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Is abnormal axonal transport a cause, a contributing factor or a consequence of the neuronal pathology in Alzheimer's disease?

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Abstract

Axonal transport, the process by which membrane-bound organelles and soluble protein complexes are transported into and out of axons, ensures proper function of the neuron, including that of the synapse. As such, abnormalities in axonal transport could lead to neuronal pathology and disease. Similar to many neurodegenerative diseases, axonal transport is deficient in Alzheimer's disease (AD), a neurodegenerative brain disorder that affects old-age humans and is characterized by the deterioration of cognitive function and progressive memory loss. It was proposed that the synaptic pathology and neuronal degeneration that develops in AD could be caused by an abnormal axonal transport, and that the mutated proteins that cause early-onset AD, as well as the genetic variants that confer predisposition to late-onset AD might somehow impede axonal transport. This paper analyzes the data that support or contradict this hypothesis. Together, they indicate that, although abnormalities in axonal transport are part of the disease, additional studies are required to clearly establish to what extent deficient axonal transport is the cause or the effect of the neuronal pathology in AD, and to identify mechanisms that lead to its perturbation.

Keywords

Alzheimer's disease; amyloid-β peptide; amyloid-β precursor protein; apolipoprotein E4; axonal transport; JNK-interacting protein-1; kinesin; neurodegeneration; presenilin-1; sortilin-related receptor

Alzheimer's disease (AD), the most prevalent neurodegenerative disorder causing mental failure in the elderly [1–3], is a multifactorial syndrome of as yet unknown etiology [4],

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predicted to affect 14 million individuals by 2050 [5]. In a small number of cases (~1%), AD starts early in life, before the age of 60–65 years; these forms of early-onset AD (EOAD, or familial AD) are caused by rare autosomal dominant mutations in three genes: *APP*, *PS1* or *PS2*, all of which alter the metabolism of the amyloid- β (A β) precursor protein (APP) and the production of the amyloidogenic peptide, A β [6–9]. For most cases of AD – those termed late-onset AD (LOAD, or sporadic AD) – the first signs of disease appear at much older ages, and are thought to be caused by yet unknown conditions associated with aging, long-term exposure to detrimental environmental factors and specific genetic predisposition. The latter includes certain common genetic variants with low penetrance [10]. So far, only one clearly established (*APOE4*), and one potential (*SORL1*) susceptibility gene for LOAD have been identified [6, 7,11–13].

The clinical symptomatology of AD is complex, and is highlighted by the deterioration of cognitive function and progressive memory loss. Pathologically, AD has been associated with synaptic dysfunction and neuronal loss, as well as with the presence of neuritic plaques and neurofibrilary tangles in specific brain regions, both of which consist of protein aggregates. Neuritic plaques are extracellular deposits of material (i.e., proteins, proteoglycans and metal ions) that contain aggregated A β at their core, generated by the proteolytic processing of APP. Neurofibrillary tangles are intraneuronal, cytoplasmic lesions consisting of aggregated, hyperphosphorylated tau (a microtubule binding protein). The clinical symptomatology in AD is likely caused by synaptic dysfunction followed by neuronal loss [14,15]. However, what triggers this neuronal pathology, and its relation to the plaques and tangles, is still not clear.

Disruption of axonal transport is part of the pathogenic process in AD, but is the disruption the cause or the result of it?

An interesting idea that has recently gained support is that the synaptic pathology and neuronal degeneration in AD might be linked to abnormal axonal transport [16–18], the process by which membrane-bound organelles and soluble protein complexes are transported into and out of axons [19]. Transport of membranous cargo within axons, known as fast axonal transport, occurs along microtubule tracks and is powered by molecular motors: kinesins (for movement in the anterograde direction) and cytoplasmic dynein (for movement in the retrograde direction) [19]. Kinesin-1, consisting of two heavy chains, which contain the motor domains, and two light chains (KLCs), which bind to the cargo, is the major anterograde motor responsible for transport of many cargoes, including APP. The recruitment of the motors to the transport vesicles, and their release – usually occurring after completion of the transport – are regulated events that determine the rate of transport.

Axonal transport is likely to be abnormal in degenerating neurons of AD brains, judged from the axonal pathology detected in affected brain regions in humans and in mouse models of the disease (see [20–22] for examples; reviewed in [23–25]). However, whether the abnormality is the cause, a facilitating factor, or the result of the disease is unknown. In principle, a block or reduction in axonal transport could cause the depletion of mitochondria, peroxisomes and synaptic vesicle precursors from axons, leading to decreased energy production, increased oxidative stress and dysregulation of synaptic function. An impaired axonal transport could also cause the accumulation of material destined for the axon within the cell body, which could lead to perikaryal degeneration.

