

# Intracellular and Extracellular A $\beta$ , a Tale of Two Neuropathologies

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**The central pathological cause of Alzheimer disease (AD) is hypothesized to be an excess of  $\beta$ -amyloid (A $\beta$ ) which accumulates into toxic fibrillar deposits within extracellular areas of the brain. These deposits disrupt neural and synaptic function and ultimately lead to neuronal degeneration and dementia. In addition to the pathological roles attributed to A $\beta$ , evidence from our laboratory would suggest that A $\beta$  serves a physiological role in the modulation of CRE-directed gene expression. This commentary also highlights some of the pathological consequences of the accumulation of intracellular A $\beta$ . Finally it discusses the impact of cortical A $\beta$  burden on transmitter-specific synaptic numbers as well as the generation of dystrophic neurites. The fundamental thesis of my proposal is that the A $\beta$  pathology seen in AD is a continuous process from an initial abnormal A $\beta$  intracellular accumulation to the well-established extracellular A $\beta$  aggregation, culminating in the formation of amyloid plaques and dystrophic neurites.**

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## INTRODUCTION

There is little doubt that overproduction of the A $\beta$  peptide in the CNS plays a central causative role in unleashing the elaborate Alzheimer's neuropathology. The most compelling evidence stems from findings that the overproduction of its precursor protein (A $\beta$ PP, the A $\beta$  amyloid precursor protein) in Down syndrome leads to an Alzheimer-like pathology (for review, see 47). Similarly, mutations in A $\beta$ PP and the presenilins result in the accelerated production of high levels of A $\beta$  peptide in the familial forms of Alzheimer disease (AD) (21) with a neuropathology similar to the later onset and more protracted development of the sporadic forms of the disease. It needs to be acknowledged at this point that AD pathology is not only restricted to A $\beta$ PP dysmetabolism and A $\beta$  overproduction, but that it is also accompanied by neurofibrillar (25, 37) and synaptic pathologies (50), oxidative stress (22) and inflammation. A $\beta$  production and deposition could be affected by a number of factors, such as high cholesterol levels (45) and brain trauma (53). As A $\beta$  is found in diverse forms, amounts and localization in the CNS, the precise contribution of each in provoking Alzheimer syndrome remains an issue of great controversy and interest in the field.

In this commentary, I would like to suggest that A $\beta$ PP fragments, including A $\beta$  itself, serve a physiological role under normal circumstances, but that this changes progressively from an intracellular to an extracellular neuropathology in the AD scenario. These ideas are based on our own observations in animal models and from the observations of others as cited in this opinion paper.

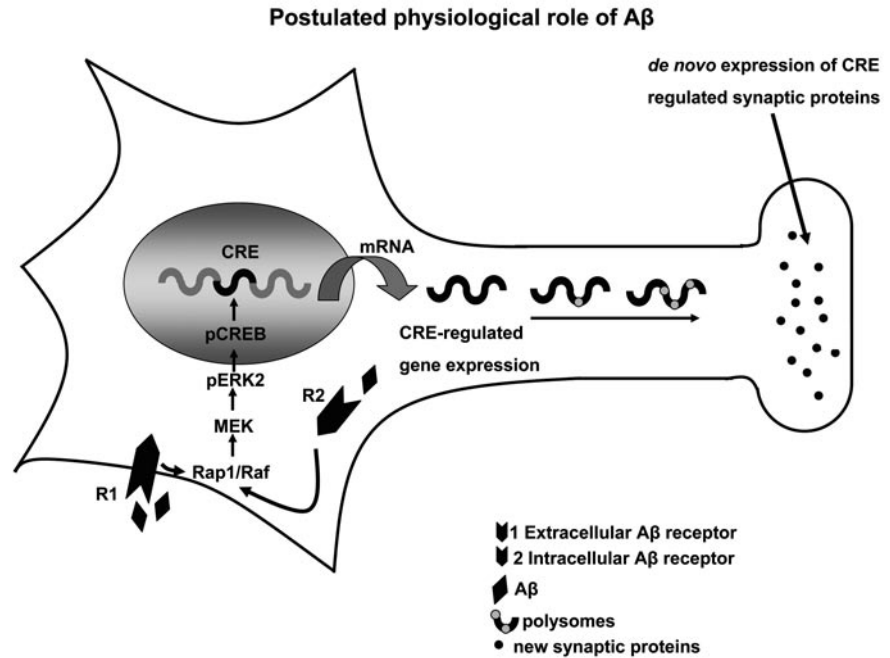
## A FUNCTIONAL ROLE FOR A $\beta$ AND A $\beta$ PP FRAGMENTS?

