Title
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Permalink
https://escholarship.org/uc/item/1gz8d2vj

Journal
Journal of cellular and molecular medicine, 13(9A)

ISSN
1582-1838

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Publication Date
2009-09-01

DOI
10.1111/j.1582-4934.2009.00861.x

Peer reviewed
Calcium in the initiation, progression and as an effector of Alzheimer’s disease pathology

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Received: 15 May 2009; Accepted: 9 July 2009

Abstract

The cause(s) of sporadic Alzheimer's disease (sAD) are complex and currently poorly understood. They likely result from a combination of genetic, environmental, proteomic and lipidomic factors that crucially occur only in the aged brain. Age-related changes in calcium levels and dynamics have the potential to increase the production and accumulation of both amyloid-β peptide (Aβ) and τ pathologies in the AD brain, although these two pathologies themselves can induce calcium dyshomeostasis, particularly at synaptic membranes.

This review discusses the evidence for a role in calcium dyshomeostasis in the initiation of pathology, as well as the evidence for these pathologies themselves disrupting normal calcium homeostasis, which lead to synaptic and neuronal dysfunction, synaptotoxicity and neuronal loss, underlying the dementia associated with the disease.

Keywords: calcium • Alzheimer's disease • amyloid-beta peptide • tau • NMDA • synapse • autophagy • aging

Introduction

Alzheimer’s disease (AD) is an age-related neurodegenerative disorder, which is characterized by progressive memory loss and behavioural changes that become more apparent and more severe as the disease progresses, and accounts for up to 70% of all dementias. Underlying these phenotypic changes are the appearance of several hallmark pathologies. The first is the accumulation of the amyloid-β peptide (Aβ) into extracellular plaques, whereas the second is the formation of neurofibrillary tangles (NFTs) composed of hyperphosphorylated τ protein within neurons [1]. The consequence of the development of these two pathologies is widespread synaptic and neuronal loss, such that a brain with advanced AD can weigh 33% less than a brain from an aged-matched control. AD research has focused on two main areas – the first is in understanding what causes the accumulation of these two pathologies in the aged brain, and the second is understanding how the presence of these pathologies affects the local brain environment to cause the widespread synaptic and neuronal loss, and cognitive decline. Through understanding, it is possible that (1) the accumulation of pathologies in the aged brain can be delayed, prevented, or even reversed, and that (2) the neurodegenerative properties of the pathologies can be dampened. Over the past 20 years aberrant calcium dysregulation has been consistently implicated in AD [2, 3].
either in the initiation of the disease, the progression of AD-related pathologies, or in mediating the neuronal and synaptic loss that results from the presence of the pathologies.

Calcium is an integral signalling molecule whose local and global levels are tightly regulated at a temporal and spatial level as elevations in calcium activate a number of cellular signalling pathways including those crucial for learning and memory such as CamKII and CREB, as well as cell death pathways. In most cell types, cytosolic calcium levels are kept low (~100 nM) relative to the extracellular space and the intracellular stores by calcium-binding buffering proteins (e.g. calbindin), and via the extrusion of cytosolic calcium across the plasma membrane through calcium ATPase pumps and exchangers, and also due to sequestration into intracellular stores such as the endoplasmic reticulum (ER) and mitochondria. Calcium influx into the cytosol occurs across the plasma membrane via store-operated calcium channels, voltage-gated calcium channels or from internal stores. The ER is the largest intracellular store, maintaining a high calcium concentration (100–500 μM) via the unidirectional pumping of cytosolic calcium into the ER lumen by SERCA. Calcium release from the ER into the cytosol occurs via two types of calcium channels: IP3Rs and RyRs. IP3R-mediated release is regulated at the plasma membrane by ligand binding to specific G-protein coupled receptors that induce phospholipase C to cleave phosphatidylinositol-4,5-bisphosphate into diacylglycerol and IP3, which then binds to the IP3R in the ER membrane. Calcium influx across the plasma membrane occurs due to depolarization, through VGCCs, and glutamate receptors such as the NMDA receptor and the mGluR.

Familial and sporadic AD

The overwhelming majority of AD cases occurs with no obvious environmental or genetic cause and is known as sporadic AD (sAD). It typically manifests at a late age of onset of more than 65 years. Around 5% of all AD cases, however, have a Mendelian pattern of inheritance and these are referred to as familial AD (fAD). The age of onset of dementia in fAD can vary dramatically, but typically fAD cases occur at much younger ages (from 30 years or more), and their dementia is usually more severe and progresses more rapidly. Clinically, and neuropathologically, both sAD and fAD are very similar, with all patients exhibiting widespread plaque and tangle pathologies with extensive neuronal loss.