Another possibility is that small, more difficult to detect changes in axonal transport of specific proteins, rather than its global perturbation, initiate the neuronal pathology leading to AD. Such subtle deficiencies in the transport of a limited number of proteins may lead to their mislocalization, and loss (or gain) of function at specific intracellular sites. For example, a defective transport of APP, a protein highly relevant for AD, might perturb its proteolytic

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processing and increase the production of the amyloidogenic peptide, A β [26]. Similarly, an abnormal axonal transport of kinases (e.g., cyclin-dependent kinase 5 [Cdk5], glycogen synthase kinase 3 β [GSK3 β], casein kinase 2 [CK2], c-Jun N-terminal kinase [JNK]) could result in improper localization of the entire signaling cascades, and promote aberrant phosphorylation of proteins. Hyperphosphorylation of tau, followed by its aggregation into neurofibrillary tangles [27,28], or of APP, causing increased susceptibility to proteolytic cleavage by secretases [29], may be the result of such processes. Perturbation of the transport of a limited set of proteins has been occasionally reported for other neurodegenerative diseases. For example, a recent study demonstrated that the palette of signaling molecules transported by retrograde axonal transport is changed in neurons from a transgenic mouse model for familial amyotrophic lateral sclerosis [30].

A different view is that the AD-specific neuronal pathology is triggered by dysregulation of cellular processes that are not linked to axonal transport. For example, such conditions could prevent proper recycling of synaptic vesicles [31], or could interfere with normal mitochondrial function [32], both of which could affect the function of the synapse. These abnormalities may thus lead to axonal pathology that is independent of axonal transport. Eventually, this neuronal pathology could also perturb the axonal transport, in a subsequent phase, and the axonal transport deficiency may accelerate and accentuate the neuronal pathology. However, in this case, the alteration of transport is a consequence rather than the cause of the initial pathogenic event.

How could the AD-specific conditions lead to abnormal axonal transport? Do the proteins implicated in the pathogenesis of AD impinge on intracellular transport? This scenario seems likely, since the genetics of AD suggest a link between the proteins associated with the disease and axonal transport. For example, APP and the presenilins – proteins that are mutated in EOAD – have been implicated in the regulation of axonal transport (see later). In addition, APOE4 and SORLA, proteins encoded by genes that confer susceptibility to LOAD, also appear to participate in the regulation of intracellular transport processes (see later). Given the multiple connections between axonal transport and the pathobiology of AD, the question whether a deficient axonal transport is the cause, a facilitating factor or the consequence of AD is essential for the prevention, early detection and, possibly, treatment of the disease. In the following sections, we will briefly analyze the evidence that links the proteins associated with increased risk for AD to abnormal axonal transport.

Is APP a cargo linker or only a cargo protein for kinesin-1?

Approximately one decade ago, a report demonstrating that APP can bind directly to the microtubule motor, kinesin-1 [33] prompted a heated debate aimed at determining how relevant for the *in vivo* situation this interaction is and, if true, how it might impact the field of AD (see [34,35] for examples).

Amyloid- β precursor protein is a type I glycoprotein with receptor-like structure [36], comprising a family of ubiquitously expressed, alternatively spliced and post-translationally processed polypeptides [37]; the neuronal isoform is APP695. Part of the newly synthesized APP follows the secretory pathway and is targeted, as full-length protein, into axons, by fast axonal transport [36]. A fraction of the synthesized, full-length APP reaches the plasma membrane and re-enters the cell via endocytic vesicles. Along the secretory and endocytic pathway, APP undergoes endoproteolytic cleavages by proteases, known as secretases [38, 39], which generate three classes of biologically active peptides: the soluble N-terminal fragments (sAPPs); short middle segments, such as the amyloidogenic, 40–42-amino-acid long A β ; and the carboxy-terminal fragments (CTFs) (Figure 1) [9]. Thus, the processing and transport of APP are coupled events that influence each other.

Two findings from the Goldstein laboratory revived the interest in the axonal transport of APP, in relation to AD. The first showed that APP might function as a receptor for kinesin-1 on transport vesicles, directly binding to the tetratricopeptide repeat (TPR) domain of the KLCs [33,40]. The second finding suggested that $A\beta$ can be generated in transit, within the transport vesicles – thus, linking the transport of APP to AD [26]. Although these studies have since been challenged by several laboratories [35,41,42], they reignited the interest in investigating the intricate relationship between APP processing and transport, and the role a deficient axonal transport might play in the pathogenic process in AD. Let us take a closer look at the possibility that APP is a linker for kinesin-1 to cargo vesicles.

