ADVANCES IN NEUROPSYCHIATRY

Presenile dementia syndromes: an update on taxonomy and diagnosis

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The four major degenerative dementias that often begin in presenescence: are reviewed. These are Alzheimer's disease, frontotemporal dementia, dementia with Lewy bodies, and Creutzfeldt–Jakob disease. Their epidemiological, genetic, and clinical features are reviewed, and controversies in taxonomy arising from recent discoveries described. Particular attention is given to the pathological role of protein aggregation, which appears to be a factor in each disease.

New clinical, pathological, and basic science discoveries often extend, challenge, and eventually change the nomenclature of major medical disorders. Nearly 40 years ago in Newcastle, England, the finding of a strong correlation between dementia in elderly nursing home patients and β amyloid plaque concentration in the brain transformed our understanding of what constituted Alzheimer's disease. Before this key series of studies, Alzheimer's disease was considered a rare disorder that afflicted presenile populations, while dementia in the elderly, or “senility”, was a poorly defined condition thought to be secondary to cerebral arteriosclerosis. Linking the brain changes originally described by Alzheimer in patients with a presenile dementia to patients with senile dementia transformed the field. Subsequent studies on the genetic basis, pathogenesis, and clinical sequelae of the amyloid plaque have produced a remarkably coherent picture of Alzheimer's disease across all these areas of research. The same has been true for the varied set of genetic, infectious, and sporadic disorders that were linked together through the discovery of prions. In contrast, nomenclature issues with what was once called Pick's disease and is now called frontotemporal dementia have been more complicated, and recent discoveries challenge previous dogma regarding how we should classify patients with the widely varied clinical and pathological syndromes now linked to frontotemporal dementia. Dementia with Lewy bodies (DLB) also presents a less coherent picture, given its predominantly non-Mendelian genetics and considerable overlap with both Alzheimer's disease and Parkinson's disease.

Increasingly, neurodegenerative syndromes are considered disorders of abnormal protein aggregation—a concept discussed in an accompanying article in this issue of the journal. Alzheimer's disease is strongly linked to the accumulation of amyloid β-42 protein (Aβ42), frontotemporal dementia to abnormalities in the protein tau, and Creutzfeldt–Jakob disease to abnormal aggregation of the prion protein. With each of these dementias there are genetically determined forms of the illness caused by mutations in genes that either code for or affect the function of a protein. Similarly, for each dementia subtype there is a sporadic form of the illness in which no genetic abnormalities are evident. Beyond the single gene mutations that lead to abnormal protein aggregation there are susceptibility genes that increase the likelihood that abnormal protein aggregation will occur. An apolipoprotein gene, e4, represents a susceptibility gene for Alzheimer's disease and DLB, while homozygosity for codon 129 on the prion protein increases susceptibility to Creutzfeldt–Jakob disease (table 1).

The remarkable coherence between clinical, pathological, and molecular findings in both Alzheimer's disease and Creutzfeldt–Jakob disease has protected these disorders from major controversy over nomenclature. However, with frontotemporal dementia the field is substantially divided between “lumpers” and “splitters”. This dispute is complicated by the discovery of substantial overlap between frontotemporal dementia and disorders traditionally considered under the category of parkinsonian–dementia syndromes, including progressive supranuclear palsy and corticobasal ganglionic degeneration. Similarly, DLB has been subject

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to nomenclature controversy owing both to the prominent clinical overlap with Alzheimer's disease and Parkinson's disease and to inconsistent clinical criteria. In this article we review the four major degenerative dementias that often begin in presenescence: Alzheimer's disease, frontotemporal dementia, DLB, and Creutzfeldt–Jakob disease. We review some epidemiological, genetic, and clinical features of these disorders and describe controversies in taxonomy that have arisen from recent discoveries.

ALZHEIMER'S DISEASE

Alzheimer's disease is the most common degenerative dementia, accounting for around two thirds of all cases. Alzheimer's disease most commonly occurs in late life, but a small percentage of patients have onset before 60 years (presenile). Differences between early and late onset cases do exist, and one clear distinction is that many presenile cases are secondary to known genetic mutations. In contrast, the aetiology of late life Alzheimer's disease is more heterogeneous, with many factors—genetic, age related, and environmental—contributing to the dementia. Despite these and other differences between presenile and senile Alzheimer's disease, most consider the two to be the same illness. Even though only a small minority of Alzheimer patients show dementia in the presenium, these relatively uncommon presenile cases have had a great impact upon our understanding of the pathogenesis of the disease.