APP processing and the amyloid cascade hypothesis

Studies have shown that cases of fAD are accounted for by mutations in only three genes – the amyloid precursor protein (APP) gene [4, 5], presenilin 1 (PS1) gene [6] and presenilin 2 (PS2) gene [7]. APP is a large membrane spanning protein that contains the Aβ peptide sequence [4]. Aβ is sequentially cleaved from APP. An enzyme identified as BACE first cleaves APP [8] to produce a 99 amino acid stub within the membrane known as C99. C99 is then cleaved by a complex known as γ-secretase to release Aβ [9, 10]. Presenilins comprise the catalytically active subunit of the γ-secretase complex (which also includes nicastrin, APH-1 and PEN-2 [11]), and thus are essential for release of the Aβ peptide.

Although the BACE cleavage site of APP occurs at a specific sequence, the γ-secretase cleavage has loose sequence specificity and can cleave C99 between 38 and 43 amino acids from the N-terminal, to release Aβ peptides of variable lengths. The most common lengths are 1–40 and 1–42. Aβ1–40 is 10–20 times more abundant than Aβ1–42. Mutations in presenilins associated with fAD increase the production of Aβ1–42 at the expense of shorter Aβ peptides [12], whereas mutations in APP are associated with increased BACE cleavage [13], Aβ structural misfolding [14], or an increased prevalence of Aβ1–42 at the expense of shorter Aβ peptides [15].

APP cleavage by BACE and γ-secretase to generate Aβ is not the predominant APP processing pathway. More commonly APP is cleaved by an enzyme with α-secretase activity, at a site juxtaposed to the membrane, to release the large ectodomain known as sAPPα, as well as an 83 amino acid stub within the membrane known as C83. α-secretase cleavage occurs within the Aβ sequence to preclude Aβ production. Hence, stimulation of α-secretase cleavage leads to reduced Aβ production [16], and is a possible therapeutic target for the disease.

The production of Aβ alone is not sufficient to be toxic, and many studies have highlighted the aggregation state of Aβ as being crucial to its toxicity in vitro [17, 18], its ability to impair learning and memory [19] as well as long-term potentiation (LTP) [20, 21], in vivo [22]. Aβ1–42 aggregates more readily than Aβ1–40, and forms the majority of Aβ species present within AD plaques, explaining why mutations that affect the Aβ 40/42 ratio cause fAD without affecting total Aβ levels. More recently, soluble oligomers have been identified as the disease active state of Aβ, and include dimers and trimers [23, 24] to dodecamers [19] and beyond.

The sum of these findings led to the amyloid cascade hypothesis, which states that the accumulation of Aβ causes both sAD and fAD, and the downstream consequences of Aβ accumulation lead to pathology and ultimately synaptic and neuronal loss [25]. Although the causes of fAD are for the most part understood, the cause(s) of sAD are poorly understood. It should be noted that both sAD and fAD are age-related – despite fAD mutations in APP and PS1/2 being present from birth, dementia onset usually requires many decades to manifest suggesting that either pathology accumulation is very slow, or that the younger brain is better able to cope with the increased Aβ pathology that these mutations lead to, either via its degradation or by protecting against the downstream effects of Aβ accumulation. With sAD, the cause of pathology development and consequent dementia is not well understood. It likely results from combinations of genetic, environmental, proteaomic and lipidomic interactions resulting in the accumulation of AD-related pathologies. Understanding both the causes for the appearance of AD-related pathologies and the effects of these pathologies on the local brain environment is crucial in order to understand, prevent or treat the disease.
This review will discuss the possibilities that calcium plays in both disease initiation, progression, and as a disease effector molecule, as summarized in Fig. 1. Notably, the presenilins, in addition to being the catalytic subunit of the γ-secretase complex, are also key regulators of ER calcium homeostasis and have been reported to regulate SERCA [26], IP3 receptors [27] and Ryanodine receptors [28] and also reported to form calcium conducting ER leak channels themselves [29]. fAD linked PS mutations have profound effects on calcium signalling [30], but the significance of this to fAD disease progression is unclear as mutations that affect Aβ accumulation/increased production are sufficient to cause fAD without these additional effects on calcium caused by PS mutations. Thus it seems that, at least with respect to fAD, fAD linked PS mutations alter Aβ metabolism to give rise to fAD, and additionally affect calcium homeostasis which may or may not affect disease progression. The effects and implications of presenilin regulation of both calcium and γ-secretase activity will not be discussed in this review as it was recently covered elsewhere [31].

Calcium in the initiation of pathology

**Effects of calcium on APP processing and Aβ deposition/aggregation**

APP processing lies at the heart of AD, and a majority of therapeutics are designed to alter APP processing to prevent the production of Aβ, to prevent the disease and to halt further progression [32]. Ultimately Aβ accumulation is the tenet of the disease, and accumulation occurs due to an increase in production during aging, a decrease in degradation or a change in predisposition to aggregate. For example, changes in APP metabolism can produce more aggregate prone Aβ1–42 at the expense of shorter Aβ peptides.