Numerous studies implicate kinesin-1 in the transport of APP into axons [43–47]. However, the notion that APP could have the function of recruiting kinesin-1 to cargo vesicles, as proposed by the Goldstein laboratory, confers to APP a major role in regulating the axonal transport. Although it was initially believed that APP anchors kinesin-1 to a significant fraction of axonally transported vesicles [33], the pool size of vesicles that recruit kinesin-1 via a direct interaction with APP may be small. In addition, more recent data show that the binding of kinesin-1 to APP may not be direct, but instead mediated by scaffolding proteins capable of binding simultaneously to APP and kinesin-1 [48–50]. The JNK-interacting protein-1 (JIP-1) is such a bridging protein that binds APP, via its phosphotyrosine-binding (PTB) domain [51-53], and the KLC subunits of kinesin-1, via a C-terminal motif [54,55]. Under endogenous conditions, JIP-1 mostly recruits kinesin-1 to a phosphorylated form of APP (pAPP; generated by phosphorylation of Thr668 of APP [56,57]), which constitutes a small fraction of the total cellular APP [50]. Fe65, another APP-binding scaffolding protein may fulfill a similar function, since it is capable of binding with high affinity to the KLCs of kinesin-1 [35] and APP [58]. Importantly, since Fe65 binds nonphosphorylated APP with an affinity approximately sevenfold higher compared with pAPP [59], it may primarily recruit kinesin-1 to nonphosphorylated APP (Figure 2).

In addition to recruiting kinesin-1 via JIP-1 and, possibly, Fe65, APP may also recruit the kinesin-2 motor, KIF17, via the PTB domain-containing protein, Mint1/X11 α , which also binds to KIF17 via a PDZ domain (Figure 2) [58,60]. Since KIF17 is primarily a dendritic motor [60], it is likely to transport APP species into dendrites. We note that significant amounts of APP are localized to dendrites. While in principle, APP could serve as an anchor for kinesins to cargo vesicles via the mechanisms described above, the extent to which this actually occurs *in vivo*, under native conditions, has only started to be evaluated. While under normal conditions, pAPP does bind kinesin-1 via JIP-1, and the two proteins are transported into neurites as a complex [50], it is possible that a significant fraction of non-phosphorylated APP are only cargo proteins for kinesins and not cargo linker proteins.

A slowed down axonal transport may increase production of amyloid- β , but does this mechanism operate in AD?

Since kinesin motors may bind – mostly indirectly – to APP, APP gains an important role not only for its own transport, but also for the transport of a number of membrane and soluble proteins that are sorted with APP in the transport vesicle. The premature cleavage of APP, resulting in the generation of A β inside the transport vesicle and shedding of the cytoplasmic C-terminal fragment, could theoretically lead to the release of the motor, and terminate the transport of many axonal proteins. In this scenario, the deficiency in axonal transport is a consequence of the premature cleavage of APP. A reversed situation, where an initial slowing down of axonal transport leads to APP cleavage and abnormal generation of A β can also be envisioned. This scenario is supported by earlier studies from the Goldstein laboratory, which suggest that the APP processing machinery (i.e., the secretases) is packaged into the same transport vesicle that carries APP [26,33]. Thus, an impeded transport could lead to premature

proteolysis of APP and generation of cleavage products – including A β – in places where they may not be properly cleared (Figure 3A) [26]. According to this model, a retarded axonal transport is the initial event that leads to the neuronal pathology in AD, including the abnormal generation and aggregation of A β . The finding that the disruption of kinesin-1-driven transport in transgenic mice leads to increased axonal pathology, production of AB and amyloid deposition is consistent with the model described previously [22]. However, other studies failed to confirm key features of this model, such as the interaction of APP with kinesin-1 [35], and the cotransport of APP with BACE1, the major β -secretase [41]. Moreover, recent studies indicate that a significant fraction of APP is cleaved very early in the secretory pathway, prior to sorting into transport vesicles. Subsequently, these APP-derived fragments are sorted separately, and transported independently to various destinations within the neuron [61]. As a consequence, the axonal transport of APP becomes a vague concept, referring to transport of a large number of APP cleavage products, phosphorylated or not, in addition to full-length APP. Importantly, the cleavage of APP by secretases may or may not result in the release of the cleaved APP intracellular domain (AICD) - to which presumably the kinesin motors bind - from the cargo vesicle into the cytoplasm, as previously thought [61,62]. Thus, the processing of APP within the transport vesicle, if it occurs, would not necessarily release the kinesin, and halt transport. Although attractive, a mechanism where a slowed down axonal transport leads both to the abnormal generation of A β and to further inhibition of transport, owing to cleavage of APP and release of the kinesin motor, may not operate to a large extent in AD.

Does amyloid-β block axonal transport in AD?

Amyloid- β is clearly associated with the pathogenic process in AD. As a result, numerous studies have asked whether A β could be a cause of the axonal transport abnormalities that accompany AD. There are many reasons to suspect that A β is detrimental to axonal transport. Thus, A β is highly toxic for neurons in cell culture and in animal models of disease. When generated in excess, A β (a hydrophobic peptide with high propensity for protein–protein interactions) is likely to affect a multitude of neuronal processes, including axonal transport. While APP-induced axonopathies might not necessarily be inflicted by a toxic effect of A β [63,64], several studies indicate that A β can lead to inhibition of axonal transport via a variety of mechanisms, depending on the oligomeric state of A β (i.e., monomeric, oligomeric or fibrillar) and the site of its action (extracellular or intracellular). We note that, although in AD A β accumulates primarily in extracellular plaques that contain high levels of fibrillar A β , the current thinking is that soluble A β oligomers – acting either from the extracellular space or from the inside of the cell – are the real culprits in AD [14,65].

