



Prion-like mechanisms and potential therapeutic targets in neurodegenerative disorders☆



Masato Hasegawa*, Takashi Nonaka, Masami Masuda-Suzukake

Department of Dementia and Higher Brain Function, Tokyo Metropolitan Institute of Medical Science, 2-1-6 Kamikitazawa, Setagaya-ku, Tokyo 156-8506, Japan

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ABSTRACT

Prion-like propagation of abnormal intracytoplasmic proteins, which are the defining features of major neurodegenerative disorders, such as Alzheimer's disease (AD), Parkinson's disease (PD) and amyotrophic lateral sclerosis (ALS), has been proposed. A growing body of evidence strongly suggests that abnormal tau, α -synuclein and TDP-43 have prion-like properties, convert the corresponding normal proteins into abnormal forms, and are transmitted from cell to cell, spreading throughout the brain. This idea is extremely important not only for understanding the pathogenesis and progression of these diseases, but also for the development of molecular therapies. Since the distributions and spreading of the abnormal proteins are closely associated with disease symptoms and progression, gain-of-toxic-function of these proteins may affect the neurons and glial cells either directly or indirectly, or both. It is essential to regulate the aggregation of abnormal intracellular proteins and their cell-to-cell transmission in order to stop, or at least slow, the progression of these diseases.

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Contents

1. Introduction	22
2. Abnormalities in the pathological proteins found in neurodegenerative disorders	23
3. Experimental evidence that abnormal tau, α Syn and TDP-43 spread by prion-like propagation.	24
4. Molecular mechanisms of cell-to-cell transmission.	29
5. Therapeutic strategies to block prion-like propagation	29
Conflict of interest.	30
References	30

1. Introduction

Neurodegenerative diseases are characterized by progressive degeneration of subsets of neurons and gliosis. Many of these disorders are

Abbreviations: AD, Alzheimer's disease; PD, Parkinson's disease; DLB, dementia with Lewy bodies; MSA, multiple system atrophy; ALS, amyotrophic lateral sclerosis; α Syn, α -synuclein; TDP-43, TAR DNA binding protein of 43 kDa; FTL, frontotemporal lobar degeneration; PrP(C), normal prion protein; PrP(sc), abnormal prion protein; TTR, transthyretin.

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* Corresponding author.

E-mail address: hasegawa-ms@igakuken.or.jp (M. Hasegawa).

accompanied with distinct pathologies in the central nervous system, consisting of abnormal proteinaceous structures: neurofibrillary tangles in Alzheimer's disease (AD), Lewy bodies in Parkinson's disease (PD) and skein-like inclusions in amyotrophic lateral sclerosis (ALS) (Fig. 1A–C). Since the 1980's, attempts had been made to identify the components of these pathological structures in order to explore the molecular mechanisms of neurodegeneration and the pathogenesis of the diseases. A microtubule-associated protein, tau, was discovered to be the major component of neurofibrillary tangles in the mid-1980's (Brion, Flament-Durand, & Dustin, 1986; Delacourte & Defossez, 1986; Grundke-Iqbal et al., 1986; Kosik, Joachim, & Selkoe, 1986; Nukina & Ihara, 1986). α -Synuclein (α Syn) was found to be the protein component of Lewy bodies in PD and dementia with Lewy bodies (Spillantini et al., 1997; Baba et al., 1998; Spillantini, Crowther, Jakes, Hasegawa, & Goedert, 1998). Also, TAR DNA binding protein of 43 kDa (TDP-43) was identified as a component of intracellular ubiquitin-positive

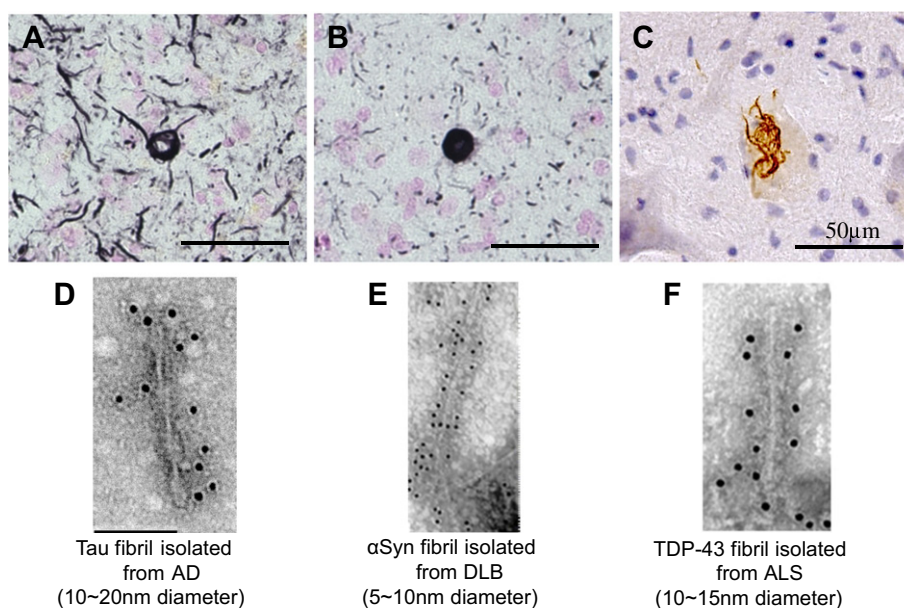


Fig. 1. Intracellular abnormal protein pathologies in major neurodegenerative diseases and electron microscopy of the isolated abnormal fibrils. A: Neurofibrillary tangles and threads in neocortex of AD (stained with an anti-phospho tau antibody AT8), B: Lewy bodies and Lewy neurites in neocortex of DLB (stained with an anti-phospho α Syn antibody, PS129), C: Skein-like inclusion in spinal cord of ALS (stained with an anti-phospho TDP-43 antibody, pS409/410). D: Paired helical filament that constitutes the building block of tau pathologies in AD (stained with AT8, 10 nm gold particles). E: α Syn fibrils isolated from DLB brain (stained with PS129, 5 nm gold particles). F: TDP-43 fibrils isolated from brains of ALS patients (stained with pS409/410, 10 nm gold particles). Bar: 100 nm.