**Genetic linkage in calcium related genes and AD**

Calcium has been implicated both in the development of sAD through genetic susceptibility, as well as altered Aβ production.
and metabolism. A number of genes are associated with increased risk of developing sAD, most notably the apoE4 allele [33]. Recently, a polymorphism in a gene encoding a novel calcium conducting channel was found to have linkage to sAD [34], although this same polymorphism was shown not to have linkage in three separate studies of Caucasian populations [35–37]. This channel is called calcium homeostasis modulator 1 (CALHM1) and is a conserved three-transmembrane domain containing glycoprotein. It localizes to both the ER and the plasma membrane. Overexpression of this channel induces a cytosolic calcium influx pathway, which is unaffected by conventional calcium channel blockers, but prevented by the removal of extracellular calcium and non-specific ion channel pore blockers. The induction of this particular calcium influx route into the cytosol results in an increase in sAPPα production, with a concomitant reduction in Aβ, suggesting an effect on one of the α-secretase enzymes. Notably, knockdown of this channel, or the presence of the identified polymorphism for sAD (rs2986017 encoding P66L substitution) decreases calcium permeability and increases Aβ production. These data provide strong evidence that calcium signalling and influx can contribute to the initiation of AD pathology in the aged brain, and that specific calcium pathways can affect APP metabolism. Although further studies are required to replicate the linkage of the CALHM1 rs2986017 polymorphism to sAD, this important finding points to how changes in calcium influx pathways can alter APP processing and Aβ production.

Since a polymorphism in the CALHM1 gene has linkage to increased risk of sAD, it begs the question whether other known risk factors could also directly involve altered calcium homeostasis. One study has shown an association between the NMDA receptor NR2B subunit gene promoter polymorphism and development of sAD in a North China population [38]. The NMDA receptor is an important focal point in AD, both in modulating the appearance of Aβ and α-pathology, and also in mediating Aβ-induced synaptic and behavioural deficits in the disease process. Offering a direct correlation between calcium and cognitive decline, serum calcium levels correlate well with cognitive decline during aging, with elevated serum calcium levels being associated with worse cognitive function [39], and also a faster rate of decline. A regulator of serum calcium levels, the calcium-sensing receptor – is associated with development of AD, particularly with a dinucleotide repeat polymorphism within the promoter sequence [40].

**ApoE4 and calcium**

Of note, the presence of the ApoE4 allele is a strong risk factor for the development of sAD [33] – with ~50% of sAD patients having at least one copy of the ApoE4 allele. Despite the significant impact of this lipid transporter on AD, we do not fully understand its role in disease progression. The ApoE genotype modulates elevated serum calcium and cognitive decline [41]. High serum calcium is associated with worse cognitive function especially in ApoE4 carriers, but this interaction is lost in ApoE2 carriers [41]. Furthermore, the ApoE4 allele is associated with increased calcium rises in neuronal cultures, which can mediate cell death [42], and also in SHSY5Y cultures where increased calcium influxes increase GSK3β activity leading to reduced cell viability [43]. ApoE4 appears to affect calcium influx into neurons via plasma membrane calcium channels [42], which, as discussed below, can alter APP processing leading to increased production of Aβ.

**Calcium and APP processing**

In addition to CALHM1, a number of other studies have reported altered calcium homeostasis affecting the α-secretase dependent processing of APP. Increasing calcium influx into the cytosol with the ionophore ionomycin leads to increased sAPPα production [44], whereas calcium influx through the NMDA receptor increases sAPPα production and decreases Aβ generation [45]. However, despite calcium influx through certain channels stimulating α-secretase processing of APP, the vast majority of studies have shown that increased calcium influx, and elevated cytosolic calcium levels, leads to a net increase in Aβ production [46–48].

More recently, data have highlighted the ER calcium stores as playing an important role in the regulation of Aβ production. We have shown that inhibition of SERCA, either with thapsigargin or by siRNA knockdown, leads to a robust reduction in Aβ generation whereas overexpression of SERCA increases Aβ generation [26]. In agreement with ER calcium stores playing a regulatory role in Aβ generation, knockdown of the IP3 receptor also reduces Aβ generation [27]. In concert, calcium release from the ER via the ryanodine receptor increases Aβ production [49], whereas capacitative calcium entry (CCE), a calcium influx pathway across the plasma membrane, affects APP metabolism with inhibition of CCE leading to increased Aβ 1–42 production [50]. Taken together these studies point to ER calcium as a key regulator of Aβ generation, as well as calcium influx across the plasma membrane through channels such as CALHM1 and CCE channels, and demonstrates how age- or environment-dependent changes in neuronal calcium homeostasis could alter APP processing to drive the increased production of Aβ, or altered metabolism to preferentially generate Aβ1–42.