The effect of extracellular $A\beta$ on axonal transport was investigated in mice *in vivo* [66], and in neuronal cultures [67]; in all cases, axonal transport of vesicular cargo was severely impaired. The impairment of transport was acute, suggesting that it occurs through activation of specific signaling pathways. Interestingly, the trafficking of mitochondria was particularly sensitive to treatment with $A\beta$, and the signaling pathway leading to the blockade of mitochondrial motility was mediated by GSK3 β [68]. Whether this signaling alters the microtubules, the motor itself, or the attachment of the motor to the cargo or to the microtubules, remains to be established. We note that the inhibition of transport appears to be more pronounced for fibrillar compared with monomeric $A\beta$, although there is no consensus on this among the different studies.

The effect of intracellular $A\beta$ on axonal transport was studied in isolated squid axoplasm [69] – an *in situ* system that shows robust vesicular transport [70]. Perfusion of the axoplasm with soluble $A\beta$ oligomers, but not monomeric or fibrillar $A\beta$, disrupted the fast axonal transport via a pathway that led to the phosphorylation of KLCs by casein kinase 2, and release of kinesin-1 from the transport vesicles (Figure 3B) [69]. We note that the perfusion of squid axoplasm with $A\beta$, as performed in this study, does not reflect the expected topology of $A\beta$,

which normally should be confined within membrane compartments rather than being free in the cytoplasm. Even if A β is taken up from the extracellular space via endocytosis, it should still remain encapsulated inside the endocytic vesicle. However, occasionally, intraneuronal A β was detected in the cytoplasm [71], where it could have escaped by translocation across the encapsulating membrane, or through leak-age from a damaged membrane compartment. While these remain real possibilities (since A β is a hydrophobic peptide and could, in principle, cross or translocate through membranes), they need to be confirmed experimentally.

Unlike the soluble cytoplasmic A β , the intraneuronal accumulations of A β , present occasionally within membrane compartments in the neurites of cultured neurons, do not globally perturb the axonal transport [72]. However, transport and distribution of mitochondria within the neurites that contain such accumulations are perturbed (Figure 4), and the neurites themselves appear dystrophic [73]. These and other results indicate that transport of mitochondria is particularly sensitive to AD-related insults and may thus be a major cause of the detected axonopathy.

The studies previously summarized clearly show that $A\beta$ has a detrimental effect on axonal transport, acting via a multitude of pathways. However, whether this is a mechanism for impairment of axonal transport in AD is still unknown.

Presenilin-1 regulates vesicle transport, but its enzymatic activity may not be required

The idea that APP processing and generation of $A\beta$ could regulate axonal transport prompted the re-investigation of presenilins – proteins encoded by genes mutated in EOAD, which participate in APP processing – as possible regulators of axonal transport. Presenilin (PS)-1 had been previously shown to play a role in intracellular trafficking of various proteins (see [74–76] for examples). Several recent studies have indeed demonstrated that PS1 regulates axonal transport, and that EOAD-linked mutations in PS1 lead to the impairment of fast axonal transport [77,78]. The exact mechanisms by which PS1 controls axonal transport are not fully understood, but may not involve the proteolytic activity of PS1. It was proposed that EOADlinked PS1 mutants might somehow activate GSK3 β , leading to increased phosphorylation of kinesin-1 and its release from the cargo vesicles (Figure 3B) [77,78]. The PS1 mutants would thus trigger axonal pathology by acting directly on axonal transport. However, this may be an oversimplification, since a recent study demonstrated that EOAD-linked PS1 mutants may actually alleviate axonal transport defects caused by EOAD-linked APP mutants [64].

Opposing effects of tau on kinesin-1-based transport

Although no mutations in *MAPT* (the gene encoding tau) were found in AD, tau is clearly linked to the pathology of AD [7]. The binding of tau to microtubules leads to their stabilization [79], and should thus enhance the transport of cargo along microtubules. In principle, this positive effect on axonal transport could be counterbalanced by the inhibition of kinesin-1-driven transport [80], likely due to blocking of kinesin-1 binding to microtubules by tau [81–83]. Thus, the net effect of increased tau levels, presumably occurring in AD [84], on axonal transport may not be very detrimental. For the same reason, a reduced affinity for microtubules of hyperphosphorylated tau [85] – also observed in AD [86–88] – may not necessarily perturb axonal transport through the diminished binding of phosphorylated tau to microtubules (causing thus decreased microtubule stability), but rather via other mechanisms. However, tau overexpression and abnormal phosphorylation likely contribute to the formation of disease-related intraneuronal structures, such as the neuropile threads and neurofibrillary tangles, which themselves may physically damage the microtubules and obstruct transport within the neurites [89,90]. Moreover, the extent by which the opposing effects of tau on axonal transport may

balance each other out depends on the concentration of tau and on its affinity for microtubules (regulated, in part, through phosphorylation of tau). This could explain the apparently contradictory results obtained using different experimental systems, with regard to the effect of tau on axonal transport. Indeed, while overexpression of tau in *Drosophila* motor neurons [91,92] or mammalian neurons [80] was shown to interfere with organelle transport (including transport of mitochondria), other studies, using the squid axon system or transgenic mice, found that axonal transport is unaffected by increased concentrations of tau [93,94].

