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INTRACELLULAR CHOLESTEROL HOMEOSTASIS AND AMYLOID PRECURSOR PROTEIN PROCESSING

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Abstract

Many preclinical and clinical studies have implied a role for cholesterol in the pathogenesis of Alzheimer's disease (AD). In this review we will discuss the movement of intracellular cholesterol and how normal distribution, transport, and export of cholesterol is vital for regulation of the AD related protein, A β . We focus on cholesterol distribution in the plasma membrane, transport through the endosomal/lysosomal system, control of cholesterol intracellular signaling at the endoplasmic reticulum and Golgi, the HMG-CoA reductase pathway and finally export of cholesterol from the cell.

Keywords

Alzheimer's disease; amyloid; apolipoprotein E (apoE); APP; cholesterol; cholesterol ester; lipid rafts; sterol regulatory element binding protein (SREBP); SREBP cleavage activating protein (Scap); niemann-pick type C disease (NPC)

Introduction

Alzheimer's disease (AD) is a progressive neurodegenerative disease characterized histologically by the presence of amyloid plaques and neurofibrillary tangles in the brain. The plaques consist of aggregated proteinaceous material, a major component of which is β -amyloid (A β). The neurofibrillary tangles are composed of paired helical filaments of the microtubule associated phosphoprotein, tau. For more than a decade, research has focused on how A β is generated, how it impacts cellular function, and how it promotes the pathobiology of tau.

The Amyloid Precursor Protein (APP), located on chromosome 21 was implicated as being a critical player in AD as the A β peptide that is found in amyloid plaques is derived from APP following a number of cleavage events, and Down syndrome patients, who have three copies of Chromosome 21, always develop AD. In 1991, the first mutations that cause familial forms of AD were identified in APP (reviewed in [1]). In 1995, a second genetic locus, presenilin 1

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(PS-1) was found to be associated with familial AD. This was soon followed by the identification of mutations in the presenilin 2 (PS-2) gene. Mutations in APP and the presenilins have as their unifying feature the ability to alter the processing of APP such that more A β peptides are produced (reviewed in [1]). It is therefore thought that the accumulation and/or aggregation of A β underlies the etiology of the disease, and that preventing this accumulation is a valid therapeutic target. Multiple studies have indicated a role for cholesterol in maintaining normal levels of A β in vitro and in vivo. In this review we will discuss the movement of intracellular cholesterol and how normal distribution, transport, and export of cholesterol is vital for homeostatic regulation of A β . The major stages of intracellular cholesterol transport are shown in Figure 1.

Evidence for a role of cholesterol in Alzheimer's disease

Sporadic, late onset AD accounts for greater than 95% of all AD cases [2]. Within these sporadic cases, by far the greatest genetic risk factor is possession of the apolipoprotein E ϵ 4 (APOE4) allele [3;4]. Not only does APOE genotype act as a risk factor for developing the disease, but also the age of onset of the disease. In an almost dose-responder manner, the average age of onset for patients with 2 ϵ 4 alleles is less than 70 years of age, with 1 ϵ 4 allele, 80 years, whereas for those expressing no ϵ 4 alleles it is 90 years [3].

How APOE genotype alters risk of developing AD remains an open question, but a primary role for apolipoproteins is cholesterol transport, and thus studies of AD began to examine the effects of cholesterol on APP processing. The brain is the most cholesterol rich organ in the body, but following development, CNS cholesterol is generated solely through *de novo* synthesis, with transfer from the periphery contributing little [5]. Cholesterol is essential in the CNS for, amongst other things, synapse formation, neuronal repair, myelin and neurosteroids production. ApoE plays an important role in the transport of cholesterol around the CNS, for repair, neurogenesis and myelin maintenance. Despite its important role, knockout of apoE does not cause severe impact on neurological function. This may be due to the fact that other apolipoproteins, such as apoD, can compensate when the APOE gene is removed [6].

