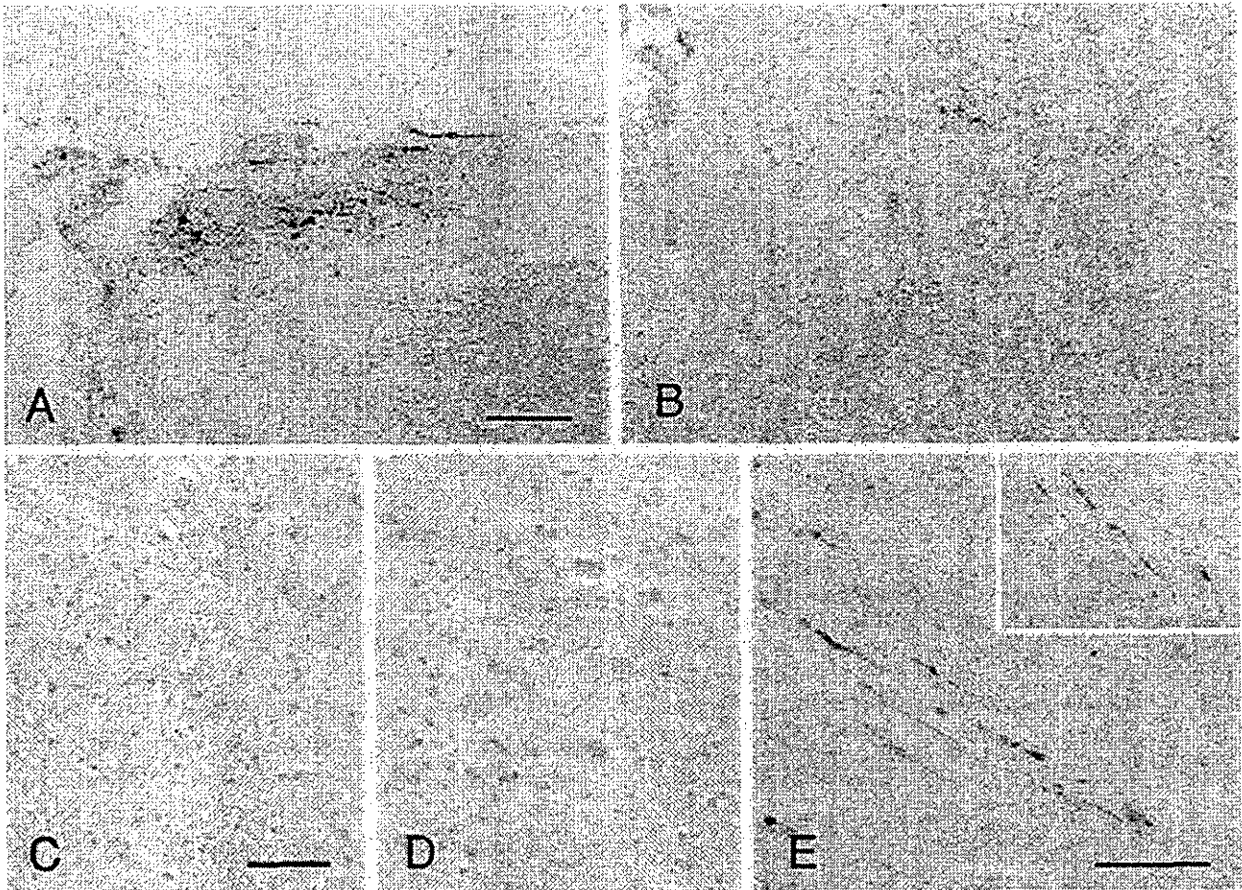

Figure 5. Immunohistochemistry for prion protein (PrP) deposition. A: cerebellum. B: frontal cortex. C: centromedial nucleus of the thalamus. D: inferior olivary nucleus. PrP deposition can not be detected in the thalamus or the inferior olivary nucleus, but coarse or fine granular PrP deposition is visible within the cerebral cortex and the cerebellar molecular layer. E: axons with swollen and/or frizzled features can be detected in the white matter at the border of the thalamus. These axons are also immunostained with anti-APP (amyloid precursor protein) antibody (inset). Bars: 50 μm (A, B, E), 100 μm (C, D).

Although the conventional method of Western blot analysis for PrP^{res} failed to detect any particular signal (data not shown), by application of phosphotungstic acid precipitation that preferably concentrates PrP^{res} but not cellular PrP [12], Western blot analysis of the extract from the frontal cortex of this case revealed a detectable amount of PrP^{res} (Figure 6). The molecular weight of non-glycosylated form of PrP was about 19 kDa (PrP^{res} type 2 pattern) and also the PrP^{res} glycoform ratio was compatible with that of FFI, which has been previously reported [14]. The extract from the cerebellum showed no significant signal in Western blot analysis even with phosphotungstic acid precipitation (data not shown).

