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The role of overexpressed DYRK1A protein in the early onset of neurofibrillary degeneration in Down syndrome

Jerzy Wegiel · Karol Dowjat · Wojciech Kaczmarski · Izabela Kuchna · Krzysztof Nowicki · Janusz Frackowiak · Bozena Mazur Kolecka · Jarek Wegiel · Wayne P. Silverman · Barry Reisberg · Mony deLeon · Thomas Wisniewski · Cheng-Xin Gong · Fei Liu · Tatyana Adayev · Mo-Chou Chen-Hwang · Yu-Wen Hwang

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Abstract The gene encoding the minibrain kinase/dualspecificity tyrosine phosphorylated and regulated kinase 1A (*DYRK1A*) is located in the Down syndrome (DS) critical region of chromosome 21. The third copy of *DYRK1A* is believed to contribute to abnormal brain development in patients with DS. In vitro studies showing that DYRK1A phosphorylates tau protein suggest that this kinase is also involved in tau protein phosphorylation in the human brain and contributes to neurofibrillary degeneration, and that this contribution might be enhanced in patients with DS. To explore this hypothesis, the brain tissue from 57 subjects

J. Wegiel (⊠) · K. Dowjat · W. Kaczmarski · I. Kuchna · K. Nowicki · J. Frackowiak · B. Mazur Kolecka · J. Wegiel Department of Developmental Neurobiology, NYS Institute for Basic Research in Developmental Disabilities, 1050 Forest Hill Road, Staten Island, NY 10314, USA e-mail: J_Wegiel@msn.com

W. P. Silverman Department of Psychiatry and Behavioral Medicine, John Hopkins University Medical School, Baltimore, MD, USA

B. Reisberg · M. deLeon · T. Wisniewski
Department of Psychiatry,
New York University School of Medicine,
Silberstein Aging and Dementia Research and Treatment Center,
New York, NY, USA

C.-X. Gong · F. Liu Department of Neurochemistry, NYS Institute for Basic Research in Developmental Disabilities, 1050 Forest Hill Road, Staten Island, NY, USA

T. Adayev · M.-C. Chen-Hwang · Y.-W. Hwang Department of Molecular Biology, NYS Institute for Basic Research in Developmental Disabilities, 1050 Forest Hill Road, Staten Island, NY, USA including 16 control subjects, 21 patients with DS, and 20 patients with sporadic Alzheimer's disease (AD) was examined with two antibodies to the amino-terminus of DYRK1A (7F3 and G-19), as well as two polyclonal antibodies to its carboxy-terminus (X1079 and 324446). Western blots demonstrated higher levels of full-length DYRK1A in the brains of patients with DS when compared to control brains. Immunocytochemistry revealed that DYRK1A accumulates in neurofibrillary tangles (NFTs) in subjects with sporadic AD and in subjects with DS/AD. Overexpression of DYRK1A in patients with DS was associated with an increase in DYRK1A-positive NFTs in a gene dosage-dependent manner. Results support the hypothesis that overexpressed DYRK1A contributes to neurofibrillary degeneration in DS more significantly than in subjects with two copies of the DYRK1A gene and sporadic AD. Immunoreactivity with antibodies against DYRK1A not only in NFTs but also in granules in granulovacuolar degeneration and in corpora amylacea suggests that DYRK1A is involved in all three forms of degeneration and that overexpression of this kinase may contribute to the early onset of these pathologies in DS.

Keywords Minibrain kinase DYRK1A · Neurofibrillary degeneration · Granulovacuolar degeneration · Corpora amylacea · Down syndrome · Alzheimer's disease

Introduction

A broad spectrum of developmental and age-associated changes in patients with Down syndrome (DS) is considered to be a result of the overexpression of genes localized in the DS critical region of chromosome 21. The extra copy of the gene encoding amyloid precursor protein (*APP*) located on chromosome 21 appears to be the main cause of the early onset of brain amyloidosis- β in patients with DS. Overexpression of APP has been associated with an increase in amyloid β (A β) 42 level in the brain of fetuses with trisomy of chromosome 21 [64], the development of diffuse A β -positive plaques in approximately 50% of individuals with DS younger than 30 years of age [35, 37, 72, 73], Alzheimer-type pathology in the majority of DS subjects older than 40 years of age [55, 69, 73], and an elevated risk of Alzheimer's disease (AD)-associated dementia [24, 33, 41, 78].

While early brain amyloidosis- β appears to be directly associated with an extra copy of the APP gene, the mechanism leading to neurofibrillary degeneration in patients with DS is unknown. Recent studies suggest that the Minibrain kinase/dual-specificity tyrosine phosphorylated and regulated kinase 1A (DYRK1A) gene, located in the critical region of chromosome 21 [20, 50, 62] and overexpressed in DS [21, 27], may play a significant role not only in developmental brain defects associated with DS [46, 59] but also in neurofibrillary degeneration. The discovery that DYRK1A phosphorylates microtubule-associated protein tau at several phosphorylation sites and in cultured cells [40, 74] implicates DYRK1A in abnormal tau phosphorylation and neurofibrillary degeneration in patients with AD. The observation that Thr²¹² is phosphorylated in fetal tau and hyperphosphorylated in filamentous tau from subjects with AD [43] supports this hypothesis. Moreover, phosphorylation of Thr²¹² by DYRK1A stimulates in vitro the rate of phosphorylation of tau about eightfold through priming tau phosphorylation by glycogen synthase kinase 3 (GSK3) at Ser²⁰⁸ [74]. Phosphorylation of Ser²⁰⁸ has also been reported in tau in patients with AD [43]. DYRK1Ainduced tau phosphorylation inhibits tau activity to stimulate microtubule assembly and promotes its self-assembly into filaments [40]. These findings strengthen the position that DYRK1A may be involved in tau protein phosphorylation in the human brain and potentially in neurofibrillary degeneration.

The aim of this study was to explore the hypothesis that DYRK1A contributes to neurofibrillary degeneration in patients with sporadic AD and that this contribution is enhanced in DS subjects with trisomy of chromosome 21 and AD. Western blots demonstrated higher levels of DYRK1A in the brains of patients with DS, whereas immunocytochemistry with antibodies to DYRK1A showed the presence of this kinase in neurofibrillary tangles (NFTs) in patients with sporadic AD. A several-fold increase in the number of DYRK1A-positive NFTs in patients with DS/AD supports the hypothesis that DYRK1A contributes to the early onset of neurofibrillary degeneration in DS.