In around 2% of cases, Alzheimer's disease is transmitted as an autosomal dominant gene with strong penetrance. Most of these autosomal dominant cases present before the age of 60. Three genes account for these familial cases, and all three result in excessive production of β amyloid 42 (Aβ42), the most neurotoxic of the three common forms of amyloid. The original studies on familial Alzheimer's disease focused on chromosome 21 because the amyloid precursor protein is located on this chromosome and because all patients with Down's syndrome (trisomy 21) develop amyloid plaques by the age of 40. The amyloid precursor protein gene on chromosome 21 was the first Alzheimer's disease related gene to be identified and characterised, and new mutations in the gene continue to be identified. Most of the early onset cases have a documented genetic basis are caused by mutations of the presenilin 1 gene found on chromosome 14, and most of these cases occur before the age of 50. However, there are a few cases caused by mutations in a homologous gene called presenilin 2 on chromosome 1. Presenilin genes code for a pair of proteins that share considerable biochemical overlap, which, when mutated, favour the production of Aβ42. A fourth gene, apolipoprotein ε, coding for apolipoprotein E (apoE), has been implicated in the more common late onset, sporadic form of Alzheimer's disease. ApoE has three common alleles—E2, E3, and E4—and it is the E4 allele that increases susceptibility to Alzheimer's disease. The apoE4 allele is neither necessary nor sufficient for the development of Alzheimer's disease. However, the majority of cases of Alzheimer's disease between the ages of 50 and 60 years are homozygous or heterozygous for apoE4. Other potential susceptibility genes have been proposed.

It is not possible to differentiate presenile from senile Alzheimer's disease on the basis of neuropathological findings. Alzheimer's disease has three important pathological findings: amyloid plaques, neurofibrillary tangles, and neuronal loss. The mechanisms through which plaques, tangles, and neuronal loss develop can be traced by studying genetic forms of the disease and by evaluating mice carrying genetic mutations. Amyloid plaques are extraneuronal aggregates of Aβ protein. Neurofibrillary tangles are aggregations of tau and neurofilaments found in neuronal cell bodies. The plaques and tangles lead to neuronal loss. One staging system for Alzheimer's disease emphasises the density of neurofibrillary tangles. This system relies upon the fact that neurofibrillary tangle distribution and density are the best correlates of cognitive deficits; it stages disease severity from 1 to VI, with stages I and II confined to the entorhinal and hippocampal cortex, and stages V and VI involving the neocortex.

A consensus on the clinical diagnosis of Alzheimer's disease was reached in 1984 with the development of the NINCDS–ADRDA criteria. These criteria remain valid today. To meet the diagnostic criteria for probable Alzheimer's disease, a patient must have deficits in at least two areas of cognition, progressive worsening of memory and other cognitive functions, and onset between ages 40 and 90. Many individuals who carry presenilin 1 mutations become ill before age 40, persuading the value of this item in the criteria. There now appears to be no other neurological disorder that could explain the cognitive decline, and therefore the clinician must be able to differentiate Alzheimer's disease from frontotemporal dementia, DLB, Creutzfeldt–Jakob disease, or other causes of dementia. The NINCDS–ADRDA criteria do not offer much guidance for differentiating other dementias from Alzheimer's disease. Therefore they show good sensitivity (90–95%) but only modest specificity (60–70%).

Efforts have been made to increase the specificity by using adjunctive investigations such as blood tests, cerebrospinal fluid analysis of Aβ42, and measurements of apoE status, yet none of these tests has reached the level of a gold standard for diagnosis.

In the first stages of Alzheimer's disease, when it is pathologically confined to the medial temporal region, patients have episodic memory deficits. Forgetfulness, difficulty in learning new spatial routes, problems with remembering where items have been placed, and difficulty in remembering new names or faces are common symptoms. As the disease moves from the medial temporal lobe to the posterior temporoparietal cortex, language and visuospatial deficits emerge. A thorough mental state examination can therefore often provide a rough estimate of the pathological stage of disease, such that isolated memory impairment may correlate with stage II or III, whereas memory impairment coupled with prominent visuospatial or language difficulties suggests progression to stages V or VI. In studies of patients carrying the genetic mutations that predispose to Alzheimer's disease, a decline in verbal memory and performance intelligence quotient seem to be the earliest manifestations of the illness.

Focal variants of Alzheimer's disease are seen, where memory loss is not a prominent early feature. These focal presentations are common in the presenile setting. In such cases, isolated cognitive impairments occur based upon focal cortical brain degeneration. Naming, praxis, and calculation deficits are seen with focal degeneration in the left temporoparietal regions, loss of executive function and behaviour deficits with focal degeneration in the frontal lobes, visuospatial loss develops with asymmetric right parietal degeneration, and visual disturbances emerge with posterior cortical degeneration. In these cases, Alzheimer's disease may mimic other focal degenerative disorders such as frontotemporal dementia presenting with primary progressive aphasia, or corticobasal degeneration presenting with progressive apraxia. Psychiatric symptoms are common with Alzheimer's disease, particularly as the illness progresses. In addition to relatively mild behavioural problems such as irritability and sleep disturbance, major depression occurs in up to 20% of patients and in the later stages up to 50% will have delusions. Alzheimer's disease is slowly progressive but ultimately fatal. Median survival following the onset of symptoms has been estimated to be in the range of five to nine years. In presenile dementia the course is sometimes more rapid.