The disease of this case was successfully transmitted to some of the mice inoculated with tissue homogenate from the frontal cortex. The incubation time was 571.6 ± 61.1 days (5/7 of the inoculated mice developed TSE) in the NZW mice and 736 ± 64.4 (5/8) in the Tg7 mice, respectively. Although not all the mice developed TSE, diseased mice demonstrated lethargy in the terminal stage rather than excitability. In the TSE-developed mice pathological examination of the brain showed that spongiform change and gliosis were prominent in the cerebral cortex in addition to the thalamus (Figure 7). Immunohistochemistry for PrP revealed that diffuse granular PrP deposition was present within the deep layer of the cerebral cortex as well as in the lateral portion of the thalamus (Figure 7B,E).

Discussion

It is established that there is an overlapping spectrum between classical FFI and CJD in association with PRNP

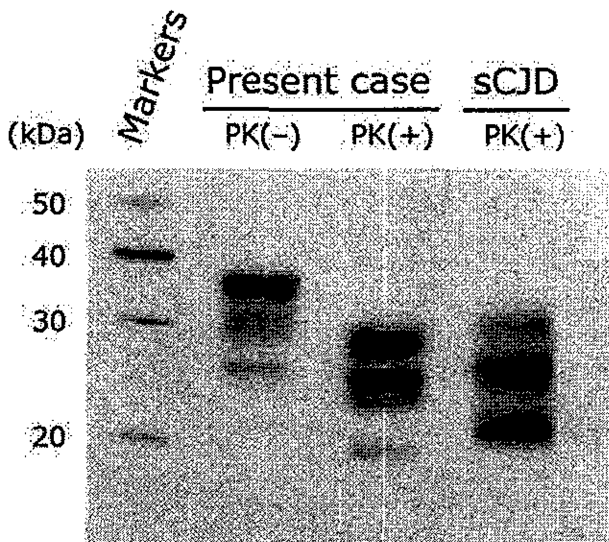

Figure 6. Western blot analysis for protease-resistant prion protein (PrP[res]). Brain homogenate of the frontal cortex of this case is treated with or without proteinase K (PK), and then PK-digested sample is followed by the 40-times concentration with phosphotungstic acid precipitation for PrP[res]. The abnormal PrP molecules in the frontal cortex of this case migrate as PrP[res] type 2. PrP molecules in the lane sporadic Creutzfeldt-Jakob disease (sCJD) are also shown as a standard type 1 PrP (MM1). Molecular sizes (kDa) are indicated on the left.

D178N [6]; however, this case adds to our knowledge about this disease. Although the present case had FFI genotype, the clinical features were initially characterized by prominent cerebellar ataxia, and the neuropathological findings were also atypical in the following respects. First, PrP deposition and spongiform change in the cerebral cortex were more conspicuous than in the thalamus or inferior olivary nucleus, both of which are extremely vulnerable sites for FFI. It has been previously reported that heterozygotes Met/Val at codon 129 result in a longer clinical course than homozygotes [4], and it is therefore possible that the lesions seen in the cerebral cortex were more prominent simply because of the longer course of illness in this patient. However, a further noteworthy point about this case is rather that there was no PrP deposition either in the thalamus or in the inferior olivary nucleus.

Second, immunohistochemical examination detected a peculiar deposition of PrP within the molecular layer of the cerebellum. The localized lesions of granular deposits of PrP and spongiform change in the cerebellar molecular layer seemed to be similar to those reported in a patient from an Austrian FFI family [15]. The cerebellar ataxia of this case could have attributed to the loss of granular neurones and degeneration of Purkinje's cells, in addition to the lesions of the inferior olivary nucleus, although the pathology related to PrP deposition could have also been responsible.

A third atypical feature is that the neuronal loss in the thalamus was most noticeable in the centromedial nucleus. A previous study revealed that severe atrophy of the anterior ventral and dorsomedial thalamic nuclei was consistently observed, whereas that of other thalamic nuclei was less severe and they were inconsistently affected [2]. In this case, the medial portion of the thalamus was indeed damaged crucially, but the principal lesion was different from the typical pathology of FFI.

In addition, an interruption of axonal transport was suggested. Some of the axons were swollen and associated with PrP accumulation, and both APP and SNAP-25 were also accumulated in those axons. APP and SNAP-25 are presynaptic protein and APP is considered as the most effective marker for axonal injury [16]. Aberration in recruitment of PrP might be involved in the pathogenesis of TSE, as described previously [17,18].