The first evidence that cholesterol may impact A β production in the brain was provided in 1994, when Sparks and colleagues demonstrated that dietary cholesterol increases amyloid production in rabbits [7]. They demonstrated that a high cholesterol diet for as little as 4 weeks caused increased β -amyloid immunoreactivity in rabbit hippocampal neurons. This was followed by work on mice genetically modified to deposit cerebral β -amyloid, again showing that a cholesterol-enriched diet resulted in increased amyloid plaque deposition, increased A β and β -cleaved APP c-terminal fragment (CTF) production, and decreased α -cleaved secreted APP (sAPP α) [8]. Similar studies have confirmed these observations [9-12]. As mentioned above, very little cholesterol is transferred from the periphery to the CNS, so the findings that increased dietary cholesterol can contribute to brain A β levels is puzzling. One possible explanation is the issue of blood brain barrier permeability, which is reported to be impaired in transgenic models of AD [13]. We have fed 5% cholesterol diets to the PS/APP transgenic mouse, and to nontransgenic mice. We found that only PS/APP mice had elevated brain A β , while A β levels in nontransgenic mice were unaltered by the diet (unpublished data).

Further evidence for a role for cholesterol in AD was found when cholesterol lowering drugs, statins, were shown to reduce both intracellular and extracellular levels of A β 40 and A β 42 peptides in primary cultures of hippocampal neurons transfected with human APP [14]. Studies by our group and others have shown that these drugs can also significantly reduce the levels of A β 40 and A β 42 in transgenic mice, wild-type mice and guinea pigs [14-16]. In addition, epidemiological studies indicated that statin use may decrease the risk of developing AD by up to 70% [17-19], and studies in humans have shown that statin use can reduce levels of

A β in the plasma [20] and the β -cleaved fragment sAPP β in CSF [17;21]. However prospective cohort studies have failed to demonstrate the protective effects of statins on dementia [22-24] and other studies have not replicated the A β lowering effect of statins in the CSF [25-27].

Plasma membrane

Cholesterol has many functions in animal cells, including a vital role in the plasma membrane. Lipids account for approximately 40% of the dry weight of plasma membranes, the remainder consisting of proteins. Phospholipids are the most abundant of these lipids (~70%) with cholesterol being the majority of the remainder (~21%). These proteins and lipids are arranged into bilayer leaflets, which display extraordinary fluidity. One of the major factors influencing this property is the cholesterol content of the membrane. An increased cholesterol concentration results in stiffening of the plasma membrane and reduced lateral motion of the membrane. This can influence basic membrane functions by preventing the translocation of substrates to proteins embedded in the membrane, and increasing endocytosis.

Early elegant experiments demonstrated the importance of membrane cholesterol for activity of the APP secretase enzymes, α - and β -secretase. Bodovitz and Klein were the first to demonstrate that increased cholesterol could impact α -secretase cleavage of APP [28]. They used a rapid delivery system that transfers cholesterol to the cell membrane in under 60 seconds [29]. This was followed by work that demonstrated that methyl- β -cyclodextrin, which removes cholesterol rapidly from the cell membrane, could decrease β -secretase cleavage of APP when combined with lovastatin [30]. Our lab has confirmed this work with methyl- β -cyclodextrin alone, and found that as cholesterol is reduced there is an inverse relationship between β - and α -secretase activities, with β -secretase cleavage decreasing and α -secretase cleavage increasing [31]. This leads to a reduction in A β 40 and A β 42.

While the different lipid classes are distributed asymmetrically in the leaflets of the plasma membrane they are not always assigned randomly. Even though cholesterol is an integral part of the plasma membrane, it is not evenly distributed in and between the two leaflets of the bilayer [32;33]. There is a marked asymmetry in cholesterol distribution which strongly favors the cytofacial leaflet, with approximately 85% of total membrane cholesterol residing in this region [32;34]. The importance of transbilayer distribution of cholesterol is not fully understood, but modifications to this distribution are associated with aging, statin use, alcohol and activity of membrane proteins [15;32-37]. The aging mouse membrane is particularly relevant for studies of AD, which is primarily a disease of aging. In synaptosomal plasma membranes, the normal cholesterol levels in the exofacial leaflet of a young male mouse are ~15% of total membrane cholesterol; by 15 months this has increased to ~25%; and by 24 months of age to ~32% [32]. The fluidity profile of the membrane was also altered, with decreased fluidity of the membrane in the older mice [32]. Membrane fluidity is important for basic membrane functions, such as endocytosis of the plasma membrane. Changes in exofacial cholesterol have been demonstrated with the cholesterol modifying agents methyl- β -cyclodextrin, apoE and statins [15;32-37]. All of these compounds are known to impact A β production *in vivo* and *in vitro*, and we have shown that the changes in neuronal plasma membrane cholesterol distribution caused by statins directly correlate with A β levels *in vivo* [15].