Materials and methods

Tissue for morphological studies

Brains of 57 subjects were examined, including 16 control subjects 32-92 years of age (12 males and 4 females), 21 subjects with DS from 38 to 72 years of age (12 males and 9 females), and 20 subjects with sporadic AD 73-97 years of age (11 males and 9 females). Functional assessment staging (FAST) [51] was used to evaluate the progression of the clinical course of sporadic AD. Control subjects were classified as FAST stages 1 and 2 (normal adult without functional decrement documented in available medical records, and normal aged adult with subjective deficit in word finding, respectively). At the time of their demise, three AD subjects were at FAST stage 3 (incipient AD with deficits noted in demanding employment settings); three subjects at FAST stage 4 (mild AD with deficits in performance of complex tasks of daily life); two subjects at FAST stage 5 (moderate AD manifested with deficient performance in choosing proper attire); four subjects at FAST stage 6 (moderately severe AD manifested with five substages, including decreased ability to dress properly, to handle mechanics of bathing and toileting, and finally with urinary and fecal incontinence), and eight subjects at FAST stage 7 (severe AD with several substages, including incipient averbalism, limitation of speech to a single word, and loss of ambulation). Patients were examined by clinicians in the Aging and Dementia Research Center at New York University Medical Center.

Fixation, processing, embedding, and neuropathological evaluation

One brain hemisphere from each subject was cut into 1-cmthick frontal slabs. Diagnostic tissue samples were cut off and fixed in formalin for routine neuropathological examination, while the majority of tissue was frozen at -80° C for biochemistry. Neuropathological evaluation was based on gross brain description and examination of 8-µm-thick sections from 16 brain regions. Alzheimer-type pathology was evaluated using criteria developed at a consensus conference organized by the National Institute on Aging and the Reagan Institute of the National Alzheimer Disease Association [44]. According to Braak and Braak criteria [5], all control subjects younger than 42 years of age were free of AD pathology. All older control subjects were affected with incipient neurofibrillary degeneration corresponding to Braak and Braak criteria limbic stage I. They were free of amyloid plaques. AD subjects represented a broad spectrum of neuropathological changes from Braak and Braak limbic stage II to neocortical stage VI. In the examined cohort, all subjects with DS older than 38 years of age were affected

by both neurofibrillary degeneration and β -amyloidosis with neuritic plaques corresponding to Braak and Braak II–VI.

The other hemisphere was fixed in 10% buffered formalin for 6 weeks to several months and then dissected into 1-cm-thick frontal slabs. The tissue blocks were dehydrated in a graded series of ethanol: for 5 days, in 70% ethanol; then 2 days in 80% ethanol; and finally, 1 week in 96% ethanol. Dehydrated tissue was infiltrated with polyethylene glycol (PEG) 400 (Merck #807 485) for 6 days (two changes of 3 days each at room temperature) and with PEG 1000 for another 6 days (two changes of 3 days each at 42°C). Slabs were embedded in fresh PEG 1000 [25] and stored in +4°C. Tissue blocks were then cut at a temperature of 18°C into 50-µm-thick serial sections. They were stored in 70% ethyl alcohol at room temperature and used for immunocytochemical and morphometric studies.

The methods applied to this study were approved by the Institutional Review Board at the New York State Institute for Basic Research in Developmental Disabilities. All brain tissue samples were identified by case number and examined blind to clinical and demographic information.

Immunostaining

Serial sections stored in 70% ethyl alcohol were washed with phosphate buffer solution (PBS) and immunostained as free-floating sections. Several antibodies were used to detect DYRK1A (Table 1), including two antibodies to the amino-terminus of DYRK1A: monoclonal antibody (mAb) 7F3 (IBR) [70] and goat polyclonal antibody (pAb) Dyrk1A G-19 (Santa Cruz Biotechnology Inc.). The carboxy-terminus of DYRK1A was detected with sheep pAbs X1079 from Exalpha Biologicals and sheep antibody 324446 from Calbiochem. Abnormally phosphorylated tau protein of NFTs was detected with mAb Tau-1. Tau-1 recognizes an epitope between amino acids 189 and 207 of the human tau sequence [17]. To obtain optimum staining with Tau-1, sections were dephosphorylated with alkaline phosphatase prior to the application of primary antibody [19] (Sigma, Type VII-L, 400 μ g/ml in PBS, pH 7.4). A β -positive plaques were immunodetected with mouse mAb 4G8 (NYS, IBR), which recognizes amino acids 17-24 of the amyloid β -peptide [29]. To enhance the immunoreactivity of A β , sections were treated with concentrated formic acid [31].

The endogenous peroxidase in the sections was blocked with 0.2% hydrogen peroxide in methanol. Non-specific bindings were blocked with 10% fetal bovine serum in PBS for 30 min. The antibodies were diluted in 10% fetal bovine serum in PBS and were incubated with sections overnight at 4°C. The sections were washed and treated for 30 min with either biotinylated sheep anti-mouse IgG antibody or Ë

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Table 1 Anti	bodies used for Western blotting	g and immu	nostainings			
Symbol	Company and catalog number	Host	Type, isoform, purification, concentration	Immunogen (epitope)	Dilution for immunocyto- chemistry	Secondary antibody
7F3	IBR, Dr. Y-W. Hwang	Mouse	Monoclonal, IgG, purified, 1 mg/ml	Immunogen: 1–160 aa of rat Dyrkl A.Epitope: 74–78 aa of 754 aa isoform [70]	1:50	RPN 1001Anti-mouse Ig biotinylated from sheep, 1:200Amersham Biosc.
Dyrk1A (G-19)	Santa Cruz sc-12568	Goat	Polyclonal, IgG, purified, 1 mg/ml	20–50 aa of N-terminus of human Dyrk1A	1:75	RPN 1025 anti-goat Ig biotinylated from donkeyAmersham Biosc.
Dyrk	Exalpha Biologicals X1079	Sheep	Polyclonal, IgG, purified, 1 mg/ml	486–763 aa of human DYRK1A	1:50	RPN 1025 anti-goat Ig biotinylated from donkeyAmersham Biosc.
Anti-DYRK	Calbiochem 324446	Sheep	Polyclonal, IgG, purified, 1 mg/ml	Unique non-catalytic region of C-terminus of DYRK1A	1:50	RPN 1025 anti-goat Ig biotinylated from donkeyAmersham Biosc.
Tau-1	IBR; Dr Richard Kascsak	Mouse	Monoclonal, IgG, purified	189–207 aa of human tau [17]	1:100,000	RPN 1001 anti-mouse Ig biotinylated fr sheep, 1:200Amersham Biosc.
4G8	IBR; Dr Richard Kascsak	Mouse	Monoclonal, IgG, purified	17–24 aa of A β [29]	1:8,000	RPN 1001Anti-mouse Ig biotinylated fraction sheep, 1:200Amersham Biosc.