A $\beta$  fragments appear to be produced under physiological conditions by the concerted action of 2 convertases referred to as  $\beta$ - and  $\gamma$ -secretases (for review, see 10, 62). The identities of these convertases have been elucidated. The main  $\beta$ -secretase is BACE 1 (beta amyloid convertase enzyme) (55). On the other hand, the  $\gamma$ -secretase is a membrane protein complex of which presenilin provides the catalytic function (for review, see 29). Activation of the  $\gamma$ -secretase generates intracellular C-terminal fragments from 2 membrane-resident proteins: Notch and A $\beta$ PP. These resultant fragments are called NICD (Notch Intracellular C-terminal Domain) and AICD (A $\beta$ PP Intracellular C-terminal Domain), respectively. There is evidence that these fragments migrate to

the nucleus and can directly or indirectly modulate gene expression (for review, see 31, 32). The Notch pathway is implicated in the development of the CNS while the functions of the AICD fragment are less clear. Yet, the existence of such pathways suggests that therapies aimed towards the suppression of  $\gamma$ -secretase activity may eliminate vital neuronal regulatory functions. Moreover, cleavage of A $\beta$ PP by  $\alpha$ -secretases within the A $\beta$  domain prevents the formation of A $\beta$  amyloidogenic fragments and releases into the extracellular domain the A $\beta$ PP N-terminal fragment referred to as soluble A $\beta$ PP alpha (sA $\beta$ PP $\alpha$ ). Presently, the metallopeptidase, ADAM 10, is thought to be the best candidate for this role (1). This  $\alpha$ -secretase activity is apparently regulated by the CNS activation of muscarinic receptors (41), leading to the production of sA $\beta$ PP $\beta$  fragments which apparently display neurotrophic effects in vitro (38). It is therefore plausible that in the early AD stages an overproduction of A $\beta$ PP may generate sA $\beta$ PP $\alpha$ , thus favoring the transient de novo formation of cortical-hippocampal synapses. If this were the case, it could explain the initial up-regulation in the number of cholinergic synapses in pre-plaque stages (63) as well as the transient up-regulation of cortical glutamatergic and GABAergic synapses in early stages of plaque formation observed in cortical areas of AD transgenic models (3). In line with this, De Kosky and collaborators (9, 24) have found an upregulation of choline acetyl transferase activity in the cortex and hippocampus of patients with mild cognitive impairments (9) in which a trophic response to sA $\beta$ PP $\beta$  should not be rule out.

A pressing and somewhat neglected question is: Is there a physiological role for A $\beta$  in the CNS? We know that a certain amount of A $\beta$  appears to be generated intracellularly and is transported axonally un-

der normal conditions, however, its functions at the synaptic level is not yet clear (27, 30, 46, 61). The prevailing notion is that pathological amounts of A $\beta$  accumulate in extracellular spaces of the brain and participate in AD's neuropathology as well as the demise of synapses. However, under basal physiological conditions, it is possible that A $\beta$  plays a quite different role in reinforcing synaptic plasticity. For example, our laboratory has observed that undifferentiated rat pheochromocytome PC12 cells display an activity-dependent (high molarity K<sup>+</sup> and forskolin-induced) up-regulation of phospho-ERK2 along with an elevation in CRE-regulated gene expression. These responses were markedly greater in PC12 cells transiently transfected with mutated forms of human A $\beta$ PP and expressing modest amounts of endogenous A $\beta$  (7, 13) (also Echeverria et al, unpublished data). In these transfected cells the stimulus-dependent increase in CRE-directed gene expression was blocked by  $\gamma$ -secretase inhibitors, suggesting that this response was mediated by either A $\beta$  or the AICD peptide. Conversely, the over-expression of the AICD fragment did not replicate the increase in CRE-directed gene expression, suggesting that A $\beta$  was the responsible agent. Our observations, thus, would indicate that endogenously produced A $\beta$  peptides, acting on either intracellular or extracellular receptors, can facilitate CRE-directed gene expression. These observations are of functional relevance as CRE-regulated gene expression plays a key role in the development of late LTP formation requiring *de novo* protein synthesis (for review, see 28, 58). We speculate that, at this stage, the A $\beta$  peptides are produced transiently and in relatively low amounts, and that they facilitate rather than inhibit the CRE-directed gene expression. This regulatory mechanism apparently requires the phosphorylation of the MAP kinase ERK 2, and the application of suitable inhibitors has shown that this signalling mechanism involves the Rap1-MEK-ERK pathway (7, 13) (also Echeverria et al, unpublished data). It is therefore feasible that in normal circumstances, low levels (pM-nM concentrations) of A $\beta$  would stimulate CRE-directed gene expression and neural plasticity in the CNS. On the other hand, at higher, non-physiological levels, A $\beta$  would inhibit CRE-directed gene expression. It is still unclear whether



