LacCer and GluCer synthases have been utilized to identify the involvement of these sphingolipids in terms of atherogenesis. Thereby, it was shown that LacCer and GluCer increase vascular dysfunction, since aortic wall thickening, presence of Ca<sup>2+</sup> deposits and vascular stiffness were decreased upon blockade of glycosphingolipid synthesis in ApoE<sup>-/-</sup> mice [189].

This finding is strengthened by previous studies showing that LacCer exerts an influence on many mechanisms relevant to atherosclerosis. For example, LacCer mediates TNF-induced NF- $\kappa$ B expression and ICAM-1 expression in endothelial cells by activation of a redox-dependent transcriptional pathway [190,191]. In the same manner, it has atheroprogresive effects by stimulating the expression of MAC1 on monocytes or neutrophils, presumably facilitating their adhesion to endothelial cells and initiating atherosclerosis [189]. Based on the results of various studies, Chatterjee and colleagues postulated the following pathway to mechanistically explain LacCer-induced atherosclerosis progression: OxLDL increases the production of endogenous LacCer, which, in turn, activates NADPH oxidase [118,191–193]. The resulting production of superoxide [190,194,195] induces GTP loading of P21ras and thus activation of a kinase cascade from Raf-2, Mek2 and p44MAPK. Phosphorylation of p44MAPK results in a local shift of p44MAPK from the cytoplasm to the nucleus [118,191]. This step determines the final expression of c-fos, proliferating nuclear antigen and cell proliferation.

GluCer has previously been implicated in arterial stiffness and vascular cell wall thickening [189] but, in addition, appears to have a direct impact on atherosclerotic plaque development, as inhibition of glucosylceramide synthase attenuates atherosclerotic plaque development and the expression of inflammatory genes [196]. These glycosylceramide synthase-associated effects were found even more pronounced in ApoE\*leiden mice, in which pharmacological inhibition of glucosylceramide synthase also led to a drastic reduction of atherosclerotic plaques. This effect was accompanied by a decreased cholesterol level in the liver and an increased excretion of cholesterol by feces and an increased secretion of bile [196]. The effects could also be replicated in LDL receptor KO mice. In vitro, glucosylceramide per se initiates apoptosis in HCASMC and induces an inflammatory response, evident as increased expression of IL-6, MCP- and macrophage inflammatory protein-1 $\beta$  [59]. In view of these reported functions, LacCer and GluCer seem to exhibit primarily pro-atherogenic effects.

## 7. Conclusions

The large quantity of sphingolipids identified in atherosclerotic plaques supports a possible link between sphingolipids and atherosclerosis. The key question, however, that remains to be clarified is whether sphingolipids are the cause or the consequence of atherogenesis. Various studies have demonstrated a specific effect of sphingolipids on cellular processes relevant to the development of atherosclerosis, such as impaired NO production [109,115–119], apoptosis [59,71,86,107], plaque development [70,79,80,120,127,196], or LDL aggregation [59,80,125]. As a function of their cellular and tissue context or their respective sphingoid bases, however, the sphingolipids Cer, DhCer, GluCer, LacCer, SM and S1P can exert chimeric and often opposing functions in the pathogenesis and progression of atherosclerosis (Table 1). While the existing evidence reviewed herein suggests that SM, DhCer, LacCer and GluCer exclusively mediate atheroprogresive effects, Cer and S1P may exert both protective as well as progressive properties in atherosclerosis (Table 1). As such, S1P mediates anti-apoptotic [139,159] and anti-inflammatory [162,170] processes as well as enhancing vasoconstriction [149,155] while maintaining endothelial barrier function [145,154] (Table 1). In contrast, its pro-atherogenic functions are evident in its ability to activate lymphocytes [153] and to promote primary hemostasis and thrombus formation [163] (Table 1). Furthermore, it should be considered that the synthetic pathway of sphingolipids is intertwined, and enzymes that can synthesize multiple species of a sphingolipid class with unique properties mediate the generation of one from another. Very long chain Cer is pro-thrombotic, induces cell proliferation and TNF $\alpha$  secretion, and correlates with LDL aggregation, whereas long chain Cer inhibits proliferation and induces apoptosis [128,130]—highlighting the chimeric properties of different Cer species depending on their sphingoid bases (Table 1). This opposing mode of action within the same class of sphingolipids is influenced by their biosynthesis, as for, e.g., Cer, the activity of CerS1-6 results in the generation of Cer with a distinct chain length, which has unique progressive or protective functions on atherogenesis [128,130]. Furthermore, sphingolipid receptors can essentially determine whether the mediated effect is atheroprotective or

atherogenic, as exemplified by the differential expression and upstream signaling facilitated by S1PR1-5 [139,141–143]. In addition, an increasingly recognized level of regulation is the bioavailability of S1P mediated by its specific carrier molecules. While ApoM-associated S1P mediates atheroprotective effects, S1P bound to albumin can mediate either atheroprotective or atheroprotective effects [145,150,152].