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biotinylated donkey anti-goat IgG antibody diluted 1:200. The sections were treated with an extravidin peroxidase conjugate (1:200) for 1 h, and the product of reaction was visualized with diaminobenzidine (0.5 mg/ml with 1.5% hydrogen peroxide in PBS). After immunostaining, the sections were lightly counterstained with cresyl violet.

To confirm the specificity of the immunostaining, the reaction was tested with antibodies not absorbed as well as those absorbed with appropriate peptides. For absorption, antibody at the dilution 1:50 or 1:75 (the dilution at which consistently positive results are achieved on sections) was combined with a fivefold (by weight) excess of blocking peptide. The mixture was incubated for 2 h at room temperature (or overnight at 4°C) and used for immunocytochemical tests. The effect of tau protein dephosphorylation was examined on sections pretreated and those not treated with alkaline phosphatase.

Co-localization of Mnb/Dyrk1A and NFTs by double immunofluorescence

Fifty-µm-thick brain sections were double-immunostained for DYRK1A using pAb G19, and for NFTs, using mAb Tau-1. Sections treated with alkaline phosphatase were incubated with G19 antibody, followed by a secondary donkey antiserum against goat IgG conjugated with biotin (Amersham), and then streptAvidin-FITC (Vector). The second primary antibody was mAb Tau-1. The reaction was detected with antibody made in sheep: $F(ab)_2$ fragments against mouse IgG conjugated to Cy3 (Sigma). The absence of cross-reactions was confirmed as described earlier [15, 16]. Images were generated using a Nikon Eclipse E800 microscope and the Nikon PCM 2000 dual-laser scanning confocal microscopy system and with the C-Imaging-SIM-PLE32TM image analysis system (Compix Inc.).

Detection of DYRK1A using Western blotting

Frozen tissue samples from the cerebral frontal cortex of five control subjects 31–71 years of age, 12 DS subjects from 28 to 63 years of age, and four subjects with AD from 71 to 85 years of age were used for the detection of DYRK1A using Western blotting. Tissue was mortarground in liquid nitrogen. 0.1 g of powdered brain tissue was lysed in 1 ml of 6 M guanidine chloride buffer (6 M GuHCl, 10 mM NaH₂PO₄, 10 mM Tris, 20 mM imidazole, pH 8) and sonicated. The protein concentration was determined by bicinchoninic acid assay (BCA) (Pierce Rockford, IL, USA). The total amount of protein extracted from each brain sample was calculated, and volumes corresponding to 4 mg of protein were mixed with 50 μ l of Ni-Sepharose High Performance (Amersham) slurry and incubated for 2 h at 4°C with constant rotation. Ni-Sepharose beads were then washed three times with 1 ml 8 M urea buffer (8 M urea, 100 mM NaH₂PO₄, 10 mM Tris, pH 5.9) for each wash. Elution was carried out by boiling for 2 min in 50 μ l of 1× tricine sample buffer (BioRad) with 50 mM EDTA. Ten microliter of each eluate was electrophoresed on 8% tricine SDS-polyacrylamide gel and transferred onto PVDF membrane. The membranes were blocked in 5% non-fat milk and incubated overnight with the appropriate dilution of monoclonal 7F3 [70] and goat polyclonal G19 (Santa Cruz Biotech, Inc.) antibodies. After incubation with alkaline phosphatase-conjugated secondary antibody (Amersham), the membranes were developed by color reaction using BCIP/NBT as a substrate. The immunoreactive bands on digitalized images were densitometrically quantified using the 1D Scan EX3.1 software program (Scanalytics Corp., Rockville, MD, USA).

Morphometry

On consecutive sections from the brains of subjects with DS/AD and sporadic AD, the numerical density (n/mm³) of neurons stained with cresyl violet (CV), and of neurons with NFTs immunostained with antibodies Tau-1 and G-19 was evaluated. NFTs were counted in the four layers of the entorhinal cortex known to be affected early and severely with neurofibrillary degeneration (transentorhinal/entorhinal stage of neurofibrillary degeneration) [5], including layers II (islands of stellate cells), III, V, and VI. Sectors CA1 and CA4 of the cornu Ammonis and the pyramidal layer in the subiculum proper were selected as representative for the limbic stage of neurofibrillary degeneration. Neurons with NFTs were counted in a frame of 0.0035 mm² in the entorhinal cortex and in a frame of 0.007 mm² in three subdivisions of the hippocampal formation. The depth of the examined dissector was 30 µm. Based only on neurons containing Tau-1-positive NFTs, the proportion of neurons with G-19-positive NFTs was calculated. NFTs were counted in three histological sections representing the frontal, middle, and caudal portions of the entorhinal cortex and hippocampal formation. Twenty-five test areas were evaluated for each examined subregion at a final magnification of 3,100×.

Results

All samples were subjected to partial purification by immobilized metal (Ni²⁺) ion affinity chromatography (IMAC). DYRK1A binds with high affinity to nickel ions chelated to sepharose matrix through the histidine-rich domain in the C-terminal part of the molecule [20, 23]. The samples were made out of pools of cerebral frontal cortex tissue of five control, 12 DS, and four AD subjects. Figure 1 shows the

103 kDa	CTR DS AD			CTR DS AD		
	boat boat south			aning).	10.00	
	7F3 mAB			Dyrk1A (G-19) AB		

Fig. 1 Representative Western blot of Ni-affinity purified Dyrk1A from the pools of the cerebral frontal cortex tissue of control (CTR five cases; 31-71 years), Down syndrome (DS 12 cases; 28-63 years), and Alzheimer disease (AD four cases; 71-85 years) subjects. Mixing equal amounts of powdered brain tissue from each case made pools, which were then processed for Western blotting and densitometric quantification. Equal loads of two identical sets of samples were separated on 8% tricine SDS-polyacrylamide gel and immunoblotted with a 1:5,000 dilution of monoclonal 7F3 antibody at a 1:10,000 dilution of polyclonal G19 antibody. The blot was developed by color reaction using AP-conjugated secondary antibodies and BCIP/NBP substrate. The density of immunoreactive bands on digitalized images was measured by the 1D Scan EX3.1 software program. The value of the DS:CTR ratio was the mean \pm SD of three independent Western blots. Densitometry of control and DS samples revealed an enhancement of DYRK1A levels in the DS pool by a factor of 1.47 for 7F3, and 1.9 for G19 antibody. Both antibodies revealed less DYRK1A in the AD pool when compared to controls