inclusions in ALS and frontotemporal lobar degeneration (FTLD) (Arai et al., 2006; Neumann et al., 2006). Immunocytochemical and biochemical studies using specific antibodies to the normal and abnormal proteins revealed widespread abnormal protein pathologies in these diseases (Fig. 1A–C), and the distributions and spread of the abnormal proteins were closely correlated with clinical presentation and disease progression. At an early stage of AD, tau pathology is observed in transentorhinal cortex, and at more advanced stages, widespread tau pathologies are seen in limbic regions and cortex (Braak & Braak, 1991; Braak & Del Tredici, 2011). In PD or DLB, α Syn pathology is detected in olfactory bulb, and/or dorsal motor nucleus in medulla at an early stage, and it is seen in amygdala, cholinergic nuclei of the basal forebrain and substantia nigra at later stages (Braak et al., 2003), or spread from the temporal lobe to the frontal lobe and further spread to the parietal and occipital lobes in DLB (Saito et al., 2003). It was also reported that TDP-43 pathology in ALS appears to disseminate in a sequential pattern that permits the recognition of 4 neuropathological stages consistent with the hypothesis that TDP-43 pathology is propagated along axonal pathways (Braak et al., 2013; Brettschneider et al., 2013; Brettschneider et al., 2014; Kassubek et al., 2014; Smethurst, Sidle, & Hardy, 2015), although these studies were done after the theory had been proposed. Thus, these diseases are “progressive” disorders, and the intracellular pathologies seem to spread during the course of the diseases. However, little attention had been given to the questions of why these diseases are progressive, and why the pathologies spread to different brain regions during the course of the diseases. This prion-like propagation is the principal molecular mechanism of these major neurodegenerative diseases with amyloid-like abnormal protein pathologies, and can account for the degeneration of subsets of neurons, the diverse but characteristic pathologies, and the disease progression (Brettschneider, Del Tredici, Lee, & Trojanowski, 2015; Frost & Diamond, 2010; Goedert, 2015; Halliday, Radford, & Mallucci, 2014; M. Hasegawa, 2016; Jucker & Walker, 2013; Prusiner, 2012). Therefore, regulation of the propagation of abnormal proteins is an important goal for disease-modifying therapy of these major neurodegenerative disorders.

2. Abnormalities in the pathological proteins found in neurodegenerative disorders

The pathological inclusions and their principal protein components (tau, α Syn and TDP-43) (Fig. 2) deposited in the brains of patients have been extensively studied by means of biochemical, immunochemical and protein chemical analyses. Although these proteins are different in size, structure (Fig. 2), subcellular localization and physiological functions, there are some common features (Table 1). Immunoelectron-microscopic studies demonstrated that the proteins accumulate as abnormal fibrils or filaments (Fig. 1D–E) with some cellular organelles, irregularly shaped structures and lipofuscins (M. Hasegawa et al., 2007; Lee, Balin, Otvos, & Trojanowski, 1991; Lin & Dickson, 2008; Spillantini, Crowther, Jakes, Cairns, et al., 1998; Spillantini, Crowther, Jakes, Hasegawa, et al., 1998). Most of these structures are Thioflavin S/T-positive (Delaere, Duyckaerts, Brion, Poulain, & Hauw, 1989; Hashimoto et al., 1998; Bigio et al., 2013), suggesting that the fibrils are amyloid-like and well-ordered, being clearly distinct from amorphous disordered aggregates. The term “amyloid-like” is used for intracellular proteins similar to amyloids, because the classical histopathological definition of “amyloid” is an extracellular, proteinaceous deposit exhibiting beta-sheet structure. All these abnormal proteins are sarkosyl-insoluble (Fujiwara et al., 2002; Greenberg & Davies, 1990; M. Hasegawa et al., 2008), and can be partially purified or enriched by utilizing this property in combination with ultracentrifugation. Mass spectrometric and immunohistochemical analyses revealed that these proteins deposited in brains are abnormally phosphorylated (Hasegawa et al., 1993; Morishima-Kawashima et al., 1995; Fujiwara et al., 2002; Hasegawa et al., 2008; Kametani et al., 2016), partially ubiquitinated (Morishima-Kawashima et al., 1993; Hasegawa et al., 2002; Kametani et al., 2016), deamidated and fragmented (Hasegawa et al., 1992; Arai et al., 2004; Hasegawa et al., 2011; Dan et al., 2013; Kametani et al., 2016; Taniguchi-Watanabe et al., 2016), in contrast to the situation in the brains of control cases. So, what is the most important feature for the pathogenesis among these abnormalities? Ubiquitination of α Syn

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