

FRONTOTEMPORAL DEMENTIAS: UPDATE ON RECENT DEVELOPMENTS IN MOLECULAR GENETICS AND NEUROPATHOLOGY

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Frontotemporal dementias (FTD) are the second most common type of presenile dementias, considered to be clinically and pathologically different from Alzheimer's dementia (AD). FTD differs clinically from AD because memory loss is rarely an early symptom. Instead, FTD is usually denoted by behavioural and language difficulties, and may co-occur with motor neuron disease (MND). Frontotemporal lobar degeneration (FTLD) with ubiquitin-positive, tau-negative inclusions (FTLD-U) is the most common underlying pathology with and without MND. TAR DNA-binding protein 43 (TDP-43), encoded by the TARDBP gene, has been identified as the major pathological protein of FTLD-U with or without MND, demonstrating that abnormal TDP-43 alone is sufficient to cause neurodegeneration. FTLD is a genetically complex disorder. A proportion of cases of FTLD-U have various pathogenic mutations in the *progranulin* (GRN) gene. Other FTLD-U entities with TDP-43 proteinopathy include FTLD-U with valosin-containing protein (VCP) gene mutation and FTLD with MND linked to chromosome 9p. In contrast, chromosome 3-linked dementia, a FTLD-U with chromatin modifying protein 2B (CHMP2B) mutation, has TDP-43 negative inclusions. Thus, TDP-43 defines a novel class of neurodegenerative diseases called *TDP-43 proteinopathies*. These recent discoveries will contribute to an accurate diagnosis, and facilitate the development of diagnosis and therapy.

KEY WORDS: *frontotemporal lobar degeneration, granulin, motor neuron disease, mutation, TARDBP, TDP-43 protein*

Frontotemporal dementias (FTD) are a clinically, genetically, and neuropathologically heterogeneous group of diseases accounting for up to 20 % of presenile dementia cases. FTD is a focal, non-Alzheimer form of dementia, clinically characterised as behavioural and/or language dysfunction, and may co-occur with motor neuron disease (MND) (1). Typically, the patient with FTD does not have an amnesic syndrome, at least in the early stage of the disease, which distinguishes FTD clinically from Alzheimer's disease (AD) (2, 3), but there are exceptions (4). However, the clinical diagnosis of FTD may only be considered after other potential causes of dementia (e.g. small and/or large vessel disease), systemic conditions (e.g., hypothyroidism, vitamin B deficiency), tumours, and substance abuse have been excluded. Frontotemporal

lobar degeneration (FTLD) with ubiquitin-positive, tau-negative inclusions (FTLD-U or FTLD-MND-type) is the most common underlying pathology in FTD with and without MND (5). TAR DNA-binding protein 43 (TDP-43), a nuclear protein implicated in exon skipping and transcription regulation (6, 7, 8) was recently identified as the major pathological protein of sporadic and familial FTLD-U, with and without MND, as well as in sporadic amyotrophic lateral sclerosis (ALS), (9, 10), and rapidly confirmed by others (11, 12). Pathologic TDP-43 in these disorders is abnormally phosphorylated, ubiquitinated and cleaved to generate C-terminal fragments, and is recovered only from pathologically affected CNS regions including hippocampus, neocortex, and spinal cord (10). Therefore, the presence of abnormal

aggregates of phosphorylated and ubiquitinated TDP-43 defines a novel class of neurodegenerative diseases called *TDP-43 proteinopathies* that includes FTLN-U, FTLN, MND, and ALS. The increasing life expectancy will result in increased prevalence of neurodegenerative diseases. This review focuses on exciting findings providing a number of important advances in our understanding of the neuropathology, molecular genetics, and biochemistry of FTD, as one of neurodegenerative diseases that have occurred recently in this rapidly developing field of dementia research.

NEUROPATHOLOGY OF FTLN

FTLN comprises a neuropathologically heterogeneous group of neurodegenerative diseases which share the common feature of predominant degeneration of the frontal and temporal lobes

(13-15). FTLN can broadly be divided into two main classes, based on abnormal accumulation of hyperphosphorylated tau protein: those with tau-positive and with tau-negative inclusions. Whereas, in the past, most attention focused on FTLN associated with tau-based pathology and microtubule-associated protein tau gene (*MAPT*) mutations (*tauopathies*), there has recently been greater attention paid to non-tau or tau-negative FTLN (*non-tauopathies*) (14). FTLN-U accounts for 5 % to 15 % of all dementia disorders (16).