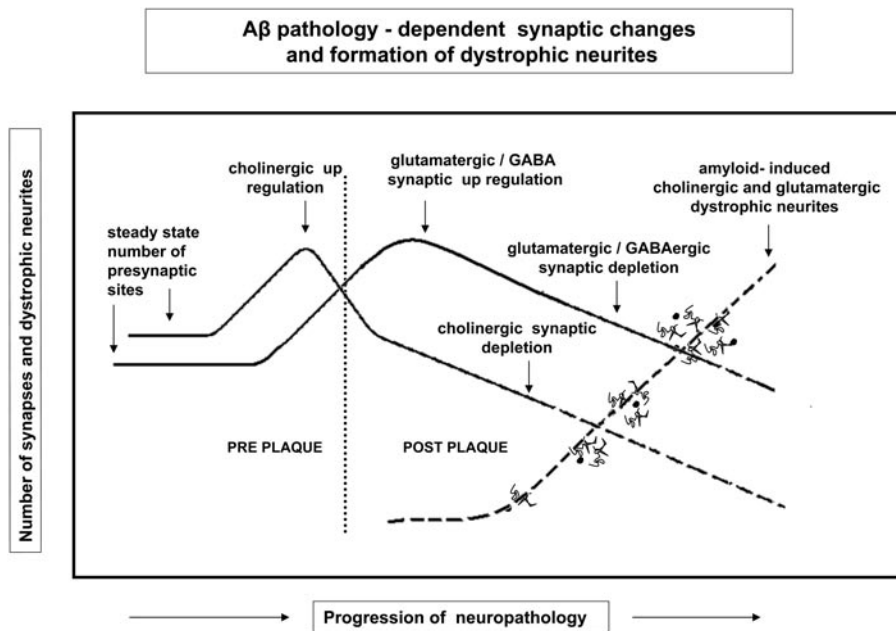
**Figure 1.** Schematic representation of A $\beta$ -induced up-regulation of the activity-dependent expression of CRE-regulated genes and production of synaptic related proteins.

a similar mechanism would occur *in vivo*. In this regard it is interesting to note that some transgenic animal models expressing the A $\beta$ PP<sub>Swe,IndP</sub>, unexpectedly display larger than normal LTP responses while in most models a down regulation of LTP formation is observed (for discussion, see 26). To explain these discrepancies, I would like to suggest that moderate levels of A $\beta$  would, at least initially, facilitate synaptic plasticity through its effects on CRE-directed gene expression (Figure 1). As the levels of A $\beta$  increase and accumulate extracellularly, the level of A $\beta$  oligomers become detrimental to synaptic plasticity and to CRE-directed gene expression. Such changes are known to exert a negative effect on the formation of LTP *in vitro* (57) and can diminish the rate of the activation of CREB in isolated cortical neurons (51). Likewise, fibrillar A $\beta$  inhibits the activity-dependent CRE-gene expression in PC12 cells (Echeverria et al, unpublished data).

#### INTRACELLULAR A $\beta$ ACCUMULATION AND AD-RELATED NEUROPATHOLOGY

There is evidence that A $\beta$  is generated and sequestered intracellularly early in AD's pathology (for review, see 11, 60, 61). In a review article (11), we referred to the accumulation of intracellular A $\beta$  as "a sign of worse things to come." This assertion is based on our own observations in a rat

transgenic model in addition to abundant findings in the literature. We and others (notably the laboratories of Gunnar Gouras, Frank La Ferla, Andrea Leblanc, Thomas Bayer and Mike D'Andrea) maintain the general hypothesis that the accumulation of intracellular A $\beta$  within neurons plays a role in the early pathogenesis of AD. For example, Gouras and collaborators (49) proposed that while a certain amount of A $\beta$  in intracellular compartments may always be present in the rodent and human CNS, the amounts of intracellular A $\beta$  elevate markedly in Down syndrome (39) and in AD (2, 18). In addition, by injecting A $\beta$  peptides in dissociated cortical human embryonic neurons, Leblanc and collaborators (65) have demonstrated that relatively low concentrations of intracellular A $\beta$  peptides lead to apoptotic cell death with the involvement of a caspase cascade. Tabira and collaborators have also linked the accumulation of A $\beta$ -immunoreactive material within cortical neurons with apoptotic signals in the AD human brain and in transgenic mice over-expression AD-related PS1 mutations (5). The possibility that A $\beta$  plaques originate from the remnants of dead neurons that contain elevated amounts of A $\beta$  has also been suggested from microscopical observations in AD brains (8, 17). It is therefore likely that the accumulation of intracellular A $\beta$  leads



