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It's a lipid's world: Bioactive lipid metabolism and signaling in neural stem cell differentiation

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Abstract

Lipids are often considered membrane components whose function is to embed proteins into cell membranes. In the last two decades, studies on brain lipids have unequivocally demonstrated that many lipids have critical cell signaling functions; they are called “bioactive lipids”. Pioneering work in Dr. Robert Ledeen’s laboratory has shown that two bioactive brain sphingolipids, sphingomyelin and the ganglioside GM1 are major signaling lipids in the nuclear envelope. In addition to derivatives of the sphingolipid ceramide, the bioactive lipids discussed here belong to the classes of terpenoids and steroids, eicosanoids, and lysophospholipids. These lipids act mainly through two mechanisms: 1) direct interaction between the bioactive lipid and a specific protein binding partner such as a lipid receptor, protein kinase or phosphatase, ion exchanger, or other cell signaling protein; and 2) formation of lipid microdomains or rafts that regulate the activity of a group of raft-associated cell signaling proteins. In recent years, a third mechanism has emerged, which invokes lipid second messengers as a regulator for the energy and redox balance of differentiating neural stem cells (NSCs). Interestingly, developmental niches such as the stem cell niche for adult NSC differentiation may also be metabolic compartments that respond to a distinct combination of bioactive lipids. The biological function of these lipids as regulators of NSC differentiation will be reviewed and their application in stem cell therapy discussed.

Keywords

sphingolipids; eicosanoids; lysophospholipids; steroids; embryonic stem cells; neural progenitor cells; differentiation

Overview

The signal transduction mechanism of bioactive lipids is not always exclusively defined by one pathway. There are lipids that specifically act through receptors such as endocannabinoids, but then there are others that can bind to receptors and intracellular protein targets such as S1P. Some lipids can affect the same receptor or signaling protein even twice, by direct binding and affecting the composition of lipid rafts such as ceramide (Fig. 1). Therefore, the classification of signaling lipids into three categories (bioactive lipids signaling through receptors; bioactive lipids binding to protein kinases, phosphatases, and other signaling proteins; and bioactive lipids regulating lipid rafts and raft-associated proteins) and is not always based on a clear-cut distinction. Likewise, individual lipids do not have exclusive functions in NSC differentiation. For example, LPA and S1P can promote proliferation and self-renewal in one population of stem cells, while their main

function is induction of differentiation in another one. These differences, which may very well be species-dependent (mice vs. men) will also be discussed in greater detail. Finally, bioactive lipids may not be sufficiently effective by themselves, but they significantly enhance the function of other growth factors and differentiation inducers, e.g., in the case of raft-associated receptors. Therefore, it is rather a specific “cocktail” of bioactive lipids and growth factors that ultimately determines the fate of differentiating stem cells. It is the goal of the review to discuss the underlying mechanisms for this lipid-dependent NSC differentiation and how lipids can then be used to steer stem cell differentiation toward one lineage.

1. Bioactive lipids signaling through receptors

A variety of bioactive lipids is known to specifically interact with receptors in the cell membrane, the endoplasmic reticulum (ER), and the nucleus (Fig. 1). The lipid classes acting through receptors in the cell membrane are lysophospholipids (e.g., LPA and S1P), eicosanoids (e.g. prostaglandins), and endocannabinoids (e.g., anandamide). These lipids activate G-protein coupled receptors (GPCRs) of the G_i, G_q, and G12/13 type. Downstream of these lipid-activated GPCRS are the PI3K-to-Akt (G_i), Ras-to-ERK (G_i, G_q), and PLC-to-PKC (G12/13) cell signaling pathways for neural stem cell proliferation and differentiation. In most cases, the specific cellular response (e.g., proliferation vs. differentiation) is orchestrated by a particular combination of bioactive lipids. Likewise, several lipids bind to nuclear receptors such as the terpenoid retinoic acid, and steroid hormones and glucocorticoids. Recently, lipid receptors in the ER have gained attention because they can affect ER stress and redox balance, two metabolic regulators of neural stem cell differentiation. In particular, sigma receptors are interesting in that they can bind to sphingolipids and neurosteroids as specific ligands, but are also lipid raft-associated. Therefore, many bioactive lipids can regulate stem cell differentiation in two different ways, by binding to specific receptors and modulating the membrane environment of receptors associated with lipid rafts. In this chapter, we will first discuss the receptor specific binding of these lipids and elaborate on the metabolic effect (ER stress and redox potential) and lipid raft-dependent stem cell differentiation in the subsequent chapters.

1.1. Nuclear receptors: Terpenoids and steroids

Terpenoids or isoprenoids are a structurally rich family of lipids based on the ene-addition of isoprene units, which are known to give rise to linear, branched, and cyclic compounds including cholesterol and other steroids. Retinoic acid (RA) is one of the terpenoids with a long tradition of usage in neural stem cell differentiation. In 1982, it was for first time described that RA induces neuronal and glial differentiation of P19 embryonic carcinoma cells [1]. Five years later, it was shown that RA induces motoneuron differentiation of mouse embryonic stem (ES) cells and it was speculated that it is a physiological inducer of the Hox2.3 gene [2]. It is now well established that RA is critical for the spatially regulated expression of Hox genes during embryonic brain development. This is achieved by the tissue-specific expression of retinal aldehyde dehydrogenase 2 (Raldh2), an enzyme converting retinaldehyde (retinal) into RA. In contrast to Aldh1, a commonly used marker for undifferentiated stem cells, RALdh2 is missing in undifferentiated ES cells preventing the undue expression of RA as differentiation inducer when stem cells are in contact with retinal or retinol (vitamin A) [3]. Interestingly, incubation of mouse ES cells with retinol/vitamin A resulted in the promotion of self-renewal, suggesting a novel cell signaling pathway for maintenance of pluripotency by vitamin A. While it has been found that this signaling pathway involves the insulin growth factor 1 receptor (IGF-1R)-to-PI3K-to-Akt-to-mTORC1 signal transduction pathway, it is not clear how vitamin A activates this pathway [3, 4].

In contrast to vitamin A, its derivative RA triggers the expression of Hox genes by activation of nuclear RA receptors or RARs. RA binds to RAR in heterodimer with RXR, the retinoid X receptor, and releases repression of a histone demethylase, which in turn activates the expression of genes for neural differentiation such as the homeobox transcription factor gene *dlx5* [5]. Another heterodimer binding partner of RXR, the oxysterol receptor LXR, has also been shown to promote neural differentiation, in particular of dopaminergic hindbrain neurons, while it prevents glial differentiation [5].

In addition to terpenoids, steroids bind to nuclear receptors that promote neural differentiation, for example the estrogen receptor (ER). Estrogen is a female sex steroid hormone, which originates in the same biosynthetic pathway as terpenoids, the mevalonate or HMG-CoA reductase pathway. It is not surprising that certain statins, inhibitors of HMG-CoA reductase and cholesterol-lowering drugs, inhibit self-renewal of ES cells, although the mechanism is not via nuclear steroid receptors but by preventing isoprenylation (geranylgeranylation) of RhoA [6]. Interestingly, simvastatin not only prevents ES cell self-renewal, but also induces osteogenic differentiation [7]. Estrogen, on the other hand, promotes differentiation and survival of dopaminergic neurons and therefore, it belongs to a group of neurogenic and neuroprotective steroids (neurosteroids) with great potential to be used in the clinics [8]. Another steroid of this group is the pregnancy hormone progesterone, which has been shown to alleviate neuronal damage in traumatic brain and spinal cord injury, to attenuate astrogliosis, and to enhance oligodendrocyte differentiation and myelination [9, 10]. While it is not clear whether these neurosteroid effects are solely induced by nuclear receptors or other non-nuclear receptors are involved, it is known that the majority of lipids in stem cell differentiation act via receptors in other subcellular compartments, in particular the cell membrane. This holds also true for cholesterol, the common precursor for the biosynthesis of steroid hormones.

Cholesterol is not only interesting as a metabolic precursor, but directly affects stem cell differentiation via two distinct mechanisms: it constitutes a post-translational modification covalently linked to one of the key morphogens in embryogenesis, sonic hedgehog (Shh), and it is a critical component of lipid rafts [11–14]. Sonic hedgehog has many important functions during embryonic and adult neurogenesis. In the embryo, it acts as a morphogen regulating the ventral patterning of the brain, while in the adult brain; it specifies the ventral position of neural stem cells. It has been shown that during embryogenesis, cholesterylation of sonic hedgehog is critical for establishing a morphogenetic gradient [15]. When administered to ES cells, Shh can stimulate the differentiation of two distinct progenitor populations: when combined with RA it promotes differentiation of ES cells to motor neurons, while subsequent incubation of RA+Shh-treated ES cells with fibroblast growth factor-2 (FGF-2) induces the generation of oligodendrocyte precursors (OPCs) [16, 17]. It is not clear whether this remarkably precise editing of *in vitro* ES cell differentiation recapitulates a physiologically significant process and which role cholesterylation of Shh may play. However, it becomes clear from this example that bioactive lipids not only act on distinct receptors, but can also modulate the activity of receptors for growth factors that are not specific for lipids. In particular, cholesterol is an important modulator of raft-associated growth factor receptors, which will be discussed in chapter 3.

