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Lipid metabolism, remodeling and intercellular transfer in the CNS

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Abstract

Lipid metabolism encompasses the catabolism and anabolism of lipids, and is fundamental for the maintenance of cellular homeostasis, particularly within the lipid-rich central nervous system (CNS). Increasing evidence further underscores the importance of lipid remodeling and transfer within and between glial cells and neurons as key orchestrators of CNS lipid homeostasis. In this Review, we summarize and discuss the complex landscape of processes involved in lipid metabolism, remodeling and intercellular transfer in the CNS. Highlighted are key pathways, including those mediating lipid (and lipid droplet) biogenesis and breakdown, lipid oxidation and phospholipid metabolism, as well as cell–cell lipid transfer mediated via lipoproteins, extracellular vesicles and tunneling nanotubes. We further explore how the dysregulation of these pathways contributes to the onset and progression of neurodegenerative diseases, and examine the homeostatic and pathogenic impacts of environment, diet, and lifestyle on CNS lipid metabolism.

[H1] Introduction

The mammalian CNS is highly enriched in lipids, which constitute at least 50% of its dry weight, surpassing the lipid content of the liver and heart, but remaining lower than that of adipose tissue^{1,2}. Within the CNS, lipids such as **cholesterol** [G] and **fatty acid (FA)** [G] -containing lipid classes (such as **phospholipids** [G] and sphingolipids) have essential structural, functional, protective and metabolic roles that contribute to the maintenance of neuronal and glial cell physiology (**Supplementary Table 1**)(reviewed in REFs ^{1,3-6}). At the subcellular level, lipids are essential for maintaining cell membrane integrity and function, regulating signaling pathways, supporting organelle activity and organizing lipid rafts (dynamic lipid-rich microdomains in the plasma membrane that act as hubs for processes such as signal transduction and cell communication)⁷⁻¹⁰. At the cellular level, they play essential roles in energy storage and utilization and the stabilization of myelin sheaths, and facilitate processes such as oligodendrogenesis, neurogenesis and neuroprotection^{1,11-14}. At the tissular level, lipids are vital for synaptogenesis, efficient neuronal communication and the preservation of the integrity of the blood-brain barrier (BBB)^{5,15,16}. Finally, at the organ level, lipids are indispensable for brain development, cognitive function and processes involved in neuroinflammation and repair¹⁷⁻²⁰. Not surprisingly, perturbations in lipid metabolism are increasingly recognized to disrupt cellular physiology and cause cytodegenerative effects in the CNS^{1,21}.

The CNS has evolved complex molecular pathways to maintain lipid homeostasis in glial and neuronal cells, or to restore it in response to pathological disruptions (reviewed in REFs ^{21,22}). In this context, the metabolism, remodeling (intracellular modification and redistribution) and intercellular transfer of lipids have emerged as important processes for ensuring the efficient utilization and turnover of lipids in the CNS. In this Review, we first explore key pathways involved in intracellular lipid metabolism and remodeling, including the biogenesis and breakdown of the most abundant lipids in the CNS (namely cholesterol, FAs and phospholipids) and **lipid droplets** [G], as well as the oxidation of lipids. We then discuss the intercellular transfer of lipids, mediated by **lipoproteins** [G] and

extracellular vesicles [G]. The impact of disease on these metabolic processes, as well as how disruptions in these processes contribute to the development and progression of neurodegenerative diseases, is also addressed. Finally, we examine how environment, diet, and lifestyle influence lipid metabolism, remodeling and transfer in the CNS and how these changes may affect CNS disease. We focus in particular on multiple sclerosis (MS), Alzheimer disease (AD), Parkinson disease (PD), and Huntington disease (HD) because of the strong experimental and genetic evidence linking disrupted lipid metabolism to the onset of these diseases and the similarities in their lipid-related pathophysiology.

[H1] CNS lipid metabolism and remodeling

The CNS lipidome [G] is tightly regulated through coordinated bidirectional lipid fluxes between the periphery and the CNS, along with complex local metabolic and remodeling pathways. This section explores the metabolism and remodeling of cholesterol and FAs, which are essential components of the cellular membranes of neurons, glia and myelin. These lipids follow well-defined and key metabolic pathways within the CNS, making them fundamental to both CNS health and disease. We examine their synthesis, modification and degradation, as well the formation and breakdown of lipid droplets and the synthesis and remodeling of phospholipids. Disruptions in these processes are thought to contribute to the progression of CNS disorders.

[H2] Cholesterol and fatty acid synthesis

The brain is the most cholesterol-rich organ in the body²³, and its cholesterol content is tightly regulated to safeguard normal brain function. Given that cholesterol is unable to freely traverse from the bloodstream into the CNS, the brain and spinal cord rely on de novo cholesterol synthesis in glial cells and, to a lesser extent, neurons (**Fig. 1**)²⁴. The synthesis of cholesterol in the CNS is regulated by the sterol regulatory element-binding proteins (SREBP) and SREBP cleavage-activating protein (SCAP), which control the expression of genes involved in cholesterol production based on cellular

cholesterol levels (reviewed in REF ²⁵). In mammals, cholesterol biosynthesis mainly occurs between the perinatal period and adolescence, coinciding with the developmental ensheathment of axons by cholesterol-rich myelin²⁶. Accordingly, deficits in cholesterol synthesis in mice result in a marked delay in myelination^{12,27}. Once myelination is complete, cholesterol synthesis decreases and turnover slows^{28,29}, resulting in relatively stable brain cholesterol levels in adults²⁴. Despite reduced turnover, cholesterol synthesis (primarily in astrocytes, which supply cholesterol to adjacent cells³⁰) remains essential for CNS homeostasis in adults, with disruptions in this process linked to CNS complications. For example, reduced cholesterol synthesis is observed in postmortem brain tissue of HD patients³¹. Furthermore, in the aging human and mouse brain, a reduced cholesterol synthesis and progressively declining cholesterol content of myelin are associated with low-grade inflammation^{30,32-35}. Finally, inhibition of cholesterol biosynthesis via statins has been shown to impair remyelination in adult mice following cuprizone-induced demyelination³⁶⁻³⁸, and negatively impact neuronal physiology in vitro by reducing their viability and promoting neurite loss^{37,38}. However, other studies have reported neuroprotective and pro-regenerative effects of statins in humans and mice³⁹⁻⁴², and statins have been shown to reduce amyloidogenic processing of the amyloid precursor protein and formation of tau-containing neurofibrillary tangles in mouse models of AD⁴³⁻⁴⁷. The reasons for this discrepancy remain unclear but may include differences in model systems, statin concentrations, treatment windows and in the mechanisms of action of different statins^{25,42,48}.

Alongside cholesterol, and despite FAs being able to cross the BBB (**Box 1**), the CNS is capable of synthesizing **saturated fatty acids (SFAs)** **[G]** and **monounsaturated fatty acids (MUFAs)** **[G]** (**Fig. 1**), processes that (like cholesterol synthesis) are transcriptionally regulated by SREBP and SCAP (reviewed in REF ⁴⁹). The biosynthesis of SFAs and MUFAs occurs in the cytosol of both glial cells and neurons, with the SFA palmitate being a major intermediate product⁴⁹. Successive elongation and desaturation steps, catalyzed by elongases (ELOVLs) and desaturases ($\Delta 5$, $\Delta 6$, and $\Delta 9$), can subsequently generate FAs with varying FA chain lengths and degrees of unsaturation (reviewed in REF ²¹). ELOVLs add two-carbon units to the FA chain, while desaturases introduce double bonds at

specific positions within the carbon chain. While mammals cannot synthesize **polyunsaturated fatty acids (PUFAs)** [G] through de novo biogenesis, dietary PUFAs, such as linoleic acid and alpha-linolenic acid can be remodeled through elongation and desaturation. Supporting the importance of proper FA synthesis for CNS physiology, studies using genetic knockout models and pharmacological inhibitors have shown that CNS remyelination, adult neurogenesis, and developmental neuronal dendrite expansion are highly dependent on cell-autonomous FA synthesis by oligodendrocytes, neural stem cells (NSCs), and neurons, respectively⁵⁰⁻⁵³. On the other hand, studies have demonstrated that CNS demyelination increases the activity of ELOVL6 and the $\Delta 9$ FA desaturase stearoyl-CoA desaturase-1 (SCD1) in microglia and that this perturbs their pro-regenerative capacity and impairs remyelination^{19,54}. Similarly, disease-associated elevations in SCD1 activity have been reported to impair regulatory T cell differentiation^{19,55}, suppress NSC proliferation⁵⁶, and promote the pathogenic processing of neuronal α -synuclein (α -syn; a protein closely linked to PD, where its abnormal aggregation drives neuronal dysfunction and cell death)⁵⁷. These detrimental changes were found to enhance neuroinflammation and neurodegeneration in animal models of MS, AD, and PD⁵⁵⁻⁵⁷. Why SCD1 and ELOVL6 abundance is enhanced in these disorders is unclear but the increase may reflect compensation for the loss of long-chain MUFAs or altered lipid homeostasis due to disease-related metabolic changes (reviewed in REFs ^{21,58}). Overall, these studies underscore the important role of cholesterol and FA synthesis in maintaining and restoring CNS homeostasis, while demonstrating that disease-associated increases in FA elongation and desaturation are detrimental.