This case showed a small amount of specific PrP[res] in the cerebral cortex but not in the cerebellum as detected by Western blotting. The ratio of PrP[res] quantity in those regions was visually correlated with that of immunohistochemical reactivity for PrP. Although fresh frozen samples from the thalamus or the inferior olivary nucleus were not obtained for Western blot analyses, we suspect that PrP[res] in such regions would be too sparse to be detected by Western blotting even in combination with phosphotungstic acid precipitation. The type 2 migration pattern and the glycoform ratio of PrP[res] in this case were compatible with those in the typical FFI [14]; however, it remains to be elucidated whether these abnormal proteins that can be classified in the same PrP[res] type may have different influences on the neurodegeneration processes.

Finally, the transmission study revealed that a pathogen in the frontal cortex of this case might be different from that of an FFI case previously reported by Dr Tateishi and his colleagues [13]. NZW mice infected with a thalamic tissue sample of a typical FFI case exhibited excitability as the principal clinical sign and demonstrated PrP deposition predominantly localized within the thalamus. On the other hand, NZW mice infected with a frontal cortical tissue sample from the present case showed lethargy as a clinical sign, and demonstrated diffuse PrP deposition within the deep layer of the cerebral cortex, as well as in the lateral portion of the thalamus. The PrP deposition

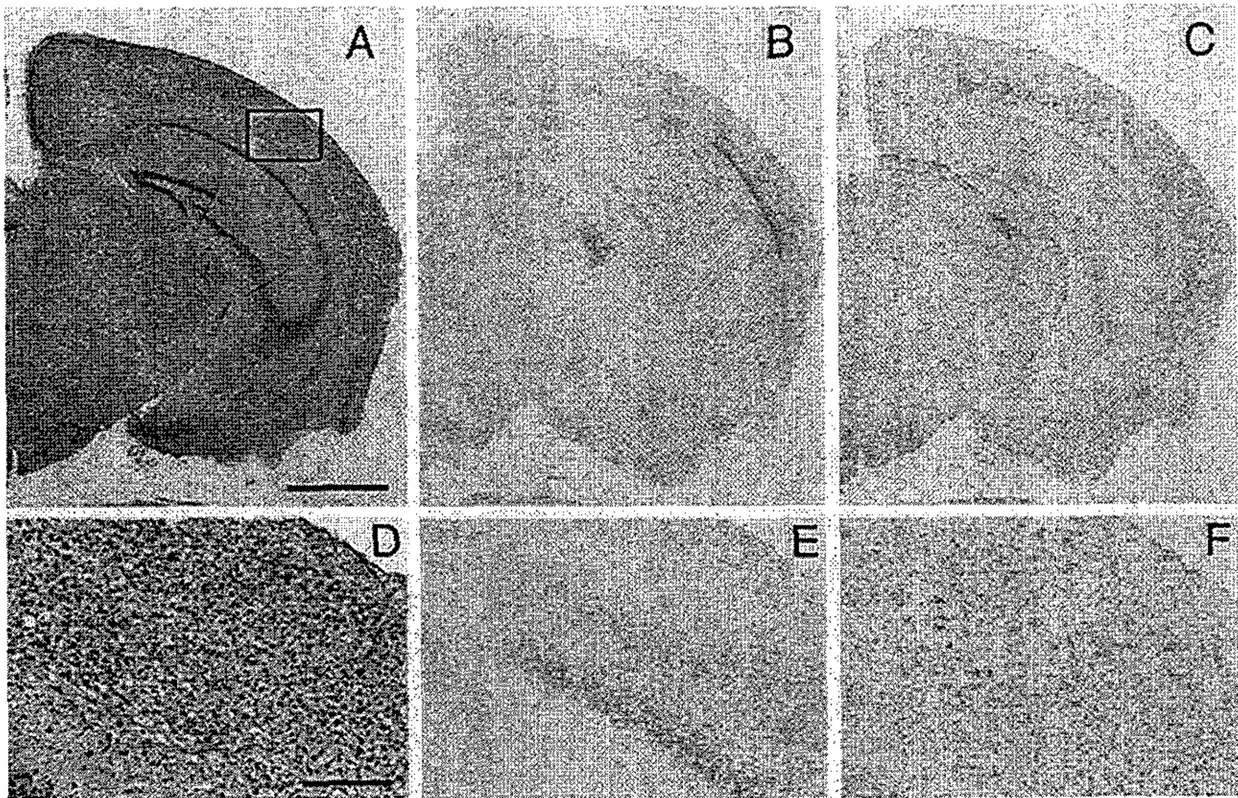

Figure 7. Histological profiles of the mice inoculated with the patient's brain material. A, D: hematoxylin and eosin stain. B, E: prion protein (PrP). C, F: glial fibrillary acidic protein. Spongiform change, PrP deposition and astrocytic gliosis can be observed within the deep layer of the cerebral cortex as well as in the lateral portion of the thalamus. D–F: high power magnifications of the cortical lesions represent the rectangular area depicted in panel A. Bars: 1 mm (A–C), 150 μm (D–E).