The effect of tau on axonal transport in AD may be difficult to assess in experimental systems that do not reproduce the full palette of disease-related changes, such as the presence of $A\beta$. In this respect, one study demonstrated that tau, instead of stabilizing microtubules, facilitates their disassembly, initiated by the addition of extracellular, prefibrillar A β [95]. Thus, tau may have opposing effects on the stability of microtubules, depending on specific extracellular and intracellular conditions. Interestingly, aberrant activation of Cdk5 in transgenic mice leads to a typical AD brain pathology, including the development of neurofibrillary tangles (aggregated and hyperphosphorylated tau), enhanced amyloidogenic processing of APP, intraneuronal accumulation of A β , and neuronal degeneration and loss [96,97]. Although the neuronal degeneration observed in these mice is compatible with axonal transport defects, it is difficult to determine if this is a cause or the result of the AD-like pathology. However, a recent study demonstrated that the selective disruption of kinesin-1-driven transport in a transgenic mouse triggers a cascade of events that leads to tau hyperphosphorylation and accumulation [98]. Whether such a mechanism may operate in AD is difficult to predict, since no tau pathology develops in most mouse models of AD, in spite of extensive axonal pathology. We note that axonal transport defects are seen in familial forms of frontotemporal dementia with parkinsonism linked to chromosome 17, as well as other tauopathies, which are caused by mutations in the MAPT gene, and show extensive tau pathology, similar to AD [24,99,100]. Future studies are needed to fully elucidate how tau might regulate axonal transport in vivo, and how this regulatory role might be compromised in AD.

APOE4 allele: a major risk factor for LOAD that may cause disruption of axonal transport

ApoE, the major apolipoprotein in the brain, with an essential role in cholesterol uptake and transport, is encoded by the *APOE* gene, which exists as a polymorphic locus represented by three alleles, ϵ_2 - ϵ_4 . Of these, the ϵ_4 allele confers high risk for LOAD. The mechanism by which APOE4 predisposes to AD is not fully understood, but may include modulation of APP metabolism and A β aggregation, regulation of brain lipid metabolism and synaptic function, as well as an effect on axonal transport [23,101]. In support of the latter, it was found that transgenic expression of human APOE4 causes significant tau hyperphosphorylation, axonal pathology and disruption of axonal transport, in brain and spinal cord neurons [102,103]. The disruption of axonal transport may be caused by a direct effect on the cytoskeleton (possibly by the phosphorylated tau), or on the interaction of kinesin-1 with the cargo. Indeed, the APOE4 receptor, APOER2, is known to bind to JIP-1 [104], and this interaction may compromise the role of JIP-1 as anchor of kinesin-1 to the cargo vesicle (Figure 2).

Sortilin-related receptor, SORLA, a possible risk factor for LOAD, regulates sorting & intracellular trafficking pathways

Systematic meta-analyses of AD genetic association studies have identified additional loci that confer AD risk, but the risk is modest and inconsistent among different studies [6,7]. Of these, the sortilin-related receptor, SORLA, with a role in sorting and intracellular trafficking pathways [105], was shown to interact with APOE and APP, and to affect $A\beta$ generation

[106–108], and thus, it could be, via these proteins, indirectly involved in the regulation of axonal transport. However, the association of *SORL1* (the gene encoding SORLA) with LOAD is still debated [7].

The chicken or the egg dilemma of AD: which comes first, axonal transport deficiency or neuronal pathology?

There is no doubt that an abnormal axonal transport is part of the pathogenic process in AD, as well as the evolution of the disease. However, it is still not clear whether a deficient axonal transport constitutes the very initial event that triggers the neuronal pathology in AD (Figure 5). As previously described, there is considerable evidence suggesting that the EOAD-linked genes, APP and PS1, encode proteins with a role in the regulation of axonal transport. There is also some evidence that the mutations or allelic variations in the genes that predispose for LOAD may adversely affect axonal transport. However, there is no definitive evidence demonstrating that the disruption of axonal transport through these AD-linked proteins is the cause of the axonal pathology and synaptic failure that are characteristic for the disease. The proteins that confer risk to AD, although having an effect on axonal transport, could also have other effects on processes that are not related to axonal transport. For example, they could target the activity of mitochondria, the generation of toxic reactive oxygen species or the composition of intracellular membranes. In this case, the abnormalities of axonal transport could be only a secondary factor, which does not itself trigger, but contributes to the disease process. Finally, it is possible that deficiencies in axonal transport, caused by factors that normally appear in old age, initiate the cascade of events that lead to neuronal pathology. For example, environmental stress (e.g., oxidative and osmotic) is known to perturb axonal transport via specific signaling [109]. This cascade may involve proteins that are linked to AD, such as APP and the presenilins.