immunoblot of IMAC-purified samples representing each pool, developed with monoclonal 7F3 and polyclonal G19 antibodies, both directed against the N-terminus of DYRK1A protein. In all three pools, 7F3 and G19 antibody detected, with similar relative intensity, two closely spaced bands with the approximate molecular weights of 97 and 94 kDa. Densitometrical quantification of immunoreactive bands revealed an enhancement of DYRK1A levels in the DS pool by the factor of 1.47 and 1.9 for 7F3 and G19 antibody, respectively. When compared with control subjects, the AD pool showed visibly less DYRK1A by both antibodies. The data shown in Fig. 1 are part of our published study [13] in which cases comprising each pool were tested and evaluated individually. In the course of this study, we measured the levels of DYRK1A in six brain structures and found a statistically significant (P < 0.05) increase in the DS group. The overall increase calculated for all six structures was 1.43-fold, very close to the predicted gene dosage value of 1.5.

Immunodetection of DYRK1A in neurons free of neurofibrillary degeneration

Immunostaining with mAb 7F3 revealed Mnbk/Dyrk1A in the nucleus and cytoplasm of neurons, including neuronal processes and synapses. However, the reaction varied substantially, ranging from heavy granular staining in the cell nucleus and cytoplasm to just traces or even a total lack of reaction product. The immunoreactivity was neuron typeand brain structure-specific. Cytoplasmic immunoreactivity with mAb 7F3 in normal brain was characterized in our previous article [70]. Antibodies to DYRK1A G-19, X1076, and 324446 also immunolabeled neurons in the brains of controls, patients with DS, and patients with sporadic AD. Immunoreactivity was observed mainly in the cell body, but some reactivity was present in apical dendrites, especially in sections stained with mAb 7F3 (not shown) and pAb G-19 (Fig. 2a–c).

DYRK1A immunoreactivity in NFTs

Antibody Tau-1 to tau protein and three antibodies to DYRK1A (G-19, X1079, and 324446) immunolabeled neurons in different stages of NFT formation (Fig. 2d-i). In the early stage of neurofibrillary degeneration, formation of NFTs was associated with enlargement of the cell body and deformation and displacement of the cell nucleus to the cell periphery. In this stage, the entire cytoplasm, including NFTs, showed weak to moderate immunoreactivity to DYRK1A (Fig. 2d-f). The progression of changes was associated with a more selective enhancement of reaction in bundles of cytoplasmic fibers with the morphology of NFTs and the proximal portion of dendrites, including apical dendrites of pyramidal neurons (Fig. 2g-i). In contrast to the common and strong pAb G-19 immunoreactivity of intraneuronal NFTs, and the weaker immunoreactivity with pAbs to the C-terminus of DYRK1A (X1079 and 324448), NFTs released from dead neurons (ghost tangles) were always immunonegative (Fig. 2j-l). Mab 7F3 did not detect intracellular or extracellular NFTs (ghost tangles, Table 2).

Immunostaining with pAb G-19 revealed numerous DYRK1A-positive NFTs in the entorhinal cortex, limbic structures, and isocortex in subjects with DS, but many fewer in patients with sporadic AD (Fig. 3).

The appearance of Tau-1 immunoreactive NFTs in neuronal bodies was associated with the appearance of Tau-1-positive deposits in dendrites (neuropil threads). The increase in the number of Tau-1-positive neuronal bodies was paralleled with an increase in the number of Tau-1immunopositive neuropil threads in DS/AD and sporadic AD. However, in sections stained with antibodies G-19, X1079, and 324448, immunolabeling of NFTs in cell bodies was observed in the almost total absence of immunoreaction in neuropil threads for both groups (Fig. 3). The increase in the number of DYRK1A-positive NFTs in persons with DS and severe Alzheimer-type pathology was not associated with an increase in the number of immunopositive neuropil threads.

Colocalization of abnormally phosphorylated tau and DYRK1A in NFTs

Intraneuronal co-localization of DYRK1A and NFTs was confirmed by double-immunofluorescence and confocal microscopy. For Tau-1-negative neurons, the staining for DYRK1A was granular and the product of immunoreactivity

Fig. 2 Antibodies to the N-terminal (G-19) and the C-terminal portion (X1079 and 324446) of DYRK1A reveal immunopositive material in the cytoplasm in neurons in CA1 (a), the nucleus basalis of Meynert (b), and CA4 (c) in a 31-year-old control subject (arrowheads). Early stage of neurofibrillary degeneration was associated with an increase in the size of neuronal body filled with G-19-, X1079-, and 324446-positive NFTs; displacement of the nucleus to the cell periphery; and diffuse cytoplasmic immunoreactivity in cell cytoplasm (arrowheads CA4 sector in a 52-year-old subject with DS/AD; **d**-**f**). In the advanced stage of NFT formation, immunopositive material marks mainly bundles of NFTs (arrowheads CA1 sector in a 52-yearold subject with DS/AD: g-i). However, ghost tangles (arrows) are immunonegative when stained with these three antibodies (CA1 sector in a 52-year-old subject with DS/AD; j-l). Magnification: $\mathbf{a} \times 390$; **b–l**×690



was evenly distributed in the cytoplasm in both the neuronal body and cell nucleus (Fig. 4a; and higher magnification, Fig. 4aa). In some neurons, the accumulation of abnormally phosphorylated tau in the early stage of neurofibrillary degeneration (pretangle stage) was associated with the formation of numerous Tau-1-immunopositive cytoplasmic granules in the cell body and dendritic tree (Fig. 4b, bb). Some Tau-1-positive deposits were co-localized with G-19-immunofluorescent

Table 2 Immunoreactivity in neurons, glial cells, and corpora amylacea with antibodies to DYRK1A

Structures	Antibody						
	7F3 N-terminus	G-19 N-terminus	X1079 C-terminus	324446 C-terminus			
Neuron							
Nucleus	+++	+	+	+			
Cytoplasm	+++	++	++	++			
NFTs	-	++++	++	++			
Neuropil threads	-	±	±	±			
Ghost tangles	_	_	-	_			
Dystrophic processes in amyloid plaques	+	+	+	+			
GVD (cores)	_	_	++	++			
Astrocyte							
Nucleus	_	_	_	_			
Cytoplasm	++	++	++	++			
Microglia and oligodendrocytes							
Nucleus, cytoplasm	_	_	-	_			
Corpora amylacea	+++	++	++	++			

NFTs neurofibrillary tangles, GVD granulovacuolar degeneration

material, but some were not (Fig. 4c, cc). In spite of the strong immunoreactivity of DYRK1A in both the cell body and processes, co-localization of Tau-1- and G-19-positive material was limited almost exclusively to the cell body. The progression of neurofibrillary changes was reflected in an increase in the amount of Tau-1-positive material and the formation of bundles of NFTs immunoreactive only with Tau-1 or both Tau-1 and G-19 (Fig. 4d–f). In some neurons, the portion of bundles of NFTs was immunoreactive with Tau-1 or G-19 only, or with both antibodies (Fig. 4g–i). Tau-1-positive neuropil threads were almost always G-19-negative. In contrast to the uniform distribution of a fine granular product of reaction with pAb G-19 observed in normal neurons, G-19-immunopositive material formed several clusters of rough granules in astrocytes (Fig. 4d, f).