Within the membrane itself, specific lipid domains exist such as annular lipids that closely border integral proteins [38;39]; fast, slow and non-exchangeable cholesterol pools [40]; and dynamic assemblies of cholesterol and sphingolipids into moving platforms in the exoplasmic leaflet (lipid rafts) [41]. Lipid rafts are thought to be a more static, liquid-ordered phase within the phospholipid-rich liquid disordered phase of the membrane. Sphingolipids laterally associate with each other with cholesterol filling any remaining voids. The observance that

cholesterol-sphingolipid rafts are insoluble in detergents led to the observation that these rafts associate with many membrane bound proteins. Glycosylphosphatidylinositol (GPI)-anchored proteins found in rafts suggest that lipid rafts are important entities in membrane signaling [42;43]. Rafts are not the dominating lipid phase in the exoplasmic leaflet of the membrane except in the case of myelin in the oligodendrocytes where concentrations of cholesterol and sphingolipids are much higher than in other cells [44]. Rafts are also implicated in membrane trafficking of proteins involved in biosynthetic and endocytic pathways [41].

The discovery that APP, β -secretase, PS-1, and $A\beta$ are all present in lipid rafts [45;46] has led to speculation that the lipid raft fraction is a putative site of membranous APP cleavage. Depletion of membrane cholesterol affects the association of APP with rafts [30], and studies that used antibody cross linking to isolate β -secretase from lipid rafts succeeded in reducing $A\beta$ production [47]. This led to the suggestion that APP in lipid rafts is primarily processed via the β -secretase pathway, and APP outside of rafts is processed via the α -secretase pathway [47]. Further studies have implicated γ -secretase activity in lipid rafts [48]. The localization of these proteolytic proteins suggests that rafts are a prime target for $A\beta$ reducing compounds, and as cholesterol is a key component of these rafts, membrane cholesterol is an early pathway target for reducing $A\beta$ production.

Interestingly, we have found that cytoplasmic APP interacting proteins show differential distribution within cells, sometimes localizing to the regions of lipid rafts. While FE65 and Mint/X11, two APP adaptor proteins [49], are found outside of lipid rafts, Disabled-1 (Dab1) partially fractionates with lipid rafts. Fyn tyrosine kinase, which phosphorylates both APP and Dab1 [50], is found exclusively in lipid rafts [51], as are the tyrosine phosphorylated forms of APP and Dab1 (unpublished data). We further found that phosphorylation of Dab1 promotes its association with APP, which increases its presence on the cell surface and its cleavage by α secretase [50;52]. Thus, the APP found in lipid rafts differs in its phosphorylation state, its interaction with adaptor proteins, and its proteolytic processing.

Intracellular Cholesterol Transport – the endosomal – lysosomal – ER pathway

The endosomal-lysosomal pathway is involved in the proteolytic processing of APP to $A\beta$ [53-58]. Endosomal abnormalities have been found in AD, where they precede amyloid and tau pathology in the neocortex. Enlarged neuronal endosomes have also been recorded in Down's syndrome (Trisomy 16) prior to dementia symptoms [59], and are thought to be caused by the excess production of APP β -CTF [60]. Thus, factors affecting APP trafficking in endosomal compartments are important for understanding AD pathogenesis.

One disorder that is valuable in determining a role for intracellular cholesterol trafficking in neurodegenerative disease is Niemann-Pick Type C (NPC) disease. The defect in NPC disease can be caused by mutations in either the NPC1 or the NPC2 gene, however NPC1 mutations account for 95% of all cases. NPC1 is a membrane protein found in late endosomes/lysosomes [61]. NPC2 is smaller soluble protein found in the lysosomal lumen [62]. They are both involved in the transport of cholesterol from late endosomes / lysosomes to the endoplasmic reticulum (ER), and are thought to work in tandem with NPC2 facilitating the egress of cholesterol from lysosomes [63]. This explains why mutations in either gene cause a similar phenotype. NPC1 shares sequence homology with the cholesterol-sensing domains of several other proteins, including 3-hydroxy-methylglutaryl-CoA (HMG-CoA), which are implicated in cholesterol homeostatic mechanisms. NPC disease is an autosomal recessive disorder caused by a mutation in the NPC1 gene. It is characterized by a fatal build up of endocytosed, unesterified cholesterol and sphingolipids in late endocytic organelles, leading to demyelination, progressive neurodegeneration and death [61;64]. The Balb/c npc^{nih} mouse

