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1 Serial Review: Redox-Regulated Phospholipase Signal Transduction  
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3 Sphingolipid signaling and redox regulation<sup>☆</sup>

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9 **Abstract**

10 Sphingolipids including ceramide and its derivatives such as ceramide-1-phosphate, glycosyl-ceramide, and sphingosine (-1-phosphate) are  
 11 now recognized as novel intracellular signal mediators for regulation of inflammation, apoptosis, proliferation, and differentiation. One of the  
 12 important and regulated steps in these events is the generation of these sphingolipids via hydrolysis of sphingomyelin through the action of  
 13 sphingomyelinases (SMase). Several lines of evidence suggest that reactive oxygen species (ROS; O<sub>2</sub><sup>-</sup>, H<sub>2</sub>O<sub>2</sub>, and OH<sup>-</sup>), and reactive nitrogen  
 14 species (RNS; NO, and ONOO<sup>-</sup>) and cellular redox potential, which is mainly regulated by cellular glutathione (GSH), are tightly linked to the  
 15 regulation of SMase activation. On the other hand, sphingolipids are also known to play an important role in maintaining cellular redox  
 16 homeostasis through regulation of NADPH oxidase, mitochondrial integrity, and antioxidant enzymes. Therefore, this paper reviews the  
 17 relationship between cellular redox and sphingolipid metabolism and its biological significance.

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19 *Keywords:* Catalase; Ceramidase; Glycosylceramide; Mitochondria; NADPH oxidase; Nitric oxide synthase; Reactive nitrogen species; Reactive oxygen species;  
 20 Sphingolipid; Sphingomyelinase; Superoxide dismutase

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*Abbreviations:* A-SMase, acidic sphingomyelinase; BH<sub>4</sub>, (6R)-5,6,7,8-tetrahydro-L-biopterin; DTT, dithiothreitol; eNOS, endothelial nitric oxide synthase; FAN, factor associated with N-SMase activation; GalT-2, galactosyl transferase-2; GPX, glutathione peroxidase; GSH, reduced form of glutathione; iNOS, inducible nitric oxide synthase; lyso-PAF, lyso-platelet-activating factor; nNOS, neuronal nitric oxide synthase; N-SMase, neutral sphingomyelinase; NO, nitric oxide; Phox, phagocytic NADPH oxidase; PI3K, phosphatidylinositol-3'-kinase; PKC, protein kinase C; PKG, cGMP-dependent protein kinase; PPAR, peroxisome proliferator-activated receptor; RNS, reactive nitrogen species; ROS, reactive oxygen species; SM, sphingomyelin; SMase, sphingomyelinases; SOD, superoxide dismutase; TNF-R1, tumor necrosis factor-receptor I; TPX, thioredoxin peroxidase.

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40

## 41 Introduction

42 Under normal conditions, reactive oxygen species (ROS)  
 43 and reactive nitrogen species (RNS) can be generated as a by-  
 44 product of normal metabolic processes and function as  
 45 physiological signaling molecules [1,2]. However, in patholog-  
 46 ical conditions, the excessive increase in ROS, such as super  
 47 oxide anion ( $O_2^-$ ), hydrogen peroxide ( $H_2O_2$ ), and hydroxyl  
 48 radicals ( $\cdot OH$ ), and RNS, such as nitric oxide (NO) and  
 49 peroxynitrite ( $ONOO^-$ ), by mitochondrial dysfunction, activa-  
 50 tion of xanthine, and NADPH oxidases and increased gene  
 51 expression of inducible nitric oxide synthase (iNOS) can cause  
 52 cell death and tissue damage [3]. Under normal conditions, the  
 53  $O_2^-$  is scavenged by superoxide dismutase (SOD) which  
 54 specifically processes  $O_2^-$  and produces  $H_2O_2$ . The  $H_2O_2$  is in  
 55 turn detoxified by catalase and glutathione peroxidase (GPX)  
 56 because otherwise  $H_2O_2$  would react with transition metals to  
 57 generate highly toxic hydroxyl radicals through the Fenton  
 58 reaction ( $H_2O_2 + Fe^{2+} \rightarrow \cdot OH + Fe^{3+} + \cdot OH$ ). Moreover, RNS  
 59 are also able to affect cellular redox homeostasis [2], such as the  
 60 physiologically relevant action of NO in the activation of the  
 61 guanylate cyclase and the subsequent activation of cGMP-  
 62 mediated signaling cascades, whereas an excessive amount of  
 63 NO can cause cell and tissue damage [3,4]. NO can scavenge  $O_2^-$   
 64 and other free radicals and inhibit the  $O_2^-$  driven Fenton reaction  
 65 and lipid peroxidation. On the other hand, large amounts of NO  
 66 as generated by the iNOS isoform in inflammatory disease  
 67 conditions are often accompanied by a large production of ROS,  
 68 and will shift NO chemistry toward indirect effects such as  
 69 nitrosation, nitration, and oxidation [2,5]. The interaction of NO  
 70 with molecular oxygen ( $O_2$ ) or  $O_2^-$  gives rise to the formation of  
 71 the potent nitrosating agent  $N_2O_3$  and peroxynitrite ( $ONOO^-$ ),  
 72 respectively. S-Nitrosothiol adducts are formed by the interac-  
 73 tion between  $N_2O_3$  and certain protein thiol groups and evoke  
 74 signaling by altering protein kinases and phosphatases, G-  
 75 proteins, ion channels, protein tyrosine kinases, and redox-  
 76 sensitive transcription factors [5,6].

77 Sphingolipids are ubiquitous constituents of membrane  
 78 lipids in mammalian cells. Along with their structural role,  
 79 sphingolipids have received attention due to their role as second  
 80 messengers in proliferation, differentiation, apoptosis, and  
 81 inflammation [7–12]. In mammalian cells, the majority of  
 82 sphingolipids are colocalized with cholesterol in specific  
 83 membrane domains called “lipid rafts” also known as  
 84 “detergent-resistant membrane domains” due to their insoluble

property in non-ionic detergents such as Triton X-100 [13,14]. 85  
 These specialized membrane microdomains contain a variety of 86  
 sphingolipid-metabolizing enzymes such as sphingomyelinase 87  
 (SMase) [15,16], ceramidase [17], sphingosine kinase [18], and 88  
 ceramide kinase [19]. Dobrowsky reported that depletion of 89  
 membrane cholesterol abolished p75<sup>NTR</sup>-dependent sphingo- 90  
 myelin hydrolysis and ceramide generation [13]. Due to the 91  
 localization of sphingolipid-metabolizing enzyme in lipid rafts 92  
 and the strong association of sphingolipids with cholesterol, 93  
 cholesterol appears to affect sphingomyelin metabolism 94  
 through modulation of lipid raft integrity. The lipid rafts also 95  
 contain a variety of receptors and signaling enzymes, such as 96  
 GTPases and kinases, and mediate receptor-mediated intracel- 97  
 lular signaling cascades and membrane trafficking [20,21]. 98  
 Therefore, it is now believed that the regulation of sphingolipid 99  
 metabolism in these membrane domains may be linked to 100  
 various cellular signaling events as well as cellular cholesterol 101  
 levels. 102

103 Recently, it has been reported that ROS and RNS are in-  
 104 volved in sphingolipid metabolism. For example, the depletion  
 105 of cellular reduced glutathione (GSH) by increased ROS and  
 106 RNS regulates enzymatic activities of SMases and ceramidase  
 107 [22–25]. Conversely, sphingolipids including ceramide, sphin-  
 108 gosine, and sphingosine-1-phosphate have the ability to regu-  
 109 late cellular redox homeostasis through regulation of NADPH  
 110 oxidase [26], mitochondrial integrity [27], NOS [10,28,29], and  
 111 antioxidant enzymes [30,31]. Therefore, in this review, we  
 112 discuss the mechanisms of activation and regulation of enzymes  
 113 which are involved in sphingomyelin metabolism and redox  
 114 regulation.