[H2] Cholesterol and fatty acid oxidation

As compared to other tissues, the CNS is highly enriched in **PUFAs** (reviewed in REF ²¹). This abundance can be explained by the essential role of PUFAs in supporting the structure and function of neuronal and glial cell membranes, and acting as precursors to signaling molecules such as endocannabinoids (reviewed in REF ⁵⁹). The function of PUFAs in the CNS is highly dependent on their oxidation, a form of lipid remodeling. This can occur non-enzymatically, when oxidants such as reactive oxygen species (ROS) interact with the carbon-carbon double bonds in PUFAs, a process

known as lipid peroxidation, or enzymatically, through the regulated conversion of PUFAs to their oxidized forms by enzymes such as lipoxygenases, cyclooxygenases or cytochrome P450 enzymes (reviewed in REF ²¹) (**Fig. 1**). Enzymatic oxidation of PUFAs results in the formation of lipid mediators such as prostaglandins, eicosanoids and specialized pro-resolving mediators (SPMs) — so called bioactive lipids^{60,61} — that play important roles in cell signaling events related to CNS disease progression and resolution. In this context, oxygenated ω 6 and ω 3 PUFAs, which differ in having their first double bond at the sixth and third carbon atom from the methyl end, are generally regarded as harmful and benign, respectively^{21,60,61}. However, exceptions to this simplified categorization exist; for instance, disease- or age-related reductions in lipoxin A4 — a bioactive lipid formed through the oxidation of the ω 6 PUFA arachidonic acid (ARA) — exacerbate astrocyte, microglia, and monocyte activation, impair neuronal function and promote T-cell infiltration and encephalitogenicity within the CNS of mice and humans⁶²⁻⁶⁴. The interaction of oxidized PUFAs with specific receptors, along with factors such as their concentration, tissue and cellular context, and the disease state and stage during which they are generated, collectively determine their eventual impact in the CNS (reviewed in REFs ^{61,65}). In contrast to the controlled enzymatic oxidation of PUFAs, the uncontrolled peroxidation of membrane PUFAs in phospholipids driven by excessive ROS production in CNS disorders, leads to membrane dysfunction and iron-dependent necrotic cell death, a process known as ferroptosis (reviewed in REF ⁶⁶). Such oxidized phospholipids have been shown to be associated with neuroinflammation and to compromise the viability of neurons and oligodendrocytes in various CNS disorders^{67,68}.

Similar to PUFAs, cholesterol is highly susceptible to oxidative modification through both enzymatic and non-enzymatic pathways, leading to the formation of oxysterols that have pleiotropic functions (**Fig. 1**). Cytochrome P450 46A1 (CYP46A1)-mediated oxidation of cholesterol is a key homeostatic pathway in neurons, leading to the formation of 24(S)-hydroxycholesterol (24S-OHC), which readily diffuses across the BBB for disposal^{24,69,70}. Since unmodified cholesterol cannot freely cross the BBB, this pathway is essential for cholesterol turnover in the CNS. Brain cholesterol can also be oxidized by

CYP27A1, expressed in neurons and glial cells, to form 27-OHC, which is also capable of crossing the BBB⁷¹. Alongside regulating cholesterol flux out of the brain, 24S-OHC and 27-OHC control the metabolism and intracellular transfer of other CNS lipids through the activation of astrocytic nuclear liver X receptors (LXRs)^{71,72}, as discussed in later sections. Even though these oxysterols are essential for CNS homeostasis, substantial evidence indicates that they can also promote disease progression in individuals with AD and AD models by enhancing neuroinflammation and increasing amyloid β (A β) and hyperphosphorylated tau levels⁷³⁻⁷⁶. They have also been shown to promote lysosomal membrane permeabilization-mediated pyroptosis and to negatively affect morphology and synaptic function in hippocampal neurons⁷⁷⁻⁷⁹. A recent study further revealed that 27-OHC promotes oligodendrocyte precursor cell (OPC) differentiation while simultaneously exhibiting toxicity towards immature oligodendrocytes, which was found to impair cognitive performance in mice⁸⁰. The contrasting homeostatic and disease-promoting effects of these oxysterols are poorly understood, but it is hypothesized that disease-associated elevations contribute to the detrimental impact of oxysterols in CNS disorders. In addition to 24S-OHC and 27-OHC, various other oxysterols are present in the brain, arising from the auto-oxidation of cholesterol induced by compounds such as free radical species. This results in the formation of oxysterols such as 7 α -OHC, 7 β -OHC, 25-OHC, and 7-ketocholesterol (7-KC). Among these, 7-KC stands out as one of the most abundant, biologically active, and toxic oxysterols in the CNS (reviewed in REFs ^{81,82}).

In conclusion, the remodeling of FAs and cholesterol through oxidation generates oxygenated lipids, whose effects on CNS diseases are shaped by their abundance and by the mode of oxidation (enzymatic vs. non-enzymatic).

[H2] Lipid droplet anabolism and catabolism

In addition to their oxidation, FAs and cholesterol can be remodeled to produce a variety of other lipids with important cellular functions. These include neutral lipids, such as cholesteryl esters (CEs), diacylglycerols (DAGs) and triacylglycerols (TAGs), which can be contained within lipid droplets. Lipid

droplets are dynamic intracellular organelles composed of a hydrophobic core of neutral lipids, surrounded by a phospholipid monolayer. This monolayer is embedded with lipid droplet-associated proteins (LDAPs) that regulate lipid droplet function, of which perilipins (PLINs) represent the largest family⁸³. Although the primary role of lipid droplets is the storage of energy (in the form of lipids), emerging research has revealed that lipid droplets play an essential role in intracellular signaling and metabolism by establishing functional contact sites with other organelles, such as mitochondria and the endoplasmic reticulum (ER), to facilitate metabolic crosstalk and regulation^{84,85}. While the quantities of lipid droplets present in the healthy CNS are typically low, substantial accumulations resulting from an imbalance between lipid uptake, synthesis, and mobilization, are observed in the aging brain and in neurodegenerative disorders such as MS, AD, PD and HD, primarily in glial cells⁸⁶⁻⁸⁸. This dysregulation can arise from disease-associated changes in lipid droplet formation and breakdown^{19,89,90}.

Lipid droplet biogenesis begins at the ER, where neutral lipids are synthesized primarily through the actions of acyltransferases⁹¹. These enzymes catalyze the esterification of DAGs, which are derived from the breakdown of phospholipids or through de novo synthesis, and cholesterol with fatty acyl-CoA, forming TAGs and CEs, respectively. As these neutral lipids accumulate in the ER membrane, they promote localized lipid droplet formation, resulting in the budding and release of lipid droplets into the cytosol. Once in the cytoplasm, lipid droplets can expand by incorporating additional neutral lipids or by fusing with other lipid droplets. Aberrant lipid droplet formation in the CNS has been suggested to be linked to neurodegenerative disease, with the most compelling evidence for this in AD. Specifically, the apolipoprotein E4 (*APOE4*) allele, the strongest genetic risk factor for AD, promotes lipid droplet formation in both glial cells and neurons^{92,93}. This increase in lipid droplet formation is thought to be driven, at least in part, by the upregulation of genes encoding proteins involved in lipid synthesis, such as acyl-CoA synthetase, which plays an essential role in the generation of fatty acyl-CoA⁹³.