The diagnostic value of neuroimaging is now under active study. Positron emission tomography (PET) and single photon emission tomography (SPECT) have been used as adjunctive tests in patients with probable Alzheimer's disease, with the
fairly consistent finding of decreased perfusion or metabolism in the temporoparietal regions. Measures of hippocampal and entorhinal cortex atrophy with structural magnetic resonance imaging (MRI) have been used in efforts to increase diagnostic accuracy. Studies of functional imaging using MRI or PET are still relatively sparse but suggest a pattern of increased task-related activation in the earliest clinical and perhaps even preclinical state, progressing to decreased task-related activation once patients become more symptomatic. With the prospect of preventive interventions for Alzheimer's disease, enhanced understanding of genetic risks, and growing awareness that many patients with minimal cognitive impairment progress to Alzheimer's disease, techniques are being investigated that increase diagnostic specificity in demented and at risk individuals. To date, neuroimaging has been able to distinguish group differences but still lacks specificity for Alzheimer's disease at the individual patient level.

Treatment options for Alzheimer's disease, while still modest in their effect, have continued to emerge and there are now treatments to improve or maintain cognition, prolong independence in activities of daily living, and minimise behavioural problems. The mainstay of treatment currently involves the use of an acetylcholinesterase inhibitor. Several different inhibitors are available with similar efficacy but varying half lives and side effect profiles. Studies suggest that the efficacy of these compounds is modest but can persist for years with continued treatment. Vitamin E has shown efficacy in slowing progression of Alzheimer's disease and has no important side effects. There are many agents that show promise as treatments to prevent Alzheimer's disease, but none has yet been proven to be effective. In selected patients antidepressants or atypical antipsychotic compounds may improve behaviour, though one report has suggested that antipsychotic agents may hasten cognitive decline.

One advance in the study of Alzheimer's disease has been the emergence of transgenic mouse models. Mice carrying mutations in the amyloid precursor protein gene or presenilin 1 gene develop amyloid plaques by six months of age. Various treatments have proven effective in these mice, including lowering cholesterol, treating with inhibitors of the protease γ-secretase, and—perhaps most promisingly—immunising against Aβ42. ApoE4 mice also develop cognitive impairment by six months when compared with mice with wild type apoe4 or apoe3. These mice represent a powerful model for studying the mechanisms leading to brain degeneration associated with apoE4 and provide a testing ground for potential treatments.

FRONTOTEMPORAL DEMENTIA

The nomenclature for frontotemporal dementia has been a modern quagmire, imposed in part by evolving clinical and genetic data. This process began with better clinical and pathological studies of patients with frontotemporal dementia by European investigators, but was further transformed by the finding of tau mutations in familial forms of frontotemporal dementia. No degenerative dementia has seen more names for the same disorder than frontotemporal dementia, which has been called Pick's disease, progressive subcortical gliosis, dementia of the frontal type, frontal lobe dementia of the non-Alzheimer's disease type, thalamic dementia, dementia, disinhibition amytrophic syndrome, frontotemporal dementia with parkinsonism linked to chromosome 17, multi-system tauopathy, dementia lacking distinctive histology, frontotemporal dementia, and frontotemporal lobar degeneration.

First described by Pick in 1892, and later called Pick's disease in 1924, much has been made of whether the Pick body constitutes a core feature of this illness. Even though Pick himself had little interest in the cellular inclusions that bear his name, for many decades pathologists were reluctant to make a specific diagnosis in patients with frontal and temporal atrophy unless these cellular inclusions were present. The majority of patients with frontotemporal atrophy do not have cellular inclusions. Therefore, by insisting upon the presence of Pick bodies and by ignoring frontotemporal cases without Pick bodies, the disease originally described by Pick became rare.

With the emergence of Alzheimer's disease as a major research focus during the 1980s, Pick's disease and other related disorders were badly neglected, particularly in the USA. However, a series of studies by the Lund and Manchester groups brought frontotemporal dementia back into the mainstream of dementia research. In consecutive patients with progressive dementia, these groups showed that somewhere between 12% and 16% of patients with degenerative dementia suffered from a non-Alzheimer's disease pathology characterised by frontotemporal atrophy, neuronal loss, gliosis, and sometimes Pick bodies. A series of studies by these investigators on patients with presenile dementia suggested that frontotemporal dementia was a common presenile illness that could be differentiated from Alzheimer's disease. On the basis of clinical, pathological, and imaging studies, these groups estimated that 12–25% of presenile dementia could be characterised by frontal lobe atrophy without Alzheimer's disease pathology. Fewer than 20% of these cases showed the classical pathological findings of Pick's disease, thus giving rise to a large definition-based increase in prevalence.