[65] synthesizes abnormal NPC1 protein due to an insertion in the NPC1 gene; it develops progressive neurodegeneration and dies at 8-10 weeks of age.

Using this mouse, as well as in vitro models, we and others have shown that the movement of cholesterol through the cell has profound effects on how APP is processed. In vivo, we have shown that mutated NPC1 in mice causes an accumulation of β -CTF, A β 40 and A β 42. This coincided with an accumulation of presenilins in early endocytic compartments [66]. Very similar accumulations of A β and β -CTF were also found in human NPC1 brain, with accumulations again occurring in early endosomes [67]. In CHO cells deficient in NPC1 protein, and in cells treated with U18666A (a drug that prevents the translocation of cholesterol from lysosomes to the ER), A β and presenilin accumulations were found in late endosomes [68].

Cholesterol at the ER and Golgi

Cholesterol that enters the cell via the endocytic pathway is transported to the endoplasmic reticulum (ER) for processing. The ER is a cholesterol-poor environment where regulation of the cells cholesterol balance is maintained. Within the ER resides the sterol regulatory element binding protein (SREBP). Under cholesterol poor conditions, SREBP interacts with SREBP cleavage activating protein (Scap). It binds to CopII proteins, which cluster the Scap/SREBP complex into vesicles for transport to the Golgi apparatus [69]. Once in the Golgi, SREBP undergoes proteolytic cleavage and the N-terminus is released, acting as a nuclear transcription factor, with a consequential increase in cholesterol production. Recent studies demonstrate the delicate balance of ER cholesterol in regulating the cholesterol homeostatic pathway. When ER cholesterol levels make up less than 5% of total ER lipids, it causes translocation of the sterol regulatory binding element to the Golgi; once levels increase past 5%, the Insig protein binds to Scap and vesicle budding from the ER is blocked [70]. A role for SREBP in brain cholesterol homeostasis can be appreciated by the dramatic decrease in SREBP2 cleavage observed after traumatic brain injury [71] when the brain is exposed to high levels of cholesterol from degenerating cells [72].

Excess intracellular cholesterol is either stored as unesterified cholesterol (UC) in cell membranes, or as cholesterol ester (CE) in cytoplasmic lipid droplets. The balance between UC and CE pools is regulated by an ER enzyme called acyl-coA:cholesterol acyltransferase (ACAT). Intracellular concentrations of ACAT are tightly linked to UC levels. Increased UC results in ACAT activation and increased cholesterol esterification. When UC levels decrease, CE hydrolysis increases and UC pools are renewed. Mutant cell lines with inactive SREBP (M19 cells), Scap (25RA cells) or ACAT (AC29 cells) have been used to examine the role of intracellular cholesterol compartmentalization on A β production [73]. M19 cells, in which SREBP never becomes active, had significantly decreased UC and unaltered CE. A β levels were unaltered in M19 cells. 25RA cells, in which SREBP is constitutively active, had normal UC and increased CE. These cells had elevated A β levels. Finally in AC29 cells, in which ACAT is inactive and CE cannot be formed, there is a fourfold increase of UC, but an almost complete lack of CE. AC29 cells produced almost no detectable A β [73]. This work suggested that not only did the distribution of cholesterol within the cell matter, but so did the ratio of UC and CE within the cell. The importance of ACAT was confirmed in animal studies that showed that pharmacological inhibition of ACAT can reduce amyloid deposition and reduce cognitive deficits in APP overexpressing mice [74]. However, when cholesterol modulation occurs at the cell surface there is an inverse relationship between α - and β - secretase activity [31]. An intriguing aspect of this ACAT work is that both α - and β - cleaved APP products are simultaneously reduced [73;74]. However, the Kovacs group has recently found that ACAT inhibition causes the delayed maturation of full length APP in the early secretory pathway,

both in vivo and in vitro, reducing the availability of APP at the cell surface for processing by the amyloidogenic or non-amyloidogenic pathways [75].