pattern of this mouse was distinctive against that of mouse models with other scrapie strains, thus refuting the possibility of contamination. It is not clear whether there were more than two pathogen strains in the brain and whether the strains were dependent on the brain areas. Because we have not examined transmissibility of this case systematically and not obtained frozen materials for Western blot analysis, this aspect still awaits further clarification.

In conclusion, the present case which had FFI genotype showed atypical features, especially with regard to the PrP deposition pattern; there was no deposition within the thalamus or inferior olivary nucleus. Diversity in disease phenotype among patients with the same genotype suggests that some other unidentified factors as well as abnormal PrP deposits or other as yet unknown genetic factors may be responsible for the pathogenesis of the disease. In this study we have shown that variation in pathogen strains may also be one such factor and this factor could have greatly affected the pathogenesis in the present case of FFI.

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Diffusion-weighted MRI abnormalities as an early diagnostic marker for Creutzfeldt–Jakob disease

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Abstract—*Objective:* To evaluate the usefulness of diffusion-weighted MRI (DWI) for the early diagnosis of Creutzfeldt–Jakob disease (CJD). *Methods:* Thirty-six consecutive patients (age 56 to 82 years) were enrolled, and 26 were examined by DWI. Nine were definite based on the World Health Organization criteria, and 27 were probable. The percentages of DWI abnormalities, periodic sharp wave complexes (PSWCs) on the EEG, detection of CSF 14-3-3 protein, and increase of CSF neuron-specific enolase (>25 ng/mL) on the first examination were compared. For DWI, 32 patients (age 31 to 84 years) who showed progressive dementia or impaired consciousness served as disease controls. *Results:* The percentage of DWI abnormalities was 92.3%, of PSWCs 50.0%, of 14-3-3 protein detection 84.0%, and of NSE increase 73.3%. Two of the 32 control subjects were falsely positive on DWI. The sensitivity of DWI was 92.3% (95% CI 74.8 to 99.5%) and specificity 93.8% (95% CI 79.2 to 99.2%). In 17 patients who did not show PSWCs on the first EEG, abnormal DWI findings were still clearly detected. Four patients who were negative for 14-3-3 protein also showed DWI abnormalities. DWI abnormalities were detected as early as at 3 weeks of symptom duration in four patients in whom PSWCs were not yet evident. *Conclusions:* DWI can detect characteristic lesions in the majority of patients with CJD regardless of the presence of PSWCs. DWI was the most sensitive test for the early clinical diagnosis of CJD; consideration should be given to its inclusion in the clinical diagnostic criteria of CJD.

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Creutzfeldt–Jakob disease (CJD) is a transmissible, progressive, fatal spongiform encephalopathy.[1] The transmission of bovine spongiform encephalopathy to humans as variant CJD[2] has focused increased attention on CJD. The cardinal manifestations of the disease are rapidly progressive dementia, generalized myoclonus, and periodic sharp wave complexes (PSWCs) on EEG. However, cases that do not consistently show such typical manifestations have been recognized, and the spectrum of disease manifestations has been extending.[3,4] An early and accurate diagnosis is important to prevent disease transmission, but diagnosis is not easy, especially in the early stage of the disease.

PSWCs on EEG have been used as one of the central diagnostic tests for CJD.[5] However, PSWCs are observed in only 60% of patients[3,4] and usually appear after the middle stage of the disease. In addition, PSWCs are not always specific for CJD.[6] PSWCs are therefore of limited use for the early diagnosis of CJD. The detection of brain-specific proteins such as 14-3-3 protein[6] and neuron-specific enolase[7] (NSE) in CSF also supports the diagnosis of CJD. Although the sensitivity and specificity of 14-3-3 protein[6] and NSE[7] are higher than those of PSWCs,[8] false-positive results are observed in several neurologic diseases such as herpes simplex encephalitis, cerebrovascular disease,[6] Hashimoto encephalopathy,[9] and paraneoplastic neurologic disorders.[10]

Recently, several reports described that diffusion-weighted MRI (DWI) could demonstrate early brain lesions in CJD patients when scans were negative on T2-weighted MRI examination (T2I).[11] In this study, we evaluated the usefulness of DWI for the early clinical diagnosis of CJD by comparing it with other MR sequences such as T2I and fluid-attenuated inversion recovery imaging (FLAIR) and with other diagnos-