DYRK1A-immunoreactive NFTs in morphometric study

Figure 5 illustrates the result of the immunocytochemistrybased quantitative evaluation of G-19-positive NFTs in subjects with DS/AD and sporadic AD. In subjects with sporadic AD, the percentage of G-19-positive NFTs was structure-specific: 10%, 16%, and 14% in layers II, III, and V in the entorhinal cortex, respectively; 24% in the pyramidal layer in the CA1 sector; but only 7% in the CA4 sector and 12% in the pyramidal layer of the subiculum proper. In subjects with DS/AD, the numerical density of G-19-positive NFTs in entorhinal cortex layers II, III, and V was increased by about $3.7 \times$, $2.0 \times$, and $3.0 \times$, respectively. In the CA1 and CA4 sectors, their numbers were $1.5 \times$ and $3.6 \times$ higher, respectively, and in the subiculum, the number was increased by $2.6 \times$.

Neuronal loss and neurofibrillary degeneration in the second layer of the entorhinal cortex (EC) of subjects with DS/AD and sporadic AD is illustrated in Fig. 6. The similar age at the onset of AD pathology in almost all DS subjects results in the rather uniform neuronal loss in the second layer of the EC of DS subjects and the correlation between the numerical density of CV-positive neurons and age (r = 0.75). The neuronal loss is paralleled by a decrease in the numerical density of Tau-1-positive NFTs (r = 0.65). The immunoreactivity of NFTs with G19 in all DS subjects, including the youngest in the examined cohort (38 years), suggests that DYRK1A contributes to the early onset of neurofibrillary degeneration in DS subjects. The increase in the percentage of G19-positive NFTs (considering Tau-1-positive neurons as 100%) with age (r = 0.51)suggests that DYRK1A contribution to neurofibrillary degeneration in DS increases with age. In three DS subjects, 58, 63, and 67 years, almost all tau-positive NFTs were G19-positive.

In the AD cohort, the numerical density of CV-positive neurons in the second layer of the EC decreases from 30,000–40,000 mm⁻³ in the early stages to about 15,000 in the late stages (Fig. 7b–f). In contrast to DS, sporadic AD starts at a different age, and the correlation between the number of neurons and age is not significant (r = 0.29), but the number of neurons correlates with FAST stage (r = -0.87). The decrease in the numerical density of Tau-1-positive neurons correlates with the stage of AD (r = -0.76), but not with age (r = 0.22). In sporadic AD, G19-positive neurons represent a low percentage of Tau-1-positive NFTs. They were not found in eight subjects with sporadic AD. In six subjects, the

Fig. 3 In a 67-year-old subject with DS, the presence of numerous Tau-1-immunoreactive NFTs in neuronal bodies and apical dendrites (arrows) is paralleled with the appearance of numerous Tau-1-immunopositive neuropil threads (arrowheads) in the amygdala (AMY), the second layer of the entorhinal cortex (EC), and the pyramidal layer in the cornu Ammonis (CA1). In a subject with sporadic AD and moderately severe clinical pathology at the time of demise (FAST 6d), only a few NFTs (arrow) are G-19-positive. In contrast to Tau-1, which stains numerous neuropil threads, antibody G-19 does not detect neuropil threads. Magnification \mathbf{a} - $\mathbf{c} \times 55$, \mathbf{d} - $\mathbf{l} \times 430$



number was less than 10%. In only six AD subjects, it varied from 25 to 67%. In contrast to DS, in sporadic AD, in the later stages of disease (FAST stages 6 and 7), the number of G19-positive NFTs was low, or they were not found.

Dephosphorylation with alkaline phosphatase did not change the pattern of DYRK1A distribution in NFTs (not

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shown) in the brains of people diagnosed with DS/AD or sporadic AD.

DYRK1A immunoreactivity in astrocytes

Two types of reaction were observed in astrocytes. In the majority of the subjects with DS, almost all astrocytes

Fig. 4 Section from the cornu Ammonis of the 67-year-old DS subject immunostained with pAb G-19 (green color) and mAb Tau-1 (red color) and analyzed in confocal microscope. Fine-granular, evenly distributed immunoreactivity is detected with pAb G-19 in the body of two neurons marked with arrowheads (a, aa higher magnification of one of these neurons). One neuron contains roughgranular deposits of Tau-1positive material (arrow) corresponding to an early stage of neurofibrillary degeneration (b, bb). Yellow color corresponds to partial co-localization of the product of reaction for DYRK1A and abnormally phosphorylated tau protein (c, cc). d-f A normal neuron positive for DYRK1A and negative for Tau-1 (1); a neuron with NFT immunoreactive with Tau-1 only (2); and a neuron with NFT immunoreactive with both antibodies (3; G-19, green; Tau-1, red; co-localization, yellow). The cytoplasm of astrocytes is filled with clusters of G-19positive rough granules (arrowheads). A neuron in the late stage of neurofibrillary degeneration (g-i, arrowhead) contains some bundles of NFTs immunoreactive with both G-19 and Tau-1 (vellow). Magnification **a**–**c** and **d**–**f** \times 410, **aa**–**cc** ×1,430, g-i ×1,025



contained a variable amount of clusters of rough granules that were 7F3- and G-19-positive but Tau-1-negative. Immunoreactivity with X1079 and 324448 was less consistent and usually weaker than immunolabeling with 7F3 and G19. In patients with sporadic AD, granular staining in astrocytes was less pronounced, but astrocytes in the subependymal zone and in the molecular layer of allocortex and isocortex often showed mild to strong diffuse staining both in the astrocyte body and the processes. This pattern of staining was especially common and strong in astrocytes in neuritic plaque corona (Fig. 7a–c). Immunoreactivity with 7F3, G-19, X1079, and 324448 was undetectable in microglial cells and oligodendrocytes. DYRK1A immunoreactivity in neuritic plaques

All subjects with DS/AD older than 38 years of age and all patients with sporadic AD were affected by amyloidosis, with numerous neuritic plaques detected with mAb 4G8 in both cortical and subcortical gray matter (not shown). In a large proportion of the neuritic plaques detected with mAb 4G8, mAb Tau-1-labeled numerous swollen dystrophic neuronal processes containing abnormally phosphorylated tau. Mild/moderate reactions with G-19, X1079, and 324448 were observed in about 20% of these plaques in areas corresponding to dystrophic neuronal processes (Fig. 7d–f).