During cell growth or in situations of energy deprivation, lipid droplets can be mobilized and broken down for energy production and the synthesis of membrane components and signaling molecules (reviewed in REF ⁹⁴). Breakdown of lipid droplets occurs through two main pathways: lipolysis and lipophagy (**Fig. 1**). Lipolysis involves the hydrolysis of TAGs into free FAs and glycerol by the consecutive actions of adipose triglyceride lipase (ATGL), hormone-sensitive lipase (HSL), and monoacylglycerol lipase (MGL). The importance of lipolysis for CNS function is demonstrated by recent findings that microglia in actively demyelinating lesions from people with MS show increased abundance of PLIN2, which inhibits lipolysis-mediated degradation of lipid droplets in these cells⁹⁵. In preclinical models of MS, loss of PLIN2 reduced lipid droplet accumulation and induced a less-inflammatory microglia phenotype, thereby improving remyelination. These findings position PLIN2 as an LDAP that prevents lipid droplet degradation, similar to PLIN5, whose interaction with ATGL inhibits lipid droplet lipolysis⁹⁶. The molecular drivers of the increased PLIN2 abundance in microglial lipid droplets during CNS demyelination remain unclear, but it is possible that CNS inflammation increases microglial PLIN2 expression⁹⁷. This may be a protective response, because blocking ATGL-mediated breakdown of lipid droplets in microglia has recently been shown to alleviate neuroinflammatory and behavioural responses to lipopolysaccharide⁹⁸, suggesting that PLIN2-mediated inhibition of lipolysis during demyelination protects against excessive neuroinflammation. This feedback mechanism may be beneficial early after demyelination but could become detrimental during chronic demyelination, when abundant microglial lipid droplet formation is evident and associated with the induction of a harmful foamy microglia phenotype^{19,54,89,99}. Neurons also express an additional specific lipid hydrolase, DDHD domain-containing 2 (DDHD2), which initiates lipolysis independently of ATGL and whose deficiency is linked to neuronal TAG deposition and hereditary spastic paraplegia¹⁰⁰. However, the functional relevance of neuronal lipolysis remains unclear, as neurons typically do not form abundant lipid droplets¹⁰¹.

In addition to lipolysis, lipid droplets can be degraded by a specific form of **autophagy** **[G]**, called lipophagy¹⁰², a process that has gained substantial attention for its role in driving lipid metabolism in

the CNS (reviewed in REFs ¹⁰³⁻¹⁰⁵). The autophagic degradation of lipid droplets involves their sequestration into autophagosomes, which subsequently fuse with lysosomes for degradation by lysosomal acid lipases (**Fig. 1**). Disruption of autophagy is an important pathological hallmark of various neurodegenerative conditions and aging, and increasing evidence suggests that impaired lipophagy may underlie abnormal lipid droplet accumulations in these conditions. For example, it was recently demonstrated that macrophages and microglia in active lesions of people with MS, which contain abundant remnants of intracellular myelin debris, exhibit signs of impaired lipophagy¹⁰⁶, leading to excessive lipid droplet accumulation and the adoption of a highly inflammatory, repair-inhibitory phagocyte phenotype. Inducing lipophagy with the autophagy activator trehalose reversed this harmful metabolic and inflammatory microglia phenotype, and promoted remyelination in preclinical models of MS. A direct link between impaired lipophagy and HD has also been reported: Huntingtin (HTT), the gene product responsible for HD (and which in its mutated form creates toxic aggregates that cause neuronal damage¹⁰⁷) has been shown to function as a scaffold protein for the selective autophagy of lipid droplets¹⁰⁸. By physically interacting with the autophagy cargo receptor p62 to facilitate its association with the autophagosome, HTT was found to be essential for at least three types of selective autophagy: lipophagy, mitophagy, and aggrephagy. Accordingly, loss of *Htt* in mice led to increased sensitivity to lipid challenges and a more pronounced accumulation of lipid droplets.

The fate of the lipids released by lipolysis and lipophagy is multifaceted. Once lipid droplet-associated neutral lipids are converted to free FAs, these can be degraded through mitochondrial **β -oxidation** **[G]** to meet cellular energy needs^{102,109}. The acetyl-CoA that is produced from this β -oxidation could subsequently be utilized for de novo lipogenesis¹¹⁰. Alternatively, lipid droplet processing via lipophagy and lipolysis can facilitate the intercellular transfer of free cholesterol and FAs^{111,112}. Notably, recent findings indicate that enhancing lipophagy promotes microglia lipid efflux within demyelinating brain lesions¹⁰⁶. Catabolism of lipid droplets can also release or lead to the generation of signaling lipids. For example, unmodified free FAs may serve as endogenous ligands for

peroxisome proliferator-activated receptors (PPARs)^{55,113}, while oxidized PUFAs are potent signaling molecules^{60,61}. Finally, lipid droplets can act as a readily available lipid reservoir, from which lipids can be released and used for membrane formation¹¹⁴. Collectively, these findings underscore the important roles of lipid droplet catabolism and anabolism in controlling lipid remodeling in the CNS. **Tracer lipidomics [G]** studies are, however, required to fully define the fate of the lipids released by lipolysis and lipophagy.

[H2] Phospholipid synthesis and remodeling

Phospholipids, including phosphatidylcholines (PCs) and phosphatidylethanolamines (PEs), are key components of cellular membranes in the CNS, ensuring structural integrity, fluidity, and proper membrane functionality (reviewed in REF ¹¹⁵). Their de novo synthesis occurs in all cell types via the Kennedy pathway, which converts choline and ethanolamine through successive enzymatic steps into PC and PE, respectively (**Fig. 1**)¹¹⁶. The final enzymatic step in this pathway involves the addition of a DAG, which can be synthesized de novo from glycerol-3-phosphate (G3P) or derived from the catabolism of existing phospholipids or TAGs stored in lipid droplets, among other sources (reviewed in REF ¹¹⁷). The composition of the DAG molecule that is added, which is defined by the two FAs that are esterified to its glycerol backbone, directly influences the type of phospholipid that is synthesized, providing a form of lipid remodeling and diversifying the lipidome at the phospholipid level. Despite the abundance and tremendous compositional diversity of phospholipids in the CNS⁶, direct evidence linking disturbances in the Kennedy pathway to glial cell and neuronal dysfunction in neurological disorders is currently lacking. Choline supplementation studies, however, suggest that limited availability of the substrates of the Kennedy pathway may be a key factor driving chronic neuroinflammation and failure of repair in preclinical MS models¹¹⁸. The quantity and composition of PCs and PEs are further known to impact glial cell and neuronal activity, plasticity, and survival^{1,119-122}, and alterations in the levels and composition of these phospholipids are associated with the pathology of most neurodegenerative disorders^{1,68,123}. Notably, phospholipid synthesis also facilitates the formation of the autophagosomes that engulf organelles during autophagy¹²⁴. Hence, in addition

to the active remodeling of the phospholipidome, the Kennedy pathway may indirectly affect lipid remodeling by influencing lipophagy.

The highly diverse and asymmetrical distribution of FA chains in phospholipids cannot be fully achieved by de novo synthesis via the Kennedy pathway. Instead, phospholipids in cellular membranes are postulated to be metabolically active and — in a process known as the Lands cycle — to continuously undergo deacylation and reacylation reactions. This results in the dynamic incorporation of PUFAs, particularly ARA and docosahexaenoic (DHA), into the phospholipids (reviewed in REF ¹²⁵). Mechanistically, the FA chains of de novo-synthesized phospholipids are hydrolyzed by divergent phospholipase A2 (PLA2) isoforms, generating lysophospholipids. These lysophospholipids are then reacylated by lysophospholipid acyltransferases (LPLATs), resulting in the incorporation of a new FA and the formation of a new phospholipid (**Fig. 1**). Reflecting its high activity level in the brain¹²⁶⁻¹²⁸, alterations in the Lands cycle have increasingly been linked to disruptions in CNS homeostasis. For example, elevated PLA2 activity is associated with neuroinflammation, neurodegeneration, and cognitive deficits in mouse models of AD, PD, and MS, and in people with MS¹²⁹⁻¹³⁵. Interestingly, A β has been shown to enhance PLA2 levels, which may, at least in part, contribute to A β toxicity^{133,134}. Conversely, loss of the *PLA2G6* isoform shortens lifespan, impairs synaptic transmission, promotes α -syn aggregation, and induces neurodegeneration in cell culture and animal models of PD^{136,137}, potentially by causing ER stress¹³⁸, while, *PLA2G6* mutations are associated with infantile neuroaxonal dystrophy (INAD), neurodegeneration with brain iron accumulation (NBIA), and familial type 14 PD (PARK14)^{139,140}. The dual harmful and benign roles of PLA2 isoforms remain poorly understood, but could reflect disease stage-dependent changes in PLA2 isoform abundance and isoform substrate specificity^{141,142}. In contrast to PLA2 isoforms, research on LPLATs in CNS disease is sparse. *LPCAT1* is elevated in postmortem brain tissue from people with PD, and its suppression reduces α -syn accumulation and toxicity in a human PD cell culture model¹⁴³. Furthermore, overexpression of *LPEAT2*, which uses DHA as a substrate, induces necrotic cell death in neuronal cells in vitro¹⁴⁴. Collectively, ample evidence suggests that the Lands cycle is dysregulated

in various CNS disorders, though the mechanisms and implications of this remain unclear. Future studies should explore whether altered Lands cycle activity arises from disease-related changes in the compositional requirements of membrane phospholipids, serves to detoxify peroxidated membrane phospholipids, or reflects a need to generate lipid signaling metabolites. Notably, while outside the scope of this review, lipid remodeling similar to that seen in phospholipids is involved in the metabolism of many other lipid classes, underscoring the complexity and importance of these processes in maintaining cellular function in the CNS¹⁴⁵.