**Figure 2.** Schematic representation of the involvement of cortical transmitter systems as the A $\beta$  burden. AD pathology evolves from a pre-plaque phase to the advanced formation of mature, neuritic plaques.

to neuronal dysfunction including changes in synaptic plasticity. We have observed that the overexpression of APP in neurally differentiated P19 cells can provoke considerable intracellular accumulation of A $\beta$  peptides (19), and that this accumulation is accompanied by mitochondrial changes that include a shift to lower mitochondrial membrane potentials (20). This mitochondrial involvement is of interest in the context of the AD pathogenesis given that it has been shown that intracellular A $\beta$  can specifically bind and distort the active site of mitochondrial alcohol dehydrogenase, a change that favors the formation of free radicals and neuronal cell death (36).

Other organelles may also be negatively affected by the intracellular accumulation of A $\beta$ . We have recently developed a number of transgenic rat models that over-express mutated forms of A $\beta$ PP in which the most prominent phenotype has been the accumulation of intracellular A $\beta$  in most neurons of the CA2 and CA3 regions of the hippocampus and within many large pyramidal neurons of the cerebral cortex (12). These A $\beta$ -accumulating neurons displayed a significant increase in both the proportional area occupied by the Golgi apparatus elements as well as the size of individual saccules in the hippocampus of transgenic rats as compared to controls. We also found an elevated number of lysosomes and lipo-

fuscin bodies in the hippocampi of transgenic rats, as well as an increase in the mean individual cross sectional area of lipofuscin bodies in the cortex of transgenic rats as compared to controls (35). These findings support the hypothesis that intracellular A $\beta$  accumulation has a wide impact on subcellular compartments. Indeed, this rise in lipofuscin bodies signals the premature aging of A $\beta$ -accumulating neurons. The up-regulation of lysosomes is also in line with observations for the involvement of this subcellular compartment in early stages of AD (42).

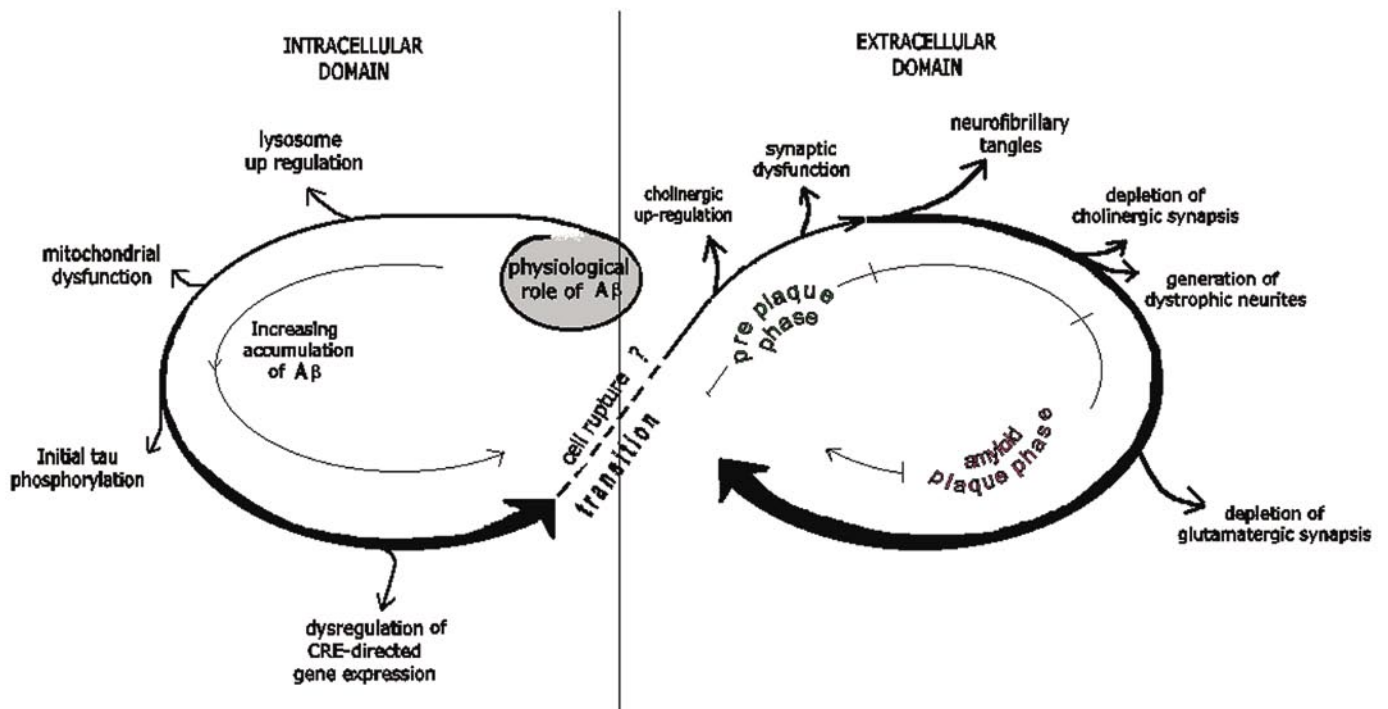
Besides alterations in organelles, the early intracellular accumulation of A $\beta$  leads to changes in protein expression (56) (also Vercauteren et al, unpublished data) and, most importantly, in the activity of key proteins involved in cell signalling. In our transgenic rats with a phenotype of intracellular A $\beta$  accumulation, we observed a robust up-regulation of the MAP kinase ERK2 (12). This increase in ERK2 was accompanied by an augmented phosphorylation of tau, specifically in its MAP kinase phosphorylation sites (14). The dysregulation caused by persistent ERK2 activation, as opposed to activity dependent ERK2 activation (as discussed above), may have a negative impact on cell and synaptic function. Persistent ERK2 activation appears to diminish the degree of phosphorylation