1.2. Cell surface receptors: Lysophospholipids, eicosanoids, and endocannabinoids

Receptors for bioactive lipids can reside within all membranes or subcellular organelles of neural stem cells. So far, nuclear receptors for terpenoids and steroids have been discussed as inducers or suppressors of gene expression in neural differentiation. However, nuclear receptors are “blind” toward spatial cues. In contrast, receptors at the cell membrane can be activated by gradients of bioactive lipids (Fig. 1). This is important for the synchronization of neural differentiation with cell polarity, e.g., during neural migration or process

formation. The most prominent examples for lipids associated with cell polarity are lysophospholipids (LPLs) and phosphatidylinositolphosphates (PIPs). This section focuses on LPLs, while PIPs will be discussed in chapter 2.

LPLs are bioactive lipids that can be generated by hydrolytic cleavage of fatty acid from glycerophospholipids, a reaction catalyzed by phospholipases. Distinct phospholipases cleave off either one of the two (PLA1 and PLA2) or both (PLB) fatty acid residues, or they cleave off the phosphate-containing head group (PLC) or the alcohol (PLD) [18–26]. For example, PLD cleaves off the choline residue from phosphatidylcholine (PC) giving rise to phosphatidic acid (PA). PLA1 and PLA2 then generate lysophosphatidic acid (LPA) by cleaving off one of the two fatty acids from PA. Another enzyme, lysophospholipase D or autotaxin catalyzes a reaction similar to PLD; it generates LPA from lyso-PC, which is derived from PC by cleaving off one fatty acid residue, a reaction catalyzed by PLA2 [23, 27, 28]. While the enzymatic reactions generating LPA are well known, it is less clear which cells release LPA that can affect NSCs. In blood, activated platelets are a rich source for LPA and lesions or injuries, including those of the brain, may acutely release LPA to stimulate cell growth or migration. However, a systemic supply with LPA through the bloodstream as in the developing embryo would only affect NSC differentiation independent of spatial cues. In contrast, LPLs including LPA and S1P have been shown to act as chemotaxins, which is clearly demanding for defined albeit unidentified cellular source for neural development. LPA receptors are critical in cell proliferation and tumorigenesis and have recently been shown to promote proliferation of human neural precursor cells [22, 29–33]. In fact, LPA seems to inhibit neuronal lineage specification in favor of oligodendroglial differentiation. LPA receptors are GPCRs, which is a receptor family that also encompasses receptors for other bioactive lipids such as S1P, eicosanoids, and endocannabinoids (Fig. 2).

The biosynthesis of eicosanoids is initiated by the activation of PLA2, an enzyme that has a dual function: the generation of lyso-PC from PC and LPA from PA, but also the release of arachidonic acid, a fatty acid that is the precursor for the generation of eicosanoids [34–39]. Arachidonic acid is converted to a variety of pro-inflammatory eicosanoids among which prostaglandins, thromboxanes, and leukotrienes are the most important signaling lipids. The effect of eicosanoids on ES cells is not well understood and research is mostly limited to results with mouse ES cells. Due to their activity on GPCRs, LPA and eicosanoids such as prostaglandin E2 (PGE2) appear to activate similar downstream cell signaling pathways, mainly the PI3K-to-Akt/PKB, MAPK, and Wnt/GSK-3 β pathways [30, 32, 33, 40–43] (Fig. 2). The eicosanoid and LPL (LPA and S1P) receptors belong to the family of Class A Rhodopsin-like GPCRs [20, 22, 24, 32, 33, 44]. They mediate the activation of downstream cell signaling pathways through different types of GTPases, mainly Gi, Gq, and G12/13, acting upon PI3K-to-Akt/PKB (Gi), Ras-to-ERK (Gi, Gq) Rho (G12/13), and PLC-to-PKC (Gq) cell signaling pathways for pluripotency and cell survival (Akt/PKB), proliferation (Rho and PKC), and differentiation/specification (MAPK) pathways (Fig. 2).

The combination of cell signaling lipids with cytokines or growth factors such as LIF or FGF-2 activating similar effectors has been found to be useful in directing stem cell fate toward pluripotency, proliferation, or differentiation, respectively [24, 29, 30, 45, 46]. In contrast to activation of the receptors for LIF and FGF-2, LIFR α or FGFR2, however, stimulation of Akt/PKB by PGE2 has not been reported to sustain pluripotency, but is rather anti-apoptotic/cell protective and promotes stem cell proliferation. This may not be surprising since generation and conversion of arachidonic acid is often a response to hypoxic insults, which can damage mitochondria and induce apoptosis. Notably, inhibition of eicosanoid biosynthesis reduces the potential of mouse and human ES cells to self-renew, indicating a role of eicosanoids in stem cell maintenance or pluripotency [47]. Thromboxane has not been described to play a role in stem cell differentiation, maybe because its main

function is rather confined to platelet aggregation. In contrast, prostacyclin, a similar eicosanoid in platelet aggregation has been shown to promote cardiogenic differentiation from human ES cells [48, 49]. In addition to prostacyclin, leukotriene of the LTD₄ type has been used in several studies to promote proliferation and cardiovascular differentiation of mouse ES cells [39, 50, 51].

Most recently, two additional classes of bioactive lipids, endocannabinoids and particular poly-unsaturated fatty acids have been shown to stimulate gliogenesis and neuroblast migration (endocannabinoids) or promote neuronal differentiation from NSCs (docosahexanoic acid (DHA)), respectively [52–55]. Endocannabinoids such as anandamide (N-arachidonylethanolamine) and 2-arachidonoylglycerol bind to the CB1 receptor and activate the G α i subunit upstream of PI3K-to-Akt, while DHA binds to GPR40, a G α q subunit upstream of PLC γ (Fig. 2). Its activation generates DAG and IP₃, a signaling lipid mediating Ca²⁺ release from the ER. At this point, it becomes clear that many GPCRs, although binding specifically distinct bioactive lipids will activate very similar downstream cell signaling pathways, which may not be completely surprising considering that LPA, eicosanoids, endocannabinoids, and PUFAs are generated by interconnected metabolic pathways initiated by phospholipases. In this view, metabolically interconnected pathways constitute cell signaling niches since conversion of metabolic intermediates generates bioactive lipids with similar function for NSC differentiation. This view cannot be generalized when applied to S1P, another LPL triggering GPCRs. S1P is not generated in the phospholipase pathway, but it is derived from ceramide or sphingosine in sphingolipid metabolism. S1P and other bioactive sphingolipids will be discussed in chapter 2.

1.3. Receptors in the ER and other subcellular organelles

In addition to nuclear and cell membrane-resident receptors, bioactive lipids may also bind to receptors in other subcellular organelles. One of these receptors discussed in the context of NSC differentiation is the sigma receptor in the ER. The sigma receptor is highly enriched in mitochondria-associated membranes (MAM) of the ER and associates tightly with cholesterol, a lipid usually found at rather low concentration in the ER [56, 57]. It has been shown that the interaction with cholesterol is amino acid specific. However, since cholesterol is embedded in lipid microdomains or ER rafts this interaction is considered a lipid raft association. In addition, the sigma receptor can also bind to other steroids and sphingosine, but not to S1P [58]. Recently, the sigma receptor has gained attention because it binds to a multitude of psychopharmacological agents including opiates, antipsychotics, anti-depressants, and neurosteroids. This is consistent with the high abundance of this receptor in neurons and oligodendrocytes. Most surprisingly, however, is the ability of the sigma receptor to elevate and redistribute cholesterol and gangliosides, thereby reconstituting lipid rafts [59]. It is currently being investigated how this receptor-dependent remodeling of lipid rafts can regulate neuronal or glial function. One of these functions has been shown to directly involve oligodendrocyte differentiation and myelination [60]. The observation that cholesterol- and galactosylceramide-associated sigma 1 receptor is critical for the differentiation of oligodendrocyte precursors may lead to a novel drug target for treatment of dys- or demyelination diseases. The role of sigma receptors in the developing brain is not known.

2. Lipid mediators that affect protein kinase cascades and energy metabolism

The bioactive lipids discussed so far bind to specific lipid receptors and therefore, are mostly distinct from those binding to intracellular cell signaling proteins. The two lipids discussed here are phosphatidylinositols (PIPs) and ceramide. PIPs, in particular PI(3,4)P₂ and

PI(3,4,5)P₃ bind to specific protein domains such as the pleckstrin homology (PH) domain. These domains are present on a variety of protein kinases and other proteins that are regulated by PIPs. There also binding domains for ceramide, however, these have not been described as consensus sequences in protein kinases or phosphatases yet. Therefore, the specific effect of ceramide on these proteins, although well documented in many studies, is still lacking a clear molecular mechanism. Ceramide is distinct from PIPs in that it is a metabolic precursor for many other bioactive lipids that act via different mechanisms such as receptor binding (S1P) or lipid raft formation (glycosphingolipids or GSLs). This metabolic conversion makes it difficult to distinguish the biological effect of ceramide from that of its derivatives. One of these derivatives, S1P is unique in that it activates cell membrane receptors (S1P-specific GPCRs) as discussed in the previous chapter, but it can also bind to a variety of intracellular targets that will be discussed in this chapter.