Fig. 5 Graph illustrates a several-fold higher percentage of G-19-positive NFTs (when the number of Tau-1-positive NFTs is considered 100%) in the entorhinal cortex (*EC*) layers II, III, and V, in the *CA1* and *CA4*, and in the subiculum proper (*SUB*) in the brains of persons with DS compared to subjects with sporadic *AD*

Granulovacuolar degeneration and corpora amylacea

In subjects with DS/AD and sporadic AD, numerous neurons in the pyramidal layer of the cornu Ammonis were affected with granulovacuolar degeneration (GVD). The cell body contained several large vacuoles, each with a dense core. The number of affected neurons was increased in the middle stages of AD pathology. However, in subjects with severe AD, loss of neurons was associated with a reduction in the number of neurons with GVD. In the cornu Ammonis, immunostaining with C-terminal antibodies X1079 and 324448 revealed moderate to strong reactions in the cores in cytoplasmic vacuoles of neurons affected with GVD; however, reactions with N-terminal antibodies 7F3 and G-19 were undetectable (Fig. 7g–i).

Corpora amylacea were a common finding in control subjects as well as in subjects with sporadic AD and DS/AD. They were strongly immunopositive in sections stained with 7F3 and moderately immunopositive in sections stained with G-19, X1079, and 324448 (Fig. 7j–l). A strong reaction was present not only in large corpora amylacea in the subpial, subependymal, and perivascular location but also in tiny corpora amylacea dispersed in the cortex and subcortical structures undetectable with CV.

Discussion

Recent studies indicate that DYRK1A is involved in several molecular mechanisms leading to tau phosphorylation and that overexpression of DYRK1A in patients with trisomy of chromosome 21 and DS may contribute to the early onset of neurofibrillary degeneration. DYRK1A is present in the neuronal cytoplasm and nucleus [14, 70] and has been reported to phosphorylate both cytoplasmic and nuclear substrates [30, 40, 53, 74], suggesting that the subcellular localization of the protein reflects the cytoplasmic and nuclear pathway leading to neurofibrillary degeneration.

Cytoplasmic pathway

DYRK1A phosphorylates the human microtubule-associated protein tau at Thr²¹² in vitro, a residue that is hyperphosphorylated in AD and tauopathies, including Pick disease [14, 74], and permits phosphorylation of Ser²⁰⁸ by glycogen synthase kinase 3 [74]. Ferrer et al. [14] observed modifications in the expression of the constitutive cytoplasmic and nuclear DYRK1A in neurodegenerative diseases associated with tau phosphorylation including DS, AD, and Pick disease. However, they could not find any evidence that DYRK1A is directly involved in neurofibrillary degeneration.

Application of two antibodies to the N-terminal portion of DYRK1A, 7F3 [70] and G-19, and of two antibodies to the C-terminal portion of DYRK1A, X1076 and 324446, shows that the distribution of DYRK1A in the normal human brain undergoes modifications associated with neurofibrillary degeneration in DS/AD and sporadic AD. In control brains, all four antibodies reveal immunoreactivity in neurons, both in the cell body and the processes. A stronger reaction is observed in sections stained with antibodies to the N-terminal compared to the C-terminal portion of DYRK1A. DYRK1A contains a nuclear targeting sequence and is distantly related to the mitogen-activated protein kinase family [27]. Intranuclear localization of DYRK1A was shown in human neurons with mAb 7F3 [70], but three antisera, including G-19, detect less immunoreactivity in the cell nucleus.

Our study offers evidence that DYRK1A is directly involved in neurofibrillary degeneration. Antiserum G-19 shows a strong reaction in the majority of NFTs in patients with DS, and antisera X1079 and 324446 react with only about 10% of G-19-positive NFTs.

On the other hand, mAb 7F3 does not react with NFTs. The observed differences in immunoreactivity of NFTs with antibodies and antisera against different sequences of DYRK1A suggest that epitopes detected by these antibodies/antisera are masked in complexes of DYRK1A with tau and other proteins in intracellular NFTs.

Nuclear substrates phosphorylated by DYRK1A

We documented that DYRK1A phosphorylates alternative splicing factor (ASF) at three sites (unpublished data).



Fig. 6 Graphs illustrate changes in the numerical density of neurons stained with CV and NFTs immunoreactive with Tau-1 antibody and G19-antibody detecting DYRK1A in NFTs in DS subjects (*left column*) and in AD (*right column*) in the second layer of the entorhinal cortex. The early onset of neurofibrillary degeneration results in neuronal loss. In the DS cohort, the decrease in the numerical density of CV-positive neurons and Tau-1-positive NFTs correlates with age (r = -0.75 and -0.65, respectively; P < 0.05). DYRK1A-positive NFTs (G19-immunoreactive NFTs) are present in the entorhinal cortex of all DS subjects from 38 to 72 years. The percentage of DYRK1A-positive NFTs (tangles immunoreactive with G19) increases with age

Phosphorylation of ASF by SRPK1 in cytosol results in ASF relocation to the nucleus, whereas phosphorylation by Clk/Sty releases ASF from speckles [32, 45] and recruits it into nascent transcripts, where ASF regulates alternative splicing. Phosphorylation of nuclear ASF by DYRK1A drives ASF back to speckles, decreases involvement in regulation of alternative splicing, and inhibits ASF's ability to promote tau exon 10 inclusion. The result is an increase in 3R tau level and an imbalance of 3R/4R tau ratio. The imbalance could be the factor initiating and accelerating neurofibrillary degeneration in the brain of patients with DS.