## 115 Regulation of sphingolipid metabolism by oxidative stress

116 Sphingomyelin (SM) is a constituent of membrane lipids and  
 117 mainly localized in the plasma membrane. At least two different  
 118 subtypes of SMases are involved in SM hydrolysis which is  
 119 regulated by intra- or extracellular stimuli. These are  
 120 identified based on pH optima, subcellular localization, and  
 121 cation dependence. The acidic, or lysosomal, sphingomyelinase  
 122 (A-SMase or SMPD1) was the first sphingomyelinase to be  
 123 identified and subjected to intensive investigation due to its role  
 124 in ceramide generation [32,33]. Later, the neutral, membrane-  
 125 bound  $Mg^{2+}$ -dependent SMases 1 and 2 (SMPD2 and 3) were  
 126 cloned [34] and characterized [35,36]. Although the detailed  
 127 mechanism for the activation of these SMases is still under

128 investigation, cellular redox potential is regarded as one of the  
129 key regulators for the activation of these enzymes.

### 130 Acid sphingomyelinase

131 A-SMase, which is deficient in patients affected with type A  
132 and B Niemann-Pick disease, has been known to play a role in  
133 stress signaling and apoptosis [37]. So far three types of human  
134 A-SMase have been cloned (types I, II, and III). These are  
135 generated by alternative splicing from a single transcript and  
136 only type I, which is major A-SMase species, has functional A-  
137 SMase activity [38]. A-SMase is generated from a 75-kDa  
138 proprecursor by proteolytic processing to a 72-kDa protein in  
139 ER/Golgi and further processing in endosome-lysosome to the  
140 fully active 70-kDa enzyme [39]. A-SMase is also found as an  
141 extracellular form and this secretory A-SMase is encoded from  
142 the same gene that encodes the lysosomal form of type I A-  
143 SMase but with different posttranslational modification [40].

144 The mechanism for A-SMase activation is not fully  
145 understood at present but several factors are identified to be  
146 involved in its activation or inhibition. These include tumor  
147 necrosis factor-receptor I (TNF-R1) [41,42], ApoC-III, an  
148 apolipoprotein [43], phosphatidylinositol-3'-kinase (PI3K)/  
149 protein kinase B (Akt) [44], and certain lipids such as mono-,  
150 di-, and triacylglycerols [45], 1,2-diacylglycerol [46], and  
151 sphingosine-1-phosphate [47]. A-SMase is also known to be

152 regulated by thiol oxidation. Although the enzymatic activity of  
153 A-SMase is not affected by GSH, it is inhibited by DTT in a  
154 dose-dependent manner [48,49]. The inactivation by DTT may  
155 not simply be due to disulfide reduction because effects of  
156 DTT on activity were reported to be unrelated to disulfide  
157 reduction [50]. Interestingly, Qiu and co-workers [23] have  
158 shown that the C-terminal cysteine (Cys<sup>629</sup>) is an unbridged  
159 free form and modification of this thiol group by dimerization,  
160 chemical modification, or deletion increases the enzymatic  
161 activity of A-SMase. Notably, restoration of the thiol group with  
162 DTT inhibits copper-mediated dimerization as well as activation  
163 of A-SMase [23]. Therefore, it is possible that oxidative  
164 modification of the C-terminal cysteine may be required for  
165 full activation of A-SMase and that certain antioxidants such as  
166 DTT may counteract by restoring the thiol group (Fig. 1).

167 In vitro studies have shown that NO plays a role in the  
168 regulation of A-SMase activity. NO from exogenous or  
169 endogenous sources has been known to inhibit TNF- $\alpha$ -  
170 mediated apoptosis via inhibition of ceramide generation  
171 [24,51]. Although the mechanism for NO-mediated inhibition  
172 of A-SMase is not fully understood, a NO-mediated increase in  
173 cGMP and the activation of cGMP-dependent protein kinase  
174 (PKG) have been suggested to be involved in the inhibition of  
175 A-SMase [52,53]. However, excessive amounts of NO are also  
176 known to activate apoptosis via activation of ceramide  
177 generation which is sensitive to A-SMase inhibitor [54]. NO-