See also pages 410, 436, and 450

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Table 1 Profiles of CJD patients and examination results

Patient no.	Type	Age/sex	Duration, wk	DL	*PRNP*	PSWC	DWI	14-3-3	NSE, ng/mL
1	Sp	71/M	3	P	MM	−/+	+		
2	Sp	63/M	4	P	MM	+	+		26
3	Sp	78/M	6	P	MM	+	+	+	79
4	Sp	76/F	6	P	MM	+	+	−	22
5	Sp	61/M	7	P	MM	+	+	+	37.3
6	Sp	66/M	8	P	MM	−/+	+	+	29
7	Sp	76/F	8	P	MM	+	+	+	177
8	Sp	69/M	8	P	MM	+			
9	Sp	54/M	8	P	MM	−/+	+	+	18.4
10	Sp	69/M	8	D	VV2	−	+	+	110
11	Sp	68/M	9	P	MM	−/+	+	+	24
12	Sp	63/F	9	P	MM	+		+	56
13	Sp	74/M	10	D	MM	+			31
14	Sp	71/M	12	P	MM	+		+	25.2
15	Sp	63/F	12	P	MM	−/+	+	+	
16	Sp	79/M	13	D	MM	+		+	36
17	Sp	75/F	21	D	VV2	−	+	+	48
18	Sp	74/F	23	P	MM	+	+	+	51.4
19	Sp	59/M	24	D	MM2-T	−	−	−	15.4
20	Sp	73/F	25	P	MV	−	+	+	56.2
21	Sp	73/F	3	P		−/+	+		
22	Sp	69/M	3	P		−/+	+		50
23	Sp	67/F	6	P		+		+	95
24	Sp	70/M	8	P		+	−/+		
25	Sp	72/F	8	P		+			120
26	Sp	59/M	9	P		+	+		62
27	Sp	74/F	17	D		+		+	72
28	Sp	66/F	25	P		+		+	300
29	Fa	76/M	4	P	V180I	−	+	+	19.5
30	Fa	56/F	8	P	M232R	−/+	+	+	110
31	Fa	58/M	9	P	E200K	−/+	+		
32	Fa	72/M	12	D	V180I	−	+		60.4
33	Fa	82/F	13	P	V180I	−	+	+	32.1
34	Fa	79/M	24	D	V180I	−	+	−	13
35	Ia (Dura)	57/M	3	D	MM	−/+	+	−	18
36	Ia (Dura)	70/F	8	P		+		+	15.8

CJD = Creutzfeldt–Jakob disease; Duration, wk = duration from the onset to diagnostic examinations; DL = diagnostic level based on World Health Organization criteria; PSWC = periodic sharp wave complex; DWI = diffusion-weighted imaging; NSE = neuron-specific enolase; Sp = sporadic CJD; P = probable; MM = homozygosity for methionine at codon 129; D = definite; VV = homozygosity for valine at codon 129; MV = methionine/valine heterozygosity at codon 129; Fa = familial CJD; V180I = a point mutation of Val to Ile at codon 180; M232R = a point mutation of Met to Arg at codon 232; E200K = a point mutation of Glu to Lys at codon 200; Ia = iatrogenic CJD; Dura = a recipient of cadaveric dura mater; (−/+) = negative on the first examination but positive on the sequential examinations.

tic tests including PSWC, CSF 14-3-3 protein, and CSF NSE, which are used as the World Health Organization (WHO) CJD diagnostic criteria.[12]

Patients and methods. *Study group.* Thirty-six consecutive patients with CJD seen from January 1, 1994, to June 30, 2003, at the Department of Neurology, Tohoku University Hospital, and its related hospitals (age 56 to 82 years with a mean age of 68.9 years; 21 men) were enrolled in this study. These patients included the patients in our previous reports.[13-15] According to the WHO criteria,[12] 9 were definite and 27 were probable. A genetic study of human prion protein gene (*PRNP*) was performed in 27 patients, and 20 were sporadic CJD (17 had methionine homozygosity at codon 129 of *PRNP*, and, among those, 3 were definite; 14 were probable at the diagnostic level, and 2 who were definite had valine homozygosity and 1 who was probable had methionine/valine heterozygosity). Six had familial CJD (two definite, four probable, in which two had V180I[15] and one had E200K[16]; one had M232R).[17] Of two patients who were recipients of cadaveric dura mater (iatrogenic CJD), one was definite and one was probable (one had methionine homozygosity at codon 129 of *PRNP*). Our patients composed various clinical phenotypes of CJD including uncommon variants with rather longer clinical courses. The profiles of these patients are listed in table 1.