Since axonal transport both depends on, and affects many intracellular processes, it will be difficult to clearly establish the sequence of events that lead to its perturbation. According to a model of initiation of the neuronal pathology proposed by Kamal *et al.* [26], the abnormal axonal transport is both causative and consequential of the pathology (Figure 3A). In this case, the slowed down transport of the APP-carrying vesicle would lead to the premature cleavage of APP and the generation of A β , which in turn will further block axonal transport (due to the concomitant release of the kinesin motor). Vice versa, a premature cleavage of APP and generation of A β would lead to slowed down or blocked transport, which in turn, will facilitate further cleavage of APP. Although more recent studies no longer support this mechanism, the alternative models that were recently advanced also envision an abnormal axonal transport that is caused by the release of the kinesin motor from the cargo vesicle (Figure 3B). Whether this is triggered by a signaling downstream of intracellular A β , or from mutant presenilin, or by yet another mechanism, is a question of mechanistic details that needs to be solved by future studies.

Future perspective

The dilemma of whether an abnormal axonal transport is a cause or a result of the AD neuronal pathology will not be easily solved. This will require very sensitive assays to detect subtle defects in axonal transport and APP processing. In addition, these parameters should be accurately measured *in vivo*, at least in animal models of AD. Methods to image and noninvasively quantify axonal transport and axonal degeneration are already being developed [20,110–113]. For now, it appears that the deficiencies in axonal transport represent important components of AD, as they very likely appear relevant for a variety of other neurodegenerative diseases [24,114]. It is certain that, owing to the importance of axonal transport for the

physiology of the neuron, any dysfunctionality in axonal transport will lead to some form of disease.

From a clinical point of view, it will become necessary to identify ways to enhance axonal transport for preventive and therapeutical reasons. Implicitly, this requires the identification of the molecular species that trigger the abnormal transport, and of the proteins involved in axonal transport, whose functions are affected by old age or by the disease process. The interventions should also aim to stabilize the axonal microtubule network. This could be carried out by increasing expression of microtubule-associated proteins, capable to reduce the microtubule dynamics without interfering with kinesin motility, or by promoting posttranslational modifications of tubulin that increase microtubule stability, such as acetylation and detyrosination (see [115,116] for examples). Since these modifications are the result of enzymatic reactions, the responsible enzymes could be targeted with specific drugs, pending the solving problems of delivery and interference with other intracellular processes. Eventually, correcting deficiencies of axonal transport associated with AD will also allow for the clear establishment of whether these deficiencies are a cause or consequence of the neuronal pathology. If the neuronal pathology is corrected by the intervention, then the abnormal axonal transport is most probably the cause or a facilitating factor of the pathology, but not a consequence of it.

Executive summary

Is abnormal axonal transport a common mechanism of disease in early-onset & late-onset Alzheimer's disease?

- Several lines of evidence suggest an affirmative answer to this question, and implicate proteins relevant to the disease process in the regulation of axonal transport.
- The axonopathy observed in post-mortem Alzheimer's disease (AD) brains and in mouse models of AD suggests impeded transport. Quantitatively, decreased rates of transport are detected in mouse models of AD.
- Increased expression of Amyloid β precursor protein (APP) in transgenic mice causes abnormal transport of select proteins.
- Amyloid-β, applied either intracellularly or extracellularly, disrupts axonal transport.
- Mice expressing early-onset AD-specific presenilin 1 variant show impaired axonal transport.
- Presenilin 1, a key enzyme in APP processing, can regulate axonal transport via mechanisms that target the binding of kinesin-1 to the cargo vesicles.
- Tau can regulate kinesin-driven transport, but the mechanisms could involve more than regulating the stability of microtubules and the accessibility of kinesin-1 to microtubules.
- Alteration of axonal transport in a KLC1-null mouse leads to tau hyperphosphorylation.
- Aberrant activation of Cdk5 in a transgenic mouse featuring AD-related amyloid and tau pathology leads to axonal pathology consistent with transport deficiencies.
- Mice expressing APOE4, which in humans predisposes for late-onset AD, display pathology consistent with disruption of axonal transport.

Conclusion

- Overall, these observations are consistent with the notion that axonal transport slows down in old age, and suggest that an abnormal axonal transport could contribute to AD.
- Nevertheless, the AD-related conditions could cause axonal pathology via mechanisms not involving axonal transport, but at the same time, could indirectly affect axonal transport.
- Thus, the alterations in axonal transport could be both the cause and the consequence of AD-specific neuronal pathology.