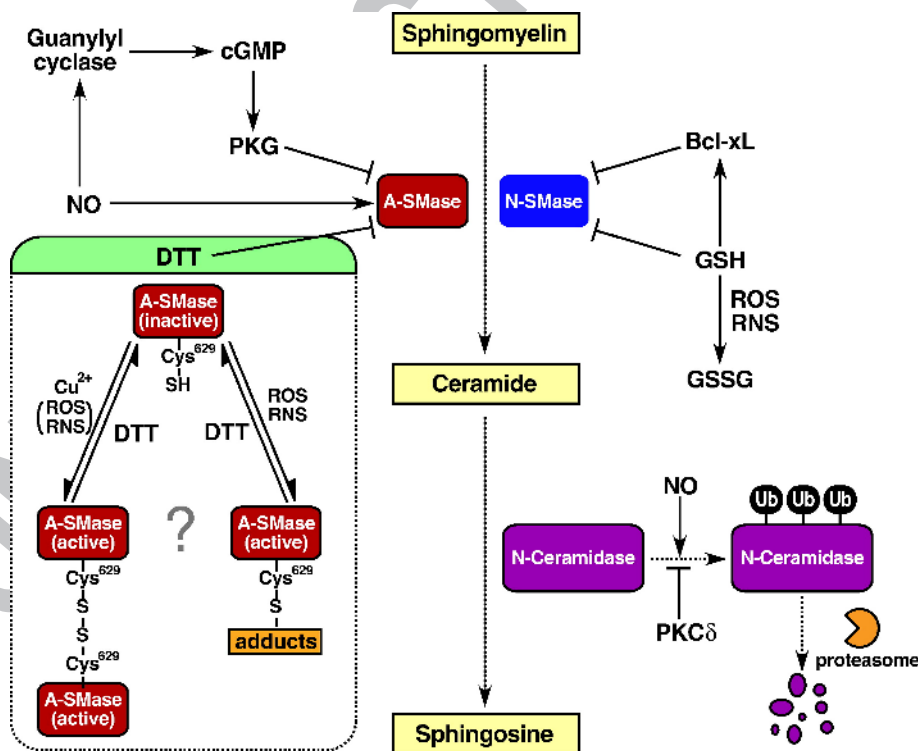


Fig. 1. The possible regulatory mechanism of sphingolipid metabolism by cellular redox potential. In mammalian cells, sphingomyelin is hydrolyzed by neutral (N-SMase) and acid sphingomyelinase (A-SMase). N-SMase is regulated by cellular level of reduced and oxidized forms of glutathione (GSH and GSSH) which are affected by antioxidant protein, Bcl-xL, and reactive oxygen or nitrogen species (ROS and RNS). Although the activity of A-SMase is not affected by GSH, it is inhibited by dithiothreitol (DTT) and cGMP-mediated pathway. The adduct formation on cysteine<sup>629</sup> is known to enhance the activity of A-SMase. However, whether ROS and RNS are implicated in the formation of adduct on cysteine<sup>629</sup> and whether DTT counteracts ROS/RNS-mediated adduct formation are not known at present. Ceramide accumulation is also regulated by the activity of neutral ceramidase (N-ceramidase). Nitric oxide (NO) is known to activate N-ceramidase degradation through the ubiquitine/proteasome pathway. Solid arrow and T-shaped heads represent stimulatory and inhibitory effects, respectively.

mediated interaction between A-SMase and procaspase-3 [55] suggests the possible regulation of A-SMase activity by NO through interaction with other proteins. Indeed, several protein molecules have been identified to be regulated posttranslationally by NO through S-nitrosylation-mediated adduct formation [56]. Therefore, posttranslational modification of A-SMase by S-nitrosylation, in particular on C-terminal cysteine (Cys<sup>629</sup>), and its role in A-SMase activity should be an interesting investigation (Fig. 1).

### Neutral sphingomyelinase

The presence of N-SMase which has neutral pH optimum (~7.4) and magnesium dependency was firstly described in 1967 by Schneider and Kennedy [57]. Later N-SMase1 was cloned and characterized in mice and humans by Tomiuk and co-workers [36]. The cloned human N-SMase1 (sphingomyelin phosphodiesterases; *smpd2*) gene is localized on chromosome 6 and encodes proteins with a predicted molecular mass of 47.6 kDa with two putative transmembrane domains at the C terminus [36]. Although it shows activity for SM hydrolysis in vitro [58], it shows no change of SM hydrolysis as compared to increased hydrolysis of 1-O-alkyl-lyso-phosphatidylcholine (lyso-platelet-activating factor or lyso-PAF) in N-SMase1-overexpressing cells, suggesting that the cloned enzyme is actually a lyso-PAF phospholipase C, but not N-SMase [58]. More recently, Hofmann and co-workers cloned and characterized another mammalian brain-specific magnesium-dependent N-SMase (N-SMase2; *smpd3*) from humans and mice [35]. Human N-SMase 2 gene is localized in chromosome 16 and encodes proteins of 655 amino acids, resulting in a predicted molecular mass of 71 kDa. The N terminus contains two predicted transmembrane domains, whereas the C terminus contains the putative catalytic domain. In contrast to N-SMase1, the N-SMase2 showed properties similar to the previously purified rat brain N-SMase but had no activity against lyso-PAF [35]. The primary subcellular localization of N-SMase2 was described to be in the Golgi [35]; however, later, its localization was reported to be in the plasma membrane [59].

The mechanism of N-SMase activation has been intensely studied during the past decade. In addition to bioactive lipids, such as arachidonic acid [35,36], anionic phospholipids (i.e., cardiolipin and phosphatidylglycerol) [60], and phosphatidylserine [61], N-SMase is also reported to be regulated by caspases 3 [62] and 9 [63] and TNF-R1 through FAN (factor associated with N-SMase activation) protein [64]. Upon ligation of TNF- $\alpha$ , TNF-R1 recruits FAN through its N-SMase activation domain (NSD) and activates N-SMase [64]. The role of FAN in N-SMase activation was further supported by other groups using FAN-deficient mice and cells which overexpressed dominant-negative FAN [65,66]. Since N-SMase1 does not exhibit sphingomyelinase activity in vivo, the increased N-SMase activity by these activators appears to be mediated by N-SMase2. However, the detailed mechanism for the activation of N-SMases2 by these activators is not known at present. Moreover, a possible role of other isoforms of N-SMase cannot be excluded.

Indeed, the existence of multiple forms of N-SMase in bovine brain was demonstrated previously based on different chromatographic and biochemical properties [67].

Previously, we and other groups have observed that proinflammatory cytokines (i.e., IL-1 $\beta$  and TNF- $\alpha$ ) or hypoxia induced SM hydrolysis and ceramide generation in a redox-sensitive event [12,68]. Moreover, A $\beta$ <sub>1-42</sub> or its synthetic peptide A $\beta$ <sub>23-35</sub>, which produces pathologies of Alzheimer's disease, induces ceramide generation by activation of N-SMase in a redox-sensitive manner without altering A-SMase [69,70]. Therefore, these studies suggest that oxidative stress-mediated N-SMase activation and ceramide generation may play a key role(s) in the pathobiology of various disease conditions. However, the mechanism of regulation of N-SMase by oxidative stress is not completely known. Liu and Hannun showed that GSH, but not DTT and  $\beta$ -mercaptoethanol, dose dependently inhibited partially purified N-SMase activity [49]. They reported that  $\gamma$ -glutamyl-cysteine, but not the free sulfhydryl group, in GSH may function as an allosteric regulator of N-SMase. GSH was also known to regulate N-SMase activity through Bcl-xL or Bcl-2, an antiapoptotic protein, by inhibiting oxidative stress-mediated SM hydrolysis and ceramide generation [71,72]. Moreover, tyrosine kinases such as Lyn and PKC $\zeta$  are also implicated in oxidative stress-mediated regulation of N-SMase activity and ceramide generation [73,74]. In addition to GSH, Takeda and associates reported that sodium nitroprusside (SNP), a NO donor, also increases cellular sphingomyelin hydrolysis and ceramide generation through activation of N-SMase [75]. However, whether NO mediates N-SMase activation through direct interaction or depletion of cellular GSH levels is not currently known (Fig. 1).

### Ceramidase

In addition to the increase in ceramide levels via activation of A-SMase or N-SMase, the increased ceramide levels may also be due to concomitant inhibition of ceramidases [22]. In renal mesangial cells, NO from an exogenous donor causes a chronic upregulation of ceramide levels by activating sphingomyelinases and concomitantly inhibiting ceramidases, and particularly in the late phase ceramide generation may be responsible for the further processing of a proapoptotic signal [22]. Later, this effect was shown to be due to the action of NO on ubiquitin/proteasome-mediated proteolysis of neutral ceramidase and counterregulated by protein kinase C (PKC), especially the  $\delta$ -isoform [76,77]. Therefore, neutral ceramidase may represent another novel target for interference with the cellular stress response and modulate programmed cell death, a typical feature of many inflammatory diseases (Fig. 1).