**Synaptic regulation of Aβ production**

Adding a new dimension to these in vitro experiments, in vivo studies have allowed a new insight into how the workings of neuronal networks can regulate Aβ production, in both mouse models of AD and also in the human brain [51]. Synaptic activity is dependent upon calcium, both at the pre- and post-synaptic terminals, for exocytosis of neurotransmitters across the synaptic cleft, as well as initiation of action potentials at the post-synaptic membrane. In slice cultures from APP overexpressing mice, increased synaptic activity led to increased production of Aβ, whereas inhibition of synaptic activity suppressed Aβ generation [52]. Synaptic activity affects APP processing at either the α- or β-secretase levels, as levels of C99 were modulated. Using microdialysis, and subsequent analysis of interstitial Aβ, it was shown that synaptic activity also regulated Aβ production in vivo [53]. Decreased synaptic activity, elicited by tetrodotoxin (TTX) administration, reduced Aβ and partially corresponded to increased C83 levels,
suggestions an increase in α-secretase or a decrease in BACE processing of APP with decreased synaptic activity. TTX inhibits action potentials by blocking Na⁺ channels, to prevent calcium influx at the pre-synaptic terminal and subsequent exocytosis of neurotransmitters and concomitant endocytosis of synaptic vesicles. Hence, TTX application is likely associated with a decrease in synaptic calcium and a decrease in Aβ generation/secretion. Furthermore, Aβ production and secretion are dependent upon synaptic vesicle membrane recycling; increasing synaptic activity leads to synaptic APP being recycled into endosomes where Aβ generation occurs which is then secreted from the neuron [54]. Though synaptic activity modulates both endocytosis of APP into Aβ-generating endosomes and also exocytosis of neurotransmitter containing vesicles, it is unclear if Aβ is secreted from synapses alongside neurotransmission or in a more passive manner. Furthermore, synaptic activity increases Aβ oligomer formation in a synaptic cleft zinc dependent manner [55], suggesting that synaptic activity can not only increase Aβ generation/secretion but it can also drive it to aggregate into toxic oligomeric species. Of interest, it seems that secreted Aβ at synapses is able to regulate synaptic function through depression of excitatory transmission by an NMDA receptor-mediated pathway [52], suggesting that Aβ is purposely secreted from the pre-synaptic terminal in order to play a role in synaptic plasticity. How this role differs between healthy controls and AD patients is unknown, but may suggest a calcium dependent physiological role for Aβ at the synapse.

Normal age-related changes in calcium homeostasis and AD pathology

Normal brain aging involves subtle changes in cognition and brain functioning, attributed in part to age-related changes in how neurons shuttle calcium from their various internal stores, and altered calcium influx across the plasma membrane. These changes are likely due, partially, to an increased presence of reactive oxygen species, as well as transcriptional and post-translational changes in the calcium channels and receptors themselves. Age-related changes in neuronal calcium handling have been elegantly reviewed elsewhere [3, 56, 57], but the consensus is that aged neurons tend to have increased resting cytosolic calcium levels [58–60] as well as increased cytosolic calcium fluxes due to increased calcium entry through voltage gated calcium channels [61, 62]. This increased influx, in turn, stimulates calcium release from the ER via the ryanodine receptor [63, 64] resulting in further elevated cytosolic calcium fluxes. Voltage gated calcium channels, such as the L-type, are known to be modulated by reactive oxygen species [65], as well as by Aβ [66], both of which are increased through the aging process. Furthermore, aged neurons appear to have reduced calcium buffering capabilities due to reductions in the levels of cytosolic calcium binding proteins such as calbindin D28K [67] and calreticulin [68], reduced mitochondrial buffering capabilities [69], and a decrease in SERCA function [70], at least in peripheral nerves, resulting in prolonged calcium fluxes. Interestingly, IP₃ evoked calcium fluxes appear to decrease in aged neurons [71], perhaps due to a diminished chemical gradient due to elevated cytosolic calcium. Hence, age related increases in cytosolic calcium levels could conceivably alter APP processing as discussed above, and shift APP metabolism to produce increased amounts of Aβ.

Along these lines, aging is associated with a decrease in calcium currents through NMDA receptors in neurons [56]. Since current through the NMDA receptor is associated with increased α-secretase processing of APP [45], it suggests that age-related reductions in NMDA receptor calcium would no longer stimulate the α-secretase enzymes to the same degree, resulting in a shift to Aβ generation. Due to the presence of age-related changes in calcium, coupled with the observations that calcium regulates synaptic plasticity and increases in calcium leads to neuronal cell death, the calcium homeostasis disruption theory of AD was first postulated over 20 years ago [2], and remains a viable theory for the appearance of AD in the aging brain.

