

MINI-SYMPOSIUM: "Prion-Like" Templated Misfolding in Neurodegenerative Disorders

From Soluble A β to Progressive A β Aggregation: Could Prion-Like Templated Misfolding Play a Role?

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INTRODUCTION

More than 100 years ago, neuropathologist, Alois Alzheimer, discovered peculiar changes in the brains of individuals who had died from dementia: amyloid plaques, neurofibrillary tangles and, to varying extent, vascular amyloidosis (3, 31, 53, 58). Today, we know that extracellular amyloid plaques and vascular amyloid are proteinaceous deposits mainly composed of A β peptides (51). In contrast, neurofibrillary tangles are intracellular aggregates of hyperphosphorylated tau proteins (50). Normally retaining a soluble state, A β peptides generated through proteolytic cleavage from the amyloid precursor protein (APP), as well as tau proteins, are integrated into cellular physiology (33, 50). However, in Alzheimer's disease (AD), A β peptides and tau proteins adopt alternative, misfolded conformations rich in β -sheet structure that renders them aggregation prone and promotes association into oligomeric as well as fibrillar structures. The abundance of these pathological deposits increases with disease severity and compromises synaptic and neuronal function. In advanced stages of AD, vast areas of the brain are affected (13), particularly in the so-called default mode network (78), and significant synaptic and neuronal loss causes progressive dementia (79).

The amyloid cascade hypothesis of AD proposes that A β misfolding and aggregation are the central and earliest events in the disease and trigger a sequence of pathological changes that result in tau hyperphosphorylation and aggregation, brain inflammation and ultimately synapse as well as neuronal loss, and dementia (35).

Abstract

Accumulation, aggregation and deposition of A β peptides are pathological hallmarks in the brains of individuals affected by Alzheimer's disease (AD) or by cerebral β -amyloid angiopathy (A β -CAA). While A β is a peptide of yet largely unknown function, it is constantly produced in the human brain where it normally remains in a soluble state. However, A β peptides are aggregation prone by their intrinsic ability to adopt alternative conformations rich in β -sheet structure that aggregate into oligomeric as well as fibrillar formations. This transition from soluble to aggregated state has been hypothesized to initiate the pathological cascade and is therefore subject to intensive research. Mounting evidence suggests prion-like templated misfolding as the biochemical phenomenon responsible for promoting progressive A β aggregation. Here, we review studies *in vitro* and *in vivo* that suggest that cerebral A β aggregation may indeed progress via prion-like templated misfolding. The implications of these findings are discussed with respect to understanding initiation and progression of the disease and to developing therapeutics.

Conclusive evidence is derived largely from rare forms of familial AD (<3%) caused by mutations in APP or in the catalytic subunits of the γ -secretase complex, that is, presenilin 1 (PS1) and presenilin 2 (PS2), responsible for releasing A β peptides from APP (33). These mutations lead to subtle changes in APP metabolism with increased production of A β peptides and/or to generation of more aggregation prone variants (7, 12, 16, 19, 20, 52, 76). Similarly, an increase in aggregation prone A β peptides caused by increased production and/or reduced degradation and clearance is hypothesized to initiate the more common sporadic cases of AD. A β aggregates, especially small units termed oligomers, have been shown to be neurotoxic and to alter synaptic function (57), although by themselves they may not be sufficient to lead to dementia, correlating only insufficiently with cognitive decline (14, 47, 54, 59, 91). Current state of knowledge suggests that A β toxicity in AD is at least partially mediated via tau hyperphosphorylation and aggregation that promote neuronal dysfunction (38, 72, 88). This part of the mini-symposium focuses on the mechanisms and consequences of A β aggregation. Initiation and spreading of tau aggregation and concomitant pathological changes are reviewed in Clavaguera *et al* (this issue).

AD is not the only disorder with signature A β aggregation. In cerebral β -amyloid angiopathy (A β -CAA), a sporadic or rare hereditary pathological condition in the elderly, characteristic A β deposits are found almost exclusively in the walls of small- to medium-sized blood vessels in the leptomeninges and in the cortex (9). These vascular A β deposits can weaken vessel walls

and compromise vascular function. Abundant vascular A β deposits can cause intracerebral microbleeds or hemorrhages as well as neurodegeneration and cognitive impairment. Pathological accumulation of A β peptides has also been described in sporadic inclusion body myositis (sIBM), an inflammatory and degenerative muscle disease (5, 60, 87); however, the contribution of A β aggregation to sIBM is controversial (32).