### Regulation of redox potential by sphingolipids

As discussed in the preceding section, oxidative stress regulates sphingolipid metabolism to generate sphingolipid molecules which participate in intracellular signaling. On the other hand, a growing body of evidence also suggests that

286 certain sphingolipids, such as ceramide or its glycosyl deri- 341  
 287 vatives, are able to induce cellular oxidative stress through 342  
 288 activation of NADPH oxidase [26], mitochondrial dysfunction 343  
 289 [27,78], and NOS [10,28,29], and/or downregulation of  
 290 antioxidant enzymes [30,31]. Therefore, sphingolipid metabo-  
 291 lism and redox homeostasis are regulated in a bidirectional  
 292 manner.

### 293 Regulation of NADPH oxidase

294 In phagocytes, ROS are generated by a membrane-associated 346  
 295 phagocytic NADPH oxidase (Phox; also known as NADPH 347  
 296 oxidase-2, Nox2), with its catalytic moiety gp91<sup>phox</sup>, which is 348  
 297 activated by assembly with regulatory proteins such as 349  
 298 p47<sup>phox</sup>, p67<sup>phox</sup>, and Rac [79,80]. Recently, other oxidases 350  
 299 similar to the Phox complex have also been identified in other 351  
 300 cell types and show different expression patterns depending on 352  
 301 cell or tissue types [81,82]. Similar to Nox, dual oxidase (Duox) 353  
 302 isoforms (i.e., Duox 1 and 2) include molecular mass gp91<sup>phox</sup> 354  
 303 homologs with an N-terminal peroxidase domain in addition to 355  
 304 the C-terminal NADPH oxidase activity. 356

305 The involvement of sphingolipids in the regulation of 346  
 306 NADPH oxidase activity was first discussed in Gaucher disease 347  
 307 type I [83]. Liel and co-workers reported that monocyte 348  
 308 dysfunction in Gaucher disease type I patients is caused by 349  
 309 suppression of NADPH oxidase-mediated superoxide genera- 350  
 310 tion as glucosylceramide (glucocerebroside) accumulates [83]. 351  
 311 Recent studies by Moskwa and co-workers further support the 352  
 312 role of glucosylceramide in the regulation of NADPH oxidase 353  
 313 activity, in which glucosylceramide is able to inhibit NADPH 354  
 314 oxidase in a cell-free system [84]. However, other related lipids 355  
 315 such as lactosylceramide (CDw17) and ganglioside GD3 have 356  
 316 been reported to upregulate NADPH oxidase activity [85–88]. 357  
 317 The lactosylceramide-mediated activation of NADPH oxidase 358  
 318 and ROS generation is involved in endothelial and neutrophil 359  
 319 cell functions through regulation of endothelial cell prolifera- 360  
 320 tion, adhesion molecule expression, and phagocytosis [85–87]. 361  
 321 Recently, our group also reported the involvement of lacto- 362  
 322 sylceramide-mediated ROS generation in sequential activation 363  
 323 of hRas/NFκB and inflammatory gene expression [89]. The 364  
 324 exact mechanism of lactosylceramide-mediated activation of 365  
 325 NADPH oxidase is not clear, but it is believed that lacto- 366  
 326 sylceramide interactions with a Src family kinase (i.e., Lyn) in 367  
 327 the lipid rafts may lead to generation of ROS through phos- 368  
 328 phatidylinositol-3-kinase-, p38 MAPK-, and PKC-dependent 369  
 329 signal transduction pathways [90]. 370

330 Along with lactosylceramide, ceramide has also been 381  
 331 implicated in the regulation of NADPH oxidase. Recently, 382  
 332 ceramide-mediated activation of NADPH oxidase and resultant 383  
 333 oxidative stress were reported to be involved in endostatin- 384  
 334 induced endothelial dysfunction [91]. The mechanism for 385  
 335 ceramide-mediated activation of NADPH oxidase is not fully 386  
 336 understood but involvement of ceramide-mediated activation of 387  
 337 Rac small GTPase, a regulatory component of NADPH oxidase, 388  
 338 was suggested recently [92]. Moreover, ceramide activates the 389  
 339 NADPH oxidase through activation of PKCζ [93]. PKCζ may 390  
 340 also activate p47<sup>phox</sup> adapter protein via phosphorylation [93],

followed by translocation of activated p47<sup>phox</sup> to membrane to 341  
 facilitate stimulus-induced binding of p67<sup>phox</sup> to the holo 342  
 NADPH enzyme complex [26]. 343

### Mitochondrial dysfunction: mitochondrial redox regulation 344 and apoptosis 345

Mitochondria play a central role in cellular metabolism. 346  
 They are the site of fatty acid catabolism and the citric acid 347  
 cycle, which produces NADH and FADH<sub>2</sub>. These molecules 348  
 transfer electrons to the respiratory chain, and finally to oxygen, 349  
 a process that generates ATP. It has long been recognized that 350  
 the mitochondrial electron transport chain is a site of free radical 351  
 generation [94]. The two sites where this occurs are complex I 352  
 (NADH-coQ reductase) and complex III (cytochrome *c* 353  
 oxidase). The electron leaks from mitochondria and formation 354  
 of O<sub>2</sub><sup>-</sup> have been identified in normal as well as pathological 355  
 conditions. Mitochondria are also known to play a central role 356  
 in regulating apoptosis [95]. Mitochondria sense the catastro- 357  
 phic cellular changes and irreversibly commit cells to apoptosis 358  
 by releasing death factors into the cytosol, such as cytochrome *c* 359  
 [95], Smac 2/DIABLO [96], AIF [97], and EndoG [98]. 360

Ceramide was reported as a regulator for the generation of 361  
 ROS and activation of the mitochondrial irreversible apoptotic 362  
 process. Mitochondria isolated from TNF-α-treated hepatocytes 363  
 showed a higher content of ceramide, compared to control [27], 364  
 and addition of C<sub>2</sub>-ceramide to mitochondria from untreated 365  
 cells increased ROS production [27]. Moreover, naturally 366  
 occurring C<sub>16</sub> ceramide was shown to cause an increase in 367  
 ROS generation through mitochondria [78]. Ceramide may 368  
 function to generate ROS from mitochondria as a consequence 369  
 of cytochrome *c* release, an electron carrier of the respiration 370  
 chain between complexes II and III in mitochondria [99]. 371  
 Ghafourifar et al. have shown that C<sub>2</sub>- and C<sub>6</sub>-ceramide induce 372  
 release of cytochrome *c* from isolated mitochondria [100]. 373  
 Since cytochrome *c* release causes a decrease in mitochondrial 374  
 oxygen consumption, mitochondrial inner transmembrane 375  
 potential (ΔΨ<sub>m</sub>), and Ca<sup>2+</sup> retention and all of which lead to 376  
 mitochondrial dysfunction and ROS generation [100], cera- 377  
 mide-mediated release of cytochrome *c* may be one of the key 378  
 events in the induction of ROS generation from mitochondria 379  
 (Fig. 2). 380