Age-related reductions in autophagy

Alterations in neuronal calcium may also affect AD-related pathology through indirect means. Of interest, AD and many other neurodegenerative diseases are age dependent even when mutations in key genes force the aggregation of proteins into toxic conformations. What is the protection that the youthful brain provides against these proteinaceous aggregates seen in neurodegenerative diseases that the aged brain does not afford? Protein turnover is crucial both in neurodegenerative diseases as well as normal and successful aging. Cytosolic proteins are degraded by either the ubiquitin proteasome system, or by the lysosome system through macroautophagy or chaperone mediated autophagy. The UPS system is impaired in AD patients [72], and proteasome inhibition increases levels of both τ and Aβ in vivo [73]. Notably, autophagy is crucial for the removal of cytosolic aggregates especially those implicated in neurodegenerative diseases [74]. Age-related reductions in autophagy are implicated in allowing the aged brain to develop neurodegenerative diseases, whereas stimulators of macroautophagy have proven successful in mouse models of Huntington’s disease, by removing cytosolic aggregates of expanded huntingtin protein from neurons [75]. Cytosolic calcium levels are potent regulators of macroautophagy. Inhibition or depletion of the IP₃ receptor, and L-type voltage gated calcium channel blockers [76] are associated with increased autophagy. Age-related increases in L-type calcium currents and overall increases in cytosolic calcium levels may negatively influence macroautophagy, allowing the buildup of cytosolic proteins in aged neurons such as hyperphosphorylated τ, α-synuclein and huntingtin.

Calcium in disease progression

Linking Aβ, τ and

Although no mutations or polymorphisms in the τ gene (MAPT) are associated with sAD or fAD, τ protein is the component of the
second hallmark pathology of AD – the NFTs [77]. NFTs are composed of aggregated hyperphosphorylated τ and are found in the soma and dendrites of neurons. τ is a microtubule-associated protein that acts to stabilize microtubules enabling efficient axonal transport. τ can be phosphorylated on a number of threonine and serine residues by an expanding list of kinases, most notably cyclin dependent kinase 5 (Cdk5) [78] and glycogen synthase kinase 3β [79]. Abnormal hyperphosphorylation occurs in AD and leads to the disruption of axonal transport and the subsequent relocation and aggregation of the modified τ proteins into structures known as paired helical filaments, which comprise the NFTs. Of note, NFT formation is sufficient to cause extensive neurodegeneration on its own. Mutations in the τ gene are associated with a group of dementias known as tauopathies, including frontotemporal dementia with parkinsonism linked to chromosome 17 [80]. Due to the genetic data that gives rise to fAD, τ hyperphosphorylation is generally considered to be a downstream event of Aβ accumulation, but is thought to be a crucial effector molecule that causes synaptic and neuronal loss [81, 82]. This hypothesis is supported by data showing that Aβ accumulating transgenic mice develop cognitive impairments, but when the τ gene is knocked out the Aβ pathology remains stable but the mice no longer develop cognitive impairments [83]. We have also demonstrated recently that treatment of 3×Tg-AD mice with the sirtuin inhibitor nicotinamide prevents cognitive deficits by removing τ phosphorylated at threonine 231, with no effects on Aβ [84]. Any factor that impacts τ phosphorylation has the potential to play an important role in disease progression in AD. Given that many kinase and phosphatase pathways are activated by calcium, there has been a great deal of interest in studying the pathways that lead to τ hyperphosphorylation through both Aβ dependent and independent pathways. In the AD brain, post-mortem studies have highlighted changes in calcium channels and pumps that correlate with severity of τ pathology, including a reduction in ryanoide receptor binding [85] and IP3 receptor binding [86], in cases with high pathology, suggesting that either NFT formation down-regulates these calcium pumps which may contribute to neurodegeneration, or that a down-regulation of these pumps (perhaps by Aβ pathologies) encourages NFT formation.

Calcium, calpains and τ pathology
Aβ induced increases in calcium lead to τ hyperphosphorylation via both GSK3β and Cdk5 [87]. Cdk5 activity is regulated through its interactions with cyclin-related activator molecules such as p25, p35, p29 and p39. Aβ increases the activity of Cdk5 by promoting the Cdk5-p25 and p35 complexes [88–90]. p39/p35 is cleaved to p29/25 by calpains [91], releasing it from a membrane tether where it forms unregulated functional complexes with Cdk5 leading to τ hyperphosphorylation [92]. Activation of calpain is calcium dependent, and aberrant activation of calpains 1 and 2 has been linked to AD [93, 94]. Calpain 2 has increased activation in NFT bearing neurons and neurites [95], whereas a calpain substrate, calcineurin, is truncated in the AD brain [96]. Hence, Aβ-mediated and age related increases in cytosolic calcium can lead to the activation of calpains, which then lead to hyperphosphorylation of τ via Cdk5 [97]. Of note, a recent study highlighted how calpain activation in transgenic mouse models of AD leads to synaptic dysfunction and behavioural impairments [98]. Calpain 1 is expressed in dendritic spines [99], which have abnormally elevated basal calcium levels in close proximity to Aβ plaques [100] which would have the potential to activate these synaptic calpains. The application of a calpain inhibitor restores synaptic deficits and spatial memory to wild-type levels, demonstrating the role activated calpain plays in mediating Aβ-induced synaptic dysfunction and cognitive decline [98]. Calpain inhibitors also prevent Aβ-mediated cleavage of τ protein into toxic fragments, in particular a 17 kD fragment [101], suggesting that Aβ-induced calcium influx leads to calpain dependent neurotoxicity via cleavage of τ protein, and others including dynamin 1 [101, 102]. These effects could be prevented by blocking calcium influx through the NMDA receptor [102] implicating Aβ, NMDA receptor mediated calcium influx, calpains, τ and neurotoxicity. Along these lines, overexpression of τ itself can lead to increased NMDA receptor activation, leading to increased calcium influx and subsequent activation of calpain, cleavage of τ to produce the aforementioned toxic 17 kD fragment, and subsequent neuronal loss [103]. This suggests that Aβ-induced increases in synaptic calcium could lead to τ pathologies via calpain activation, but that τ pathologies themselves could then influence further increases in synaptic calcium levels through the NMDA receptor.