Future perspective

- Careful studies investigating the appearance and development of neuronal pathology in correlation with live imaging of axonal transport in animal models of AD are required. Since no animal models fully reproduce the pathology of the human disease, noninvasive methods to monitor the axonal transport in human brain neurons *in situ* are required.
- Identifying the means to correct transport deficiencies by enhancing axonal transport will soon become a priority for AD therapeutics.

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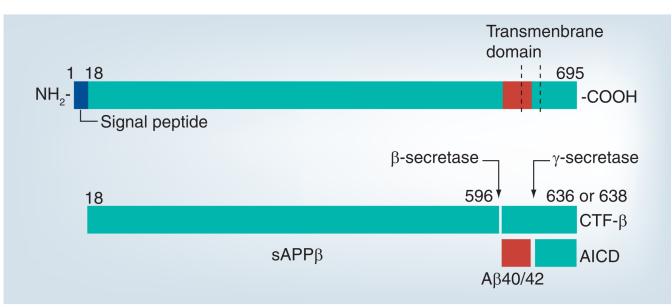


Figure 1. Processing of amyloid- β precursor protein via the β -secretase pathway

The positions of the transmembrane domain, of A β (red), and of the cleavage sites are marked. The APP-derived proteolytic fragments, sAPP β , A β , AICD, and the intermediary cleavage product, CTF- β , are also shown. The cleavage by γ -secretase at positions 636/638 generates A β 40/42.

A β : Amyloid β ; APP: Amyloid- β precursor protein; AICD: APP intracellular domain; CTF- β : Carboxy-terminal fragment- β . Modified from [4,61].

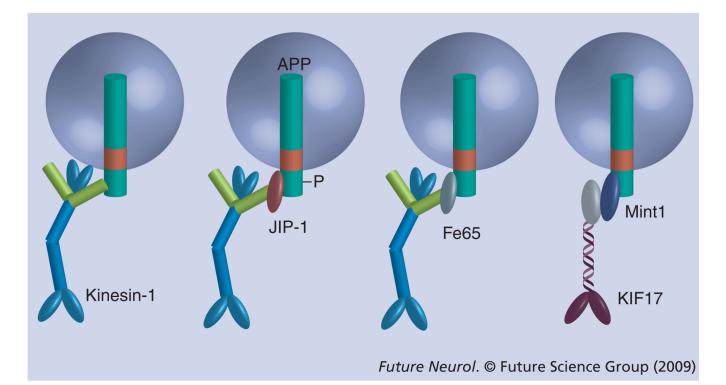


Figure 2. Amyloid-β precursor protein recruits kinesin motors to cargo vesicles via adaptor proteins JIP-1, Fe65 and Mint1/X11a bind to APP via their phosphotyrosin-binding domains. JIP-1 and Fe65 also bind to the light chains of kinesin-1; Mint1/X11α binds to KIF17, a kinesin-2 motor with dendritic targeting. In the cellular context, JIP-1 binds preferentially to phosphorylated APP, while Fe65 binds to nonphosphorylated APP. The direct binding of kinesin-1 to APP, as shown to the left, could be less relevant for the *in vivo* situation.

APP: Amyloid-β precursor protein; JIP: JNK-interacting protein.

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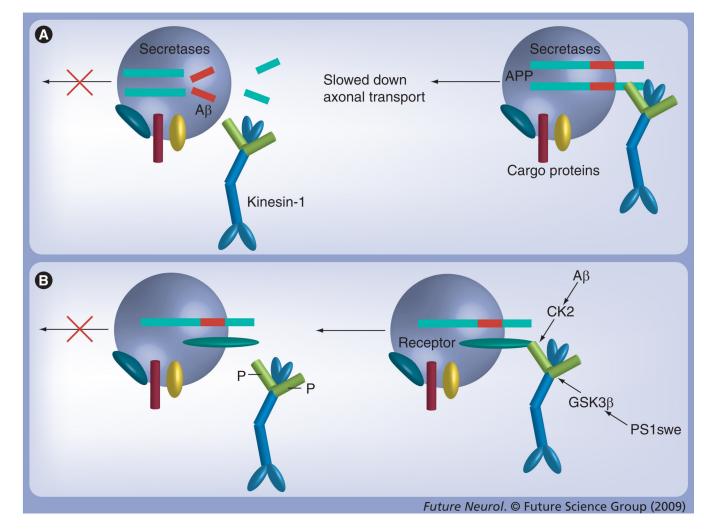


Figure 3. Hypothetical models for the release of kinesin-1 from transport vesicles, leading to disruption of the fast axonal transport in Alzheimer's disease