The cholesterol biosynthetic pathway

Following cleavage of SREBP in the Golgi, the N-terminus is released and acts as a transcription factor, entering the nucleus and inducing mRNA for HMG-CoA reductase and low density lipoprotein (LDL) receptors. The LDL receptors will allow more exogenous cholesterol to enter the cell, and HMG-CoA reductase production will induce more intracellular cholesterol production.

The cholesterol biosynthetic pathway has five major stages (Figure 2). Acetyl-CoA is converted to HMG-CoA and mevalonate. Mevalonate is phosphorylated to isopentenyl pyrophosphate and other active isoprenoid units, which condense and combine to form squalene. Squalene is converted to lanosterol, which is finally converted to cholesterol. Enzymes dictate the rate of each of these stages, with HMG-CoA reductase being the rate-limiting enzyme for the entire process. The commonly used statin drugs target this enzyme leading to inhibition of de novo synthesis of cholesterol.

As mentioned earlier, epidemiological studies have indicated a role for statins in preventing the incidence of AD [17-19], and studies on transgenic and non-transgenic mice have demonstrated that statin treatment can reduce levels of A β in mice [14-16]. Statins reduce both intracellular and extracellular levels of A β 40 and A β 42 peptides in primary cultures of hippocampal neurons transfected with human APP [14]. Studies on PS/APP transgenic mice using atorvastatin have shown that when administered to mice at an early stage in their disease progression, the statin can significantly reduce the levels of A β 40 and A β 42 [16]. However, as statins impact an early point on the cholesterol synthetic pathway, it is possible that inhibition of the dependent mevalonate pathway may be also be responsible for the effect on APP processing, rather than cholesterol itself.

The mevalonate pathway is responsible for production of many nonsteroidal isoprenoids, which are responsible for isoprenylation of many vital proteins within the cell, including small GTPases (Ras, Rho and Rab), playing an important role in protein signaling and transport. In fact this pathway has been shown to be important in diverse effects such as activation of microglia [76] and production of nitric oxide by vascular smooth muscle cells [77]. It has been shown that atorvastatin and simvastatin can increase sAPP α production via inhibition of the Rho-associated protein kinases [78]. However, another study has shown that inhibition of the mevalonate pathway will cause increased production of sAPP β and accumulation of A β [79].

Despite this, the cholesterol lowering impact of these compounds is probably the more relevant factor in lowering A β , and a study examining the effects of a 7-dehydrocholesterol- Δ 7-reductase inhibitor (BM15.766) on PS/APP transgenic mice, showed that A β and plaque formation can be reduced by treatment with these types of drugs [80]. BM15.766 inhibits the final step of the cholesterol biosynthetic pathway, and does not alter protein prenylation as statins do. This demonstrates directly that cholesterol production is vital for APP processing via the β -secretase pathway.

Cholesterol Efflux

Once cholesterol has been synthesized at the ER, it is transported to the plasma membrane within a short time frame (half-life of ~10 min) [81]. Cholesterol within the membrane is redistributed throughout the cell, with any excess removed by efflux to extracellular acceptors. The major energy dependent mechanism for cholesterol efflux is via sterol ATP binding cassette (ABC) transporters on the cell surface. There are a number of ABC transporters known

to be important for cholesterol efflux found in the CNS, including ABCA1, ABCG1 and ABCG4 [82;83]. Although all three transporters are required to move cholesterol from the cell to extracellular acceptors, there is an important distinction between them. ABCA1 can deliver cholesterol directly to lipid free / lipid poor apolipoproteins such as apoAI and apoE, whereas the ABCG transporters are involved in the delivery of partially lipidated particles generated by the action of ABCA1 [84;85].