PIPs and ceramide are signaling intermediates that are generated in response to intrinsic or extrinsic cues (Fig. 1). Extrinsic cues elevating these two bioactive lipids are often related to the regulation of the energy metabolism of an organism, organ, or cells within a tissue. For example, PIP₃ is generated by PI3K in response to insulin or IGF-1, and therefore, a lipid signaling saturation and availability of energy and reduction equivalents. In contrast, ceramide generated by activation of sphingomyelinase is rather a signal for starvation and oxidative stress. With this thought in mind, the interdependence of lipid mediators for NSC proliferation vs. differentiation and energy metabolism will be discussed in a new way that is often paid only little attention.

2.1. Phosphatidylinositols (PIPs)

The phosphatidylinositols PI(3,4)P₂ and PI(3,4,5)P₃ generated by class I phosphatidylinositol-3-kinase (PI3K) upon induction of tyrosine receptor kinases or G-protein coupled receptors (GPCRs) are known to be the major activators of the Akt/PKB cell signaling pathway for cell survival and differentiation [33, 45, 61–64] (Fig. 2). There are three functionally distinct groups of receptors activating PI3K in neural differentiation. The first group encompasses the GPCRs binding to other bioactive lipids, namely LPL (LPA and S1P), eicosanoid, endocannabinoid, and DHA receptors. Then there are receptors for growth factors such as FGF-2 and LIF, which are well known effectors of stem cell self-renewal and differentiation. Finally, there are receptors for growth factors such as insulin and IGF-1, which are not confined to stem cell differentiation, but interconnect NSC survival and differentiation with functions in general metabolism. Since the first group has already been discussed in the previous chapter, this section will focus on the second and third group of receptors activating PI3K.

In the cultivation process of ES cells, LIF and FGF-2 are used to maintain pluripotency of mouse or human ES cells respectively. These two growth factors elevate the expression of the transcription factors Oct-4 and Nanog, which is essential for maintenance of pluripotency [65, 66]. It has been shown that two cell signaling pathways are critical for this regulation: the janus kinase/signal transducer and activator of transcription 3 (Jak/Stat3) and the Akt/PKB signaling pathways [62, 67]. In the cultivation of mouse ES cells, the most important growth factor activating Stat3 and Akt/PKB is LIF (leukemia inhibitory factor), an interleukin 6 class cytokine binding to LIF receptor α (LIFR α) [68–72]. *In vitro*, LIF is added to the medium when cultivating undifferentiated mouse ES cells on feeder fibroblasts and in feeder-free culture. *In vivo*, LIF is generated by the trophoectoderm from where it penetrates the inner cell mass, the source of pluripotent ES cells in the pre-implantation embryo. In human ES cells, the role of LIF as “guardian” of pluripotency is taken over by FGF-2 [73, 74]. Binding of FGF-2 to the FGF receptor 2 (FGFR2) activates similar cell signaling pathways in human ES cells as stimulated by LIF in mouse ES cells: in particular MAPK and Akt/PKB [73, 74]. However, FGFR-dependent signaling is very diverse and it

depends on individual receptor protein complexes which specific response is elicited by FGF. For example, in mouse ES cells, FGF-2 is used to maintain the multipotent neuroprogenitor stage and to prevent further neuronal differentiation. In human ES cells, supplementation of the serum-free cell culture medium with FGF-2 is critical to prevent apoptosis and to maintain pluripotency.

The balance between PIP₃ and PIP₂ is regulated by the enzyme “phosphatase and tensin homolog deleted on chromosome 10” (PTEN). PTEN catalyzes the hydrolysis of PIP₃ to PIP₂, which leads to inactivation of the Akt/PKB cell signaling pathway [75–77]. It is a tumor suppressor mutated in many types of cancer and it is critical for the controlled growth of embryonic tissue and ES cells. Consistent with this function, deletion of PTEN activates Akt/PKB-dependent cell signaling pathways [75]. PTEN mutations are often found in human cancers such as glioblastoma, prostate cancer, and breast cancer. Loss of function of this tumor suppressor gene results in the up-regulation of the Akt/PKB-to- β -catenin pathway (Fig. 2A) [76]. Akt/PKB phosphorylates and inactivates glycogen synthase-3 β (GSK-3 β), a protein kinase in the Wnt signaling pathway that phosphorylates β -catenin [78–80]. The oncogene β -catenin is an important adhesion protein and transcription factor for genes involved in proliferation. When phosphorylated by GSK-3 β , β -catenin (in a protein complex with adenomatous polyposis coli or APC) is proteolytically degraded and thus, adhesion lost and proliferation reduced. Consistent with this function, deletion of β -catenin results in loss of pluripotency and early embryonic death of the respective knockout mouse [81]. Likewise, deletion of PTEN results in increased β -catenin levels and increased pluripotency or malignancy [75]. Therefore, the PTEN vs. PI3K-to-Akt/PKB antagonism is interesting in two biological contexts with respect to stem cell differentiation: maintenance of pluripotent stem cells and tumorigenesis of cancer stem cells. In the first context, inhibition of PTEN, activation of PI3K and Akt/PKB, or inhibition of GSK-3 β will be useful to maintain pluripotent ES cells. In the second context, activation of PTEN, inhibition of PI3K and Akt/PKB, or activation of GSK-3 β may be a useful strategy to eliminate cancer stem cells.

In addition to using natural lipids as ligands, ES cell and NSC differentiation can also be modulated by pharmacologic reagents that are either lipid analogs, inhibitors of enzymes in lipid metabolism, or drugs targeting downstream effectors of lipid-regulated cell signaling pathways. Two drugs that are inhibitors of protein kinases in the LIFR α and FGFR2 pathways have been tested on their effect on pluripotency: LY294002 and indirubin-3-monoxime, two inhibitors specific for PI3K and GSK-3 β , respectively [62, 65, 77, 82–86]. The PI3K inhibitor LY294002 has been shown to reduce the capacity of mouse and human ES cells to self-renew and to undergo subsequent steps of lineage specification and differentiation [62]. These effects are likely to involve differentiation stage-specific (contextual) other cell signaling pathways downstream (or parallel) to the PI3K-to-Akt/PKB signaling axis. While it may not be desired to interfere with ES cell pluripotency, LY294002 and other PI3K and Akt/PKB inhibitors are currently tested for cancer treatment, in particular for targeting cancer stem cells [87, 88]. If one desires to sustain self-renewal of ES cells, GSK-3 β inhibitors such as indirubin-3-monoxime or BIO are attractive candidates. BIO has been successfully used to maintain pluripotency in human ES cells [65]. Additional effectors targeting GSK-3 β are synthetic agonists of the Wnt receptor frizzled, however, their use in stem cell differentiation is not yet sufficiently investigated [86].

Interestingly, inhibitors of the MAPK pathway such as the MAPK kinase (MEK) inhibitor PD98059 have been used with mouse ES cells to promote self-renewal or pluripotency [74, 89]. This appears paradoxical since LIFR α as well as FGFR2 are known to activate MAPK, which suggests that activation of MAPK is involved in pluripotency. However, only transient MAPK activation to promote G1 re-entry is useful for self-renewal while prolonged activation will promote differentiation. Therefore, a combination of LIF with the

MEK inhibitor PD98059 activating PI3K-to-Akt/PKB while inhibiting MAPK signaling has been successfully used to promote pluripotency in mouse ES cells, but also to enhance the generation of induced pluripotent stem (iPS) cells [74, 86]. The situation in human ES cells, however, is different. In contrast to mouse ES cells, inhibition of the MAPK cell signaling pathway reduces the potential of undifferentiated human ES cells to self-renew, indicating that FGFR2-mediated activation of Ras/Raf-to-MEK-to MAPK is critical for human ES cell pluripotency [90]. A similar role has been found for Bmp4, which promotes pluripotency in mouse and differentiation in human ES cells [91–93]. It is quite possible that this difference depends on which other pathways for pluripotency are co-activated such as Jak/Stat3 in mouse or activin in human ES cells. Bioactive lipids are important in that they co-regulate several cell signaling pathways critical for pluripotency and differentiation of ES cells and NSCs (Fig. 2), in particular MAPK and PI3K downstream of ClassA/Rhodopsin-like GPCRs, as discussed in the previous chapter.

2.3. Ceramide and S1P

Ceramide is a fascinating lipid. Its level is highly regulated by three interconnected metabolic pathways, *de novo* biosynthesis from serine and palmitoyl-CoA, the sphingomyelin cycle in the cell membrane releasing ceramide from sphingomyelin, and the salvage pathway regenerating ceramide from sphingosine that was released by endo-lysosomal hydrolysis of sphingomyelin [94–96]. Moreover, this intensive metabolic turnover allows for fast remodeling of the fatty acid species in ceramide by six (dihydro)ceramide synthases in the ER (CerS1/lass1, C18 (dihydro)ceramide; CerS2/lass2, C20, C22, C24, and C26 (dihydro)ceramide; CerS3/lass3, C24 and C26 (dihydro)ceramide; CerS4/lass4, C18, C20, and C22 (dihydro)ceramide; CerS5/lass5, C16 (dihydro)ceramide; and CerS6/lass6, C14 and C16 (dihydro)ceramide) [97]. Since C18 ceramide is a major ceramide species in the brain it is not surprising that the mRNA expression level of CerS1 is about 50% of the total CerS mRNAs. Consistently, the CerS1 knockout mouse shows severe neurodevelopmental effects, particular in the cerebellum [98]. It has been reported that the portion of C18 ceramide increases in differentiating ES cells [99].