The Lund–Manchester groups coined the term frontotemporal dementia in 1994 in an attempt to establish reliable diagnostic criteria for a clinically heterogeneous group of disorders that still shared many distinct pathological features. Recently, a consensus group delineated three main cognitive subtypes of frontotemporal dementia (described in detail below). The validity of this more inclusive perspective (and, in turn, of those early prevalence estimates) has been borne out by recent advances in molecular genetics.

While frontotemporal dementia was known for many years to have a strong genetic component, the first major stride took place in 1994 with linkage of a familial frontotemporal dementia syndrome to chromosome 17. Subsequently, mutations in the tau gene on chromosome 17 were identified in clinically distinct frontotemporal dementia kindreds. Around 40% of cases of frontotemporal dementia are familial and many of these familial cases seem to be inherited in an autosomal dominant pattern. Most cases of frontotemporal dementia linked to chromosome 17 have been associated with a specific mutation on the tau gene and abnormal accumulations of tau in the brain. Recently, it has been discovered that in one of these families an absence of brain tau was a characteristic feature, so called “no tauopathy.” Evidence from some familial and sporadic cases suggests that a subgroup of patients with frontotemporal dementia either overexpress or underexpress a subtype of tau, or express a mutated form of tau. However, there are other kindreds with linkage to chromosome 17 that have yet to yield specific mutations in the tau gene. Furthermore, tau mutations do not account for the majority of familial cases and are rarely seen in sporadic frontotemporal dementia. One report has described a family with frontotemporal dementia linked to chromosome 3. Families with frontotemporal dementia and a strong association with amyotrophic lateral sclerosis rarely show tau mutations and often have no evidence of tau pathology in the brain. Families have recently been linked to chromosomes 9 and 15 (Wilhelmsen KC, personal communication).

Pathologically there is a distinctive foci atrophy of the frontal lobes, the temporal lobes, or both. The atrophy can show a unilateral predominance or be symmetrical. In the temporal lobes the more anterior regions typically show greater pathology, with the amygdala demonstrating more involvement than the hippocampus. Posterior parietal and
temporo-occipital regions are relatively preserved. On microscopically, subcortical structures such as the substantia nigra, putamen, and globus pallidus are not shown marked involvement. Those cases with a motor neurone component also show pathological changes in the anterior horn cells. The microscopic changes in the affected regions include neuronal loss, synaptic loss, gliosis, and spongiosis, often most prominent in the first three cortical layers. Some cases of frontotemporal dementia show swollen neurones with inclusion bodies. Even among cases with inclusions, only a minority shows the classical histological feature of a Pick body—silver staining. Silver staining appears to be associated with the deposition of 3Rtau rather than the 4R isofrom. Families with frontotemporal dementia may present with a primary progressive aphasia, corticobasal ganglionic degeneration pathology has been reported. Patients with both corticobasal degeneration and progressive supra-nuclear palsy are more likely to carry the A0 tau allele, and brain pathology in these conditions always shows tau abnormalities. The presence of tau positive inclusions in progressive supranuclear palsy and corticobasal degeneration has prompted some to view progressive supranuclear palsy, corticobasal degeneration, and frontotemporal dementia simply as clinical variants of “taupathy.”

Estimates of life span following a diagnosis of frontotemporal dementia can vary considerably, ranging from three to 15 years, depending on which symptoms are considered to herald the onset of disease. It is possible that patients with progressive language disorder are more likely to get an early diagnosis than those who present with psychiatric symptoms. Additionally, some of this variability related to the speed of progression is explained by whether or not the patients develop motor neurone disease or parkinsonian features either early or late in the course of the illness. Given the pronounced variation in initial presentation and the common early prominence of behavioural symptoms, misdiagnosis is a common problem for patients and their families. Clinical criteria alone can be useful in distinguishing frontotemporal dementia from Alzheimer’s disease. In a recent study, the Land–Manchester criteria were applied retrospectively to a group of pathologically confirmed cases of Alzheimer’s disease, frontotemporal dementia, progressive supranuclear palsy, and DLB and found to have a sensitivity of 97% and a specificity of 97%. The criteria have not yet been tested in a prospective manner against pathological diagnoses, but will probably prove to be less sensitive and specific in this setting. Increasingly, therefore, neuroimaging is being relied upon to improve both the sensitivity and the specificity of the diagnosis. The addition of SPECT to clinical criteria has been shown to improve diagnostic accuracy to 90%. Patients with frontotemporal dementia tend to show bifrontal and bitemporal hypoperfusion, in contrast to Alzheimer’s disease where there are temporoparietal defects. More recently, structural MRI has shown promise as an adjunct to help in distinguishing frontotemporal dementia from Alzheimer’s disease and other dementias. For the time being, however, clinical criteria remain the mainstay of diagnosis.