Calcium as a disease effector
The discovery of a polymorphism in the CALHM1 gene, linking calcium influx to Aβ generation and an increased risk of developing AD provided concrete evidence for a role for calcium in the initiation of sAD [34], but so far three further linkage studies have failed to replicate the association to sAD [35–37]. Despite this, the evidence that age-related changes in neuronal calcium homeostasis could affect APP processing and Aβ metabolism is compelling and offers potential therapeutic targets to try and reverse these changes in calcium, and hence APP processing. However, the appearance of Aβ in old age is only the first step in neurodegeneration and dementia, and there is overwhelming evidence that (1) Aβ induces disturbances in neuronal and synaptic calcium homeostasis and (2) that such changes in neuronal and synaptic calcium would lead to synaptic and neuronal toxicity as sustained increases in cytosolic calcium render neurons susceptible to additional metabolic insults, as well as being able to induce cell death pathways itself. Additionally, given the evidence that altered neuronal calcium fluxes and synaptic activity can alter Aβ generation and secretion, Aβ-induced calcium fluxes could further augment Aβ generation in a positive feedback pathway, accelerating disease progression.

The effects of Aβ on calcium can be categorized into several overlapping groups. The first is a large body of literature
when describing calcium dependent events. Soluble Aβ fluxes and is thought to underlie learning and memory [123]. The second group consists of numerous reports that Aβ can itself form unregulated calcium conducting pores within lipid membranes [119, 120]. Thirdly, there are reports that the presence of Aβ affects calcium buffering, either by affecting intraneuronal calcium sinks/stores or by affecting steady state levels of cytosolic calcium buffering proteins, such as calbindin [121]. Finally, there are also reports that Aβ can affect synaptic transmission, which influences calcium influx at both the pre- and post-synaptic membranes, and is essential for neuronal functioning and cognitive function. The effects on synaptic transmission are likely a consequence of Aβ inducing calcium dyshomeostasis, as well as a cause of further calcium dyshomeostasis.

**Synaptic calcium dyshomeostasis and AD – a focus on NMDA receptors**

Synaptic loss is the best correlate to dementia in the AD brain, although Aβ and t-tau loads associate poorly with dementia. In addition to this, work with transgenic mouse models, which overexpress Aβ, has shown that Aβ induces memory and LTP impairments without overt neuronal loss, and can do so in an acute fashion. As such the site of neuronal plasticity, the synapse, is highlighted as a crucial target for the Aβ peptide as disturbances here would interfere with both LTP and learning and memory, but would only lead to neuronal loss over extended periods of time, possibly beyond the lifespan of rodents. Neuronal synapses contain dense concentrations of calcium conducting channels, as well as cytosolic calcium buffering proteins, and a variety of calcium dependent signalling pathways also implicated in LTP and learning and memory such as calcineurin, CaMKII and CREB, all of which have been reported to be changed in AD brains and APP overexpressing transgenic mice. Hence, in order to satisfy the observation that Aβ acutely impairs LTP and learning and memory, but chronically leads to widespread synaptic and neuronal loss, the effects of Aβ on calcium could initially be at the synaptic level, but with the ability to lead to widespread neuronal loss over time.

One of the most significant advances in recent years in AD research has been the discovery that it is the aggregation state of Aβ that is crucial to its pathophysiological effects, particularly when describing calcium dependent events. Soluble Aβ oligomers appear to mediate calcium dysfunction, as well as inhibit LTP [122], a form of synaptic plasticity, which relies upon calcium fluxes and is thought to underlie learning and memory [123]. The NMDA receptor appears to be a focal point for Aβ oligomer induced calcium dysfunction and LTP inhibition [124], as well as NMDA receptor mediated synaptic loss [125], and Aβ oligomers associate and co-localize with dendritic arbour [126, 127] – post-synaptic membranes enriched in NMDA receptors. Synaptic activity can increase generation/secretion of Aβ at synapses, but can also stimulate the aggregation of monomeric Aβ into oligomeric species. This aggregation of synaptic Aβ appears to be modulated by synaptic zinc, a known facilitator of Aβ aggregation, and by the NMDA receptor, as inhibition of NMDA calcium current influx with antagonists or with zinc chelators prevents synaptic Aβ oligomerization [55]. Intriguingly, synaptic zinc is a known competitive antagonist at the NMDA receptor [128], and is released into the synaptic cleft alongside glutamate, serving to depress synaptic transmission, yet here zinc promotes Aβ oligomerization suggesting that the inhibitory effect of endogenous synaptic zinc on the NMDA receptor is not sufficient to overcome Aβ oligomerization. What is the consequence of Aβ, or Aβ oligomers, associating with NMDA receptors at synapses? It appears that Aβ may have a physiological role in regulating synaptic plasticity via the NMDA receptor, as Aβ secreted from synapses causes NMDA receptor dependent synaptic depression [52], which could arise due to increased endocytosis of the NMDA receptor in the presence of Aβ [109, 111]. This function may go awry in the AD brain where Aβ oligomer formation is promoted, as Aβ oligomers interact with NMDA receptors to promote calcium influx into post-synaptic membranes [129], or this NMDA receptor mediated synaptic depression may be enhanced as it has been reported that Aβ oligomers decrease calcium currents through the NMDA receptor resulting in synaptic loss [125].