However, the attempt to correlate specific ceramide species such as C18 ceramide with a specific function in development is difficult since the CerS1 knockout mouse shows the up-regulation of another ceramide species (C16 ceramide), most likely as a compensatory effect. Therefore, the developmental defects may result from the loss of C18 ceramide or from the elevation of the other ceramide species. A similar effect was observed for the CerS2 knockout mouse that also shows neurodevelopmental defects and up-regulation of C16 as well as C18 ceramide [100–102]. It may be possible to get a better idea of the function of ceramide in development once double-knockouts for CerS1 and CerS2 (or other CerSs) have been generated.

In our laboratory, we have examined the function of ceramide in neural differentiation of ES cells with respect to the downstream targets of ceramide, in particular cell signaling pathways important for NSC differentiation. Three major regulators of differentiation signaling pathways have been shown to be activated by ceramide: protein phosphatase 1 (PP1) and PP2a (de-phosphorylates a variety of signaling protein kinases or other proteins), kinase suppressor of Ras (upstream of Erk1/2), and atypical protein kinase C (aPKC, upstream of Akt and GSK-3 β) [103–116]. Based on our data obtained with *in vitro* differentiating ES cells showing the critical function of the ceramide/aPKC interaction for NSC differentiation, we have focused on aPKC.

Atypical PKCs comprise a family of three protein kinase C isoforms, PKC δ (zeta), λ (lambda), and ι (iota). Knockout mice for each of these enzymes have shown that these isoforms have overlapping as well as highly specific functions in development. For example,

the conventional knockout for aPKC λ/λ is embryonic lethal, while aPKC δ -deficient mice survive, but show impaired insulin, NF- κ B, and IL-4 signaling [117, 118]. Conditional knockout of aPKC λ/λ has tissue-specific effects: in the retina this aPKC isoform is indispensable for retinal development and photoreceptor cell polarity, although another study has found that aPKC δ can functionally substitute for the λ/λ isoform [119, 120]. Interestingly, the specific function of aPKC for stem cell differentiation and embryo development seems to be cell- and tissue-specific: while a double-knockout of the δ and λ isoform does not affect hematopoietic stem cell polarity and differentiation to blood cells, inhibiting all isoforms of aPKC prevents lineage commitment of ES cells and sustains self-renewal [121, 122]. This observation is important with respect to our research since ceramide appears to bind and activate all of the aPKC isoforms (unpublished).

Prior to our ceramide-aPKC binding studies, it was known from *in vitro* enzyme assays that ceramide can activate (at low dose) or inactivate (at high dose) aPKC [112, 113]. It was not known whether this activation/inactivation reaction is mediated via a ceramide binding site and occurs in cells. It was also known that aPKC interaction with an endogenous inhibitor protein, prostate apoptosis response-4 (PAR-4) previously discovered in prostate cancer cells, induces apoptosis [123–125]. We demonstrated for the first time that ceramide activates and enhances autophosphorylation of aPKC in cells, but also induces inactivation of aPKC when PAR-4 expression is elevated [108, 111, 126–129]. Moreover, we showed that this mechanism (sensitizing cells to ceramide by expressing PAR-4) eliminates teratoma (stem cell tumor)-forming stem cells from differentiating ES cells and may also control the number of neural progenitors during brain development [126, 128]. Later on, we found that there is a ceramide-binding site in the C-terminus of aPKC, which is distinct from the modified C1 domain in the regulatory moiety of the enzyme [110, 130].

The C1b domain in aPKC is similar to the C1 domain in classical PKC α , which mediates binding to DAG and activation of PKC α . Prior to our study; it was hypothesized that the C1b domain may bind ceramide because of its structural similarity to DAG [131]. However, until recently, it has never been shown that ceramide can actually bind to the C1b domain. First evidence that this binding occurs comes from a study with KSR, which contains a C1b domain mediating activation of the enzyme by ceramide [107]. Therefore, it is likely that aPKC has two binding domains for ceramide, the C1b domain and the C-terminal domain identified in our studies. Although this idea is speculative, but two binding sites may tremendously enhance ceramide binding of aPKC since the free enthalpy for binding will be the sum of that for each binding site (or the combined dissociation constant the product of the individual constants). Binding of ceramide to aPKC may proceed in a similar way as binding occurs between DAG and PKC α or PKD: the C2 domain (in PKC α) binds to Ca²⁺/PIP₂ or phosphatidylserine (PS) and promotes binding of the C1a domain to DAG [132–134]. The pseudosubstrate domain unmask the catalytic site and client proteins can be phosphorylated. The DAG binding part of the C1a domain consists of prong-like β -sheets that form a binding pocket partially penetrating the inner leaflet of the cell membrane. While this pocket almost completely engulfs one DAG residue, it forms a tight interaction with hydrophobic and anionic lipid residues at its outside. One may envision a similar “hollow rivet”-like mechanism for binding of ceramide to the C1b domain.

We have hypothesized that that ceramide binding of aPKC is associated with PIP₂ or PIP₃ binding of Cdc42, a small GTPase interacting with the N-terminus of aPKC via the polarity protein Par6 [135–137]. Moreover, this mechanism could initiate a larger sphingolipid-protein scaffold (SLIPS) regulating cell polarity, which is known to critically depend on the aPKC-Par6-Cdc42 interaction [130, 135–142]. Finally, mutually exclusive binding of ceramide-aPKC to either PAR-4 or Par6-Cdc42 may constitute a “cell fate switch” we have proposed to control the balance between apoptosis (PAR-4) and cell polarity (Par6-Cdc42)

in NSCs [126, 135–137, 143, 144]. Therefore, it is not surprising that ceramide elevation is concurrent with apoptosis in “unwanted” or surplus stem cells and cell polarity of neural progenitor cells at the same differentiation stage [126, 128, 135–137, 144]. We have found that one mechanism distinguishing these two opposite cell fates is the unequal distribution of PAR-4 to the two daughter cells of a dividing stem cell: the one daughter cell that inherits PAR-4 is going to die, while the other one differentiates to a neural progenitor cell with ceramide sustaining cell polarity instead of inducing apoptosis [126].

How can ceramide affect cell signaling pathways in NSC differentiation apart from regulating cell polarity and apoptosis? We have proposed that two rheostats between ceramide and its derivatives are involved in this regulation: the balance between ceramide and S1P and that between ceramide and GSLs, the latter ones being discussed in chapter 3 of this review [135, 143–145]. Since elevating ceramide in polarized cells puts these cells at risk to undergo apoptosis, it is important to counteract the pro-apoptotic ceramide signal downstream of its client proteins. We have shown that ceramide activation of aPKC enhances phosphorylation and thereby inactivation of GSK-3 β , a mechanism critical for cell polarity [136]. However, ceramide is also known to inactivate Akt, which compromises GSK-3 β phosphorylation and reduces cell survival [106, 146, 147]. These two opposite effects of ceramide on GSK-3 β can be regulated by the ceramide derivative S1P. We have discussed in the previous chapter that S1P binding to GPCRs activates Akt, which then in turn phosphorylates and inactivates GSK-3 β (Fig. 2). Inactivation of GSK-3 β entails hypophosphorylation and therefore, stabilization of β -catenin, a protein with dual function. As transcription factor it is critical for neural differentiation of stem cells, while as cell polarity-associated protein it sustains adhesion of neural progenitors to the basal lamina in the subventricular zone (SVZ) of the embryonic brain.

It appears as if fine-tuning of the ceramide/S1P rheostat may actually imply that these two lipids with opposite function cooperate in favor of NSC differentiation. While ceramide sustains cell polarity and promotes differentiation, S1P increases cell survival. The critical role of ceramide for neural progenitor cell polarity has been reported in our recent studies showing that chemically induced ceramide deficiency completely abrogates the distribution of β -catenin to the basal lamina of the SVZ [136]. The β -catenin dislocation leads to aberrant neural progenitor migration and defective cortical lamination. Consistently, a water-soluble ceramide analog for the first time synthesized in our laboratory (N-oleoyl serinol or S18) rescues the β -catenin distribution, neural progenitor migration, and cortical lamination in ceramide-deficient embryos [108, 136, 148]. We have tested the combinatorial cell signaling effects of ceramide and S1P in neural progenitor cells. When differentiating ES cells are incubated with S18 and S1P or its prodrug analog FTY720, residual pluripotent, teratoma-forming stem cells are eliminated while the survival and differentiation of neural progenitor cells is enhanced [126]. Elimination of teratoma-forming stem cells is due to their high expression level of PAR-4, which is only weakly expressed in neural progenitor cells. The enhanced survival of progenitor cells is due to their cell surface expression of the S1P receptor S1P1, which is missing in the tumor stem cells [144].