[H1] Intercellular lipid transfer in the CNS

The ability of astrocytes and microglia to supply neurons and oligodendrocytes with lipids has been appreciated for many years¹⁴⁶⁻¹⁴⁸. However, emerging evidence has indicated that this intercellular lipid shuttling is not unidirectional, with neurons also being able to transfer lipids to astrocytes¹⁰¹. The most well-known process driving intercellular lipid transfer in the CNS is lipoprotein-mediated lipid flux, in which astrocytes and microglia synthesize and secrete lipoprotein particles that are composed of cholesterol and a neutral lipid core, and are surrounded by a phospholipid monolayer and apolipoproteins, primarily APOE (reviewed in REF ¹⁴⁹). Neurons and oligodendrocytes take up these particles and utilize the lipids for dendrite and axonal growth, synapse formation and communication, and myelin membrane maintenance and formation^{146-148,150}. While they can also use these lipids as an energy source, this is likely to be limited due to their relatively low capacity for β -oxidation as compared to glia¹⁵¹. The essential role of lipoprotein-mediated lipid flux for CNS function is made particularly evident by studies of AD, where genetic variants of *APOE* and triggering receptor expressed on myeloid cells 2 (*TREM2*), encoding a receptor involved in the uptake of lipidated APOE, disrupt CNS lipid metabolism and drive disease progression (**Box 2**). Alongside lipoproteins, emerging evidence suggests that other mechanisms also control intercellular lipid fluxes in the CNS, including extracellular vesicles (EVs). In this section, we will discuss the key role of these intercellular lipid transfer mechanisms in maintaining CNS homeostasis and how perturbations in their activity can drive CNS pathologies.

[H2] Lipid transfer between glia and neurons

Glia-to-neuron lipid transfer plays an essential role in ensuring proper neuronal function, with the release of lipidated APOE by glial cells, followed by its uptake by neurons, being well-documented (reviewed in REFs ^{152,153}). Consistent with this, astrocyte-specific deficiency of *Scap*, a key regulator of de novo lipid biosynthesis (**Fig. 2**), impairs presynaptic terminal development, reduces neurite outgrowth, and leads to behavioral and motor defects in mice in vivo^{154,155}. Similarly, astrocytes in a mouse model of HD exhibit impaired cholesterol biosynthesis and secrete reduced levels of APOE-bound cholesterol, which hinders neurite outgrowth and synaptic activity¹⁵⁶. Enhancing astrocytic lipid biosynthesis and efflux was found to reverse these neuronal dysfunctions¹⁵⁶.

Studies also indicate that neuron-to-glia lipid transfer is a continuous and essential process in the CNS. The transfer of neuron-derived 24S-OHC to astrocytes activates APOE-mediated cholesterol efflux in these cells through an LXR-regulated pathway⁷². This 24S-OHC flux is likely to serve as a mechanism for neurons to communicate their cholesterol requirements to astrocytes, as astrocytic APOE-associated cholesterol is a major source of cholesterol for neurons. More recent studies indicate that neurons also transfer FAs to astrocytes^{90,157}, which is important because neurons typically lack the machinery to form lipid droplets (and thus to store FAs)¹⁰¹. Genetic and pharmacological studies have shown that, under homeostatic conditions, astrocytes take up blood glucose, convert it to lactate and transfer it to neurons via the monocarboxylate transporter¹⁵⁷. Neurons metabolize the lactate to produce acetyl-CoA, a key substrate for energy production and de novo FA synthesis. The neurons maintain a stable and healthy FA lipidome by exporting excess FAs to astrocytes, which oxidize these lipids for energy¹⁵⁷. It has been shown that elevated ROS levels promote the use of glial lactate to fuel neuronal lipogenesis, resulting in the formation of peroxidated FAs that are exported to, stored in, and catabolized by astrocytes¹⁵⁷. While transient increases in ROS and peroxidated lipids can be buffered by astrocytes, the authors hypothesized that chronic and excessive ROS release, such as that seen in some CNS disorders, could overwhelm this

system, leading to astrocyte dysfunction and neurodegeneration. They further showed that neuron–glia lipid transfer and astrocytic lipid droplet formation depend on APOE. Notably, APOE4 was less efficient in facilitating neuron–glial lipid transfer and impaired the formation of lipid droplets in glial cells as compared to other APOE isoforms. This provides a molecular rationale for the increased neuronal lipid peroxidation observed in individuals with AD that carry *APOE4* alleles¹⁵⁷. Supporting this notion, another study demonstrated that APOE4 hinders neuronal FA sequestration in lipid droplets, impairs the flux of lipid droplet-associated FAs from neurons to astrocytes, and reduces FA oxidation in astrocytes¹⁵⁸.

A recent study showed that lipid fluxes between neurons and astrocytes are also essential for protection against the adverse effects of neuronal hyperactivity¹⁵⁹, a hallmark of neurological disorders such as AD¹⁶⁰. Hyperactive neurons, displaying increased lipid peroxidation^{161,162}, were found to avoid lipid toxicity by transferring toxic lipid intermediates to astrocytes (**Fig. 2**)¹⁵⁹. Specifically, the hyperactive neurons released toxic lipids, including peroxidated FAs, encapsulated within APOE lipoproteins, which adjacent astrocytes endocytosed and incorporated into lipid droplets. The transferred FAs were used as metabolic intermediates for increased mitochondrial β -oxidation and detoxification in astrocytes. Neuronal glutamate released during neuronal activity was postulated to enhance this β -oxidation and lipid detoxification by binding to N-methyl-D-aspartate (NMDA) receptors on the astrocytes.

While these findings underscore the importance of neuron-to-astrocyte lipid transfer in protecting neurons from lipid toxicity, it has also been revealed that lipid fluxes between these cells follow a circadian rhythm¹⁶³. During wakefulness, mitochondrial energetic activity is heightened in neurons¹⁶⁴, resulting in the production of ROS and formation of peroxidated lipids. These peroxidated lipids are shuttled by apolipoproteins to neighboring glia, where they were sequestered into lipid droplets, thereby effectively reducing neuronal ROS levels. During sleep, the peroxidated lipids contained within the glial lipid droplets were catabolized, removing toxic radicals and producing energy to

support future neuronal activities. All in all, these studies indicate that stressed and (hyper)active neurons protect themselves from ROS and lipid peroxidation by transferring toxic lipids to glial cells.

Alongside these protective functions, intercellular lipid coupling between astrocytes and neurons can lead to cytodeneration. For example, astrocytes primed with inflammatory mediators released by activated microglia — interleukin 1 α (IL1 α), tumor necrosis factor (TNF), and complement component 1q (C1QA) — increase the loading of APOE and APOJ lipoproteins with toxic long-chain SFAs (**Fig. 2**)¹²⁰. Hindering the synthesis of these long-chain SFAs by selectively knocking out *Elovl1* in astrocytes mitigates astrocyte-mediated toxicity. The release of these toxic long chain SFAs by reactive astrocytes is likely to contribute to the cytotoxic effects of IL1 α , TNF, and C1QA on ganglion cells following retinal injury^{165,166}. Why reactive astrocytes upregulate the loading of saturated lipids on APOE and APOJ lipoproteins, and to what extent these FAs are differentially trafficked by reactive astrocytes that express divergent APOE isoforms, remains unclear. With respect to the latter, isoform- and cell-state-specific differences in the lipidation of APOE in astrocytes have been reported¹⁶⁷. Specifically, astrocytes chronically exposed to FAs and having excess lipid droplets produce TAG-rich APOE particles, a process that is boosted by the APOE4 variant¹⁶⁷. Whether there is a differential loading of APOE2, APOE3, and APOE4 with toxic long-chain SFA in reactive astrocytes, however, remains to be determined.