Treatment of frontotemporal dementia is limited to symptom management. Serotonin selective reuptake inhibitors may be beneficial in managing the disinhibition, depressive symptoms, carbohydrate craving, or compulsions often encountered in this disease. Support for such an approach can be found in molecular studies showing low serotonin receptor binding in frontotemporal dementia. Some behavioural symptoms may require more aggressive pharmacotherapy, including the use of antipsychotic agents, in which case—as with Alzheimer’s disease—the newer atypical agents are favoured in order to minimise parkinsonian side effects. At least one study has suggested that the cholinergic system is relatively spared in frontotemporal dementia, and often acetlycholinesterase inhibitors do not appear to improve cognitive status and can worsen irritability.

DEMENTIA WITH LEWY BODIES

As with frontotemporal dementia, the epidemiology of DLB is difficult to establish owing to the variability in the diagnostic criteria. The first case description of DLB is attributed to Oka-zaki et al in 1961, but standardised, consensus, diagnostic criteria were not established until 1996. In the interim various different criteria were used to define DLB and so early prevalence estimates must be interpreted with caution. Nonetheless, DLB is increasingly considered to be the second most common degenerative dementia after Alzheimer’s disease and estimates of its prevalence among patients with dementia range as high as 20%. Other studies have reported lower estimates, ranging from 8.5% of dementias in a Scandinavian study to 15% in a Japanese study. Demographically, a male preponderance has been fairly consistently reported, with a ratio of the order of between 1.5:1 and 2:1. The genetic underpinnings of DLB include an autosomal dominant component linked to familial Parkinson’s disease and a non-Mendelian component linked to Alzheimer’s disease. Familial cases of Parkinson’s disease were initially linked to an autosomal dominant but incompletely penetrant
mutation in the α-synuclein gene on chromosome 4q.12 Subsequently, two other genetic loci on chromosomes 2p and 4p were identified in Parkinson’s disease kindreds, also showing autosomal dominant inheritance with incomplete penetrance.13 Individuals in one of the kindreds linked to chromosome 4p have been described who fulfill criteria for DLB rather than Parkinson’s disease.14 Thus cases of DLB seem to make up only a fraction of the already rare kindreds with familial Parkinson’s disease. As with Alzheimer’s disease, therefore, the bulk of the genetic load in DLB is non-Mendelian. The more striking similarity is that the susceptibility gene, apoE4, is the same in the two diseases. Several groups have reported increased apoE4 allele frequencies in DLB patients with115 and without12 Alzheimer changes. Overall it seems clear that there is an increased frequency of apoE4 in DLB but that the effect is less prominent than in Alzheimer’s disease.107

The hallmark pathological finding of DLB is, of course, the presence of cortical Lewy bodies. Lewy bodies are intracytoplasmic aggregates of α-synuclein and other proteins.7 They can be detected on routine haematoxylin and eosin staining, but immunocytochemical stains are more sensitive, α-synuclein stains being superior to ubiquitin stains. Some investigators believe that the location and density of Lewy bodies is linked to the clinical syndrome, such that brain stem Lewy bodies correlate with movement disorder, limbic Lewy bodies with psychosis, and cortical Lewy bodies with depression.136 Others have not detected a tight correlation between the clinical syndrome and the extent and location of Lewy bodies.137

Immunocytochemical stains for ubiquitin and α-synuclein have also led to the detection of Lewy neurites, which are abnormal filaments that may be Lewy body precursors.138 Because Lewy bodies may be seen in a significant percentage of patients with otherwise typical Alzheimer’s disease, the presence of Lewy neurites in CA 2/3 of the hippocampus, more specific for DLB, may be a useful distinguishing neuropathological finding.139

Distinguishing features are especially important given the extensive pathological overlap between DLB and Alzheimer’s disease. The most recent international workshop report states that 15% of DLB cases have severe Alzheimer pathology, 55% have some Alzheimer pathology, and only 30% have no more Alzheimer pathology than age matched controls.140 Insights into the properties of α-synuclein are helping to elucidate the pathogenesis of DLB. The normal function of α-synuclein remains obscure, though it may play a role in synaptic transport of vesicles or synaptic plasticity.141 It is also unclear whether cell injury and death results from loss of the protein’s normal function or from a toxic gain of function caused by protein aggregation. Of all the proteins linked to α-synuclein and other proteins:7 They can be detected on routine haematoxylin and eosin staining, but immunocytochemical stains are more sensitive, α-synuclein stains being superior to ubiquitin stains. Some investigators believe that the location and density of Lewy bodies is linked to the clinical syndrome, such that brain stem Lewy bodies correlate with movement disorder, limbic Lewy bodies with psychosis, and cortical Lewy bodies with depression.136 Others have not detected a tight correlation between the clinical syndrome and the extent and location of Lewy bodies.137