In addition to Aβ oligomers interacting with post-synaptic NMDA receptors, they have also been shown to affect pre-synaptic calcium conducting channels including the P/Q type voltage gated calcium channels [108] at very low nanomolar concentrations. Calcium currents through the P/Q type channels were found to be suppressed in the presence of aggregated, but not monomeric, Aβ resulting in decreased synaptic transmission, as exocytosis at the pre-synaptic membrane is dependent upon calcium influx. The authors suggested that this reduced synaptic activity could lead to synaptic loss, as long-term withdrawal of synaptic activity is associated with synaptic retraction. This effect would be enhanced by a concomitant decrease in NMDA receptor calcium influxes in the post-synaptic membrane, in the presence of Aβ oligomers. Thus it seems that there are at least two competing theories for the mode of action of Aβ oligomers at the synapse. It is well defined that Aβ oligomers inhibit LTP, and this appears to be through an NMDA receptor dependent mechanism [124]. One theory is that LTP is impaired due to an inhibition of calcium current at the post-synaptic membrane, due to inhibition of the NMDA receptor, and perhaps a reduction in pre-synaptic activity. A lack of synaptic activity would lead to synaptic retraction over time, in accordance with reduced synapses and synaptic proteins in AD and APP transgenic mice brains. The other would suggest that Aβ oligomers induce increased calcium influx at the synapse, leading to excitotoxicity, and synaptic dysfunction, and eventual synaptic and neuronal loss. This is in accordance with the evidence that dendritic spine calcium is elevated in close proximity to Aβ plaques [100], and the evidence of activated calcium dependent proteases and calcium dependent signalling pathways at synaptic sites. Both would impair LTP, and lead to synaptic loss, but the discrepancies could be due to spatial and temporal reasons – either different...
populations of synapses are affected differentially, or the acute effects of Aβ oligomers are different from chronic exposure.

**Aβ elevates cytosolic intracellular calcium levels in vivo**

Supporting a role that chronic Aβ oligomer exposure leads to increased calcium influx, live imaging of neurites in APP overexpressing mice, revealed elevated dendritic spine basal calcium levels correlating well with close proximity to Aβ plaques [100], indicating once again that the aggregation state of Aβ is crucial but also that the net result of Aβ interactions with synaptic channels/proteins, is elevated calcium. Although the source of increased basal cytosolic calcium is not yet elucidated, it is likely a key event in synaptic dysfunction, as spines with elevated calcium displayed altered morphology tending towards a beaded or dysmorphic appearance. These morphological changes could be rescued by application of calcineurin inhibitors; calcineurin is a calcium dependent phosphatase important for learning and memory as well as LTP, indicating that increased resting calcium levels were atypically activating this phosphatase, whereas calcineurin inhibitors partially restore learning and memory deficits in APP transgenic mice [130]. In addition to calcineurin, other spine located calcium dependent enzymes have been implicated in synaptic dysfunction due to the presence of Aβ – in particular calcium pain inhibitors, as discussed, also prevent Aβ induced LTP and learning and memory deficits caused by the presence of the Aβ peptide [98].

In addition to neurons, astrocytic networks display elevated cytosolic calcium levels, and demonstrate an increased frequency of calcium transients both intracellularly and intercellularly, in APP overexpressing mice [131]. The mechanism of action is distinct from that which causes increased dendritic spine elevations in calcium as astrocytic calcium is unaffected by proximity to Aβ plaques. This suggests that Aβ aggregation state is less important to the effects on astrocytes, and is likely an easily diffusible widespread Aβ species, such as a monomer or dimer. Furthermore, astrocytic calcium dyshomeostasis is not derived, nor related, to neuronal activity, as blocking action potentials does not ameliorate elevated calcium levels or transients in astrocytes. These observations lead to speculation about the implications of widespread astrocytic calcium dyshomeostasis on AD progression. For example, it shows that astrocytes could be severely impacted as a result of pathology in AD; yet this does not seem to lead to astrocyte loss but could be indirectly contributing to the synaptic and neuronal loss seen in the disease, as astrocytes play a large supporting role for neuronal function and survival. As APP is also expressed in astrocytes could these Aβ induced changes in calcium lead to changes in APP processing in astrocytes in human beings – a positive feedback on Aβ production? Further experiments will be needed to answer these questions, as well the source of elevated dendritic spine calcium in close proximity to Aβ plaques.