In vitro experiments using primary or secondary human neuronal cell lines showed that the majority of radiolabeled cholesterol that is excreted from a neuron to the media does so as unaltered cholesterol, with only a small fraction (<25%) being secreted as a modified polar product such as 24S-hydroxycholesterol [86]. This is in contrast to in vivo calculations in mice that estimate that the conversion of cholesterol to 24S-hydroxycholesterol by the CYP46 enzyme may account for as much as 2/3 of all cholesterol efflux from the brain, much more than any other single mechanism [87;88]. The oxysterol family members (consisting of several hydroxycholesterol molecules) act as endogenous Liver X Receptor (LXR) agonists. LXR are ligand-activated transcription factors that regulate a large number of metabolic and developmental pathways, including cholesterol homeostasis. LXR agonists induce both genes [89;90] and protein levels [91] of ABCA1, ABCG1 and apoE in the CNS. LXR regulate gene expression by forming heterodimers with the 9-cis-retinoic acid receptors (RXR) and binding to LXR-responsive elements on DNA [92]. There are two known LXR isoforms, LXR- α and LXR- β , which occur in mammals. LXR α expression is mainly limited to the liver, adrenals, intestine and spleen, while LXR β is expressed in all tissue types, including the brain [93-97]. As such, the brains of LXR β knockout display developmental problems with late neuronal migration [98].

Due to their integral role in modulating cholesterol efflux from the cell, LXR have become a target of interest for AD. In vivo, induction of LXR leads to increased production of ABCA1 and ABCG1, increased cholesterol efflux, and a reduction of synaptosomal plasma membrane cholesterol [91]. As such it is expected that LXR agonists should lower A β levels; results from in vitro studies using the LXR agonist T0901317, however, have been inconsistent. While some reports show that T0901317 does indeed decrease A β [83;99;100], others have shown that T0901317 selectively increases A β 42 without changing A β 40 levels [31;101]. This appears to be partially due to the fact that T0901317 acts as a gamma-secretase modulator in vitro, selectively raising A β 42 levels at the expense of A β 38 [31]. Interestingly, this does not appear to be an issue in vivo, with studies in mice showing that T0901317 decreases A β and may only selectively decrease A β 42 [83;99;102]. APP transgenic mice treated with LXR agonists display improved spatial memory in a Morris water maze [103], and contextual memory in a fear conditioning paradigm [102;104]. Furthermore, LXR α or LXR β knockout mice crossed with PS/APP transgenic mice both have elevated amyloid deposition than PS/APP mice alone [105], and despite the preferential expression of LXR β in the brain, LXR α knockout mice had a similar increase in A β deposition to LXR β mice.

In 2005, three independent groups concurrently published studies on the effects of ABCA1 knockout in APP transgenic mice [106-108]. Using four different model types, each of the groups found that soluble apoE levels were diminished by 75-85% in the ABCA1 knockout mice. They also found that despite no evidence of changes in total or cleaved APP products, the amount of amyloid deposited in each case was increased in the knockout mice [106-108]. Conversely, overexpression of ABCA1 in APP transgenic mouse has the opposite effects, with increased lipidation of apoE particles and a dramatic reduction in the amount of amyloid deposited in the mice [109]. These studies led to the hypothesis that the lipidation status of apoE was important in the clearance and deposition of A β in vivo. This hypothesis was advanced with a recent publication that demonstrated that apoE enhanced the degradation of A β by neprilysin and insulin-degrading enzyme [110]. This enhancement was dependent on

the lipidation status of apoE [110]. Thus, the importance of cholesterol in AD pathogenesis is not limited to the cholesterol only in the cell membranes, but extends to its presence in brain-specific lipoproteins.

Conclusions

The importance of cellular cholesterol in APP processing relies not simply on the levels of cholesterol in the cell, but also on the distribution of cholesterol in the cell. This distribution can be affected by disease processes (e.g., Down syndrome or Niemann Pick Type C disease), by drug treatments (statins, ACAT inhibitors), or by normal aging. Although an understanding of the normal distribution of cholesterol in the CNS is still just underway, we also need to appreciate the changes that may occur under conditions of acute damage (such as traumatic brain injury or stroke) or chronic damage (such as AD). The redistribution of cholesterol under these conditions could contribute to altered APP trafficking and processing, and to the rate of amyloid deposition.