of pp90RSK, which would sequester the CREB binding protein (CBP) in cell systems (59). The sequestration of CBP leads to a down regulation of the CRE-directed gene expression of explicit genes that are involved in the protein-synthesis component of late LTP (for review, see 28). Therefore elevated intracellular A $\beta$  should cause synaptic dysfunction and alterations in higher CNS functions. Evidence for the first was provided by La Ferla et al showing that a diminution of LTP responses occurs prior to the appearance of amyloid plaques, and corresponds to the accumulation intracellular A $\beta$  in their triple transgenic mouse model (43). Evidence for the second effect of A $\beta$  on higher CNS functions was provided with by our rat transgenic model which displays a phenotype of elevated intracellular A $\beta$  in the absence of plaques. These animals displayed a mild but significant learning deficit along with marked hypophosphorylation of pp90RSK, which should compromise CRE-regulated gene expression (14). How these findings correspond to AD neuropathology is not yet fully established. However, Ferrer and collaborators (15) have found that activation of ERK2 is associated with early neurofibrillary pathology in AD. Likewise, an up-regulation of the phosphorylated form of ERK1/2, p-ERK1/2, has been reported in the same brain regions as AD patients with neurofibrillary neurodegeneration Braak stages I-II, but who are devoid of amyloid deposition (44). Furthermore, evidence for diminished CREB phosphorylation in the hippocampus of AD sufferers has been observed (64), which might indicate that in AD (as in the transgenic rat) sustained ERK2 activation might lead to dysregulation of protein phosphorylation patterns leading to the disarticulation of CRE-regulated gene expression and early tau phosphorylation, even in the absence of amyloid plaques.

#### ELEVATED EXTRACELLULAR A $\beta$ AND AD-RELATED NEUROPATHOLOGY

A complete discussion of the impact of extracellular A $\beta$  is beyond the scope of this commentary, and can be found in many well noted and authoritative reviews and also within this volume. In this section, I will highlight key features regarding the relationship between toxic species of ex-

## HYPOTHETICAL PROGRESSION OF A $\beta$ NEUROPATHOLOGY IN ALZHEIMER DISEASE



**Figure 3.** Schematic representation of the evolution of the A $\beta$ -dependent AD neuropathology from an early intracellular phase to advanced stages of extracellular A $\beta$  aggregation in the form of amyloid plaques. The broken lines of transition between intracellular to extracellular A $\beta$  pathology highlight the lack of direct knowledge as to how one process relates to the other.

tracellular A $\beta$  and synaptic structure and function.