DYRK1A in NFTs

Co-localization of DYRK1A with NFTs, described in this article and by in vitro studies showing tau protein



(r = 0.51; Tau-1-positive NFTs were considered as 100%). Also in the AD cohort, a decrease in the numerical density of CV-positive and Tau-1-positive NFTs strongly correlates with stage of disease (FAST; r = -0.87, and r = -0.76, respectively). However, in contrast to the DS cohort, 40% of subjects with sporadic AD do not have DYRK1A-positive NFTs; in 30% of subjects, less than 10% of Tau-1-positive NFTs are G-19-immunoreactive, and only 30% of subjects have from 25 to 67% of G19-immunoreactive NFTs. The majority of subjects with severe sporadic AD (*FAST 6* and 7) are free of DYRK1A-positive NFTs

phosphorylation with DYRK1A priming tau phosphorylation with GSK-3 β , suggests that DYRK1A plays a role in neurofibrillary degeneration and that this process is much more pronounced in neurons of patients with DS than in patients with sporadic AD. In NFTs, tau protein is phosphorylated by several protein kinases, including GSK-3 β , cyclin-dependent kinase 5 (cdk5), JNK, ERK1/2, and p38 at more than 30 phosphorylation sites [18, 22, 34, 42, 52, 60]. Several kinases in their activated forms co-localize with NFTs in AD, including mitogen-activated protein kinase (MAPK) [66], microtubule-affinity regulating kinase (MARK) [9], GSK-3 β [36, 47, 48, 76], cdk 5 [76], cdc2related kinase [38], and casein kinase 1δ (Cki δ) [57]. This study shows that DYRK1A colocalizes with NFTs in all subjects with DS/AD but in only 60% of subjects diagnosed with sporadic AD.

Fig. 7 In a 67-year-old subject with DS, in some cored plaques, diffuse immunoreactivity with G-19, X1079, and 324446 appears in the activated astrocyte body and processes (arrow*heads*, *CA1* **a**–**c**). In about of 20% of cored plaques, antibodies to DYRK1A label dystrophic neuronal processes (arrows; d-f). In the CA1 sector, granules (arrowheads) in neurons with granulovacuolar degeneration are negative when stained with antibodies to the N-terminus of DYRK1A (7F3, not shown; G-19 shown in **g**) but positive when stained with X1079 and 324446 (h, i). Strong immunoreactivity with all three antibodies appears in corpora amylacea (arrowheads) in the hippocampus (j and l dentate gyrus; k CA4). Magnification $\mathbf{a}-\mathbf{f} \times 210$, **g**–**i** ×1,075, **j**–**l** ×210



Gene dosage effect in DS

A gene dosage effect has been proposed to explain the role of an extra chromosome 21 in the pathology of DS. Studies in fetuses with DS show reduced (HACS1) or unmodified levels of some proteins (alpha A-crystalline, FTCD, GARS-AIRS-GART, and CBS), including DYRK1A, which are encoded by genes on chromosome 21. These data call into question the link between overexpressed genes and the formation of DS phenotype in utero [8].

However, in adults with DS, Western blots revealed an almost 50% higher level of DYRK1A protein in extracts from the frontal cortex. The study of other cortical regions in the human cortex confirmed a 50% increase in expression of this kinase linked to an extra copy of Dyrk1A gene in subjects with trisomy of chromosome 21 and DS [13]. The presence of DYRK1A-positive NFTs in all subjects with DS/AD but in only 60% in sporadic AD cases, and a much higher percentage of NFTs in DS subjects than in subjects with sporadic AD appears to reflect the overexpression of the DYRK1A gene and the link between the excessive amount of this kinase and the onset and progression of neurofibrillary degeneration. Overexpression of the APP gene, increased concentrations of $A\beta$ in DS fetuses [64], and the early onset of diffuse [35, 73] and fibrillar A β deposits [73] represent additional evidence of the selective overexpression of genes throughout the entire lifespan of patients with DS.

Differences in NFTs' immunoreactivity in DS/AD and sporadic AD

GSK3 β , cyclin-dependent kinase (cdk) 5 [76], and DYRK1A were found in only some NFTs. These observations and the significant differences in NFTs' immunoreactivity in patients with DS/AD and sporadic AD imply the presence of multiple patterns of tau phosphorylation and neurofibrillary degeneration. Perhaps, the percentage of NFTs that are positive to a given kinase is disease-, stage of disease-, and brain structure-specific. The present observations may also suggest that patterns of tau phosphorylation with DYRK1A in persons with DS are different than those in patients with sporadic AD. This difference may reflect a variation in the progression of neurofibrillary degeneration and, possibly, neuronal loss. In spite of the presence of significant neurofibrillary degeneration and amyloidosis- β in almost all patients with DS 40 years of age and older [72, 73], clinically significant functional deterioration has not been reported to increase in prevalence until some 10 years later [24, 33, 76]. The prevalence of dementia in the DS cohort is estimated as 55% in the age group of 50-59 years, and 75% in subjects older than 60 years of age [33]. One may hypothesize that tau phosphorylation and neurofibrillary degeneration are modified in persons with DS/AD when compared to sporadic AD. The presence of DYRK1A in the NFTs of the youngest DS subjects with neurofibrillary degeneration indicates DYRK1A's contribution to the early onset of NFTs in DS. A lower percentage of G-19positive NFTs in DS subjects before they reach 50 years of age and a high percentage in subjects in their 50s or 60s might be an indication of an enhanced contribution of DYRK1A to tau phosphorylation and neurofibrillary degeneration in older patients with DS. The lack of immunoreactivity of NFTs with antiserum detecting DYRK1A (G19) in 40% of AD subjects and the low percentage of G19-positive NFTs in the other 30% of AD subjects reflect the difference in contribution of overexpressed DYRK1A to neurofibrillary degeneration in DS/AD and the contribution of normal level of DYRK1A in sporadic AD.

The study of chromosome 21 genetic associations with late-onset AD revealed that the DYRK1A gene has the highest significance as a genetic risk factor [30]. The dramatic difference between DYRK1A's role in neurofibrillary degeneration in DS/AD and in sporadic AD appears to be the product of several molecular mechanisms, including (a) the cytoplasmic pathway with phosphorylation of tau by DYRK1A and the priming of tau phosphorylation by GSK3b [40, 78], (b) the nuclear pathway with phosphorylation of ASF by DYRK1A and the imbalance between 3R and 4R tau (unpublished data), as well as (c) phosphorylation of APP by DYRK1A at Thr⁶⁶⁸ in vitro and in mammalian cells [54]. The experiments showing that $A\beta$ loading in neuroblastoma cells and transgenic mice increases expression of DYRK1A gene, resulting in hyperphosphorylation of tau at Thr²¹², support the hypothesis that DYRK1A could be a key molecule linking β -amyloid production with tau phosphorylation [30, 53, 54].