Ceramide may also affect cellular redox potential through 381  
 regulation of the Bcl-2 family of proteins, which are regarded as 382  
 antioxidants because they increase the GSH pool or redistribute 383  
 GSH to various cellular compartments [101]. Indeed, Bcl-2 was 384  
 known to prevent ROS production, GSH depletion, and cellular 385  
 damage caused by lipid peroxidation [102,103] through 386  
 blocking cytochrome *c* release from mitochondria [104] and/ 387  
 or inhibition of mitochondrial permeability transition pore 388  
 opening, leading to collapse of ΔΨ<sub>m</sub>, by opposing the effect of 389  
 Bax, a component of the permeability transition pore [105,106]. 390  
 Long-term treatment of human keratinocytes with C<sub>2</sub>-ceramide 391  
 induced downregulation of Bcl-2 [107]. Moreover, apoptotic 392  
 DNA fragmentation following exposure to TNF-α and C<sub>2</sub>- 393  
 ceramide was also associated with downregulation of Bcl-2 394  
 mRNA in HL-60 and U-937 cells [108], suggesting the possible 395

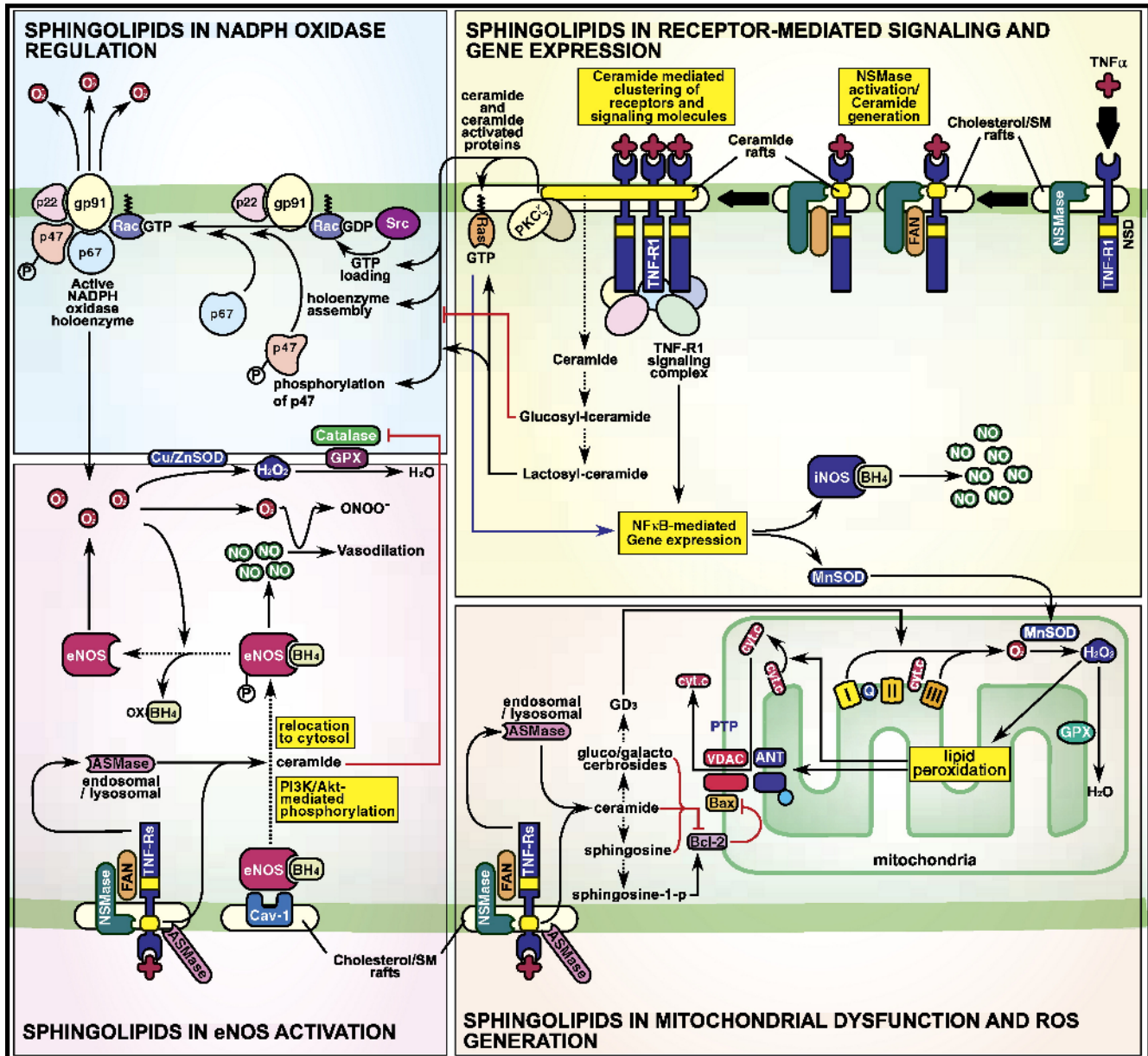


Fig. 2. The possible regulatory mechanism of cellular redox potential by sphingolipid metabolism. Ceramide is one of the key signaling mediators in receptor-mediated signaling cascades. It may activate gene expression of redox enzymes (i.e., iNOS and Mn-SOD) through receptor clustering, recruitment of signaling enzymes, and activation of ceramide-activated protein kinases (i.e., PKC $\zeta$  and KSR). The ceramide-activated signaling cascades, along with its derivatives such as glucosyl-ceramide and lactosyl-ceramide, may be implicated in the regulation of NADPH oxidase activity through regulation of Rac1 GTP loading, holoenzyme assembly, and/or p47 phosphorylation. Ceramide is also implicated in the activation of eNOS through cytosolic relocation from membrane and phosphorylation by the phosphoinositide 3-phosphate kinase (PI3K)/Akt pathway. However, ceramide is also able to inhibit action of eNOS (vasodilation) through NADPH-mediated superoxide generation (O<sub>2</sub><sup>-</sup>) leading to formation of peroxynitrite (ONOO<sup>-</sup>) or tetrahydrobiopterin (BH<sub>4</sub>) oxidation leading to uncoupling of eNOS. Ceramide, sphingosine, and GD3 are potential activators for mitochondrial dysfunction which leads to the production of massive amounts of O<sub>2</sub><sup>-</sup>. They produce interference of electron transfer, disruption of mitochondrial inner transmembrane potential ( $\Delta\Psi_m$ ), opening permeability transition pore (PTP), mitochondrial lipid peroxidation, and cytochrome *c* (cyto *c*) release. Ceramide is also known as a potent activator for mitochondrial Mn-SOD gene expression. The increased Mn-SOD by ceramide may be toxic depending on the level of glutathione peroxidase (GPX) in mitochondria. The role of ceramide on the regulation of cellular GPX is not known at present, but it was known to inhibit catalase activity. Solid arrow and T-shaped heads represent stimulatory and inhibitory effects, respectively.