While it seems evident that the transition from soluble A β peptides to misfolded and aggregated versions plays an important role in these diseases, many questions still remain open and urgently await answers in order to develop efficacious treatments. These unanswered questions include: How are misfolding and aggregation of normally soluble A β peptides initiated? With disease progression, A β aggregation affects vast areas and neuronal networks of the central nervous system (CNS)—what is the relationship of these aggregates? Do they form independently at the different sites? Or are they related?

Interestingly, new insights arise from similarities between A β aggregation and protein aggregation observed in prion disorders,

similarities with regard to a biochemical phenomenon termed templated misfolding.

NUCLEATION-DEPENDENT POLYMERIZATION, SEEDING AND PRION-LIKE TEMPLATED MISFOLDING

It has been proposed that A β aggregation as well as the aggregation of other amyloidogenic proteins follows a nucleation-dependent polymerization process similar to crystallization (36, 40). Such polymerization reactions consist of two phases: the initial nucleation phase (“lag phase”) and the growth phase (Figure 1A). During the nucleation phase, the native monomer undergoes a series of kinetically unfavorable conformational changes and misfolded or oligomeric species can be formed. These are unstable because the intra- and intermolecular interactions do not outweigh the entropic costs of misfolding and aggregation. However, under certain conditions an ordered, putatively multimeric nucleus can be formed. Once the nucleus is present, more