[H2] Glia-glia lipid transfer

Oligodendrocytes are the myelinating cells of the CNS, enveloping axons with layered, lipid-rich membranes that enable rapid saltatory axonal conduction¹⁶⁸. Cholesterol is a critical component of myelin membranes and thus adequate levels of cholesterol are essential for developmental myelination and the regeneration of myelin in response to pathological demyelination. Since peripheral cholesterol entry into the CNS is restricted by the BBB, oligodendrocytes and OPCs have adapted to synthesize their own cholesterol (**Fig. 2**)^{12,169}. Disruptions in this process can result in

oligodendrocyte apoptosis, developmental hypomyelination and impaired remyelination following demyelinating events^{12,36,170,171}. Despite FA being able to cross the BBB, oligodendrocytes also depend on de novo FA synthesis for developmental myelination and remyelination⁵⁰, which is consistent with FAs being enriched in myelin and contributing to its structural integrity¹⁷². This discrepancy may reflect the fact that SFA and MUFAs, which are major products of de novo lipid synthesis, are less efficient at traversing the BBB (**Box 2**). In addition to the cell-autonomous synthesis of cholesterol and FAs by oligodendrocytes and OPCs, intercellular lipid transfer has recently been shown to be essential for providing oligodendrocytes with lipids.

Microglia and subsets of CNS-infiltrating macrophages are increasingly being recognized for their role in meeting the demand for cholesterol by oligodendrocytes, particularly following pathological demyelination. These cells remove damaged myelin and myelin debris via phagocytosis and subsequently process and transfer myelin-derived lipids to supply oligodendroglia with the cholesterol necessary for myelin membrane formation (**Fig. 2**)¹⁷³. Mechanistically, upon receptor-mediated phagocytosis, the lysosomal degradation and intracellular processing of myelin by these cells leads to formation of oxysterols that activate cholesterol-sensing LXRs¹⁷³⁻¹⁷⁵. Recent work has shown that these receptors can also be activated by metabolites such as desmosterol that arise as a result of increased cholesterol synthesis following demyelination¹⁷¹. A similar feedback loop involving lipid-loaded macrophages, LXRs, and desmosterol is observed following low density lipoprotein (LDL) internalization in atherosclerosis¹⁷⁶. Regardless of the mode of activation, LXR activation induces an inflammatory and metabolic phenotype in macrophages and microglia that supports tissue regeneration¹⁷⁷. LXR activation also enhances the abundance of ATP-binding cassette transporter A1 ABCA1, promoting apolipoprotein lipid loading and providing a readily available pool of cholesterol for oligodendroglia to generate new myelin sheaths following demyelination²⁵.

The aforementioned studies underscore the essential role of the efficient remodeling and intercellular transfer of myelin-derived cholesterol by phagocytes in fueling remyelination in CNS

disorders. However, studies have shown that, at progressive stages of demyelinating diseases and during aging, LXR signaling becomes impaired and lipid remodeling and transfer become suboptimal in the CNS^{19,89}. Over time, the sustained intracellular accumulation of myelin-derived lipids perturbs the metabolic phenotype of macrophages and microglia^{19,54,95,106}. For example, macrophages and microglia containing abundant myelin-derived lipids exhibit a reduced capacity for lipid droplet catabolism through lipophagy and lipolysis^{95,106}. In addition, the intracellular accumulation of myelin-derived lipids leads to elevated expression of enzymes involved in FA desaturation and elongation (SCD1 and ELOVL6 in particular), resulting in the generation of ABCA1-destabilizing FAs and impaired lipid loading into lipoproteins^{19,54}. This disturbed metabolism and transfer of myelin-derived lipids contributes to the induction of a highly inflammatory macrophage and microglia phenotype that impairs OPC maturation and remyelination^{19,54,106}. Similarly, aging affects the phenotype of myelin-containing microglia in the CNS^{89,178,179}, and is a major impediment to adequate remyelination¹⁸⁰. Aging compromises the capacity of microglia to clear damaged myelin debris, which is a necessary prerequisite for remyelination¹⁸¹. Aged myelin-containing microglia further demonstrate increased lipid droplet formation and elevated ROS production^{86,178,179}. Finally, aged microglia exhibit an impaired ability to recycle large amounts of intracellular cholesterol-rich myelin debris and display reduced LXR activation, leading to lysosomal cholesterol accumulation⁸⁹. The accumulation of lysosomal cholesterol results in the formation of cholesterol crystals, which induces lysosomal rupture and the activation of the NLRP3 inflammasome, thereby impairing remyelination. To address disruptions in lipid remodeling and transfer in myelin-containing phagocytes, a recent study used an apolipoprotein A1 (APOA1) mimetic peptide to promote the stabilization and efflux capacity of ABCA1, which was effective in restoring both the intracellular processing of myelin-derived lipids in microglia and the transfer of these lipids to oligodendrocytes⁹⁹. It is important to note that dietary cholesterol can also enhance remyelination¹⁸², potentially bypassing disturbances in lipid remodeling and transfer by increasing extracellular cholesterol availability in the CNS. However, given cholesterol's inability to cross the BBB, future research will be needed to explore whether dietary cholesterol

supplementation can offer beneficial effects in CNS disorders at disease stages in which the BBB retains or regains its integrity. Collectively, these studies indicate that disrupted remodeling and efflux of myelin-derived lipids in phagocytes can contribute to the failure of remyelination.

The transfer of lipids between astrocytes and oligodendrocytes also plays an essential role in myelination and remyelination. While deficiency of *Scap* in oligodendrocytes impairs myelination only in young mice, astrocyte-specific deficiency of *Scap* leads to persistent hypomyelination¹⁸³. Interestingly, this hypomyelination could be rescued by providing a diet enriched in cholesterol and oleic acid. These findings underscore the important role of astrocyte-oligodendrocyte lipid transfer in the formation and maintenance of myelin, especially during adulthood, and suggest that dietary interventions hold promise as therapeutic strategies to correct for disturbances in the intercellular transfer of lipids in the CNS. Astrocytic lipid transfer to oligodendrocytes also promotes remyelination in models of toxin-induced demyelination¹⁸⁴. It has been demonstrated that remyelination is characterized by increased cholesterol biosynthesis in astrocytes and elevated ABCA1-mediated cholesterol transfer to oligodendrocytes. These changes inversely correlated with the activity of nuclear factor erythroid 2–related factor 2 (NRF2), a key regulator of cellular oxidative stress responses (**Fig. 2**). The molecular mechanisms underlying the regulation of cholesterol biosynthesis and efflux by NRF2 have yet to be elucidated. A similar inverse relationship between cholesterol metabolism and NRF2 activity was previously reported in the liver of *Nrf2*^{-/-} mice fed a high-fat diet (HFD)¹⁸⁵. Given the low abundance and activity of NRF2 in neurons¹⁸⁶, it is tempting to speculate that a feedback mechanism in which low NRF2 activity enhances cholesterol efflux may also facilitate the ABCA1-mediated clearance of toxic peroxidated lipids from neurons to astrocytes^{157,159,164}.

Overall, proper glia-glia lipid transfer has an essential role in developmental myelination, myelin maintenance, and remyelination. Interestingly, there appears to be a divergence in how brain and spinal cord oligodendroglia acquire cholesterol, with spinal cord oligodendroglia relying more on de

novo synthesis and brain oligodendroglia on extracellular cholesterol uptake¹⁷⁰. This distinction warrants further investigation, as it may offer valuable insights into the therapeutic potential of modulators targeting lipid disturbances in brain and spinal cord pathologies.

[H2] Lipid transfer by extracellular vesicles

EVs, including exosomes and microvesicles, are small, membrane-bound vesicles that are released from cells and facilitate intercellular communication. Alongside proteins and miRNAs, EVs transfer lipids between cells¹⁸⁷. Several studies have demonstrated that the transfer of EV-associated lipids is essential for maintaining CNS homeostasis by influencing oligodendroglia physiology. Specifically, recent research showed that EVs secreted by pro-regenerative microglia enhance OPC migration and remyelination in a lysolecithin-induced demyelination model through the intercellular transfer of sphingosine-1-phosphate¹⁸⁸. Supporting the importance of EV-associated lipids in remyelination, a study by Vanherle *et al.* found that EVs released by IL4-exposed macrophages enhance remyelination ex vivo and in vivo by promoting the differentiation of OPCs (**Fig. 2**)¹⁸. Guided by lipidomic analysis and by experiments utilizing cholesterol depletion and enrichment strategies, the authors attributed the pro-regenerative effects of EVs to elevated cholesterol levels. Mechanistically, EV-associated cholesterol was found to promote OPC differentiation primarily by supplying cholesterol to the oligodendrocyte membrane through direct membrane fusion¹⁸. By contrast, a study by Lombardi *et al.* found that EVs from IL-4-exposed macrophages do not facilitate OPC maturation, while those from macrophages exposed to inflammatory cytokines hinder OPC maturation¹⁸⁹; however, the lipid fraction of EVs isolated from inflammatory macrophages did promote OPC maturation, likely through pro-regenerative endocannabinoids such as anandamide and 2-arachidonoylglycerol present within these EVs. This suggests that other detrimental factors present in EVs can outweigh the pro-regenerative impact of endocannabinoids. The divergent outcomes of these studies may stem from differences in the experimental systems used (rat vs. mouse), the duration of OPC exposure to EVs (2 days vs. 6 days), and the EV isolation methods used^{18,189}. In light of the latter, Vanherle *et al.* incorporated an additional purification step to minimize potential contaminants¹⁹⁰.