Disease control group. We reviewed retrospectively the clinical records of our patients who were admitted to the Department of Neurology, Tohoku University Hospital, from January 1, 1998, to June 30, 2003. Excluding patients who had an abrupt onset or symptoms suggesting meningoencephalitis, such as high fever, stiff neck, etc., 81 patients demonstrated subacute dementia or impaired consciousness progressing for 1 to several months.

Table 2 Final diagnosis of control patients

Final diagnosis	Total no.
Metabolic encephalopathy, including one alcoholic encephalopathy	4
Dementia with Lewy bodies	3
Corticobasal ganglionic degeneration	3
Cerebrovascular dementia	2
Viral encephalitis, including one herpes simplex encephalitis	3
Interval form of CO poisoning	3
Cryptococcal meningoencephalitis	2
Mitochondrial cytopathy	2
Progressive dementia, not otherwise specified	4
Alzheimer disease	1
CNS lymphoma	1
Multiple sclerosis	1
Temporal lobe epilepsy	1
Leukoencephalopathy, not otherwise specified	1
Ganser syndrome	1
Total	32

Thirty-two (age 31 to 84 years with a mean age of 61.9 years; 15 men) of 81 patients had undergone DWI examination. They served as disease controls for DWI because obtaining a large group of individuals with suspected CJD but with an alternative diagnosis was difficult. Of these patients, four had seriously suspected CJD, for whom the final diagnosis in two was metabolic encephalopathy that was improved by IV vitamin administration, one was mitochondrial encephalopathy as confirmed by an enzyme assay, and one was corticobasal ganglionic degeneration verified by autopsy. These controls included dementia with Lewy bodies, Alzheimer disease (AD), cerebrovascular disease, CNS infection, metabolic or mitochondrial encephalopathy, CNS malignancy, corticobasal ganglionic degeneration, interval form of CO intoxication, etc. Dementia with Lewy bodies, AD, and cerebrovascular disease are major differential diagnoses of CJD,[18] and CNS infection, encephalopathy, and CNS malignancy sometimes demonstrate positive 14-3-3 protein test in CSF,[19] which is an important diagnostic marker for CJD.[6] The disease controls are listed in table 2.

We compared the sensitivities of the positive results of MR sequences such as DWI, T2I, and FLAIR. We also assessed interobserver agreement. We compared the sensitivities of the positive results of DWI, PSWC, and brain-specific proteins such as 14-3-3 protein and NSE in CSF for making a diagnosis of CJD. PSWC and positive assay of 14-3-3 protein are included in the WHO diagnostic criteria.[12] These examinations were carried out 3 to 25 weeks after the onset with a mean duration of 10.7 weeks after the onset (see table 1).

Methods. DWI technique. Scans were performed on a number of units. A 1.5 or 1.0 T MR unit (Signa Horizon LX, GE Medical Systems, Milwaukee, WI; or Magnetom Vision, Siemens, Erlangen, Germany) was used. DWI was performed in 26 CJD cases with single-shot spin-echo echo-planar imaging. Imaging parameters were as follows: 4,700 to 5,000/93 to 120/1 or 2 (repetition time/effective echo time/no. excitations), 10 to 15 axial sections of 5- or 6-mm section thickness with a 1.5- to 3.0-mm intersection gap, 128 × 128 matrix, 220- or 230-mm field of view, and a diffusion-encoding strength (*b* factor) of 1,000 s/mm^2. In 23 of 26 CJD cases, T2I was performed, and in 17 of 26 CJD cases, FLAIR was performed. DWI, T2I, and FLAIR were performed on the same day using the same MR unit.

MRI investigation. MRI scans were assessed retrospectively as hard copies by two well-experienced neuroradiologists blind to clinical information, who examined each type of sequence separately, without referring to the other MR sequences, independently and individually. We accepted three types of high-intensity lesions as CJD-related lesions on DWI: lesions in the striatum (caudate or putaminal or both), lesions in the thalamus including the pulvinar, and lesions along the cortical ribbon (cerebral or cerebellar). We also accepted the lesions of several types in combination. We accepted not only the symmetric lesions but also asymmetric or unilateral lesions. DWI scans of the disease control group were also assessed retrospectively as hard copies combined with five DWI scans of CJD patients by the same two neuroradiologists, completely blind to clinical information to minimize observer bias.