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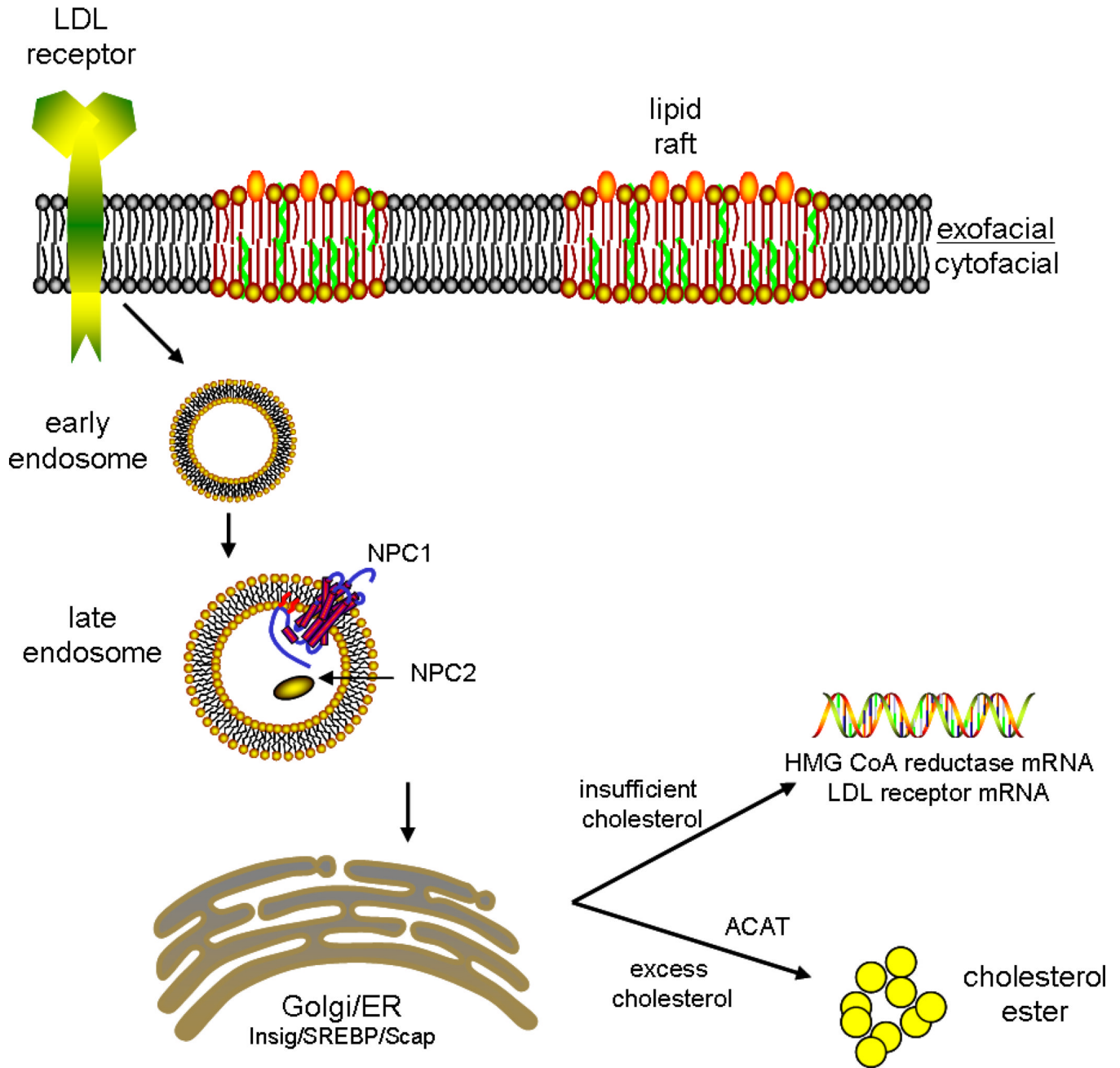


Fig 1. Schematic representation of the intracellular cholesterol transport pathway. Extracellular cholesterol enters the cell via the LDL receptor and endocytosis. Transport from the endosomal system to the endoplasmic reticulum (ER) occurs via transport proteins such as Niemann-Pick Type C (NPC) protein. Once at the ER, excess cholesterol is converted to cholesterol ester by ACAT. If intracellular cholesterol levels drop, the N-terminus of the Sterol Regulatory Element Binding Protein (SREBP) is released and enters the cell nucleus, resulting in transcription of mRNA to increase cholesterol production and increase receptors for extracellular cholesterol.