GENETIC STUDIES

FTLN is a genetically complex disorder, with multiple genetic factors contributing to the disease. A positive family history with an autosomal dominant pattern of inheritance and high penetrance is usually found in one quarter to one half of patients (17, 18).

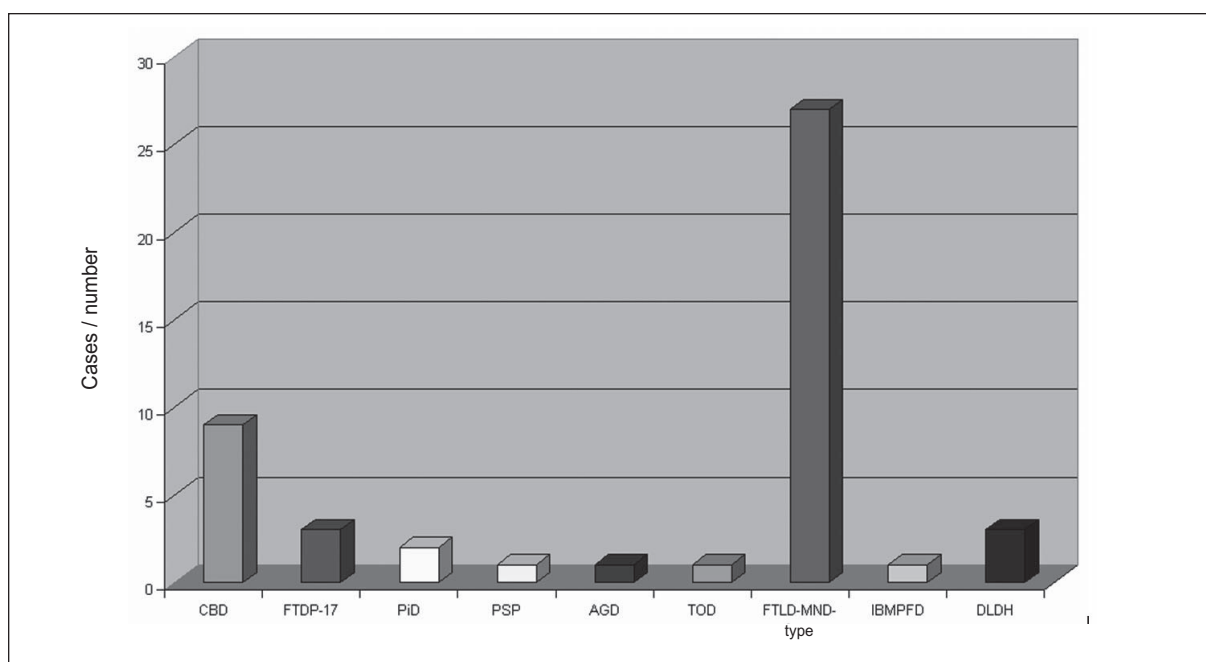


Figure 1 Spectrum of FTLN (14, 15) in a cohort of 48 cases published elsewhere (2). The FTLN-U (or FTLN-MND-type) is the most common pathology associated with clinical FTD

Abbreviations:

CBD = corticobasal degeneration

FTDP17 = frontotemporal dementia with parkinsonism linked to chromosome 17

PiD = Pick's disease

PSP = progressive supranuclear palsy

AGD = argyrophilic grain disease

TOD = tangle only dementia

FTLN-MND-type = frontotemporal dementia with motor neuron disease type inclusions

IBMPFD = inclusion body myopathy with Paget's disease of bone and frontotemporal dementia

DLDH = dementia lacking distinctive histopathology

Various pathogenic mutations in the progranulin (*GRN*) gene were recently reported in individuals with FTL-D-U linked to chromosome 17q21 (19, 20). The *GRN* gene is mutated in 5 % to 10 % of patients with FTD and in about 20 % of patients with familial FTD (21), similar to that of FTD with *MAPT* mutations (22). For the current list of pathogenic mutations in FTL-D please refer to: <http://www.molgen.ua.ac.be/FTDMutations/>. It is now recognised that FTL-D-U is the most common pathology associated with clinical FTL-D (5). Also, several genes and a locus on chromosome 9 have been linked to familial FTL-D-U. Genetic defects include mutations in the chromatin modifying protein 2B (*CHMP2B* gene), the cause of chromosome 3-linked FTL-D, and mutations in the valosin-containing protein (*VCP*) gene associated with inclusion body myopathy, with Paget's disease, and with frontotemporal dementia, which is the cause of chromosome 9-linked FTL-D (18, 23). Locus heterogeneity for FTL-D and MND is indicated by the presence of other genetic loci at 9p. The ubiquitinated pathological protein in FTL-D-U has been identified as TAR DNA-binding protein 43 (*TDP-43*) (9-11). As more entities are investigated, the pathological *TDP-43* protein is found to be a component of the inclusions of an increasing number of neurodegenerative diseases.