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The use of typical antipsychotics such as haloperidol or chlorpromazine for the treatment of psychosis in DLB is considered risky by many investigators. The use of atypical antipsychotics has also been shown to be effective in improving psychosis in DLB.114–116

**PRION DISORDERS**

A family of diseases—the transmissible spongiform encephalopathies (TSE)—are caused by an agent known as the “prion” (proteinaceous infectious particle). Prions (prion-proteins) were named and discovered by Stanley Prusiner, who was awarded the 1997 Nobel Prize for this work.117 Prions in animals cause diseases such as bovine spongiform encephalopathy (BSE or mad cow disease) in cattle, scrapie in sheep and goats, and chronic wasting disease of mule deer and elk.118 Human prion diseases include Creutzfeldt–Jakob disease, fatal familial insomnia, Gerstmann–Sträussler–Scheinker syndrome, kuru, and new variant Creutzfeldt–Jakob disease (vCJD or nvCJD). Prion diseases can be sporadic, familial, or iatrogenic/infectious.

Eighty five per cent of prion diseases are sporadic (sCJD), up to 15% are familial (fCJD, Gerstmann–Sträussler–Scheinker syndrome, fatal familial insomnia), and less than 1% are iatrogenic. The incidence of Creutzfeldt–Jakob disease is about one per million per year worldwide.119 Iatrogenic cases of Creutzfeldt–Jakob disease have resulted from insufficient decontamination of surgical instruments, corneal transplants,
dura mater grafts, and human pituitary extract treatment.\textsuperscript{133} Owing to increased awareness, there has been a decline in the iatrogenic transmission of this disease in recent years.\textsuperscript{134} Familial Creutzfeldt–Jakob disease, Gerstmann–Sträussler–Scheinker syndrome, and fatal familial insomnia are caused by gene mutations in the prion protein gene. At least 22 mutations have been identified.\textsuperscript{135} Mutations in the prion protein gene are usually found only in patients from families with clear histories of prion disease, but some mutations seem to be incompletely penetrant and may be found in patients with apparently sporadic illness.\textsuperscript{136}

Prions are infectious proteins that are abnormal isoforms of the normal human protein called PrP. Prions reproduce by transforming the normal form of PrP (PrP\textsuperscript{Sc}) into the abnormal isoform of prion (scrapie PrP or PrP\textsuperscript{C}). PrP\textsuperscript{Sc} has a very different conformation from PrP\textsuperscript{C}. The mechanism of conformational change of PrP\textsuperscript{C} to PrP\textsuperscript{Sc} is not yet known.\textsuperscript{137} In human prion disease large amounts of PrP\textsuperscript{Sc} usually accumulate in the brain. Animal models of infectious prion disease show that neurological dysfunction correlates with levels of PrP\textsuperscript{Sc} accumulation in the brain.\textsuperscript{138}

Both the normal and disease associated PrP isoforms (PrP\textsuperscript{C} and PrP\textsuperscript{Sc}) are encoded by Prnp, located on the short arm of chromosome 20. A polymorphism at codon 129 in Prnp appears to play a significant role in the host susceptibility and phenotypic expression of inherited, iatrogenic, or sporadic Creutzfeldt–Jakob disease. Homozygosity for either methionine or valine at codon 129 results in individuals having a greater susceptibility to developing sporadic or iatrogenic Creutzfeldt–Jakob disease, whereas being heterozygous at this codon seems to be protective. To date, all cases of vCJD have been homozygous for methionine at codon 129, suggesting that methionine homozygosity increases susceptibility to vCJD.\textsuperscript{139} All familial forms of prion disease are associated with a mutation (point, insertion, and stop codon mutations) in the prion protein gene. The particular mutation can drastically influence the clinical, pathological, and biochemical features of prion disease.\textsuperscript{140}

Codon 178 also appears to play a role in phenotypic expression of inherited forms of prion disease. Fatal familial insomnia is seen in persons with the haplotype of an aspartate to asparagine mutation at codon 178 (D178N) and a methionine at the polymorphic codon 129. In contrast if codon 129 is valine (and not methionine), the D178N mutation results in a clinical picture typical of Creutzfeldt–Jakob disease rather than fatal familial insomnia. Gerstmann–Sträussler–Scheinker syndrome is associated both with amino acid substitutions at codons 102, 105, 117, 145, 198, and 217 and with insertion mutations as well in Prnp.\textsuperscript{141}