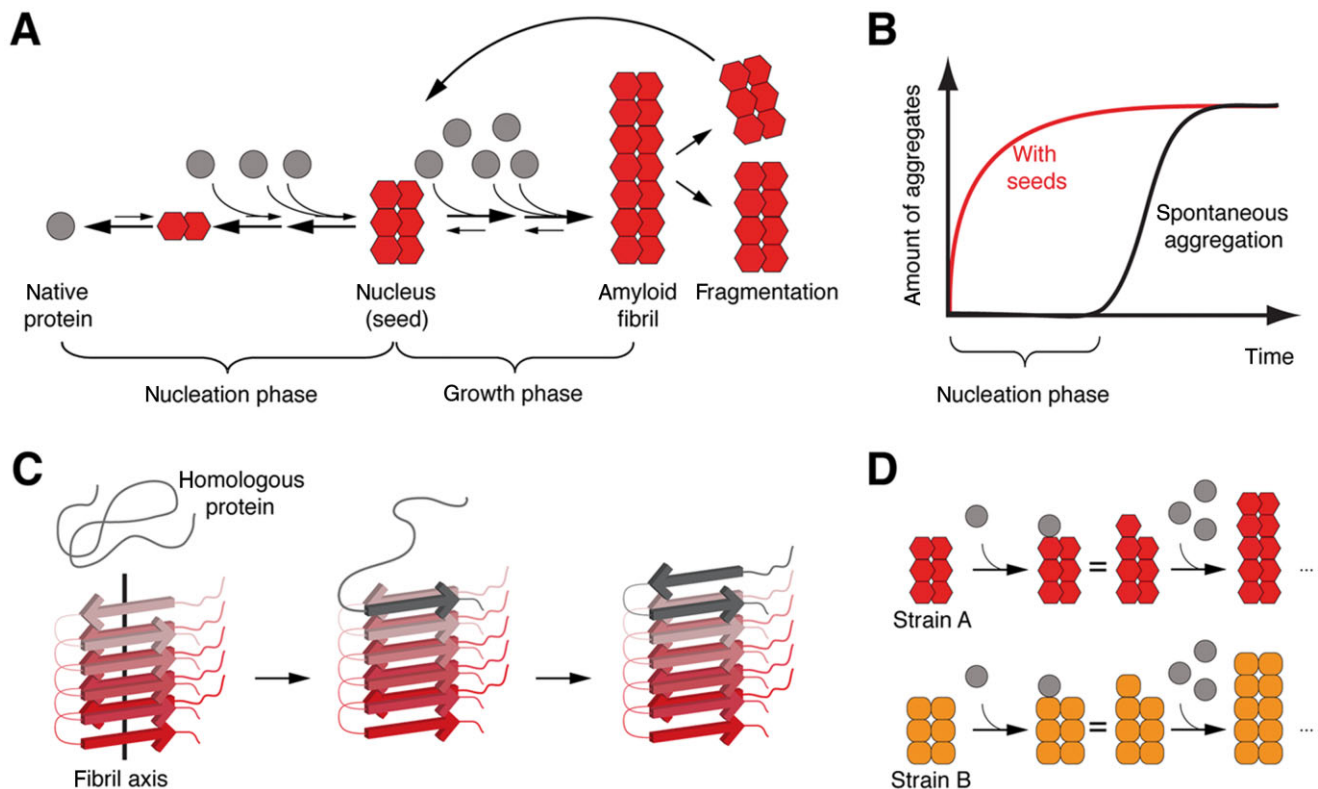


Figure 1. Nucleation-dependent polymerization, seeding and templated misfolding. **A.** Under certain circumstances, some proteins can adopt an alternative, misfolded conformation that allows for multimerization. The initial conformational changes are kinetically unfavorable and may not occur under physiological conditions. However, once a nucleus (“seed,” putatively multimeric) is formed, the properties of the nucleus allow for relatively rapid incorporation of homologous proteins into amyloid fibrils. Fragmentation of fibrils generates more nuclei resulting in a vicious cycle of progressive protein misfolding and aggregation. **B.** Schematic representation of protein aggregation kinetics. Spontaneous aggregation is delayed because of the thermodynamically unfavorable

nucleus formation. Addition of preformed seeds leads to immediate aggregate formation and circumvents the nucleation phase. **C.** Structure analysis of amyloid fibrils reveals typical sets of in-register β -sheets arranged parallel to the fibril axis. This structure has been suggested to serve as a template for the congruent incorporation of homologous, native or partially unfolded proteins that orient their peptide chains accordingly, hence termed templated misfolding. **D.** Some proteins can adopt diverse misfolded conformations that can be associated with different pathogenic properties. These conformational differences are referred to as “strains” and can be propagated by templated misfolding.

cognate proteins can be incorporated relatively rapidly into the aggregate (“growth phase”). In this model, aggregation is concentration- and time-dependent, and the rate-limiting step of fibrillization is the formation of the nucleus. Moreover, fragmentation of fibrils generates new seeds and boosts the polymerization cascade (25). Seeding, which is the addition of a preformed nucleus, circumvents the lag phase and initiates immediate aggregate formation (Figure 1B). Support for nucleation-dependent polymerization of A β peptides is derived from *in vitro* aggregation of synthetic or recombinant A β peptides in solution (48, 61, 64, 77). With regard to AD, this model takes into account that A β fibrillization in the human brain is normally a slow process: initial amyloid plaque formation may take decades in individuals who develop AD and may never even occur in histologically significant amounts at very old age (14). However, once A β aggregation has been initiated, the misfolding and deposition seems to inevitably progress.

Interestingly, structure analysis of fibrils formed by amyloidogenic proteins or fragments thereof provides a plausible and attractive explanation for the efficient incorporation of homologous molecules into the preformed nucleus. Amyloid fibrils were found to typically consist of sets of β -sheets arranged parallel to the fibril axis, with β -sheets either parallel or antiparallel and usually in-register (24). The side chains within and in between the sheets are tightly interdigitated and form a dry steric zipper devoid of water in the interface. Such structures allow for compact stacking of molecules and are stabilized by intermolecular interaction. Homologous molecules can be incorporated accordingly and this leads to progressive growth of the amyloid fibril (Figure 1C). For A β fibrils, such structures were proposed from analyses of synthetic or recombinant A β peptides aggregated *in vitro* by several groups (8, 18, 48, 63, 77, 84, 85). Although detailed structural information of A β fibrils in brain tissue is not available yet, it is conceivable that similar structures are formed *in vivo* (44, 61). Such fibrils can grow from one or both ends, depending on the polarity of the fibril. The growth rate depends (i) on the availability of soluble, homologous molecules for incorporation; (ii) on the efficiency of incorporation; and (iii) on the rate of fragmentation that generates more molecular interfaces for interaction of nucleus and homologous molecules. So, essentially proteins in their abnormal multimeric conformation seem to directly interact with native or partially unfolded homologous proteins and template them into the abnormal, misfolded form. Therefore, this phenomenon has also been termed “templated misfolding.” In case a given protein can adopt two or more differently organized aggregate formations that are reliably sustained by templated misfolding, these distinct aggregates are known as different “strains” of aggregates (2, 24) (Figure 1D). Different strains of aggregates can have different pathological properties and may differ in their growth and fragmentation rates (2).