PSWCs. EEG was recorded using the International 10–20 System. PSWCs were defined as diffuse biphasic or triphasic sharp wave complexes with a duration between 100 and 600 milliseconds and an intercomplex interval between 500 and 2,000 milliseconds.[5]

Brain-specific proteins in CSF. 14-3-3 protein immunoassay in CSF by means of western blotting was performed using a polyclonal antibody to the β isoform of 14-3-3 protein, SC 629 (Santa Cruz Biotechnology, Santa Cruz, CA). The presence of the band against the antibody, SC 629, was investigated. NSE in CSF was measured commercially using an ELISA (SRL Laboratory, Tokyo, Japan), and a value of >25 ng/mL[20] was judged as positive.

Statistical analyses of the diagnostic sensitivities of DWI, PSWCs, and NSE and 14-3-3 protein in CSF, positive rate of DWI, T2I, and FLAIR, and interobserver agreement rate were done using the Fisher exact probability test.

Results. *MRI.* DWI was examined in 26 CJD patients 3 to 25 weeks after the onset with a mean duration of 10.7 weeks. Twenty-four CJD patients showed high-intensity brain lesions by DWI examination. For both observers, the sensitivity of DWI for the CJD diagnosis was 92.3%. The interobserver agreement rate was 100%. Three patients (12.5%) showed lesions only in the caudate heads and putamen, 10 (41.7%) patients showed linear lesions only in the cerebral cortex, and 11 (45.8%) patients showed lesions in both the basal ganglia and the cerebral cortex (figure 1). Among them, only three patients (12.5%) showed lesions in the thalamus. No patients showed high-intensity lesions in the cerebellum. High-intensity lesions on DWI appeared before brain atrophy. The lesions involving the striatum were not always symmetric at the beginning but later became symmetric (figure 2), although symmetric striatal lesions are well known in CJD.[11] In some cases, the high-intensity lesions with sequential DWI did not always progress with the advance of the disease, and the signal intensity sometimes decreased with the disease progression in some lesions. In some cases, the cortical high signal varied in intensity and anatomic distribution (figure 3). In the terminal stage with profound brain atrophy, the high-intensity lesions became unclear. T2I was examined in 23 of 26 DWI-examined patients, but one T2I scan was excluded because of the low quality due to motion artifacts. One observer judged that 11 of 22 patients were positive (50.0%), and another observer judged that 8 were positive (36.4%). The interobserver agreement rate was 68.2%, and it was lower than that of DWI ($p < 0.005$). In both observers, DWI was more sensitive than T2I ($p < 0.005$ for one observer and $p < 0.0005$ for another observer). FLAIR was examined in 17 of 26 patients. One observer judged that 10 of 17 patients were positive (58.8%), and another observer judged that 7 were positive (41.2%). The interobserver agreement rate was 82.4%, and this also was lower than that of DWI ($p < 0.05$). DWI was more sensitive than FLAIR ($p < 0.01$ for one observer and $p < 0.0005$ for another observer). We show in figure 4 an example in which only DWI could detect abnormal high-intensity lesions.