Interestingly, further incubation with S1P or FTY720 promotes differentiation of neural progenitor cells toward oligodendrocyte precursor cells (OPCs), while prolonged incubation with S18 promotes neuronal differentiation [128, 143, 144]. This ceramide and S1P-regulated switch between neuronal and glial differentiation of neural progenitor cells is consistent with other studies showing that RA-induced neuronal differentiation of ES cells is concurrent with elevation of ceramide and that S1P and S1P1 is critical for survival and oligodendroglial differentiation of OPCs in primary cell culture [127, 149–152]. While our results and studies from other groups suggest that the effect of S1P (and also LPA) on ES cells and NSCs is mediated by GPCRs [29, 30, 143, 144, 153], it is quite likely that

intracellular targets of S1P are also involved in stem cell differentiation. For example, the inhibition of histone deacetylase 1 (HDAC1) by S1P synthesized by shingosine kinase 2 in the nucleus is certainly an important regulator for cancer cell proliferation, but may also promote differentiation of NSCs [154]. Therefore, it is expectable that many bioactive lipids such as ceramide and S1P regulate NSC differentiation by targeting several distinct mechanisms with similar outcome. This involves activation of lipid-specific receptors, binding of various intracellular cell signaling proteins, or forming lipid microdomains modulating the activation of growth factor receptors. Plus, lipids can be metabolically interconverted, turning them into highly dynamic and versatile regulators of various cell signaling pathways in NSC differentiation. The next section will elaborate further on this specific property of bioactive lipids.

2.4. Metabolic regulation of NSC differentiation by PIPs and ceramide

In recent years, it has become evident that the energy and redox state of stem cells, including NSCs is a critical factor controlling stem cell differentiation. As a rule of thumb: the more oxidized a stem cell the more likely it will undergo differentiation. Since the early days of ES cell culture, this was evident when pluripotent ES cells were maintained in the presence of β -mercaptoethanol thereby promoting the reduced redox state of the undifferentiated stem cell. In light of the previous sections, this curious observation can be embedded into a more scientifically grounded framework of connecting cell signaling pathways regulating stem cell differentiation with lipid metabolism. One of these pathways is the PI3K-to-Akt and Akt-to-GSK-3 β and mTOR cell signaling axis (Fig. 2). This has also been evident from the early days of ES cell culture since insulin is a critical component in N2, ITS, and B27 supplements added to serum-free media for the cultivation of ES cells and NSCs. Insulin triggers PI3K activation via the insulin receptor, which signals a saturated state: glucose is transported into the cell and converted into energy (ATP) and reduction equivalents (NADPH₂) promoting anabolic metabolism in NSCs [49, 77, 155]. In turn, cell survival (via Akt) and protein translation (via mTOR) is up-regulated, a prerequisite for the propagation of neural progenitors. IGF-1 can substitute for insulin with respect to PI3K activation and induction of neural differentiation [156, 157].

However, aerobic production of energy puts cells at risk to generate oxygen radicals or so-called reactive oxygen species (ROS). The most efficient way to eliminate ROS is the cellular production of reduced glutathione (GSH), which needs high levels of NADPH₂ for its regeneration. Therefore, low GSH levels due to high ROS or metabolic malfunction reducing NADPH₂ levels will promote aberrant differentiation of stem cells. This has been impressively demonstrated in studies showing that GSH sustains glial progenitor propagation while oxidation will promote the differentiation to reactive astrocytes, an inflammatory process commonly found in brain tissue lesion and neurodegenerative diseases such as Alzheimer's disease [158–162]. This mechanism also explains why maintenance of pluripotent stem cells or multipotent progenitor cells is often performed by cultivation of cells at low oxygen levels: while glucose supply is plenty for the production of reduction equivalents, generation of ROS is reduced. Further, astrocytes maintain a natural barrier between NSCs and neurons toward the oxygen-rich blood stream, while at the same time, providing these cells with GSH [163]. This separation is compromised in the case of brain tissue lesion, stroke, and neurodegenerative diseases leading to a sudden increase of ROS and the generation of reactive astrocytes that form a “glial scar” (gliosis).

How can lipids regulate this process? The regulation of metabolically induced proliferation vs. differentiation of stem and progenitor cells is related to two aspects of bioactive lipids: lipid metabolism and cell signaling. In particular, PIPs and other glycerophospholipids, and sphingolipids are important key players in the metabolic interconversion of bioactive cell signaling lipids. Diacylglycerides are synthesized from DAG, either by activating DAG

itself (CDP-DAG) or the alcohol/aminoalcohol such as choline (CDP-choline) or ethanolamine. For example, PIPs can be made from CDP-DAG and inositol, while phosphatidylcholine can be generated from DAG and CDP-choline. Phosphatidylcholine is then a precursor for the generation of sphingomyelin from ceramide, a reaction that is known to reduce inflammatory and pro-apoptotic cell signaling pathways [164]. Most recently, we have found that CDP-choline-driven detoxification of ceramide by its conversion to sphingomyelin can also protect neural crest stem cells and may be useful for treatment or prevention of fetal alcohol syndrome [165]. These studies and their implications for the metabolic compartmentalization of stem cell niches will be discussed in Conclusions.

At this point, one will consider these metabolic intermediates as bioactive cell signaling lipids that distinguish a saturated metabolism from a starving one. DAG as a sensor for ample supply with nutrients will directly activate PKC α and stimulate cell proliferation, while at the same time, enable the elevation of PIP₃ and reduction of ceramide, thereby increasing stem cell survival via activation of Akt. On the other hand, loss of nutrients and reduced activation of metabolic intermediates such as DAG or choline will deprive cells of precursors for phospholipid (i.e., phosphatidylcholine and sphingomyelin) biosynthesis, eventually leading to elevation of ceramide. As the consequence, stem cells will stop dividing and undergo differentiation or induce apoptosis. This assumption is in full agreement with our observations made with differentiating ES cells and embryonic mouse brain. At the transition from pluripotent EB-derived cells to multipotent neural progenitor cells or in embryonic mouse brain at gestation day E14.5, neural progenitors up-regulate ceramide, which is then converted to GSLs, in particular gangliosides (see chapter 3) [126, 127]. As discussed in the preceding section, we found that the peak of ceramide elevation does not induce apoptosis because neural progenitors show only low expression of PAR-4, the protein sensitizing cells to ceramide-induced apoptosis. Instead, ceramide is likely to function as a regulator of cell polarity in neural progenitors, thereby synchronizing NSC differentiation with the establishment of cell polarity.

The case of PIP₂ is a little bit more complicated since it is cleaved by PLC γ to DAG and IP₃, theoretically replenishing DAG in glucose-starved cells potentially activating mitogenic PKC α . However, IP₃ will bind to the IP₃ receptor (IP₃R) and induce the release of Ca²⁺ from the ER, which is a strong differentiation inducer as already discussed as one of the effect of GPCR-dependent activation of PLC γ . From this chain of events, it becomes evident that the cooperation of specific lipids with Ca²⁺ may be an important signal combination for NSC differentiation. There are at least three Ca²⁺ receptors or channels that can be regulated by bioactive lipids: IP₃R (IP₃), the ryanodine receptor (RyR, sphingosine and ceramide), the Na⁺/Ca²⁺ exchanger(s) (sphingosine, ceramide, and GM1), and the transient receptor potential channel 5 (TRPC5, LPC, S1P, and GM1) [166–169]. Most recently, ceramide has been found to regulate Ca²⁺ homeostasis in mitochondria by blocking the permeability transition pore, however, the exact mechanism is not clear yet [170]. Studies in Bob Ledeer's group have focused on TRPC5 and the Na⁺/Ca²⁺ exchanger in the nuclear membrane, which he has shown to be controlled by a specific ganglioside, GM1 [171–178]. This group of lipids and their function in neurogenesis and stem cell differentiation will be discussed in chapter 3.

3. Raft-forming lipids and raft-associated receptors in neural stem cells

The previous two chapters discussed bioactive lipids that are known to act through lipid receptors or binding proteins. Many lipids regulate cell signaling pathways through a mechanism known as “lipid rafts” or “lipid microdomains”. Lipid rafts are areas in the cell membrane (or intracellular membranes) that emerge from the self-assembly of lipids in an ordered (Lo) structure in the liquid phase of the membrane [14, 135, 179–184]. The concept

of lipid rafts goes back to studies by Gerrit van Meer and Kai Simons showing that GSLs are asymmetrically transported to the apical membrane of MDCK cells [185–187]. Simons then formulated the functional raft hypothesis in 1997 [14, 188]. This concept was preceded by that of lipid microdomains, which dates back twenty years earlier and was based on studies that are of importance for today's lipid raft research and understanding. In the mid 1970s, detergent-insoluble membranes enriched in cholesterol, sphingomyelin, and GSLs were isolated for the first time, a technique still used to prepare lipid rafts for the analysis of raft-associated lipids and proteins [189, 190]. About ten years later, fluorescence microscopy studies using filipin, a cholesterol cluster-binding reagent showed that the lipid distribution in the cell membrane is inhomogeneous with lipid microdomains of less and those of higher ordered structure [191–193].

The term “lipid microdomains” is now used interchangeably with that of lipid rafts, although rafts are commonly regarded as lipid microdomains of higher ordered structure. The minimum lipid composition of a raft is cholesterol and sphingomyelin, which is associated with GSLs or a glycerophospholipid such as phosphatidylcholine. In addition to lipid rafts, there are other highly ordered GSL-enriched microdomains (GEMs), so-called glycosynapses. Glycosynapses mediate cell adhesion and immunological responses and have been first described by Hakomori about 25 years ago [194–196]. Finally, caveolae, lipid rafts enriched with caveolin and important for endocytosis have also been found to contain GSLs. However, the functional significance of their presence is unclear since blocking GSL biosynthesis appears not to obliterate caveolae formation or function [197–199]. Although caveolae have not been found in neurons, genetic ablation of caveolin affects neural progenitor cell proliferation, indicating a role of this protein independent of caveolae formation or alternatively, a role of this type of rafts in NSCs prior to terminal neuronal differentiation [181, 200].