The essential role of EVs as transport vehicles for lipid transfer is also evident in neurodegenerative disorders such as AD and PD. In an animal model for AD, A β aggregation was found to promote the incorporation of pro-apoptotic ceramide classes into EVs secreted by astrocytes¹⁹¹. Furthermore, neuron-derived EVs have been shown to contain glycosphingolipids that facilitate the targeting of A β to microglia for clearance¹⁹²⁻¹⁹⁵. In PD patients, EVs purified from cerebrospinal fluid and frontal cortex are also heavily loaded with ceramides and neurodegeneration-linked proteins, including α -syn¹⁹⁶. In vitro studies confirmed that these EVs constitute a "pathological package" capable of inducing aggregation of α -syn¹⁹⁶. Similarly, the EV-associated gangliosides GM1 and GM3 have been identified as key factors in promoting α -syn oligomerization¹⁹⁷. Overall, these studies point towards EVs playing an important role in the transfer and distribution of lipids in the CNS. By transferring lipids between cells, EVs help to maintain the complex lipid landscape essential for glial and neuronal function. However, the relative importance of EVs in driving intercellular lipid transfer in the healthy and diseased CNS, compared to lipoproteins and the poorly understood tunneling nanotubes (**Box 3**), remains to be determined.

[H1] External influences on lipid metabolism

Emerging evidence highlights the influence of environmental factors, diet, and lifestyle on CNS lipid metabolism. Specifically, avoiding a **Western diet** [G], engaging in regular physical activity, managing stress and sleep patterns, and minimizing exposure to harmful substances are pivotal for maintaining effective lipid remodeling and intercellular transfer in the CNS.

[H2] Environmental factors

Environmental factors, including pollutants, climate, and pathogens, are increasingly acknowledged to impact lipid remodeling in the CNS. Pollutants, including micro- and nanoplastics as well as air contaminants, can cross the BBB, induce lipid peroxidation, and cause mitochondrial dysfunction and lysosomal disruption in glial cells and neurons¹⁹⁸⁻²⁰⁰. With regard to climate, a recent preprint has reported that cold exposure (4 °C; 4–6 hours) enhances neuronal activity-dependent lipid metabolic

processes in the brain, including lipid peroxidation, lipid droplet formation, and lipolysis²⁰¹, in agreement with its reported effects on lipid metabolism in adipose and skeletal tissues²⁰²⁻²⁰⁶. By contrast, a reduction in ambient temperature (17 °C; 5 weeks) was found to reduce obesity-induced lipid peroxidation in the hypothalamus of mice, which was associated with a decreased inflammatory burden²⁰⁷. This discrepancy may reflect differences between acute, severe cold exposure and prolonged, mild cold exposure. Finally, a recent study found that the Zika virus modulates neuronal lipid metabolism to favor its replication and survival, leading to significant lipid droplet accumulation in human neuroblastoma cells and neural stem cells (NSCs) in vitro²⁰⁸. Consistent with this, astrocytic lipid droplet accumulation was observed following exposure to Zika virus, Herpes Simplex virus 1, and dengue virus in vitro, with the latter also increasing lipid droplet accumulation within the brain of mice²⁰⁹.

[H2] Diet

The typical Western diet, rich in SFAs, trans FAs (unsaturated FAs with at least one double bond in the less common trans configuration that increases rigidity), ω -6 PUFAs, sugars, and cholesterol, is increasingly being linked to CNS pathologies²¹⁰. Dietary lipids can influence CNS-resident cell physiology either directly or indirectly through their effects on the microbiome, vasculature, and adipose tissue^{21,210,211}, which complicates the interpretation of studies. Despite these complexities, a few studies have demonstrated that a Western diet affects lipid remodeling and intercellular lipid transfer in the CNS. For example, Bosch-Queralt *et al.* showed that a Western Diet triggers a dysfunctional metabolic response associated with impaired myelin debris clearance in microglia, thereby attenuating lesion recovery in a mouse model of demyelination²¹². Mechanistically, this was attributed to a reduced activation of LXR-regulated genes encoding proteins involved in cholesterol efflux, including *Apoe*, *Abca1*, and *Abcg1*, which perturbed microglial processing and efflux of myelin and cholesterol. By contrast, in humanized mice homozygous for APOE2, APOE3, and APOE4, a Western diet was found to increase total brain APOE levels in an allele-dependent fashion, with the most significant increase observed in APOE2²¹³. The discrepancy in the effects of a Western diet on

APOE abundance could arise from the existence of an already perturbed foamy microglia metabolic phenotype during demyelination^{19,89}. Alternatively, it may have arisen because Bosch-Queralt *et al.* compared APOE levels only across different lesion stages, without assessing levels at baseline prior to toxin-induced demyelination.

In addition to affecting intercellular lipid fluxes in the CNS, studies have shown that exposing mice to a high fat diet (HFD), a subtype of the Western diet that is enriched in SFAs and unsaturated fatty acids (UFAs) but lacks substantial amounts of cholesterol and sugars, disturbs lipid droplet breakdown by reducing lipophagy^{214,215}. Hippocampal microglia in a mouse model of type 2 diabetes induced by an HFD exhibit lipid droplet accumulation and significant impairment of lipophagy²¹⁴. Lipid droplets were found to colocalize with and hinder the degradation of TREM1, thereby perpetuating neuroinflammation and impairing cognitive function. Interestingly, another study found that chronic HFD consumption enhances lipid droplet accumulation and impairs mito- and lipophagy in the hypothalamus of male but not female mice²¹⁵. This suggests a sexually dimorphic response to chronic HFD exposure, with females exhibiting protection against the adverse metabolic effects of the diet. An HFD is further associated with increased brain ROS levels and lipid peroxidation (reviewed in ²¹⁶), and disruptions in the quantity and quality of (lyso)phospholipids during HFD feeding²¹⁷, reflecting alterations in the Kennedy pathway and Lands cycle. Collectively, these findings highlight the detrimental effects of Western diets on lipid remodeling and intercellular transfer within the CNS. Establishing causality between these diet-induced alterations in lipid metabolism and CNS pathology in neurodegenerative disorders, while considering sex differences, will be an important goal of future research. Comparing these effects to those of intermittent fasting, which exerts opposing influences on these processes (reviewed in ²¹⁸), would also be of interest.

[H2] Lifestyle

Disruptions in circadian rhythm are increasingly linked to CNS pathologies, including AD and PD²¹⁹, but the molecular drivers remain poorly understood. Recent studies have shown that the circadian clock

regulates lipid remodeling and intercellular transfer in the CNS^{163,220,221}. The transfer of peroxidated lipids from hyperactive neurons to astrocytes, along with their subsequent storage and breakdown within astrocytes, follows a circadian pattern and is essential to retain CNS homeostasis¹⁶³. Mirroring these findings, most apolipoprotein genes show rhythmic expression in the CNS (reviewed in ²²⁰). Hence, it is tempting to speculate that disruptions in sleep patterns or disease-related alterations in circadian rhythms could contribute to lipid dyshomeostasis in CNS disorders. Regarding the latter, a recent preprint has reported that, while showing rhythmicity in controls, *Srebf1*, the gene encoding SREBP1, which is involved in lipid synthesis, loses its rhythmicity in an experimental model of AD²²¹. The extent to which intracellular accumulation of myelin or other pathogenic protein aggregates disrupts the rhythmicity of lipid metabolism, leading to glial and neuronal dysfunction, remains to be determined.

Alongside disturbances in circadian rhythms, some studies suggest that acute and chronic physiological stress are associated with alterations in lipid remodeling and intercellular transfer in the CNS. Chronic physiological stress increases CNS lipid peroxidation, which can be mitigated by treatment with non-selective reactive nitrogen species (RNS) and ROS inhibitors²²². By using a multi-omics approach, a recent study found that acute physiological stress correlates with an increase in the levels of lysophospholipids lacking acyl chains at the sn-2 position, indicative of increased PLA2 activity, and increased levels of eicosanoid and endocannabinoids in the mouse brain²²³. Similarly, single-nuclei RNA sequencing demonstrated that acute stress induces rapid and marked differences in the expression of lipid transfer and remodeling genes in astrocytes, microglia, neurons, or oligodendrocytes, including but not limited to *Apoe*, *Abca1*, *Plin2*, *Nrf2*, and diverse lipases and acyltransferases²²⁴. The relevance of these metabolic changes in driving disease pathology in neurodegenerative disorders remains unclear, but is highly pertinent in an increasingly stressed society.