Cholesterol Synthesis

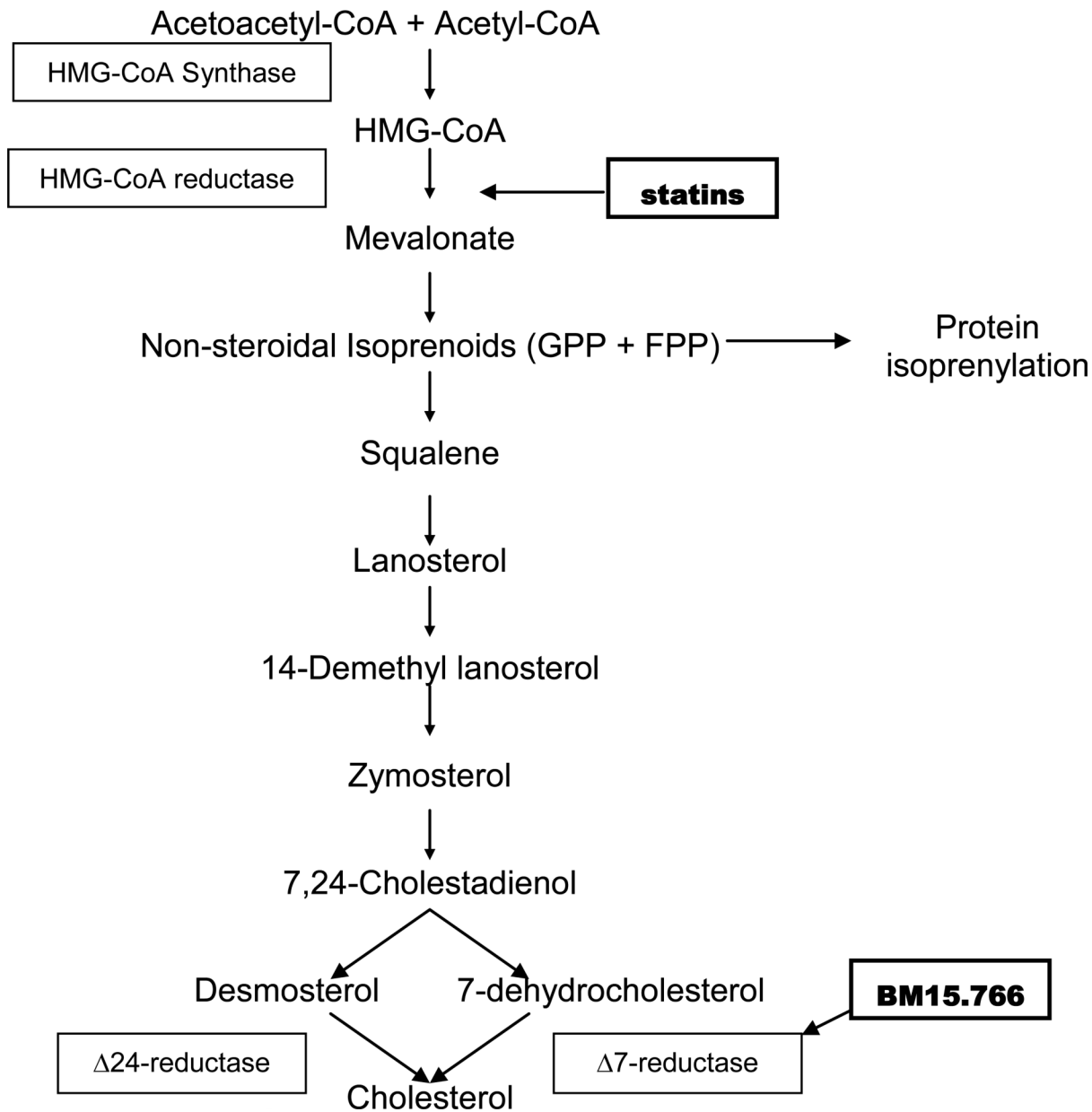


Fig 2. Schematic diagram displaying cholesterol synthesis in human cells showing target enzymes for statins and BM15.766. GPP = geranyl pyrophosphate. FPP = farnesyl pyrophosphate.