Classically, all prion diseases have a triad of neuropathological features that include vacuolar (spongiform) change, astrogliosis, and neuronal loss. However, each type of human prion disease has its own distinguishing pathological features. Creutzfeldt–Jakob disease is defined pathologically by diffuse vacuolar changes in the grey matter, gliosis, and neuronal loss with few PrP-amyloid plaques. In Gerstmann–Sträussler–Scheinker syndrome there is much spongiform change but there are extensive PrP-amyloid plaques; neurofibrillary tangles are found in some forms of this condition. Grossly, there may be mild cerebral or cerebellar atrophy. In fatal familial insomnia, there is neuronal loss and gliosis in the thalamus, inferior olives, and to a lesser extent the cerebellum, but minimal if any vacuolation (spongiform change). vCJD has distinct pathology from the other human prion diseases, with diffuse vacuolation and distinctive, dense core, PrP-containing plaques surrounded by a halo of vacuolar (spongiform) change, called florid plaques.\textsuperscript{142} In several human forms of prion disease as well as in animal models there is evidence of significant early loss of specific subpopulation of GABAergic, parvalbumin positive, inhibitory interneurons.\textsuperscript{143–145}

Though Creutzfeldt is credited with the first description of Creutzfeldt–Jakob disease, his diagnosis has been questioned, as spongiform changes were not found retrospectively in his patient. Jakob described five patients several years later in 1921; of the four in whom the pathology was reviewed retrospectively, only one had spongiform changes.\textsuperscript{146} Because of this, some people refer to the disease eponymously as Jakob–Creutzfeldt instead of Creutzfeldt–Jakob disease.\textsuperscript{147}

Creutzfeldt–Jakob disease classically presents as a triad of dementia, myoclonus, and ataxia, usually between the ages of 50 and 70, with a mean age of 60. The median duration of illness is four months and the mean is 7.6 months. Death occurs within 12 months in 85–90% of patients.\textsuperscript{148} Creutzfeldt–Jakob disease affects woman and men equally, with an incidence of about one per million.\textsuperscript{149–151} Patients with Creutzfeldt–Jakob disease usually present with a rapidly progressive dementia, leading to a general decline in overall cognitive function and eventual death. Over the course of the disease, pyramidal and extrapyramidal dysfunction, cerebellar dysfunction, gait, and speech abnormalities can occur. In a minority of patients cortical visual disturbances are present. In approximately one third of patients, vague complaints of fatigue, headache, sleep disturbances, vertigo, malaise, weight loss, poorly defined pain, or behavioural changes may precede the dementia by months or weeks. Also, in about one to two thirds of cases the EEG will eventually show 1–2 Hz triphasic periodic sharp waves.\textsuperscript{144} 145 MR imaging, particularly in sCJD, often shows hyperintensity of the basal ganglia as well as of the cortex on T2, fluid attenuated inversion recovery (FLAIR), and especially on diffusion weighted sequences (DWI). DWI is probably most sensitive for detecting hyperintensity in the affected brain regions, particularly the neocortex, basal ganglia, thalamus, and cerebellum.\textsuperscript{152–159}

The most recently recognised form of Creutzfeldt–Jakob disease, occurring primarily in the United Kingdom, has been termed new variant Creutzfeldt–Jakob disease (nvCJD, variant or vCJD) and is believed to be caused by transmission of mad cow disease (bovine spongiform encephalopathy or BSE) to humans.\textsuperscript{160–162} By 1 October 2001, 109 cases had been identified—two in France and 107 in the United Kingdom.\textsuperscript{163} 164 Variant Creutzfeldt–Jakob disease is distinctly different from sCJD in its clinical presentation and pathology. Clinically it presents as a neuropsychiatric disorder and tends to affect much younger patients, typically young adults and teenagers, with an average age of 29 years.\textsuperscript{165} The youngest patient so far was 12 years old and the oldest 74.\textsuperscript{166} Diagnostic criteria for the vCJD have been divided into possible, probable, and definite cases. Probable cases require a progressive psychiatric disorder of at least six months’ duration without a history of iatrogenic exposure, an EEG not consistent with sCJD, an MRI consistent with the diagnosis, and at least four of the following five clinical symptoms: early psychiatric symptoms, persistent painful or unpleasant dysesthesias, ataxia, dementia, and a movement disorder (chorea, myoclonus, or dystonia). Possible vCJD includes all criteria for probable vCJD, except they do not have an MRI suggestive of vCJD. Definite vCJD requires pathological confirmation (brain or tonsillar tissue).\textsuperscript{167} 168

Median survival is much longer than in sCJD, at about 14.5 months.\textsuperscript{169} On neuropathology, there is often vacuolation, diffuse astrogliosis, and numerous “kuru-type” florid plaques (resembling those found in patients with kuru), which contain a PrP positive staining core surrounded by vacuolar (spongiform) change.