396 role for ceramide in the regulation of Bcl-2-mediated anti-  
 397 oxidant activity (Fig. 2).  
 398 Ceramide was also reported to disturb the respiratory chain  
 399 through direct interaction [109,110] as C<sub>2</sub>- and C<sub>6</sub>-ceramide  
 400 treatment induced large pores in phospholipid planar mem-  
 401 branes [111]. Interestingly, rat liver mitochondria contain free

ceramide [112] and sphingolipid-metabolizing enzymes such as  
 ceramidase [113] and ceramide synthase [114]. Thus, dynamic  
 changes in the ceramide content of mitochondrial membranes  
 by vesicular transport or local production could possibly  
 regulate mitochondrial integrity and ROS generation. More-  
 over, ceramide can be converted into sphingosine by ceramidase

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408 and sphingosine-1-phosphate by further action of sphingosine  
409 kinase, thus expanding the repertoire of downstream signals  
410 which might affect cell fate. Sphingosine, as a negative  
411 regulator of cell proliferation, is known to promote apoptosis  
412 [115]. Moreover, sphingosine is also involved in the down-  
413 regulation of Bcl-2 [116] and Bcl-X<sub>L</sub> [117], increase in  
414 cytochrome *c* release [118,119], mitochondrial  $\Delta\Psi$  disruption  
415 [120], and mitochondrial generation of H<sub>2</sub>O<sub>2</sub> [27]. In contrast,  
416 sphingosine 1-phosphate stimulates cell growth and is thus  
417 antiapoptotic [115] through regulation of Bcl-2/Bax rheostat  
418 [121] and inhibition of cytochrome *c* release [122]. Interest-  
419 ingly, sphingosine kinase-overexpressing cells have decreased  
420 levels of both sphingosine and ceramide [123,124] (Fig. 2).

421 A recent cDNA microarray study showed that the *bcl-2* gene  
422 is downregulated in Gaucher disease, suggesting that the  
423 accumulation of either glucocerebroside or glucosylsphingo-  
424 sine, as a result of glucocerebrosidase deficiency, affects Bcl-2-  
425 mediated redox regulation [125]. In addition, ganglioside GD3  
426 is also known to induce swelling of isolated mitochondria  
427 through opening permeability transition pores [126,127];  
428 however, no such effects on mitochondrial permeability are  
429 described for other lipids such as GM1, GD1a, GM3, and  
430 GT1b. GD3 appears to interfere at the level of complex III of the  
431 electron transport chain [88] and GD3-mediated permeability  
432 transition pore opening is secondary to reactive oxygen species  
433 generation [128]. Therefore, the burst of ROS generation by  
434 GD3 could also induce opening of the permeability transition  
435 pores leading to cytochrome *c* release (Fig. 2).

#### 436 Regulation of nitric oxide synthases

437 Since its discovery, nitric oxide (NO) has become the subject  
438 of both intense research and heated debate over its role in  
439 various biological and pathophysiological processes. Originally  
440 discovered as a mediator of vascular smooth muscle relaxation,  
441 NO has since been implicated in a wide range of physiological  
442 mechanisms ranging from lysis of tumor cells to neural  
443 transmission [129,130]. NO is a metabolic by-product of the  
444 conversion of L-arginine to L-citrulline by a class of enzymes  
445 dubbed as the nitric oxide synthases (NOS). To date, three  
446 isoforms of NOS have been identified. Neuronal NOS (nNOS  
447 or NOS1) is expressed constitutively by neurons in the brain  
448 and enteric nervous system, whereas endothelial NOS (eNOS or  
449 NOS3) exhibits constitutive expression which is confined to the  
450 endothelial cells lining the vasculature [129,131]. The third  
451 isoform of NOS is an inducible NOS (iNOS or NOS2) and as  
452 the name implies, it is expressed only in response to certain  
453 inflammatory stimuli such as bacterial products, cytokines, and  
454 lipid mediators [130,131]. Classically, NO is considered to be  
455 an activator for cGMP [132] in the regulation of cardiovas-  
456 cular function [133] and neurotransmission [134]. Very  
457 recently, S-nitrosylation, the covalent attachment of a nitrogen  
458 monoxide group to the thiol side chain of cysteine, has  
459 emerged as an important mechanism for dynamic posttrans-  
460 lational regulation of proteins [56]. S-Nitrosylation thereby  
461 conveys a large part of the ubiquitous influence of nitric  
462 oxide on cellular signal transduction, and provides a

mechanism for redox-based physiological regulation [56]. In  
463 addition, NO in O<sub>2</sub><sup>-</sup> producing environment reacts rapidly to  
464 form the highly toxic peroxyxynitrite anion, which then  
465 protonates and decomposes to generate <sup>•</sup>OH or some other  
466 potent oxidant with similar reactivity [135]. This is of  
467 particular importance in neurodegenerating disease conditions  
468 such as demyelinating disease and in ischemia and traumatic  
469 injuries associated with infiltrating peripheral mononuclear  
470 cells and the production of proinflammatory cytokines, where  
471 subsequent astrocytes and microglia-derived NO could  
472 contribute to oligodendrocyte degeneration and neuronal  
473 death [136,137] (Fig. 2). 474