The principle of templated misfolding was first proposed and is best characterized for the misfolding cascade of prion proteins (17, 66) (Kretschmar and Tatzelt, this issue). Only recently similar mechanisms of misfolding are discussed for other proteins associated with neurodegenerative diseases, for example, A β , tau (Clavaguera *et al*, this issue), α -synuclein (George *et al* this issue) as well as several others (65, 67). With respect to proteins for which this principle of misfolding is still under debate, it is often referred to as “prion-like” misfolding. However, an important dif-

ference exists between the prion diseases and other disorders with putative “prion-like” protein misfolding: only prion diseases have been shown to be “infectious” in a sense that the disease-causing agent can be acquired from the environment and “infect” a new organism. There is no evidence that any of the other disorders with putative “prion-like” protein aggregation are “infectious” or “transmissible” in a similar sense. This is likely because of the absence of an “infectious cycle” (1). AD is a disorder that only occurs in humans and the signature protein aggregates are confined to the individual’s brain (21, 83). Moreover, clinical characteristics of AD do not suggest an infectious nature (1). Even in patients treated with cadaver-derived human growth hormone, no evidence for iatrogenic transmissibility of AD or other protein aggregation disorders was found, except prion diseases (37). However, within the brain, templated misfolding of A β peptides may explain disease progression.

EVIDENCE FOR PRION-LIKE TEMPLATED MISFOLDING OF A β

In the human brain, A β peptides are constantly generated from APP by proteolytic cleavage by β -secretase (BACE-1) generating the N-terminus and by the γ -secretase complex generating the C-terminus (33). γ -Secretase cleavage generates A β peptides of various lengths ranging from 37 to 43 residues (68). The 40-residue peptide A β (1–40) is the most abundant species in normal brain, followed by A β (1–42) species (94). In general, longer A β peptides are more aggregation prone than shorter ones (15, 68).

In vitro synthetic or recombinant A β peptides readily form aggregates of different kinds: low to high *n* oligomers have been described as well as protofibrils, annular aggregates and mature fibrillar structures (90). Aggregation *in vitro* is concentration-dependent and the precise conformation depends on the A β species studied (eg, A β 40 or A β 42) as well as on agitation, temperature, pH and peptide concentration. A detailed comparison of the structures of A β oligomers and fibrils is beyond the scope of this review and has been described elsewhere (24, 26, 86). Interestingly, A β aggregation *in vitro* can be initiated by the addition of preformed A β aggregates (“seeds”) (36, 40, 74, 93). This circumvents the lag phase and supports the concept of nucleation-dependent aggregation. When different polymorphs of A β fibrils were used for *in vitro* seeding, the structural properties of the seeds were reliably propagated to the seeded aggregates indicating templated misfolding (64).

However, most evidence for prion-like templated misfolding of A β comes from inoculation studies in transgenic (tg) rodents. The full spectrum of AD is a human-specific disorder and only partially similar pathological changes including cerebral A β deposition were found in other mammals. Although the reasons for this are not fully understood, it is likely a combination of differences in APP and A β sequence as well as shorter lifespan of other mammals in comparison to the aging human society. Therefore, numerous tg mice and a few tg rats overexpressing human APP and generating human A β peptides were made and are often used to study cerebral β -amyloidosis (4, 41, 70). Most of these tg rodents recapitulate age-related onset of A β aggregation in the brain in the form of parenchymal plaques and vascular β -amyloid very similar to the lesions characteristic for AD. The age-of-onset varies between the different tg lines and depends on the levels of APP

overexpression and related A β levels. Insertion of APP mutations associated with familial AD (FAD) or additional overexpression of human presenilin variants with AD-causing mutations promotes earlier A β deposition.

Several laboratories have studied “seeded” induction of cerebral A β aggregation in tg rodents (43, 55, 56, 92). The experimental paradigm typically uses dilute homogenates from brains of AD patients or of aged APP tg mice with abundant cerebral β -amyloidosis. Minute amounts of these homogenates are injected into the brains of young, predepositing APP tg rodents and the development of cerebral β -amyloidosis is monitored over time.