[H1] Conclusions

In recent years, it has become clear that lipid metabolism, remodeling and intercellular transfer are essential for maintaining or restoring lipid homeostasis in the CNS, with substantial evidence highlighting the key roles of lipid synthesis, lipid droplet biogenesis and breakdown, lipid oxidation, phospholipid metabolism, lipoprotein-mediated lipid transfer, and EVs. Importantly, various genetic factors (such as variants in *APOE*, *TREM2*, and *PLA2G6*), environmental influences (such as stress, circadian rhythm, diet, climate, and pathogens), and disease-associated pathological hallmarks (such as ROS, inflammation, and cellular accumulation of A β , α -synuclein, and myelin debris) have been identified as key drivers of dysfunction in these metabolic pathways.

In the upcoming years, novel lipidomics techniques will allow researchers to further unravel the molecular blueprints of disturbed lipid metabolism in the CNS. Tracer lipidomics using isotope-labeled lipid metabolites could provide precise analyses of lipid flux and transfer dynamics in the healthy and diseased CNS²²⁵. Combined with mass spectrometry imaging and Raman microscopy, researchers could further obtain spatial information on changes in the lipidome and lipid transfer on a subcellular level²²⁶⁻²³⁰. Such experiments could explain why certain lipid types (such as long-chain SFAs, lysophospholipids, and cholesterol) that are essential membrane constituents and cell signaling molecules in the CNS become toxic under specific conditions. While this may stem from supra-homeostatic or atypical lipid accumulations in organelles or cells — such as neurons and foamy microglia — that lack the necessary machinery (or progressively become less able) to properly metabolize or store them, more complex processes may also be involved. Furthermore, these novel techniques could enable the detailed assessment of whether the lipid metabolism, remodeling, and intercellular transfer mechanisms discussed in this review are differentially active in neurons and glial cells during homeostasis and disease.

Overall, while further in-depth lipidomic studies are needed, it is clear that lipid remodeling and intercellular transfer are promising therapeutic targets for restoring CNS homeostasis in neurological disorders such as MS, AD, PD, and HD.

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Competing interests

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Fig. 1. Lipid metabolism and remodeling in the CNS. The dynamic process by which the composition and structure of CNS lipids are established and modified is driven via multiple interconnected cellular pathways, occurring across essentially all CNS cell types. During de novo lipid biosynthesis (lipogenesis), acetyl-CoA acts as a substrate for the formation of new FAs and cholesterol through the fatty acid synthase (FAS) and Kandutch-Russell & Bloch pathways, respectively²³¹. The FAs synthesized through these pathways, along with internalized FAs, can undergo enzymatic oxidation by cyclooxygenase (COX), lipoxygenase (LOX), and cytochrome P450 (CYP450), leading to the production of bioactive lipids. Cholesterol can also be oxidized via CYP46A1 and CYP27A1 to form 24S-hydroxycholesterol (24S-OHC) and 27-OHC, respectively. Additionally, both FAs and cholesterol can undergo non-enzymatic oxidation through free radical species, leading to the formation of peroxidated phospholipids (PLs) and products including 7 α -OHC, 7 β -OHC, 25-OHC, and 7-ketocholesterol (7-KC), respectively²¹. Synthesized FAs and cholesterol can also be stored in lipid droplets^{21,83}. The autophagy of organelles, more specifically ER-, golgi-, lipo-, and mito-phagy, involves the incorporation of these organelles in autophagosomes. These autophagosomes then fuse with lysosomes to form autophagolysosomes, releasing lipids that can be used for various purposes¹⁰³. Similarly, lipolysis of lipid droplet-associated neutral lipids can release substrates that can be used to synthesize and remodel other lipids (including triglycerides, plasmalogens, cardiolipins, sphingomyelin, ceramides, phosphatidylserine (PS), phosphatidylinositol (PI) and phosphatidylglycerol (PG)) in organelles and the plasma membrane²³²⁻²⁴⁰. Phospholipids are synthesized via the Kennedy pathway and remodeled through the Lands cycle to generate phospholipid diversity^{116,117}. In the Kennedy pathway, a series of enzymatic reactions, operating across multiple organelles but predominantly in the ER, convert choline and ethanolamine into the phospholipids phosphatidylcholine (PC) and phosphatidylethanolamine (PE), respectively. In the Lands cycle, phospholipase A2 (PLA2) hydrolyzes a FA moiety from these phospholipids, typically releasing polyunsaturated fatty acids (PUFAs) and producing a lysophospholipid. The reacylation of lysophospholipids back to glycerophospholipids is catalyzed by lysophospholipid acyltransferases (LPLATs). LPLATs incorporate acyl chains of varying length and degrees of saturations during reacylation, resulting in the formation of structurally divergent phospholipids. While the Lands cycle is predominantly active in the ER, the enzymes involved can also be found in the Golgi apparatus, mitochondria, lipid droplets, and the plasma membrane^{241,242}.

Fig. 2: Intercellular lipid transfer in the CNS. While all CNS-resident cells can synthesize lipids through pathways mediated by sterol regulatory element-binding proteins (SREBP) and SREBP cleavage-activating protein (SCAP), intercellular lipid transfer is thought to be key in maintaining lipid homeostasis in the CNS. In pathological conditions, especially when reactive oxygen species (ROS) are elevated, astrocytes convert blood glucose to lactate, which is then transported to neurons and converted to fatty acids (FAs; light blue boxes)^{157,159}. Reciprocally, neurons expel peroxidated lipids, formed through peroxidation by ROS, which are taken up by astrocytes and stored in lipid droplets (LDs)^{90,157,159}. This intercellular

lipid transfer, along with astrocytic lipid droplet formation and β -oxidation, depends on apolipoprotein E (APOE)-containing high-density lipoproteins (HDLs) and glutamate, and protects against neurodegeneration¹⁵⁷. Neurons also transfer 24(S)-hydroxycholesterol (24S-OHC) to astrocytes, likely under both homeostatic and diseased conditions, resulting in the activation of nuclear liver X receptors (LXRs), which control the expression of genes involved in CNS lipid homeostasis (green boxes)⁷². Astrocytes have also been reported to transfer de novo synthesized cholesterol to neurons via HDLs, which neurons use to support their functions (dark blue boxes)^{154,155}. Similarly, astrocytes transfer cholesterol to oligodendrocytes to promote remyelination following pathological demyelination¹⁸⁴, complementing the de novo biosynthesis of cholesterol in these cells¹². Interestingly, during remyelination, enhanced cholesterol biosynthesis in astrocytes, followed by increased cholesterol transfer to oligodendrocytes, is inversely correlated with the activity of nuclear factor erythroid 2-related factor 2 (NRF2). This suggests that NRF2 not only serves as a master regulator of cellular antioxidant responses but also acts as a key driver of intercellular cholesterol transfer in the CNS (red boxes). Astrocytic lipid transfer is also closely associated with CNS pathology, with inflammatory mediators released by microglia, such as interleukin 1 α (IL1 α), tumor necrosis factor (TNF), and complement component 1q (C1q), enhancing the astrocytic loading of APOE and apolipoprotein J (APOJ) HDLs with toxic long-chain saturated FAs (SFAs) (purple boxes)¹²⁰. Microglia also drive intercellular lipid transfer in the CNS (grey boxes). Following demyelinating events, they process damaged myelin debris and generate lipid droplets, oxysterols, and desmosterol, with the latter two metabolites activating LXRs^{171,173-175}. LXR activation enhances ATP-binding cassette transporter expression, thereby facilitating APOE lipidation and the formation of HDLs that can be taken up by oligodendrocytes to aid myelin formation²⁵. Similarly, microglia (and CNS-infiltrating macrophages, not shown) release extracellular vesicles (EVs) containing lipids that can be used by oligodendrocytes for the genesis of new myelin sheaths^{18,188,189}. Diverse pathological parameters and environmental factors, such as neuroinflammation, aging, APOE4, oxidative stress, lipid droplet overload, and altered circadian rhythms, can impair neuron-glia and glia-glia lipid fluxes, thereby driving CNS pathologies (yellow boxes).

Box 1: Fatty acid transport into the CNS.