Lipid rafts are believed to show high affinity to specific cell signaling proteins such as growth factor or cytokine receptors, which leads to clustering and activation of these receptors (Fig. 1). Therefore, bioactive lipids can affect stem cell differentiation in two different ways: direct interaction with lipid receptors such as GPCRs and lipid raft-dependent activation of growth factor or cytokine receptors such as LIFR α , epidermal growth factor receptor (EGFR), IGFR, or FGFR2 [13, 135, 201–206]. In addition, receptor-associated signal transduction proteins such as Ras can be modified with fatty acids (palmitoylation) or terpenoids (farnesylation, geranylation) and glycosylphosphatidylinositol (GPI anchor), which tremendously increases membrane binding and raft association [14, 207–209]. It has been shown that particular GSLs termed gangliosides can regulate ES cell differentiation by the activation of FGFR2 and other receptors in lipid rafts [145, 205]. An example for this mechanism is the corrective activity of the ganglioside GM1 on the effect of the fungus toxin fumonisin B1, which causes neural tube defects by inhibiting sphingolipid biosynthesis [210, 211]. Gangliosides such as GM1 are commonly found in lipid rafts, although there is heterogeneity in their lipid composition potentially giving rise to distinct types of lipid rafts with specific function in neural differentiation. This review will focus on lipid rafts composed of sphingomyelin, cholesterol, and GSLs.

3.1. Glucosyl- and galactosylceramide

Phospholipids such as sphingomyelin as well as cholesterol are commonly found in highly ordered lipid rafts. Sphingomyelin can be hydrolyzed to ceramide, which disturbs this order and displaces cholesterol [212]. The resulting ceramide-containing microdomains or rafts have been proposed to have specific functions in cancer and neural differentiation of stem cells [135, 137, 213, 214]. Studies from our laboratory have shown that ceramide microdomains bind to cell polarity-associated aPKC, which has been discussed in chapter 2. The structural and functional variety of different rafts often stems from the composition of

specific GSLs. GSLs are synthesized by a multitude of ER- or Golgi-resident enzymes, which sequentially assemble a glycan structure to the polar head group of ceramide [145, 215–221]. The simplest GSLs are glucosyl- and galactosylceramide, more complex GSLs contain sialic acid residues and are called gangliosides. These will be discussed in the following section.

Galactosylceramide is the principal GSL in the brain. It comprises 23% of the total mass of myelin lipids [222]. However, in addition to its function in myelin, it has also been shown to be critical prior to myelin formation. In OPCs, galactosylceramide forms lipid rafts with cholesterol and sphingomyelin in the ER, which is instrumental for the activity of the sigma receptor [60] and OPC differentiation. Galactosylceramide is also the precursor for the biosynthesis of sulfatides, which comprise another 4% of the total lipids in myelin. Sulfatide and galactosylceramide are specific markers for OPC differentiation stages, which are recognized by the antibodies O1 (anti-galactosylceramide) and O4 (anti-galactosulfatide) widely used to enrich for pre- (O4(+)/O1(-)) or immature (O4(+)/O1(+)) oligodendrocytes, respectively. Interestingly, the O4 (but not O1) antibody is able to block terminal differentiation of OPCs, which indicates a functional role of cell surface sulfatide in oligodendrocyte differentiation [223]. It should be noted, however, that the function of the differentiation-stage specific expression of most GSLs and whether they act primarily as adhesion molecules or lipid raft modifiers in cell signaling pathways is still not known.

In contrast to galactosylceramide, which is synthesized in the ER, glucosylceramide is the precursor of a multitude of GSLs originating in the Golgi, including globosides and gangliosides (see next section). Cell surface glucosylceramide has been implicated in a variety of functions with respect to lipid raft and receptor regulation [181, 205, 224–226]. For example, rafts consisting of cholesterol, sphingomyelin, and glucosylceramide (CSG rafts) have been found to be associated with the activation of multidrug resistance (MDR) proteins in cancer [225]. We have suggested a novel concept of cancer treatment by breaking this CSG raft-dependent resistance in cancer stem cells [227]. Whether CSG rafts are also important in NSC differentiation is not known. However, it is known that stem cells persist throughout the whole life, which demands for a specific protection against environmental toxins. Therefore, it is not surprising that many stem cells express high levels of MDR proteins that may then be activated by CSG rafts.

Further insight into the biological function of glucosylceramide came from the use of specific pharmacologic inhibitors of glucosyltransferase, the enzyme synthesizing glucosylceramide from ceramide, and the generation of glucosyltransferase knockout mice. These two approaches clearly indicate the critical role of glucosylceramide transferase in insulin signaling and neural development, although this is likely to depend on the reduced biosynthesis of the ganglioside GM3 [228, 229]. Glucosylceramide is converted to lactosylceramide and then further sialylated to GM3, the precursor of a multitude of simple and complex gangliosides discussed in the next section.

3.2. Globo- and ganglio-series GSLs

One of the most frequently used surface markers for undifferentiated human ES cells is the globoside Gb5, also known as stage-specific embryonic antigen 3 (SSEA-3) [230–236]. It consists of a linear pentasaccharide structure (Glc-Gal-Gal-GalNAc-Gal) linked to ceramide. It can be converted to SSEA-4 by attaching a terminal sialic acid residue. Interestingly, in mouse, SSEA-4 is a marker for more differentiated (ectodermal) stem cells, while it is characteristic for the undifferentiated state in mouse ES cells [232, 237, 238]. A more thorough analysis of differentiation-stage specific GSLs showed that during differentiation of human ES cells, surface GSLs undergo a change from lacto- or globo- to ganglio-series, indicating an extension of the oligosaccharide to more complex GSLs in more differentiated

cells [239]. It has been reported that Gb5 is associated with lipid rafts, although it has not been shown to regulate their formation [240]. For globosides as well as gangliosides, it is still unclear how they affect NSC differentiation. One potential function could be that they serve as direct attachment sites for cell-to-cell contacts, a mechanism already discussed for glycosynapses. However, more likely is the GSL-dependent modulation of growth factor receptors, potentially within lipid rafts enriched with particular GSLs.

The simplest ganglioside, GM3 is synthesized by the sialylation of lactosylceramide, which comprises a ceramide residue linked to the disaccharide lactose. GM3 and its derivative GM1 are GSLs that are characteristic for GSL-containing lipid rafts. Knocking out GM3 biosynthesis leads to enhanced insulin sensitivity, which has been suggested to indicate a direct inhibition of the insulin receptor by lipid raft-associated GM3 [228]. Double knockouts deficient in the two glycosyltransferases that convert GM3 to GM2 (GM2/GD2 synthase) and GD3 (GD3 synthase) have not been reported to show enhanced insulin resistance, which may be due to a lack of GM3 accumulation in these mice [229]. However, these mice show neurodegeneration in peripheral motor and sensory nerves progressing with age, and reduced brain weight and cortical thickness, indicating a critical role of GM3-derived gangliosides in neural tissue development, maintenance or repair.

Gangliosides show amazingly profound and rapid changes during early embryonic brain development. Between gestational day E12 and E14, a large portion of GM3 is converted to GD3 and complex b-series gangliosides, which are further sialylated derivatives of GD3 [127, 215, 218, 220, 221, 241]. Later in embryonic and early post-natal brain development, GM3 is converted to its sialylated derivatives in the a-series complex gangliosides, in particular GM1. These observations suggest that increasingly complex sialylation of gangliosides is critical for NSC differentiation and brain development. Surprisingly, knockout mice deficient in sialyltransferases that synthesize these complex gangliosides did not show the severe phenotypes expected from them, although clear symptoms of aberrant neural function and neurodegeneration have been observed. For example, Bob Ledeen's group has shown that GM2/GD2-deficient mice that lack all complex gangliosides are far more susceptible to kainite-induced seizures and neuronal apoptosis than their wildtype counterparts [242]. Interestingly, the phenotype is rescued by the administration of LIGA 20, a synthetic and cell membrane-permeant analog of GM1. In the last 20 years, Ledeen's group has focused on the role of GM1 in neuronal function and found exciting results indicating a critical role of GM1 in the regulation of Ca^{2+} influx to the nucleus [169, 171, 172, 175, 176, 178, 243].

GM1 is the classical ganglioside used for staining of GSL-containing lipid rafts on the cell surface with cholera toxin subunit B. GM1 in rafts of the nuclear envelope was a novel concept brought forward and investigated by Bob Ledeen's group. He found that GM1 tightly associates with the Na^+/Ca^{2+} exchanger in the nuclear envelope and potentiates its ability to increase Ca^{2+} influx into the nucleus. LIGA 20 can restore this function in GM1-deficient knockout mice and protects neurons against kainite-induced seizures [242]. This example clearly demonstrates that gangliosides are abundant in many subcellular compartments and may regulate the activation of membrane-resident receptors, ion exchanges, and other proteins in lipid rafts distinct from those found in the cell membrane.