The essential findings are (see also Figure 2) as follows: Brain extracts from AD patients or from aged APP tg mice containing *in vivo* generated A β aggregates are capable of inducing premature cerebral β -amyloidosis in young APP tg rodents (22, 34, 43, 45, 55, 56, 73, 80, 92). Interestingly, extracts derived from aged APP tg mice with abundant cerebral β -amyloidosis were similarly effective as extracts from AD cases ruling out that factors exclusive to the human brain are essential (55). In contrast, control extracts from healthy individuals or aged wild-type mice without A β aggregates fail to induce cerebral β -amyloidosis (22, 45, 55). Likewise, induction of cerebral β -amyloidosis is prevented when A β peptides are immunodepleted from the extract or when extracts are treated with formic acid resolving any tertiary protein structure. This indicates that misfolded A β peptides are essential components of the “seeding extract” (55). In addition, seeding is

significantly reduced when A β peptides are neutralized by anti-A β antibodies (55). Induced cerebral β -amyloidosis was found to be time-dependent and occurs in histologically detectable amounts with a delay of several months (22, 34, 45, 55, 92). Induction of cerebral β -amyloidosis was also found to depend on the concentration of seeds applied (55). This supports the concept that more seeds, that is, more interfaces for templated misfolding can more efficiently recruit soluble A β peptides for templated misfolding.

The lag time until induced A β deposits are immunohistochemically detectable is not only dependent on the amount of seeds applied but also on the level of APP overexpression in the tg line used. Lower APP overexpression and thus lower availability of soluble A β peptides for misfolding goes along with an extended lag phase (34, 55, 56). In the brains of non-tg mice, no induced A β deposition was found with the same kind and amount of extracts and same incubation time (43, 55). Murine A β sequence differs from human A β sequence by three amino acids and can possibly not easily accommodate the misfolded amyloid state of human A β peptides. This again would indicate specificity of templated misfolding. However, it should not be overlooked that, all other rodent lines inoculated generate higher amounts of A β peptides because of tg overexpression. Therefore, it is conceivable that lowest availability of soluble A β peptides in non-tg mice is not sufficient for seeded induction of histopathologically detectable A β aggregation within the lifespan of these mice although templated misfolding may occur at low levels.