Unlike cholesterol, fatty acids (FAs) can cross the blood brain barrier (BBB) from the circulation into the CNS. While virtually all FAs can be transported into the CNS, some, particularly polyunsaturated fatty acids (PUFAs), rely entirely on transport mechanisms for their uptake. For those FAs synthesized within the CNS, the relative contributions of de novo synthesis versus uptake from the periphery as the primary route of FA supply to the brain remain poorly understood. The process of FA transport occurs through both passive diffusion and active protein-mediated mechanisms^{243,244}. Specifically, FA transport and binding proteins located in the plasma membrane and cytosol are essential for mediating the transfer of albumin-, lysophospholipid-, and triacylglycerol-bound FAs across the BBB^{245,246}. Given the brain's enrichment in PUFAs, their transport across the BBB has been extensively studied in both health and disease contexts^{59,247-249}. The major facilitator superfamily

member MFSD2A has been identified as a key regulator of PUFA transport into the brain^{250,251}. Similarly, acyl-CoA synthetase long-chain family member 6 (ACSL6) regulates the activation and intracellular retention of PUFAs, docosahexaenoic acid (DHA) in particular, by attaching them to coenzyme A (CoA) to form acyl-CoA, thereby facilitating their transfer into the CNS²⁵². Like PUFAs, short-chain FAs (SCFAs) such as acetate (2 carbon atoms), propionate (4 carbon atoms), and butyrate (6 carbon atoms), which are produced by intestinal bacteria through the fermentation of non-digestible dietary polysaccharides, can cross the BBB^{253,254}. The entry of SCFAs into the CNS is hypothesized to be facilitated by endothelial monocarboxylate transporters (MCTs)²⁵⁵. The accumulation of microbiota-derived SCFAs in the brain provides a molecular mechanism for the dietary and microbiota-mediated regulation of CNS metabolism and physiology (reviewed in REF ²⁵⁶). Using an *in vitro* BBB model, a pivotal role for cluster of differentiation 36 (CD36), fatty acid transporter protein 1 (FATP1) and FATP4 in facilitating the translocation of various FAs across the BBB has been revealed²⁴⁵. Increased desaturation improved the ability of FAs to cross the BBB, suggesting that there may be preferential PUFA entry into the CNS and offering a plausible explanation for the high PUFA content of the brain.

Box 2: Apolipoprotein E lipidation and uptake in Alzheimer disease

As cholesterol is insoluble in water, it binds to lipoproteins for transportation within and between cells. Within the CNS, the lipoprotein that cholesterol mainly combines with is astrocytic apolipoprotein E (APOE)²⁵⁷. The lipidation of APOE by cholesterol is mediated by lipid efflux proteins (specifically ATP-binding cassette (ABC) transporters) on the membranes of glial cells²⁵⁸, resulting in the formation and release of high-density lipoprotein (HDL)-like particles (see the figure)²⁵⁹. APOE lipidation allows it to interact with endocytic receptors, such as triggering receptor expressed on myeloid cells 2 (TREM2), low-density lipoprotein (LDL) receptor (LDLR), and LDLR-related protein 1 (LRP1), which are expressed by neurons, microglia, and oligodendrocytes^{30,260-262}. The binding of lipoproteins to these receptors and their uptake activates signaling cascades and provides an essential source of cellular cholesterol, complementing *de novo* synthesis^{147,183,263-265}. Supporting the importance of APOE-mediated cholesterol transfer in maintaining CNS lipid homeostasis, the APOE4 isoform is a strong risk factor for late-onset Alzheimer disease (LOAD), with homozygous APOE4 carriers having a 14-fold increased risk of developing LOAD compared to APOE3 carriers²⁶⁶. Amongst other processes, APOE4 dysregulates intracellular lipid processing, lipid efflux, and the binding of APOE to and endocytosis by recipient cells^{152,267-274}. Similarly, Alzheimer disease (AD)-associated mutations in the gene encoding TREM2, particularly the R47H missense mutation, result in an increased risk for LOAD²⁷⁵. *TREM2* mutations impair the binding and uptake of amyloid- β (A β) that is complexed with lipoproteins by microglia, thereby increasing amyloid pathology²⁶⁴. In parallel, *TREM2* deficiency and the *TREM2*^{R47H} variant suppress the transition of microglia to a protective phenotype²⁷⁶⁻²⁸⁰.

Box 3: Tunneling nanotubes and lipid transfer

Tunneling nanotubes (TNTs) are thin, membrane-enclosed, F-Actin-rich protrusions that link mammalian cells and permit the direct exchange of various components or signals between non-adjacent cells over long distances²⁸¹. The material transferred via TNTs can range from ions and intracellular organelles to protein aggregates and pathogens²⁸²⁻²⁸⁷.

Several studies indicate that TNTs are essential for maintaining CNS homeostasis and limiting CNS tissue damage in neurodegenerative disorders, particularly in Parkinson disease (PD). For example, in a mouse model of PD, microglia jointly degrade fibrillar α -synuclein intercellular distribution through TNTs, thereby lowering the α -synuclein burden²⁸². TNTs also enable the movement of mitochondria and α -synuclein aggregates between microglia and neurons, with a preference for α -synuclein transfer from neurons to microglia and mitochondrial movement from microglia to α -synuclein-containing neurons²⁸³. Alongside α -synuclein transfer, the cell–cell spreading of other neurodegenerative disease-associated proteins, such as tau aggregates and prions can also be controlled by TNTs in the CNS^{284,285}. Cellular overload of these pathological aggregates is postulated to lead to a cascade of reactions — such as lysosomal stress, increased formation of reactive oxygen species (ROS), and mitochondrial dysfunction — which drive the formation of TNTs²⁸⁶. This suggests an interesting link between neuronal stress and hyperactivity, in which ROS and peroxidated lipids are transported from neurons to astrocytes to prevent lipotoxicity^{157,163}. In future studies, it would therefore be of interest to assess intercellular transfer of peroxidated lipids or other lipids via TNTs, alongside lipoprotein-mediated intercellular lipid transfer, in stressed and hyperactive neurons.

Supporting the role of TNTs in lipid transfer between cells, TNTs have been shown to transfer lipid droplets between endothelial cells and their formation in these cells has been shown to be controlled by polyunsaturated fatty acids such as arachidonic acid²⁸⁷. Hence, TNTs could serve as biologically plausible ‘highways’ for the efficient transfer of macromolecules and organelles, such as lipid droplets, within the CNS, bypassing the energy-intensive and non-specific lipid remodeling and transfer pathways that involve lipid droplet breakdown, efflux, and lipoproteins. Importantly, while growing evidence underscores the role of TNTs in intercellular communication and intercellular lipid droplet shuttling, the studies that have been conducted to date were mainly limited to *in vitro* experiments, and thus their precise contribution to CNS lipid metabolism remains unclear.

Glossary

Autophagy: A cellular process in which damaged or unnecessary components of the cell are degraded and recycled within the cell through lysosomal activity.

β -oxidation: The metabolic pathway that breaks down fatty acids into acetyl-CoA units in the mitochondria for energy production.

Cholesterol: A lipid molecule synthesized in all cell types, essential for cell membrane structure, hormone synthesis, and bile acid formation.

Extracellular vesicles: Nanosized lipid bilayer particles secreted by cells that carry lipids, proteins, and RNA to distant and nearby cells, playing a key role in intercellular communication.

Fatty acid (FA): A carboxylic acid consisting of a hydrocarbon chains and a terminal carboxyl group, essential for energy storage and membrane structure.

Lipid droplets: Intracellular storage organelles primarily composed of neutral lipids such as triglycerides, serving as energy reserves.

Lipidome: The complete set of lipids in a biological system, encompassing various lipid classes.

Lipoproteins: Complexes of lipids and proteins with a hydrophobic core that transport lipids, such as cholesterol and triglycerides, through the bloodstream.

Monounsaturated fatty acids (MUFAs): Fatty acids with one unsaturated carbon bond in their hydrocarbon chain, commonly found in olive oil and avocados.

Phospholipids: Lipids containing a phosphate group, forming the structural components of cell membranes.

Polyunsaturated fatty acids (PUFAs): Fatty acids with two or more double bonds in their hydrocarbon chain, important for membrane fluidity and signaling.

Saturated fatty acids (SFAs): Fatty acids with no double bonds in their hydrocarbon chain, typically solid at room temperature and found in animal fats.

Tracer lipidomics: A method for studying lipid metabolism by tracking the incorporation of labeled lipid precursors into biological systems.

Western diet: A dietary pattern high in processed foods, fats (such as saturated fats and cholesterol), sugars, and low in fiber, commonly associated with various chronic diseases.