Brain biopsy or necropsy has been the standard for the definitive pathological diagnosis of human prion diseases. However, tonsil biopsy specimens have been shown to be positive for prion protein (PrP\textsuperscript{Sc}) by immunohistochemistry and western blot in vCJD, but not in sCJD; this should result in an easier and less invasive method of making a definitive diagnosis of vCJD.\textsuperscript{170} Proton density, T2, FLAIR, and DWI on MRI
show bilaterally high signals in the pulvinar, giving the so-called “pulvinar sign”, or in the dorsomedial thalamic nuclei, or both, giving the “double hockey stick” sign. Symmetrical high signal changes are also often seen in the striatum. A report by Oppenheim et al that noted a symmetrically high signal on various MRI sequences also identified a true diffusion abnormality in the striatum (not a T2 weighted shine-through effect), but not in the pulvinar. Whether this diffusion abnormality will be found in other vCJD cases has yet to be determined.

Kuru was a disease of the Fore (pronounced Foray) people in Papua New Guinea; kuru means “to shake or tremble” in the Fore language. The tribe’s former practice of ritualistic cannibalism resulted in human to human transmission of prion disease. The disease is also transmissible to primates. Occasionally it presents. The ataxia is often the presenting sign and can occur earlier or late in the disease course. Myoclonus may not be present. The ataxia is often the presenting sign and can occur at an early age, often in the third to fourth decades, but up to the seventh decade. The disease runs a longer course than Creutzfeldt–Jakob disease, with death occurring in two to 10 years, with a mean of about five years. Unlike Creutzfeldt–Jakob disease, periodic synchronous discharges on the EEG are not seen in Gerstmann–Sträussler–Scheinker syndrome. The disease is also transmissible to primates. Occasionally it can present with a clinical syndrome that resembles conventional Creutzfeldt–Jakob disease or even typical Alzheimer’s disease.

Familial fatal insomnia is an inherited form of prion disease characterised by the development of untreatable insomnia, followed by dysautonomia, ataxia, and variable pyramidal and extrapyramidal signs. Cognitive function is often spared until late in the disease course. A rare sporadic form of Creutzfeldt–Jakob disease can mimic the signs and symptoms of familial fatal insomnia.

Gerstmann–Sträussler–Scheinker syndrome is a rare familial variant of Creutzfeldt–Jakob disease characterised by spinocerebellar ataxia, diminished reflexes, and usually dementia. Amyotrophy and parkinsonian signs may appear early or late in the disease course. Myoclonus may not be present. The ataxia is often the presenting sign and can occur at an early age, often in the third to fourth decades, but up to the seventh decade. The disease runs a longer course than Creutzfeldt–Jakob disease, with death occurring in two to 10 years, with a mean of about five years. Unlike Creutzfeldt–Jakob disease, periodic synchronous discharges on the EEG are not seen in Gerstmann–Sträussler–Scheinker syndrome. The disease is also transmissible to primates. Occasionally it can present with a clinical syndrome that resembles conventional Creutzfeldt–Jakob disease or even typical Alzheimer’s disease.

CONCLUSIONS

The four most common presenile neurodegenerative dementias have various features in common. The prominent role of protein aggregates in the pathology of Alzheimer’s disease, frontotemporal dementia, dementia with Lewy bodies, and Creutzfeldt–Jakob disease is the most striking similarity. Treatment strategies for Alzheimer’s disease are focusing on preventing the development or speeding up the clearance of these aggregates. As more is learned about protein aggregation and its downstream effects leading to cell injury and death, one hopes that future treatment strategies will take advantage of any pathogenic overlap and therefore be more broadly applicable across these devastating disorders. At a minimum, one can expect that effective treatments in one of these disorders should guide and facilitate the development of treatment in the others.

The focal brain regions where these proteins accumulate and kill neurones are different for Alzheimer’s disease, frontotemporal dementia, DLB, and Creutzfeldt–Jakob disease, and this anatomical variability leads to distinctive clinical syndromes that can be diagnosed during life. Once the type of dementia is determined, treatment can be initiated based upon the distinctive neurotransmitter deficits associated with each disease. However, neurotransmitter modifying agents only treat symptoms and do little to slow down, and nothing to prevent, the disorder. Yet vigorous efforts are under way to develop treatments that modify the basic causes for Alzheimer’s disease, frontotemporal dementia, DLB, and Creutzfeldt–Jakob disease; effective therapeutic approaches to these conditions will require an understanding of their molecular basis. Similarly, for issues related to nomenclature, a mechanism based approach will some day supplement a syndrome based system. However, changing the names for diseases is a slow and inherently conservative process.

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