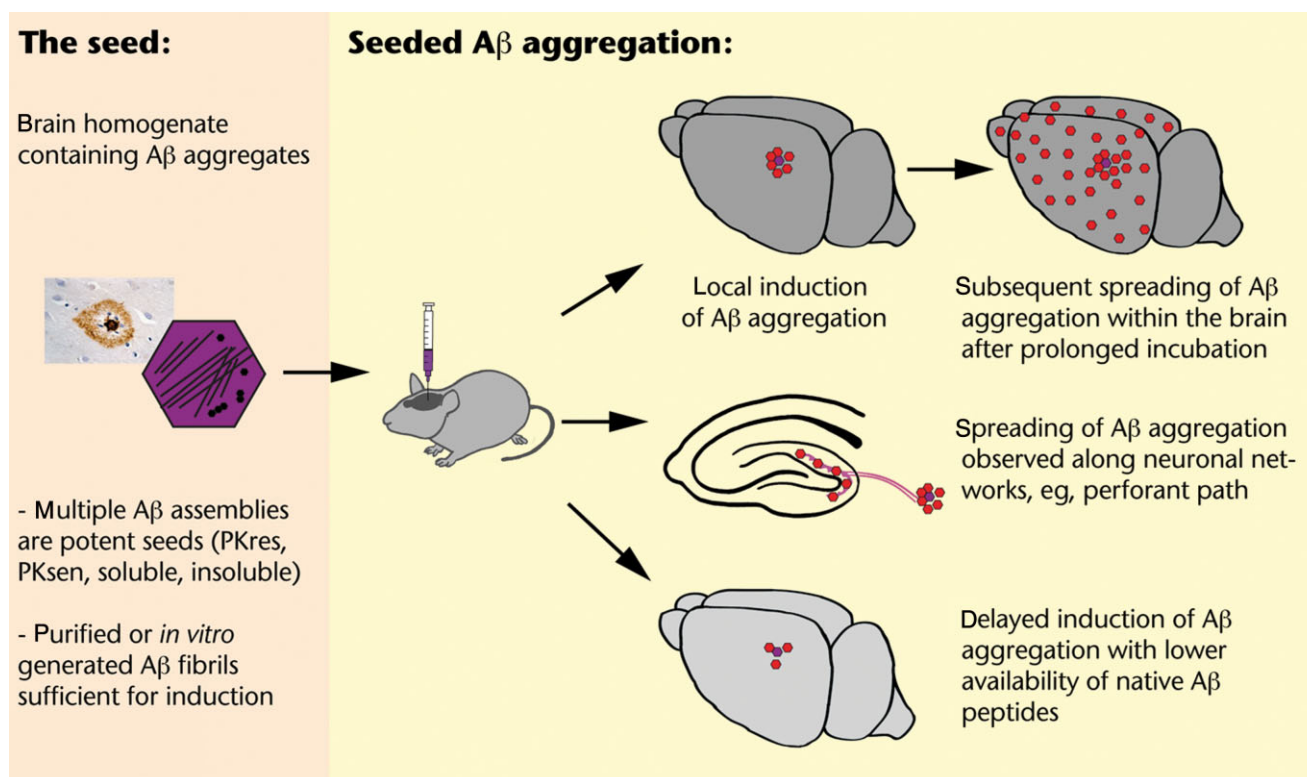


Figure 2. Schematic overview on induction and spreading of A β deposits in brains of young APP transgenic mice following intracerebral seeding with brain homogenates containing aggregated A β peptides (22, 23, 34, 45, 55, 80). [PKres = proteinase K resistant; PKsen = proteinase K sensitive under standard conditions (50 μ g PK/mL homogenate, 30 min, 37°C)].

Interestingly, multiple A β assemblies seem capable of seeding A β aggregation, that is, A β aggregates resistant or sensitive to standard proteinase K treatment, soluble or insoluble aggregates when differentiated by 100 000 \times g ultracentrifugation, and may thus represent a continuum of seeding-capable A β aggregates (45). Soluble seeds seemed to exert high activity, probably because of the presence of many molecular interfaces for templated misfolding (45). This interpretation is also supported by the observation that fragmentation increases the seeding capacity of a given extract (45).

Interestingly, induced deposits are first apparent locally at the site of injection (22, 34, 55). With prolonged incubation times, induced A β aggregates increase locally but also appear in neuroanatomically connected brain regions: inoculations into the entorhinal cortex induced A β deposits in the outer molecular layer of the dentate gyrus, that is, the terminal zone of projections of the perforant path which connects the entorhinal cortex to the hippocampus (22, 42) and vice versa (89). It was observed that local application of A β seeds can induce widespread A β deposition throughout cortex and hippocampus in APP tg mice that would otherwise not have any histochemically detectable A β deposition at this age (34). This suggests that A β aggregates may actively or passively traffic to neighboring and interconnected brain areas. Remarkably, peripheral application of A β seeds can also induce

cerebral β -amyloidosis in young APP tg mice (23). However, it is unclear how this finding would relate to the human situation.

Recently, A β fibrils purified from β -amyloid-laden mouse brains as well as fibrils generated *in vitro* from synthetic A β were shown to be capable of inducing cerebral β -amyloidosis in young APP tg mice and are discussed as the final proof that misfolded A β by itself promotes prion-like templated misfolding and induces associated pathology (80).

Perhaps the most convincing studies suggesting prion-like templated misfolding of A β peptides are observations of reliable propagation of conformational strains. It is well known that A β deposits in the AD brain are heterogeneous in histopathological appearance both within and among brain regions and patients. Amyloid plaques are diffuse or compact, smaller or larger in appearance and this is also reflected in the biochemical composition (49, 82, 83). When different extracts were used for the inoculation studies, derived from either APPPS1 tg mice with small and compact, more punctate plaques (69) or from APP23 tg mice with larger and more filamentous and diffuse plaques (81), the characteristic plaque morphology of the donor was replicated in the host (Figure 3) (55). This indicates that the type of A β aggregates present in the seeding extract has a major influence in shaping the induced deposits. This could be conveyed by prion-like templated misfolding. At present, it is not well known if polymorphic A β