DYRK1A in neuropil threads and dystrophic neurites, but not in ghost tangles

A striking feature of the immunolabeling of neurons with neurofibrillary degeneration with antibodies to GSK-3 β , MPM-2, and cdc2 is the positive reaction with PHFs in the neuronal body but the only sparse reactivity with neuropil threads and dystrophic neurites in the plaque perimeter [67, 76]. It was proposed that this difference might be attributable to a differential sensitivity of neuritic components to fixation [38]. The strong immunostaining of NFTs observed in the cell body in the majority of neurons with neurofibrillary degeneration in older subjects with DS/AD, the absence of immunoreactivity with antiserum G-19 in neuronal processes (neuropil threads), and the sparse immunoreactivity observed in dystrophic neurites in cored plaque corona may indicate that DYRK1A is involved in tau phosphorylation mainly in the cell body. These observations may also suggest that the pattern of tau phosphorylation in the cell body is different from that in the dendrites, and that G-19-positive complexes of abnormally phosphorylated tau do not extend to dendrites. However, the possibility that the lack of reaction with antibodies against DYRK1A in neurofibrillary threads in dendrites and synapses could be the result of masking of epitopes in vivo or the loss of immunoreactivity postmortem including tissue processing cannot be excluded at this phase of study.

Staining of intracellular NFTs with antibodies to cdc and MPM-2 was considered evidence that these kinases play a role in the early stages of neurofibrillary degeneration [67]. Transformation of intracellular NFTs into extracellular ghost tangles is associated with the loss of epitopes recognized by antibodies raised against the amino- and carboxytermini of tau [4]. Extracellular NFTs are immunonegative not only to DYRK1A but also to GSK-3 [76]. The majority of ghost tangles are also negative when stained with antibodies against cdc2 and MPM-2 [67]. Proteolytic processing with a loss of the N- and C-terminal portions of tau is associated with the secondary deposition on NFTs of ubiquitin [3, 63], of heparan sulfate proteoglycans [3], and, in some brain regions, of A β [4, 63, 71, 75]. The absence of DYRK1A in ghost tangles indicates that this binding reflects a contribution to neurofibrillary degeneration rather than unspecific adhesion to already formed NFTs.

Lack of DYRK1A in NFTs in aged control subjects

In general, the properties of NFTs in aged individuals without dementia recapitulate those found in NFTs in AD [10]. In contrast to the accumulation of DYRK1A in about 50% of NFTs in DS subjects and the much lower percentage of NFTs in patients with sporadic AD, NFTs in control subjects are free of DYRK1A. The lack of DYRK1A in NFTs of aged controls is not due to the absence of this kinase, because immunocytochemistry shows widespread distribution of DYRK1A in neurons. This pattern resembles the accumulation of protein kinase M ζ (PKM ζ) in the NFTs of subjects with AD but the absence in NFTs of aged control subjects [10]. The observed NFT immunoreactivity indicates that the association of kinases with NFTs is kinasespecific and aging-, AD-, and AD/DS-specific.

DYRK1A in GVD and corpora amylacea

Granulovacuolar degeneration is observed in a small percentage of neurons in the majority of normal aged subjects, and their number increases in persons with AD [65]. The granular component of vacuoles reacts with antibodies to tubulin [49] and antibodies to abnormally phosphorylated tau [11], and GSK-3 β [36, 76]. The presence of DYRK1A immunoreactivity in granules in neurons with GVD detected with C-terminal antibodies and the lack of reactivity with antibodies against the N-terminus may indicate that only N-terminally truncated products of DYRK1A processing contribute to GVD or selectively accumulate in these granules.

Corpora amylacea are considered as a sign of the earliest changes associated with aging and/or neurodegeneration in neurons [6], including axons [1, 7] and synapses [2] as well as glial cells, especially astrocytes [6, 12, 56, 61]. Strong immunoreactivity with antibody 7F3 [70] and with antibodies G-19, X1073, and 324446 suggests the involvement of DYRK1A in this form of neuron and astrocyte degeneration and the early onset of these changes in DS.

DYRK1A appears to be involved in many forms of neuronal degeneration including Lewy bodies, a pathological hallmark of Parkinson's disease. DYRK1A phosphorylates α -synuclein at Ser⁸⁷ and selectively binds to α -synuclein in transformed and primary neuronal cells. The fact that aggregates formed by phosphorylated α -synuclein are more toxic when compared with aggregates composed of unmodified wild-type α -synuclein, suggests that α -synuclein inclusion formation regulated by DYRK1A may affect neuronal cell viability [28].

Phosphorylation of tau protein by one kinase facilitates or inhibits phosphorylation by other protein kinases in vitro [26, 58, 68, 74, 77] and in vivo [39]. The reported findings and other studies [40] suggest that kinases work in a coordinated way and that overexpressed DYRK1A is a major factor contributing to the DS-specific pattern of neurodegeneration. DYRK1A's contribution to neurofibrillary degeneration in DS appears to be much more significant than in subjects with two copies of the *DYRK1A* gene and sporadic AD. The presence of DYRK1A not only in NFTs but also in granules in GVD and in corpora amylacea suggests that DYRK1A is involved in all three forms of degeneration and may contribute to the early onset of these pathologies in DS.

Protein kinase A (PKA) might be involved in the abnormal phosphorylation of tau in an upstream position to GSK-3 and possibly other proline-directed protein kinases [39]. DYRK1A phosphorylates tau at several sites and facilitates the phosphorylation of tau at several sites by GSK-3 [40, 74]. Whereas PKA is considered to be a potential therapeutic target for the treatment of neurofibrillary degeneration in patients with sporadic AD [39], overexpressed DYRK1A might be considered a major target for treatment of neuronal degeneration in patients with DS/AD.

Western blots demonstrated increase of DYRK1A in the DS group when compared to the control and AD group. The presence of DYRK1A in NFTs in sporadic AD indicates that DYRK1A contributes to tau phosphorylation even in the absence of increased gene dosage. However, an increase in DYRK1A-positive NFTs in a gene dosagedependent manner in subjects with DS, including the youngest NFT-positive DS subjects, indicates that overexpressed DYRK1A contributes to the early onset of neurofibrillary degeneration in DS. The correlation between the percentage of G19-positive neurons and the age of DS subjects suggests that the contribution of DYRK1A to neurofibrillary degeneration increases in the later stages of AD in the DS cohort. Immunoreactivity with antibodies against DYRK1A in granules in GVD and in corpora amylacea suggests that DYRK1A is involved in the early onset of all three forms of degeneration in DS.

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