Recently a link between extracellular and neurofibrillar pathology has been reported. Thus, the extracellular accumulation of endogenously expressed A $\beta$  in double transgenic mice (34) or the application of exogenous A $\beta$  (16) resulted in the generation of neurofibrillary tangles akin to those observed in AD, and with those observed in transgenic mice that carry tau transgenes with mutations observed in frontotemporal dementias. The cytoskeletal disruption arising from tangle formation should compromise axonal transport and, consequently, contribute to the severe cortical synaptic depletion observed in AD (50). It is likely, however, that the initial deleterious effects of extracellular A $\beta$  on synapses are functional in nature. In some AD transgenic models there is evidence that cognitive deficits can be observed prior to plaque appearance (33, 54) (intracellular A $\beta$ ? A $\beta$  oligopeptides?). These observations correlate with the demonstration that oligomeric A $\beta$  molecules in the extracellular space are sufficient to inhibit LTP generation in the hippocampus (57), as discussed above. Although early functional deficits can be observed in AD

transgenic models, there is well-supported evidence that the extent of the compromise of LTP formation correlates well with the amyloid plaque evolution (52).

Synaptic structural alterations in transgenic models might occur prior to the formation of amyloid plaques. Mucke and collaborators (40) have found a down-regulation of synaptophysin-immunoreactive sites while we have consistently observed an up-regulation (dysfunctional?) of cholinergic presynaptic boutons in pre-amyloid plaque phase in the cerebral cortex of murine transgenic models of AD neuropathology (23, 63). In our experience with transgenic models, these marked synaptic losses manifest themselves with the appearance and progression of amyloid plaque formation (4). We have been able to demonstrate that these structural synaptic losses follow a transmitter-specific pattern of vulnerability in which the cholinergic cortical presynaptic sites are the first to be affected followed by glutamatergic and GABAergic synapses (3, 4, 23, 63). Furthermore, the overt formation of transmitter-immunoreactive dystrophic neurites in AD transgenic models only appears with amyloid plaque formation (3, 48, 63). In our lab, Bell and col-

laborators (6) have used several transgenic mouse models to demonstrate that cholinergic fibres in the cerebral cortex are the first to be engulfed by the peri-plaque amyloid pathology and that a second wave of dystrophic neurites is composed of glutamatergic elements, while GABAergic neurites appear fairly resistant to the plaque aggregation of fibrillar A $\beta$ . These views are schematically summarized in Figure 2.

These observations from transgenic models illustrate that the presence of extracellular A $\beta$  in the CNS per se is sufficient to produce functional and structural synaptic changes, even in the absence of overt neurofibrillary tangles. While current transgenic models do not reproduce the full repertoire of AD neuropathologies, they have provided important insights into the diverse modalities by which A $\beta$ , soluble or fibrillar, contributes to the synaptic pathology of AD.

### CONCLUSIONS

The conventional hypothesis for the pathophysiology of AD—the so-called “amyloid hypothesis”—states that A $\beta$  plaques are the main cause of neurodegeneration and dementia. Consequently, the majority

of studies have focused on the neurotoxic effects of aggregated extracellular A $\beta$  fragments in AD and in AD animal models. However, in recent years the participation of intracellular A $\beta$  accumulation has been seriously considered. In this commentary I have proposed that A $\beta$ , in non-pathological concentrations and situations, plays a physiological role stimulating, post-synaptically, the expression of proteins related to synaptic plasticity via CRE-regulated gene expression. Second, I have proposed that the abnormal accumulation of A $\beta$  within intraneuronal compartments is an initial step in the AD neuropathology and leads to profound alterations in protein expression, their phosphorylation patterns (and thus activity), and thereby affects the structure and function of subcellular organelles and ultimately synaptic function. These changes are most probably insufficient to provoke a dramatic change in higher CNS function however they might “prime” it for downstream pathologies. Finally, I have briefly discussed the possible consequences of A $\beta$  fragment accumulation in the extracellular space, offering support for the notion that there is an incremental functional-to-structural compromise of synaptic elements and cortical neurites as the accumulation of A $\beta$  proceeds from soluble oligomers to fibrillar aggregates and, finally, to toxic amyloid plaques. Indeed, I believe that this extracellular A $\beta$  pathology affects CNS transmitter systems in a highly hierarchical and sequential manner. In short, I am proposing that the A $\beta$  pathology seen in AD is a tale of 2 neuropathologies, which starts at the intracellular level and proceeds into the extracellular space, and which recruits in both compartments diverse cellular and molecular components as the pathology proceeds. These ideas are schematically represented in Figure 3.

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