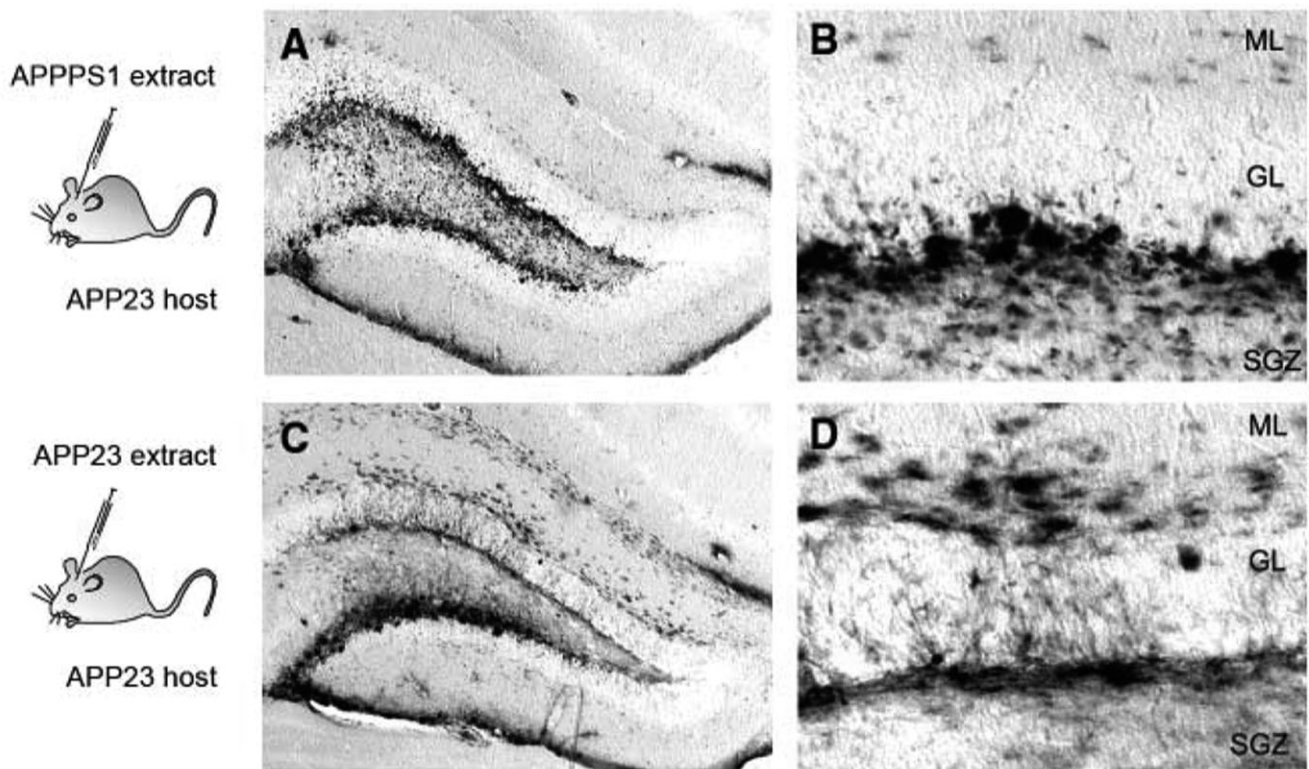


Figure 3. Seeding in young APP23 transgenic hosts induced different morphotypes of A β deposits when different extracts were used for intracerebral inoculations. Note the more punctate appearance of induced deposits with APPPS1 extract, especially in the subgranular cell layer (SGZ) (A, B) in contrast to more filamentous and diffuse A β

deposits induced with APP23 extract (C, D). The morphotypes of the induced A β deposits resemble the A β deposits found in the brains of the donors of the extracts [from Meyer-Luehmann *et al* (55). Reprinted with permission from AAAS]. ML = molecular layer; GL = granular cell layer.

“strains” are also polyfunctional and are linked to differences in toxicity or the clinical spectrum of AD.

Taken together, all these studies *in vitro* and *in vivo* support the concept of prion-like templated misfolding of A β peptides as defined earlier. Although tg rodents predisposed to developing cerebral β -amyloidosis were used, the observation of induced A β deposits in tg lines without A β deposits within the normal lifespan suggests true seeded conversion of normally soluble A β peptides and not simply acceleration of an ongoing process (56, 73). However, other possible explanations should not be neglected and will largely have to take into account that inoculation of A β aggregates may cause secondary effects that indirectly promote A β aggregation, especially in environments with unnaturally high concentrations of A β peptides. Regarding the observed spreading of A β deposits within neuronal networks, indirect effects of compromised neuronal functions that may affect APP processing or A β levels are conceivable. Moreover, these experiments in tg rodents focused on A β aggregation, and therefore do not allow a conclusion if induction of cerebral A β deposition would provoke the whole spectrum of clinical AD. However, it was repeatedly observed that the induced deposits resemble typical AD-like A β deposits and can be accompanied by dystrophic neurites with typical hyperphosphorylation of tau proteins, as well as micro- and astrogliosis (22, 23, 34, 45, 55, 92). While induced deposits may be largely diffuse initially, a more compact and congophilic appearance was noted with longer incubation times, indicating maturation toward more pathological lesions (45). In addition, region-specific differences in induced deposits may point to the importance of additional cellular factors shaping A β aggregation, for example, induced deposits in the striatum of APP23 tg mice were largely diffuse within the observation time whereas cortical or hippocampal inoculation resulted in more compact deposits (22). Last but not least, minute amounts of A β aggregates may be sufficient to trigger progressive tau aggregation and pathology (11, 29).