The generation of double knockouts for various pathways in ganglioside biosynthesis has now reached a stage that the naturally occurring redundancy in the function of sialylated GSLs preventing overt phenotypes can be avoided. A double knockout of GM3 synthase and GM2/GD2 synthase does not allow for compensation of a- or b-series complex gangliosides by the elevated biosynthesis of o-series gangliosides as observed with the GM3 single knockout mouse [244]. These mice are devoid of any ganglio-series GSLs and develop a

catastrophic neurodegenerative disease that results in death shortly after weaning. The phenotype observed, shrinkage of the brain, astrogliosis, and neuronal apoptosis in the cortex clearly indicates the essential role of gangliosides for maintenance of axon-glia interactions and other neural functions. This phenotype suggests a function of gangliosides similar to that proposed for sulfatides, as axon-glia attachment sites in the node of Ranvier, most likely by the interaction of GSLs with myelin-associated glycoprotein (MAG), which is essential for myelin stability [245, 246].

There is still much to discover for gangliosides, in particular for the function of GD3, GT3, and b-series complex gangliosides in NSC differentiation and brain development. The comparison of single with double-knockouts demonstrates that there is a very efficient compensation mechanism within the family of ganglio-series GSLs. Since during ES cell differentiation, the complexity of GSLs increases from lacto- and globo- to ganglio series GSLs, it is likely that more primitive states of NSC differentiation rely on the compensation within the families of simpler GSLs, e.g., globosides or sulfatides that could compensate for simpler gangliosides such as GD3. It is remarkable that some simpler gangliosides such as GD3 are highly specific for distinct NSC differentiation stages. Rober Yu's group discovered that GD3 is specifically expressed on the cell membrane of neural progenitor cells derived from the SVZ of embryonic mouse brain [215, 247, 248]. This group also showed that complex b-series gangliosides, in particular GD1b can protect neural cells against ceramide-induced apoptosis by regulating IGF-1 receptor-dependent Ca^{2+} influx [127]. Therefore, it is very likely that other functions of gangliosides will be found in NSC differentiation and brain development.

4. Conclusions and hypothesis: Developmental niches may be metabolic compartments for the generation of specific cell signaling lipids in stem cell differentiation

NSC differentiation can be regulated by a multitude of bioactive lipids that act on receptors, protein kinases/phosphatases and other cell signaling proteins, or lipid rafts in the cell membrane or other subcellular sites. At any given stage during differentiation, it is a cocktail of lipids that is simultaneously acting on stem cells. However, most of the research performed on NSCs neglects or overlooks the effects of lipids because of the artificial environment in which cells are cultivated *in vitro*. With the exemption of linoleic and linolenic acid, lipids are not considered essential in many media used for cell culture, although most companies specialized on stem and neural cell culture provide a lipid supplement of non-disclosed composition.

Lipids can become conditionally essential *in vitro* as well as *in vivo* depending on the metabolic capabilities of the cells. Like neurons, NSCs are living in a tissue niche, a compartment separating cells from unrestricted supply with blood-born nutrients and providing a defined environment with growth factors. For example, long-term cultivation of neurons *in vitro* requires serum-free Neurobasal medium with B27 supplement, a mixture of ingredients including lipids such as linoleic and linolenic acid, corticosterone and progesterone, and vitamin A and tocopherol. *In vivo*, however, neurons are far more needy and rely on supply with a multitude of intermediates such as citrate, lactate, serine, and glutathione provided by astrocytes [249]. A recent study found that neurons also receive their cholesterol from astrocytes, which secrete apoE type lipoproteins that are endocytosed by neurons [250]. While the metabolic compartmentalization between astrocytes and neurons has been intensely studied, only little is known about are potentially similar compartmentalization between NSCs and surrounding tissue in their developmental niche. The stem cell niches in the subventricular and subgranular zone (SVZ and SGZ) may very

well be metabolically compartmentalized: developmental niches may also be metabolic niches or compartments.

Sphingolipid metabolism is likely to be subject to compartmentalization in developmental niches such as the SVZ or SGZ niches for NSCs (Fig. 3). This hypothesis stems from our observation that neural crest-derived stem cells show ceramide elevation and apoptosis when exposed to ethanol, a potential cause for fetal alcohol syndrome and concurrent abnormalities of neural crest-derived tissues such as facial bone malformation [165]. Interestingly, *in vitro*, CDP-choline or citicoline, a precursor for phosphatidylcholine and subsequently, sphingomyelin biosynthesis from ceramide prevented ethanol-induced apoptosis in these cells. CDP-choline has also been shown to alleviate neurodegeneration and stroke damage in human trials [164, 251, 252]. Choline is a rich source for methyl groups required to synthesize phosphatidylcholine in the Kennedy pathway of glycerophospholipid biosynthesis (Fig. 3). We have published a model that integrates C1 (one carbon unit) with phospho- and sphingolipid metabolism, suggesting that stem cells such as neural crest-derived cells rely on continuous supply with methyl group donors to convert ceramide into sphingomyelin. In our model, this is mediated by the Kennedy pathway for *de novo* biosynthesis of phosphatidylcholine, which is the precursor for the generation of sphingomyelin from ceramide [135, 165] (Fig. 3). The proposed dependence of neural crest cells as a metabolically specialized compartment on the ongoing supply with methyl group donors to synthesize sphingomyelin is not obvious since one would assume that there is a sufficient amount of phosphatidylcholine coming from the bloodstream. On the other hand, our observations are consistent with previous studies showing that neuroepithelial cells in the neural tube are vulnerable toward interruption of sphingomyelin biosynthesis with the ceramide synthase inhibitor fumonisins B1 leading to neural tube defects [211].

Developmental niches such as the SVZ may induce a specialization of NSCs, e.g., through growth factor-dependent epigenetic regulation of gene expression, that is specific for the *in vivo* environment and results in a metabolic compartmentalization similar to that found between astrocytes and neurons. Figure 3 shows that this metabolic compartmentalization may very well determine the expression of bioactive lipids that have been discussed to regulate the differentiation of NSCs. Interestingly, the input metabolites such as serine or essential fatty acids are transported from astrocytes to neurons suggesting that the Kennedy pathway-to-sphingolipid metabolism integration comprises a metabolic compartment that is specifically activated in neurons (Fig. 3). Future studies will determine how similar supply with metabolites and metabolic compartmentalization may also affect bioactive lipid signaling in NSCs.

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Abbreviations

aPKC	atypical protein kinase C
DAG	diacylglycerol
EGF	epidermal growth factor
FGF	fibroblast growth factor

GSL	glycosphingolipid
IGF	insulin-like growth factor
IP₃	inositoltrisphosphate
LPA	lysophosphatidic acid
LPL	lysophospholipid
NSC	neural stem cell
PAR-4	prostate apoptosis response 4
PIP	phosphatidylinositolphosphate
PLC	phospholipase C
RA	retinoic acid
S18	N-oleoyl serinol
S1P	sphingosine-1-phosphate
SVZ	subventricular zone

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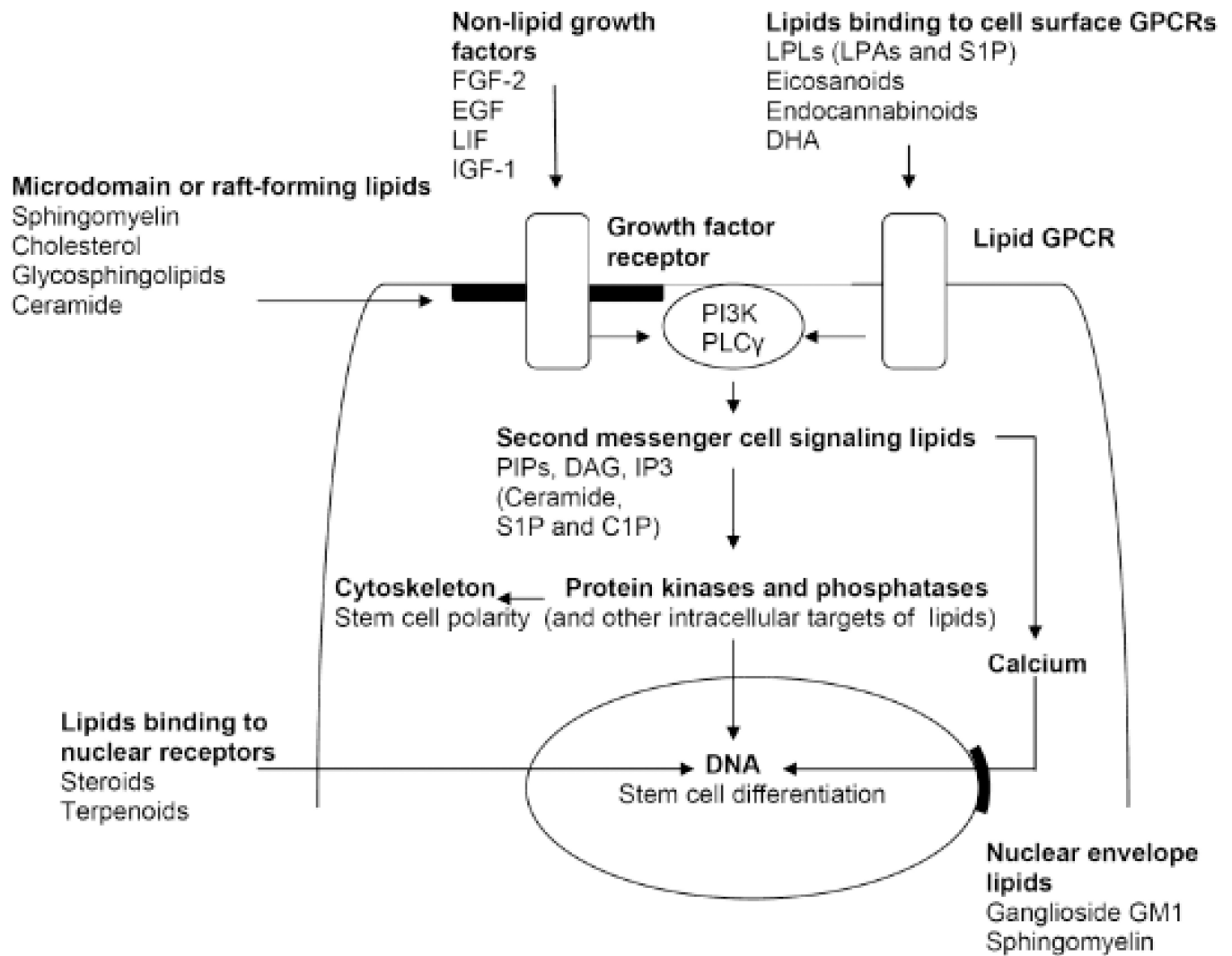