The relationship between neuronal calcium dyshomeostasis and proximity to Aβ plaques was confirmed by looking at cortical neuronal calcium transients in APP transgenic mice [132]. 21% of cortical neurons demonstrate hyperactivity, with increased frequency of spontaneous calcium transients, but only within 60 μM proximity to an Aβ plaque. A similar distance was reported for elevated basal dendritic spine calcium levels [100]. The source of calcium hyperactivity is dependent upon neuronal action potentials, as TTX prevents all spontaneous calcium transients. Intriguingly, calcium transient amplitudes do not differ between APP and wild-type mice, suggesting only the frequency of transients is affected. Spontaneous transients are presumed to be a consequence of reduced inhibition as GABA, the main inhibitory neurotransmitter, as activators decreased the frequency of spontaneous transients, whereas GABA inhibitors in wild-type mice increased spontaneous calcium transients. The consequences of these spontaneous calcium transients could be the increase in dendritic spine calcium reported in APP mice which lead to altered synaptic morphologies, but also, due to the ordered nature of the transients, could also trigger epileptic like seizures. Such seizures have been reported in other APP mouse models [133] in which the removal of τ was protective against aberrant neuronal activity, as well as in AD patients themselves [134, 135].

**Aβ and mitochondrial calcium dyshomeostasis**

 Mitochondrial dyshomeostasis has been observed in AD brains, as well as in transgenic mouse models of AD, and cell culture models [136]. Mitochondria hold the key to cellular life or death, firstly through the production of ATP, but also through the release of pro-apoptotic signals such as cytochrome C. Mitochondria buffer calcium and are involved in neuronal calcium homeostasis. Aβ, and Aβ oligomers, associate with mitochondrial membranes and have recently been shown to interact with cyclophilin D – part of the mitochondrial permeability pore [69]. Notably, this interaction makes mitochondria more susceptible to swelling in response to calcium, and increases the likelihood of pore permeability transition. Pore permeability transition is associated with a collapse of mitochondrial membrane potential and release of pro-apoptotic proteins into the cell’s cytoplasm, leading to neuronal loss. Furthermore, genetic ablation of cyclophilin D protected transgenic mice against Aβ-induced cognitive deficits, implicating Aβ and mitochondria interactions in neuronal dysfunction and highlighting the importance of normal mitochondrial function to proper synaptic transmission.

**Current therapeutics**

Given the apparent influence of calcium disturbances in the initiation, progression or effects of the disease, can calcium modulators be used as effective therapies for the disease? Currently two classes of drug are used for the treatment of AD – the first are acetylcholinesterase inhibitors, and the second consists of a single drug, memantine, which is a partial NMDA receptor antagonist. Hence, targeting NMDA receptor mediated calcium influx in AD patients is beneficial for cognition [137], as well as in mouse models of AD [138], cell culture models of Aβ toxicity via attenuation of τ.
phosphorylation [139], and is also associated with a reduction in Aβ plaque pathology [140].

Other calcium influx pathway blockers have also been evaluated in AD patients, although without sustained efficacy, including the L-type voltage gated calcium channel blocker nimodipine [141], and other calcium channel blockers associated with the cardiovascular system [142]. Dimebon, an antihistamine, has been shown to improve cognition in mild to moderate AD patients [143], and has reported activity as an NMDA receptor antagonist, as well being able to block voltage gated calcium channel currents, although at concentrations that may be too high to be responsible for the improvements in cognition [144]. Thus, targeting the NMDA receptor appears to be beneficial for the disease, and future drugs may refine this activity further. In the meantime, although calcium channel blockers have not proven effect in patients with the disease, targeting calcium influx may be able to delay or prevent the onset of AD if given at the right time, or targeting specific channels may be able to prevent the synaptotoxicity of Aβ.

Conclusions

Calcium may play a vital role in any number of aspects of AD – from modulating APP processing towards the amyloidogenic processing pathway, to underlying memory and neuronal loss. In vivo evidence has shown that Aβ increases dendritic spine calcium levels, impairs LTP and induces learning and memory deficits, highlighting the synapse as a prime Aβ target, although age related changes in calcium homeostasis may allow the buildup of pathologies associated with AD, through both direct and indirect mechanisms. Synaptic transmission is highly dependent upon tightly regulated calcium at both the pre- and post-synaptic membranes and it is at these membranes all the ion channels Aβ has been reported to disrupt are clustered, likely leading to synaptic dysfunction and more, while targeting NMDA receptors with memantine is beneficial in AD patients, as well as numerous models of the disease, placing calcium dyshomeostasis firmly at the centre of the disease.

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