**Table 1.** Sphingolipids and their associated mechanism in atherogenesis. Sphingolipids exhibit molecular mechanisms, which are either categorized as atheroprotective or atheroprotective.

Sphingolipid	Associated Mechanism	Effect on Atherosclerosis	References
Dihydroceramide	<ul style="list-style-type: none"> <li>↑ Autophagy</li> <li>↑ Oxidative stress</li> <li>↑ Inflammatory cytokines</li> <li>↑ Cell proliferation</li> <li>↑ Plaque instability</li> </ul>	Progressive	[59,63–67,69,71–73]
Cer	<ul style="list-style-type: none"> <li>↑ Inflammation</li> <li>↑ Proliferation</li> <li>↑ LDL-Aggregation</li> </ul>	Progressive	[28–30,32,59,76,90,100–102,128,130]
Long-chain	<ul style="list-style-type: none"> <li>↓ Cell proliferation</li> <li>↑ Apoptosis</li> </ul>	Protective	
Very long-chain	<ul style="list-style-type: none"> <li>↑ Cell proliferation</li> </ul>	Progressive	
S1P S1PR <sub>1</sub>	<ul style="list-style-type: none"> <li>↑ Endothelial barrier function</li> </ul>	Protective	[59,139,141,143,145,149,150,152,153,156–158,160–164]
S1PR <sub>2</sub>	<ul style="list-style-type: none"> <li>↓ Apoptosis</li> <li>↑ Chemotaxis of lymphocytes and NK cells</li> <li>↓ ICAM1 and VCAM1 expression</li> </ul>	Progressive Progressive	
	<ul style="list-style-type: none"> <li>↓ Barrier function</li> <li>↑ Recruitment of inflam. macrophages</li> <li>↑ Plaque and necrotic core formation</li> <li>↓ SMC migration</li> </ul>	Progressive Protective	
S1PR <sub>3</sub>	<ul style="list-style-type: none"> <li>↑ Endothelial barrier function</li> <li>↑ Monocyte recruitment</li> </ul>	Protective Progressive	
S1P/ApoM	<ul style="list-style-type: none"> <li>↓ Thrombus formation</li> <li>↓ Inflammation</li> <li>↓ Apoptosis</li> </ul>	Protective	
S1P/Albumin	Not shown	Protective + progressive	
Sphingomyelin	<ul style="list-style-type: none"> <li>↑ Hypercholesterolemia</li> <li>↑ Apoptosis</li> <li>↑ Inflammatory cytokines</li> <li>↑ Thrombus formation</li> <li>↑ Plaque instability</li> <li>↑ Atherosclerotic lesions</li> <li>↑ Macrophage content in lesions</li> </ul>	Progressive	[59,174–177,180–184]
Lactylceramide	<ul style="list-style-type: none"> <li>↑ TNF<math>\alpha</math>-induced NF<math>\kappa</math>B expression</li> <li>↑ ICAM-1 expression</li> <li>↑ MAC1 expression</li> <li>↑ Arterial stiffness</li> <li>↑ Aortic wall thickening</li> <li>↑ Presence of aortic Ca<sup>2+</sup> deposits</li> <li>↑ Apoptosis</li> <li>↑ Inflammatory cytokines</li> </ul>	Progressive	[59,118,189–193,197]
Glucosylceramide	<ul style="list-style-type: none"> <li>↑ Arterial stiffness</li> <li>↑ Aortic wall thickening</li> <li>↑ Presence of aortic Ca<sup>2+</sup> deposits</li> <li>↑ plaque development</li> <li>↑ cholesterol level liver</li> <li>↑ Apoptosis</li> <li>↑ Inflammatory cytokines</li> </ul>	Progressive	[1,2,5]

In a broader context, the functions of sphingolipids are determined by such a number of individual factors and steps such that one may wonder about the evolutionary purpose of this complexity. This could serve as an amplification process, so that many individual steps enhance the effect, explaining the involvement of many cell types and molecules in the context of atherosclerosis. Furthermore, this signaling network might reflect a system of mutual checks and balances, ensuring that not a single imbalance leads immediately to the formation of atherosclerosis, thus preventing atherosclerosis from developing rapidly. A deeper understanding of the complex sphingolipid network and the chimeric properties of individual sphingolipid classes and species offers new therapeutic possibilities. For example, knowledge of the biosynthesis of different species of a sphingolipid, each with chimeric functions, opens up therapeutic strategies that allow for targeted inhibition of enzymes that lead to the formation of atherosclerosis-promoting sphingolipids and, consequently, could maximize the therapeutic outcome. Taken together, this may imply that sphingolipids and their actions should be analyzed as a network rather than as individual components.

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