CLINICAL PHENOTYPE OF FTD

Recently, we retrospectively examined charts of 48 FTD individuals in order to find clinical differences between FTD and AD. All cases of FTD met pathological criteria for FTL-D (14, 15). Clinically, behavioural and language features, including impulsivity, disinhibition, and social withdrawal were significantly different between FTD and AD, as reported previously (24). The most distinctive feature of FTD, on psychometric tests, was a significant impairment of frontal lobe functioning, as reported earlier (25). The identification of different mutations in the *GRN* gene in hereditary dysphasic disinhibition dementia families 1 and 2 (HDDD 1 and HDDD 2) (26, 27) links these families to other FTL-D-U families with *GRN* mutation (28, 29). A complicating feature in both HDDD families is the presence of AD-type early memory loss which correlated with coexisting AD pathology in almost half of the cases, which distinguishes them from other families with no or little coexisting neurodegenerative disease (26, 27).

Interestingly, another family with the same *GRN* A9D mutation has been reported in an individual with corticobasal syndrome (30), indicating, again, clinical heterogeneity associated with the same mutation.

CONCLUSION

For practicing clinicians, the knowledge that changes in behaviour and language difficulties distinguish those with FTD from AD is important, although clinical and cognitive features may overlap between the two. Typically, patients with FTD do not have an amnesic syndrome, at least in the early stage of the disease, which distinguishes them from AD.

Major discoveries have been made in the recent past in the genetics, biochemistry, and neuropathology of FTD. *TDP-43*, encoded by the *TARDBP* gene, is the major pathological protein of FTL-D-U with or without MND. Thus, *TDP-43* defines a novel class of neurodegenerative diseases called *TDP-43 proteinopathies*. FTL-D-U is now recognised as the most common pathology associated with clinical FTL-D. New developments such as the discovery of *TDP-43* as disease protein have opened new view on FTD, as well as its relation to motor neuron disease.

In summary, the new genetic and pathological information now opens the way for a novel diagnostic approach. The diagnostic responsibility will increase with the development of new diagnostic tests. It is anticipated that these discoveries will contribute to an accurate diagnosis, and facilitate the development of specific therapy.

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Sažetak

FRONTOTEMPORALNE DEMENCIJE - PRIKAZ NOVIH DOSTIGNUĆA IZ PODRUČJA MOLEKULARNE GENETIKE I NEUROPATHOLOGIJE

Frontotemporalne demencije (FTD) druga su po učestalosti grupa presenilnih demencija, koja se kliničkim simptomima i patologijom razlikuje od Alzheimerove demencije (AD). FTD se razlikuje od AD budući da je gubitak memorije rijetko prvi simptom bolesti. Umjesto toga, FTD karakteriziraju smetnje u ponašanju i govoru, a mogu se javiti i simptomi bolesti motornog neurona (BMN). Frontotemporalna lobarna degeneracija (FTLD) s ubikvitin-pozitivnim, tau-negativnim inkluzijama (FTLD-U) najučestalija je patološka slika s BMN-om ili bez njega. Glavni patološki protein FTLD-U s BMN-om ili bez njega je protein 43 vezan s DNA (TDP-43), kodiran od gena TARDBP, čija prisutnost je već dovoljna za nastanak neurodegenerativnih promjena. Glede genetike, FTLD je kompleksna bolest. Dio bolesnika s patološkom slikom FTLD-U pokazuje različite patološke mutacije progranulin (GRN) gena. Ostali FTLD-U entiteti s TDP-43-proteinopatijama uključuju: FTLD-U proteinom vezanim s valozinom i FTLD s BMN-om vezanim uz kromosom 9p. Nasuprot tomu, demencija vezana na kromosom 3, FTLD-U s CHMP2B-mutacijom, sadržava TDP-43-negativne inkluzije. Stoga, TDP-43-protein definira novu grupu neurodegenerativnih bolesti koje nazivamo TDP-43-proteinopatije. Ova novija otkrića pridonijet će točnijoj dijagnozi i ubrzati razvoj dijagnostike i terapije.

KLJUČNE RIJEČI: *bolest motornog neurona, frontotemporalna lobarna degeneracija, granulin (GRN) mutacije, TARDBP, TDP-43 protein*

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