(A) A slowed down axonal transport at old age could allow premature processing of APP and release of kinesin-1 from the transport vesicle, leading to a halt in transport (crossed arrow) [26]. The cessation of transport further facilitates cleavage of APP molecules by cotransported secretases (not depicted), thus amplifying the loop where the deficiencies in axonal transport and processing of APP potentiate each other. Although this model is no longer regarded as a major mechanism operating in Alzheimer's disease (AD), the basic principle for the disruption of the axonal transport by premature release of the motor from the transport vesicle remains valid. (B) The regulation of axonal transport by phosphorylation of the light chains of kinesin-1 by casein kinase 2 (CK2) [69] or glycogen synthase kinase 3β (GSK3 β) [77,78]. The phosphorylated kinesin-1 is released from the vesicle, causing a halt in transport (crossed arrow). The signaling cascades leading to phosphorylation could be triggered by the AD-linked increased production of A β oligomers, or by a mutated presenilin-1 (PS1swe) present in some cases of early-onset AD. It should be noted that, (A) kinesin-1 binds to APP or (B) to another receptor protein. Other cargo proteins are shown.

A β : Amyloid- β ; APP: Amyloid- β precursor protein.

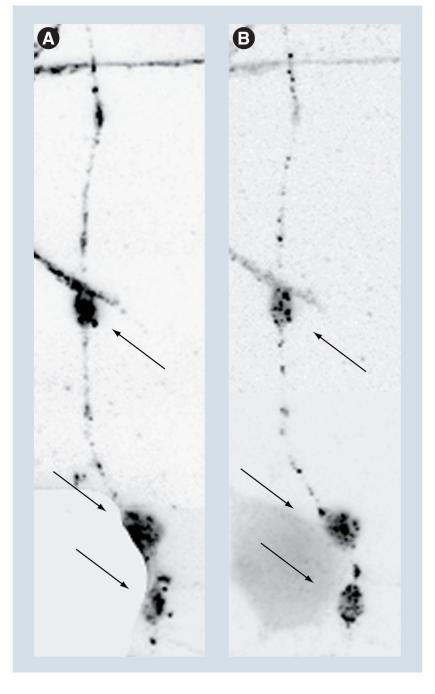


Figure 4. Clusters of mitochondria colocalize with amyloid-β deposits

Clusters of mitochondria colocalize with amyloid- β (A β) deposits (arrows) within the neurite of a CAD cell (a CNS-derived neuronal cell line [117]). Neurites containing A β deposits also contain fewer mitochondria [73]. (A) The mitochondria and (B) A β accumulations were detected with an antibody to lipoic acid and antibody 6E10 (Signet, Dedham, MA, USA), respectively. The inverted, fluorescence microscopy images were adjusted for contrast and brightness, to facilitate the evaluation of colocalization.

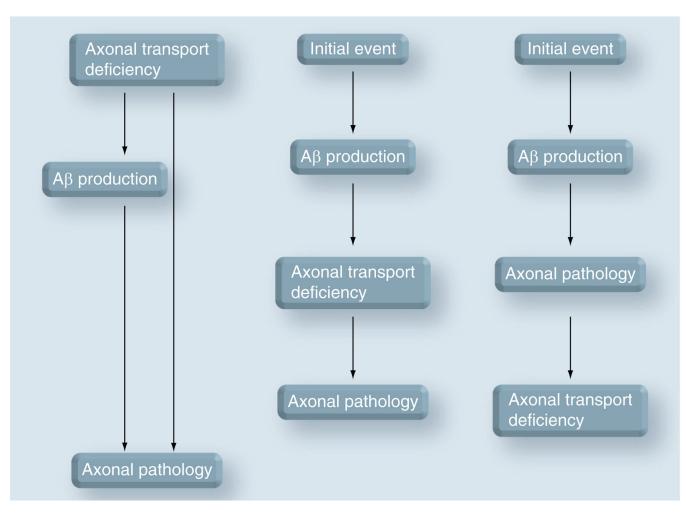


Figure 5. Relationship between abnormal axonal transport and axonal pathology in Alzheimer's disease

(A) Deficiencies in axonal transport (caused by factors that normally appear at old age or by mutated proteins with a role in axonal transport) inflict axonal pathology by depriving the axon of essential factors, and/or by causing an abnormal/increased production of toxic amyloid- β (A β). In this case, the axonal transport deficiency is the cause of the axonal pathology. (**B**) An initial event, not related to axonal transport, induces abnormal/increased production of toxic A β , which blocks axonal transport and leads to axonal pathology. Although here, the axonal transport is not itself the initial cause, it is an essential part of the mechanism that leads to axonal pathology. (**C**) Axonal pathology is caused by a toxic effect of A β on the synaptic function, independently of axonal transport (see [31] for example), triggering neuronal death by a dying back mechanism [118], when axonal transport is also perturbed. In this case, the deficiencies in axonal transport are a result of the neuronal pathology of Alzheimer's disease. The mechanisms described above are focused on A β . Other scenarios, where the abnormalities in axonal transport are a cause, a facilitating factor or a result of the neuronal pathology can be envisioned.

Aβ: Amyloid-β.