#### Endothelial nitric oxide synthase 475

eNOS identified in endothelial cells is also expressed in  
476 cardiomyocytes [138,139]. eNOS produces NO via a complex  
477 reaction which is stimulated by Ca<sup>2+</sup> and requires NADPH,  
478 along with other cofactors [138]. The role of ceramide in NO  
479 generation through eNOS was identified because ceramide  
480 affects vasorelaxation. The role of ceramide in vascular  
481 function has been extensively reviewed by Berry et al. [29].  
482 Initial studies probing the effect of ceramide on vascular  
483 contractility demonstrated that application of cell-permeable  
484 analogs of ceramide or exogenous bacterial sphingomyelinase  
485 to precontracted vascular segments results in concentration-  
486 dependent relaxation [140,141]. Subsequently, Jin and co-  
487 workers also reported that micromolar concentrations of  
488 ceramide (C<sub>2</sub>-, C<sub>6</sub>-, and C<sub>16</sub>-ceramide) induce significant  
489 relaxation in a NO-dependent manner and removal of the  
490 endothelium significantly inhibited ceramide-induced relaxa-  
491 tion [142]. Interestingly, angiotensin II type 2 receptor  
492 activation also increases intracellular concentrations of cer-  
493 amide [143,144]; therefore, ceramide may contribute to some  
494 of the physiological effects of angiotensin II through  
495 stimulation of nitric oxide production [145]. Although the  
496 precise mechanism for ceramide-induced vasodilation is not  
497 fully understood, recent studies have identified phosphatidy-  
498 linositol-3'-kinase and Akt as downstream candidate effectors  
499 for ceramide in eNOS activation [146]. Moreover, ceramide-  
500 mediated translocation of eNOS from plasma membrane,  
501 where it is bound to caveolin-1 as an inactive form, to the  
502 cytoplasm was also demonstrated as a crucial step in ceramide-  
503 induced synthesis of NO by eNOS [147,148]. However, there  
504 are opposing views on the role of ceramide in vasoregulation  
505 [11]. It was demonstrated that TNF- $\alpha$  inhibits NO-mediated  
506 endothelium-dependent vasorelaxation in small coronary  
507 arteries via sphingomyelinase activation and consequent  
508 superoxide production [149]. Indeed, ceramide was reported  
509 to inhibit endothelium-dependent vasodilation via an increase  
510 in O<sub>2</sub><sup>-</sup> and a subsequent decrease in NO availability, without  
511 altering NO synthesis [145], and this impairment of endothelial  
512 function was prevented by overexpression of Cu/Zn superoxide  
513 dismutase [150]. Therefore, ceramide appears to have a  
514 bifunctional role in the regulation of NO-mediated vaso-  
515 regulation through activation of eNOS-mediated NO produc-  
516 tion and/or ROS generation which lowers NO availability by  
517 generation of peroxyxynitrite (Fig. 2). 518

519 Interestingly, ceramide-induced ROS generation also mediates oxidation of (6R)-5,6,7,8-tetrahydro-L-biopterin (BH<sub>4</sub>, a cofactor of eNOS) that leads to BH<sub>4</sub> deficiency [151] (Fig. 2). The deficiency of BH<sub>4</sub> causes an increase in uncoupled eNOS molecules which leads to the formation of O<sub>2</sub><sup>-</sup> instead of NO [151,152]. This phenomenon, along with increased gene expression of eNOS [151] by ceramide, may be one of the mechanisms of ceramide-mediated impairment of endothelial function and increased oxidative stress under pathophysiological disease conditions, such as hypertension, experimental diabetes, and hypercholesterolemia, and in smokers [152].

### 530 *Inducible nitric oxide synthase*

531 In 1998, our group first reported the role of SMase and ceramide in iNOS gene expression and NO production [11]. The role of ceramide in the induction of iNOS gene expression was further supported by other groups [153,154]. Furthermore, recent studies demonstrated that a selective inhibitor of N-SMase downregulates LPS and/or Aβ-induced iNOS expression in macrophages and astrocytes [10,69]. On the other hand, neither pharmacological inhibition nor knockout of A-SMase affected the expression of iNOS [10,69,155]. These reports suggest a role of ceramide produced by N-SMase in the expression of iNOS as well as other inflammatory genes that are related to the regulation of cellular redox potential. The mechanism for initiation of the ceramide-mediated inflammatory signaling cascade is not clearly understood. Ceramide, generated by N-SMase, induced NFκB activation through activation of hRas signaling cascades [10]. Moreover, the requirement of tyrosine kinases in this reaction [154] suggests the possible action of tyrosine kinase in Ras/NFκB activation and iNOS gene induction (Fig. 2). Putative ceramide-interacting enzymes, such as Ser/Thr protein kinase (CAPK) [156], kinase suppressor of Ras (KSR) [157], phosphatase (PP) 2A and 1B [158], and PKCζ [159] either via direct interactions or indirectly through formation of specific membrane microdomains may play a role in these signaling events. Recent studies have shown that ceramide plays a role in clustering of TNF family receptors (i.e., TNF-R1, CD40, and CD95) [160–162]. Following ligation of these receptors with ligands, the ceramide produced by SMase around TNF receptors generates signaling microdomains. Since ceramide has the ability to self-aggregate [163], subsequent fusion of these small entities into larger membrane domains (ceramide rafts) [164] has been demonstrated to trigger the clustering of these receptor molecules [161,162]. The receptor clustering in the rafts induces close contact of receptors with other signaling molecules [165] and exclusion of inhibitory molecules (i.e., CD45 tyrosine phosphatase) [166], and thus stabilizes ligand-receptor-signaling protein interactions [161,162]. Therefore, the increase in ceramide and the formation of ceramide rafts may enhance inflammatory or death signaling events that are tightly related to cellular redox potential [161,162] (Fig. 2).

571 In addition to ceramide, its glycosylated form lactosylceramide was also demonstrated to activate iNOS and other cytokine gene expression in astrocytes and a rat spinal cord injury model [89]. The exact mechanism of this reaction is

not known but inflammatory cytokine-mediated activation of phosphatidylinositol-3-phosphate kinase appears to mediate an increase in lactosylceramide via activation of galactosyl transferase-2 (GalT-2) [167]. Both ceramide [10] and lactosylceramide induce iNOS gene expression through activation of hRas/NFκB, but whether ceramide produced by N-SMase is utilized for the synthesis of lactosylceramide is not known.

### *Neuronal nitric oxide synthase*

In contrast to eNOS and iNOS, the role of sphingolipids in the regulation of nNOS activity is relatively unknown. In neuronal cells, sphingosine treatment strongly inhibits the activity of cytosolic Ca<sup>2+</sup>-independent NOS (a putative nNOS); however, treatment with ceramide, N-acetylsphingosine, sphingosine-1P, sphinganine, and tetradecylamine had no effect on NOS activity [28]. Increasing concentrations of calmodulin led to loss of sphingosine inhibition, suggesting that sphingosine interferes with the calmodulin-dependent activation of the enzyme by a competitive mechanism [28] but without altering the intracellular Ca<sup>2+</sup> concentration [168]. These observations suggest that bioactive sphingosine plays a role in neuronal NO signaling.

### *Regulation of antioxidant enzymes*

In mammalian cells O<sub>2</sub><sup>-</sup> generated by respiration in mitochondria and by activation of NADPH-oxidase or xanthine oxidase is converted into H<sub>2</sub>O<sub>2</sub> by three forms of superoxide dismutase (extracellular and intracellular CuZn- and Mn-SODs) [169]. Extracellular SOD (EC-SOD) is mainly produced by vascular muscle cells and localized between endothelium and vascular muscle cell layers where it binds to cell surface, basal membrane, and extracellular matrix [170,171]. EC-SOD was known to be a major determinant of NO bioavailability in blood vessels through inhibition of vascular peroxynitrite generation [172]. Similarly, CuZn-SOD, which is a constitutively expressed cytosolic isoform, is also involved in the regulation of vascular functions through regulation of vascular O<sub>2</sub><sup>-</sup> level and peroxynitrite formation [173,174]. Notably, its overexpression is able to inhibit ceramide or lactosylceramide-mediated impairment of endothelial function or ICAM expression observed in pathological conditions [175]. Therefore, to protect NO over its entire diffusion route against ceramide-mediated ROS, normal expression of both CuZn-SOD and EC-SOD may be essential.