Naturally, no such inoculation studies can be undertaken in humans. Attempts in nonhuman primates were initially inconclusive (30), but subsequent work has shown that intracerebral inoculation of crude homogenates from AD brain into marmosets can induce β -amyloid deposition after incubation times of at least 5–6 years (6, 71). However, because of the high prevalence of AD and A β -CAA and the suggestive spreading within brain networks (13, 78), it will be important to further probe this concept for understanding the human disorders and for developing therapeutics.

IMPLICATIONS OF PRION-LIKE TEMPLATED MISFOLDING OF A β PEPTIDES

These findings have multiple implications with regard to future research and to the development of therapeutics for the treatment and the prevention of A β -misfolding disorders. Disease-specific A β aggregation in the brains of individuals in prodromal AD stages has been detected many years to decades before clinical symptoms manifest (39, 62). A β seeds if present in body fluids (eg, CSF) could serve as an early and disease-specific biomarker, complementing currently used measurements of total CSF A β and tau levels (10). However, it is at present unknown if any kind of A β seed exists outside the brain, and if so, how efficient its capability

to transmit the misfolded state via templated misfolding would be. Evidently, if such seeds would exist, accidental transmission between individuals must be avoided, particularly as the seeds may be chemically stable and resistant to some common inactivation procedures.

Our current understanding of the kinetics of A β aggregation and the indications for prion-like templated misfolding highlights several possibilities for therapeutic intervention: First, by keeping A β levels below the critical (local) concentration for aggregation. This could be achieved by inhibition or modulation of the enzymes releasing A β from its precursor (28, 95), by increasing A β degradation (75) or by promoting its clearance. Second, by removing existing aggregates. Some enzymes have been shown to be capable of degrading or dissociating small A β aggregates (75). In addition, immunotherapy was shown to be efficient in removing A β deposits (46). Third, by capping templated misfolding or by promoting the formation of less toxic aggregates. In principle, the cascade of progressive templated misfolding would be stopped if the molecular interfaces for templated misfolding are not available (24). Similarly, if differentially toxic A β aggregate versions exist, a shift toward less toxic and more biologically inert forms could be beneficial. Fourth, preventing the spreading of A β aggregation to neighboring or interconnected tissue may limit the detrimental effects and concomitant progressive cognitive deficits. Fifth, and maybe most importantly, compounds that preserve neuronal and synaptic integrity and function could keep dementia at bay. The prediction would be that all disease-modifying strategies are most effective as prevention or in the early stages of the disease because severe pathology and massive neuron loss in late stages are likely irreversible. Such early or preventive therapeutics may have to be taken for many years or decades, and therefore, side effects should be avoided and well-tolerated yet effective medication would be desirable. This is especially challenging as we yet lack a good understanding of the physiological functions of APP, its diverse fragments including A β and the enzymes involved in APP processing. In the end, likely not one but several of the possible therapeutic approaches outlined above may be needed for preventing cognitive decline. There is a silver lining for causative or preventive AD treatment as several compounds targeting A β -generating enzymes, anti-A β immunotherapy and others are already being tested in clinical trials and some are under consideration for preventive trials (27). While A β aggregation may be the most relevant trigger for A β -CAA, it may only be the starting point for AD. Changes in tau phosphorylation and eventually tau aggregation may be the effectors leading to AD neurodegeneration (38, 72, 88). As tau aggregation likewise seems to propagate by prion-like templated misfolding (Clavaguera *et al*, this issue), it will be important for any A β -targeting AD therapeutics to stop A β aggregation before tau aggregation proceeds beyond the point of no return.

CONCLUSIONS

Prion-like templated misfolding of A β peptides provides a plausible framework to explain the progressive accumulation and spreading of A β aggregation within the AD brain. This concept has far-reaching implications for possible therapeutic interventions. In particular, the need for early intervention before a fatal cascade of A β misfolding leads to neuropathological

damage beyond repair. However, more work is needed to fully understand the characteristics of this process and its contribution to AD and other A β -misfolding disorders. At present, prion-like templated misfolding is discussed as a driving force in several neurodegenerative disorders. This could unify experimental and translational approaches to these increasingly prevalent and devastating disorders.

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CONFLICT OF INTEREST

The author declares that she has no conflicting interest related to this article.

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