Figure 1.
Cell signaling pathways regulated by bioactive lipids

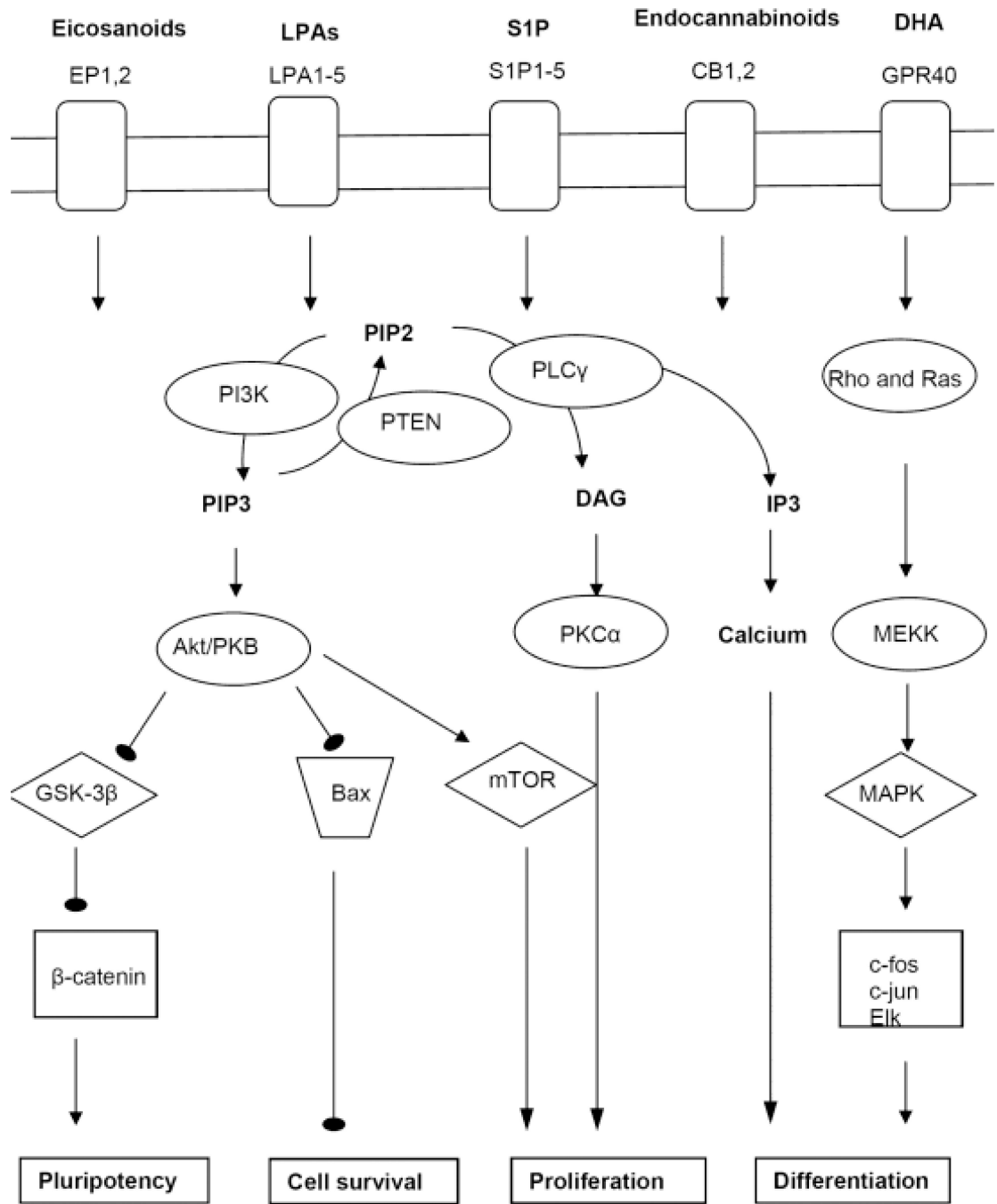


Figure 2.
Cell signaling pathways activated by lipid-specific GPCRs in NSCs

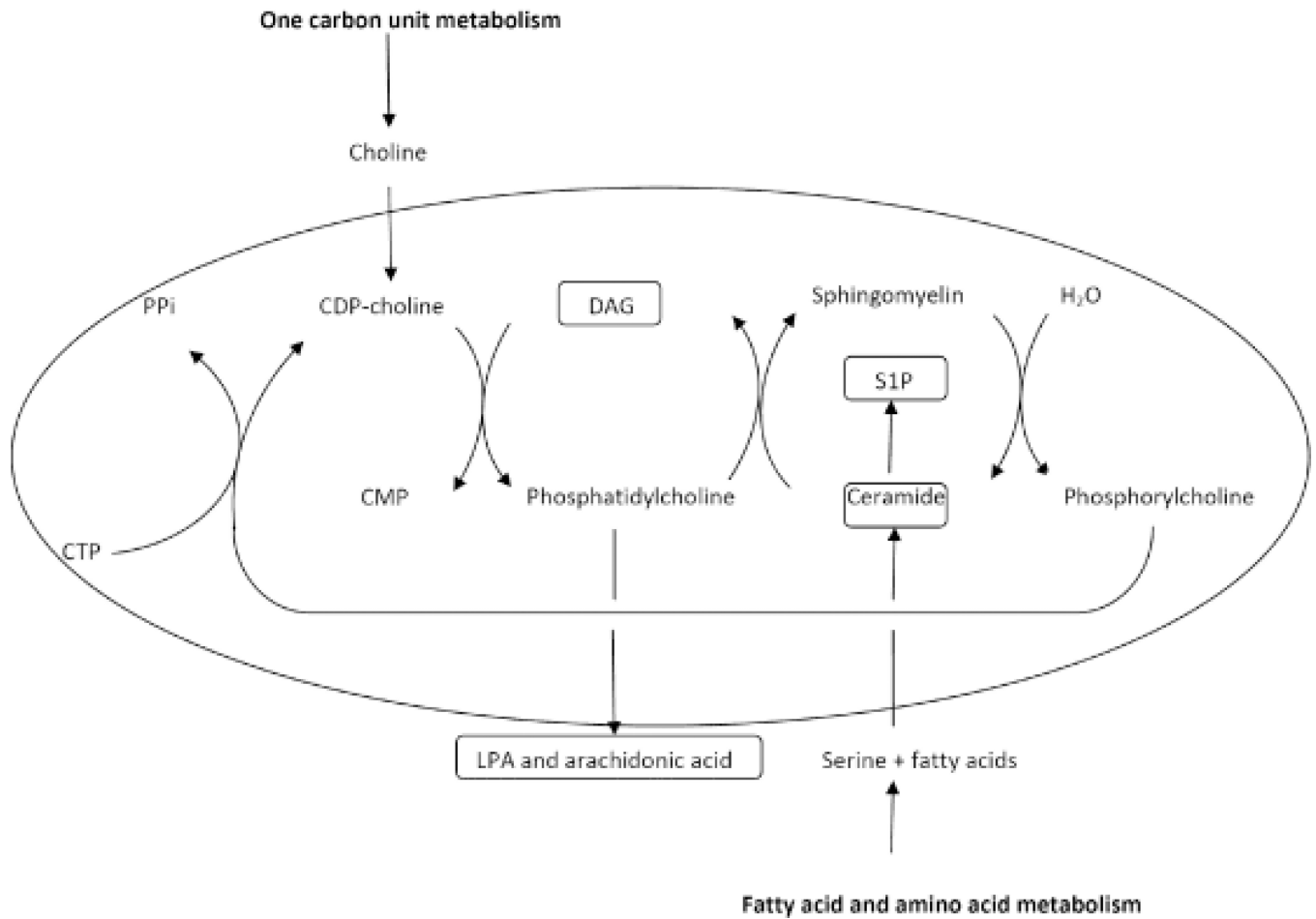


Figure 3. Metabolic compartmentalization of the Kennedy pathway for *de novo* glycerophospholipid biosynthesis and the sphingomyelin cycle in sphingolipid metabolism