Mn-SOD is an inducible isoform of SOD and mainly localized in mitochondria. Because of its localization and reported lethal phenotype in null mice, Mn-SOD is considered to be the first line of defense against oxidative stress from mitochondria [31,176]. The expression or activity of Mn-SOD or both may be altered under several physiological and pathophysiological conditions. For example, Mn-SOD is particularly responsive to and upregulated by oxidative stress caused by oxidized LDL, TNF-α, or H<sub>2</sub>O<sub>2</sub> [176]. Moreover, cell-permeable ceramide or bacterial sphingomyelinase also



629 increase the expression of Mn-SOD in various cell types such  
630 as rat primary astrocytes, rat mesangial cells, glioma, PC12  
631 cells, skin fibroblasts [168,177], and neurons [31]. Ceramide-  
632 mediated generation of ROS and subsequent activation of  
633 redox-sensitive transcription factors such as activator protein-  
634 1 (AP-1) and NF $\kappa$ B may be involved in the upregulation of  
635 Mn-SOD gene expression [178,179] (Fig. 2).

636 Following conversion of O $_2^-$  by SODs into H $_2$ O $_2$ , it is  
637 believed to play a role in various cellular signal transduction  
638 pathways associated with cellular redox [180]. In addition, in  
639 the presence of transient metals (iron or copper) it forms a  
640 hydroxyl anion which is a strong oxidant and thus participates  
641 in the pathobiology of various disease conditions. Therefore,  
642 in the absence of adequate detoxification of H $_2$ O $_2$ , increased  
643 activity of SOD may cause oxidative stress [30,181]. Two  
644 enzymatic systems are involved in the detoxification of H $_2$ O $_2$ ,  
645 catalase and peroxidases [glutathione peroxidase (GPX) and  
646 thioredoxin peroxidase (TPX)]. It is not clear how ceramide  
647 regulates activity of GPX or TPX but ceramide was reported  
648 to inhibit catalase function in various cell types [30]. The  
649 mechanism for ceramide-induced inhibition of catalase is not  
650 clear at present but the inhibitory effect of ceramide on  
651 phosphatidylinositol-3-kinase has been reported to be in-  
652 volved in this reaction [182,183] (Fig. 2).

653 Peroxisomal redox is maintained by the enzyme system  
654 for production of O $_2^-$  and H $_2$ O $_2$  and the antioxidant enzyme  
655 system (Cu/Zn-SOD, Mn-SOD, catalase, and GPX) [183].  
656 Sphingolipids may inhibit catalase activity through modulat-  
657 ing peroxisomal function. The peroxisome is a redox-  
658 sensitive organelle where H $_2$ O $_2$  produced by various oxidases  
659 is detoxified by catalase, a major peroxisomal matrix protein  
660 [182]. Drastic alteration of peroxisomal functions, as well as  
661 oxidative stress by mislocalization of catalase from peroxi-  
662 somes [184], suggests that peroxisomal integrity and function  
663 are important for the regulation of catalase activity. Recently,  
664 our group reported that galactosyl-sphingosine (psychosine),  
665 a metabolites that accumulates in the brains of globoid cell  
666 leukodystrophy (GLD) [185] or Krabbe's disease [186]  
667 patients, inhibits peroxisomal functions and increases cellular  
668 free radical production [187,188]. Although the role of other  
669 sphingolipids in peroxisomal function and catalase activity  
670 has not been studied yet, the inhibitory effect of TNF- $\alpha$  on  
671 the expression of peroxisome proliferator-activated receptors  
672 (PPARs) [189] and catalase activity [187,188] along with the  
673 concomitant increase in ceramide levels [187,188] suggests  
674 the possible role of ceramide in peroxisome function as well  
675 as catalase activity.

## 676 Summary and conclusion

677 Sphingolipids including sphingosine, sphingosine-1-phos-  
678 phate, ceramide, ceramide-1-phosphate, psychosine, gluco-  
679 sylceramide, lactosylceramide, and GD3 are known to play a  
680 key role in receptor-mediated signal cascades which regulate  
681 cell proliferation, inflammation, and endothelial function.  
682 Similarly, endogenous prooxidants such as ROS and RNS  
683 also play a key role in receptor-mediated activation of

NADPH oxidase and NOS which are also involved in 684  
various aspects of cell physiological regulation. A growing 685  
body of evidence suggests that these two pathways interact 686  
with each other. Prooxidants such as ROS and RNS regulate 687  
sphingolipid metabolism through regulating the enzymes 688  
responsible for their metabolism including SMase and 689  
ceramidase. On the other hand, sphingolipids such as 690  
ceramide, lactosylceramide, and GD3 also mediate ROS 691  
and RNS generation through regulation of NADPH oxidase, 692  
NOS, and antioxidant enzymes such as Mn-SOD and 693  
catalase. Along with the physiological signaling cascades, 694  
the interaction of these two pathways may also be involved 695  
in cytotoxic or apoptotic cascades. Ceramide and other 696  
sphingolipids such as sphingosine or GD3 were initially 697  
known as potent proapoptotic agents which produce 698  
irreversible mitochondrial dysfunction and massive ROS 699  
generation. Although events that switch the roles of 700  
prooxidants and sphingolipids from physiological to pro- 701  
apoptotic signaling cascades are still under investigation, it is 702  
believed that cellular redox potential is a crucial factor for 703  
this transition. For example, in the CNS, oligodendrocytes, 704  
which are known to have low levels of GSH compared to 705  
astrocytes or microglia, undergo apoptotic pathway activa- 706  
tion upon stimulation with neurotoxic substances or proin- 707  
flammatory cytokines, while astrocytes and microglia undergo 708  
proliferation or inflammatory activation. Under low redox 709  
buffering states, the receptor-mediated generation of ROS and 710  
RNS may produce oxidative stress and then activate redox- 711  
sensitive SMase and ceramide generation. In this event, 712  
exclusive ceramide production may be able to induce mito- 713  
chondrial dysfunction that further promotes ROS generation and 714  
apoptosis. Similarly, the interaction of ROS/RNS generation and 715  
sphingolipid metabolism may also play a crucial role in 716  
endothelial function. As discussed, ceramide exerts its role as 717  
a vasodilator through activation of eNOS. However, it may also 718  
act as a vasoconstrictor by its dual role in the activation of ROS 719  
generation when ROS are not removed due to a low redox 720  
buffering state. 721

722 During the past two decades, the regulation of  
723 sphingolipid metabolism has been under intense investiga-  
724 tion due to the involvement of these events in the  
725 pathophysiology of various disease conditions. Here, we  
726 have discussed the interregulation of ROS/RNS generation  
727 and sphingolipid metabolism as one of the crucial factors  
728 promoting the pathological outcome. Therefore, therapeutic  
729 approaches for intervention of sphingolipid-induced patho-  
730 logical signal transduction pathways and the use of  
731 antioxidants may improve the efficacy of therapeutics in  
732 these disorders.

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