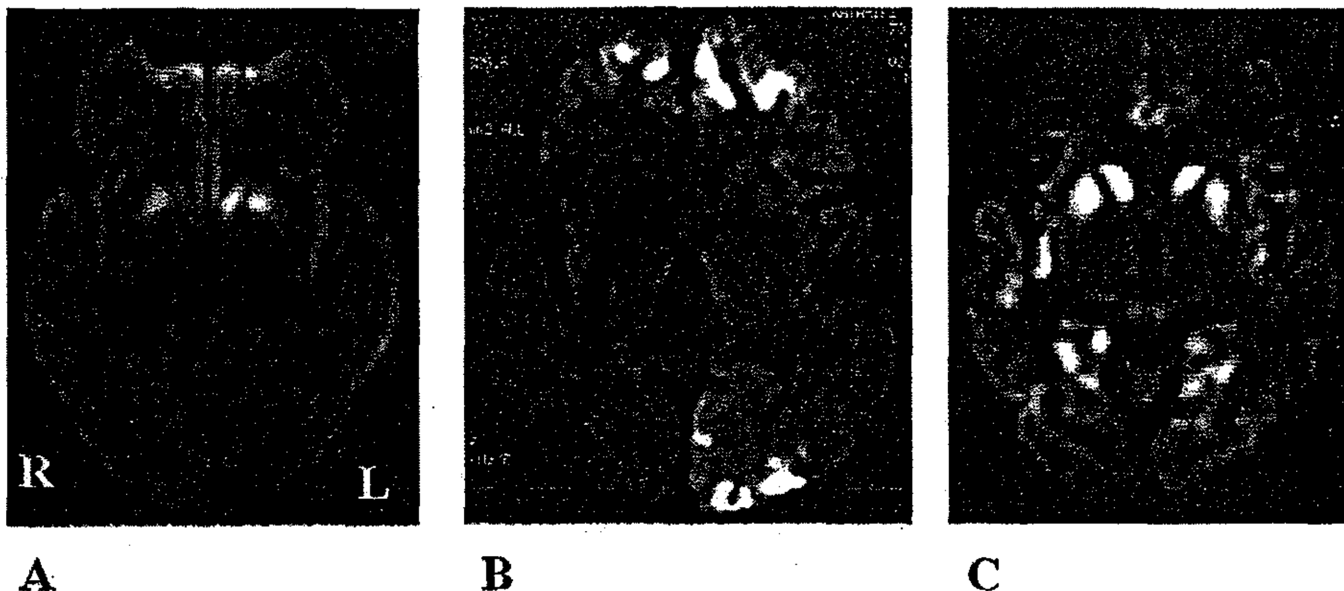

Figure 1. MRI changes seen in Creutzfeldt–Jakob disease. Three patterns of high-intensity lesions were seen: striatal lesion (A), cerebral cortical lesion (B), and a combination of both lesions (C).

DWI failed to detect any lesions in two patients at the first examination. The second DWI showed high-intensity lesions at the striatum in one of those patients. Repeated DWI scans of the other patient did not show any high-intensity lesions throughout his disease course. On the postmortem examination, protease-resistant type 2 prion protein was detected in this patient by western blot analysis using monoclonal antibody 3F4 (Signet Laboratories, Dedham, MA), and there were no spongiform changes. This case was classified as MM2-thalamic according to Parchi's classification (table 1).[3]

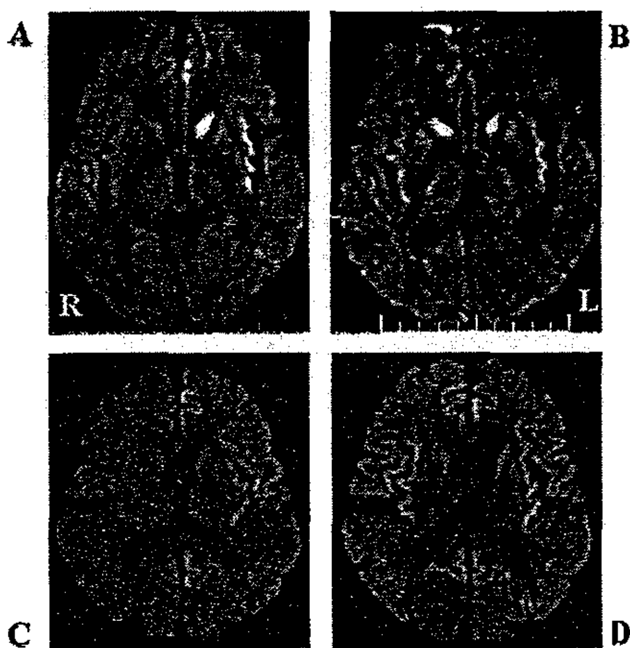

Figure 2. Chronologic change of the striatal and cortical lesions. A case of sporadic Creutzfeldt–Jakob disease (CJD) showing the progression of the basal ganglia signal changes from asymmetric (A) to symmetric (B). The interval between (A) and (B) was 2 months. A case of familial CJD with V180I mutation showing the progression of the cerebral cortex and caudate head signal changes from asymmetric (C) to symmetric (D). The interval between (C) and (D) was 4 months.

High-intensity DWI lesions that were in agreement with our criteria were observed in the disease control patients. One observer judged that a 69-year-old woman with cryptococcal meningoencephalitis and a 60-year-old woman with interval form of CO poisoning were falsely positive. Another observer judged that a 48-year-old woman with herpes simplex encephalitis and a 47-year-old with alcoholic encephalopathy were falsely positive. For both observers, the false-positive rate was 6.3% and the interobserver agreement rate was 87.5%. No highly CJD-suspected patients demonstrated high-intensity lesions. The sensitivity of DWI was 92.3% (95% CI 74.8 to 99.5%) and specificity 93.8% (95% CI 79.2 to 99.2%).

DWI detected the brain lesions before the appearance of PSWCs on EEG in 10 patients. DWI abnormalities were detected as early as at 3 weeks of symptom duration in four patients in whom PSWCs were not yet evident. In seven of eight patients who did not show PSWCs throughout their disease course, the first DWI clearly demonstrated the brain lesions (see table 1).

PSWCs on EEG. EEG was recorded from all 36 patients. Eighteen of 36 (50.0%) patients showed PSWCs that fit the criteria on the first EEG. In 10 of 18 PSWC-negative patients, sequential EEG showed PSWCs. However, eight patients (22.2%) did not show PSWCs in further sequential EEG recordings. The genetic analysis of *PRNP* demonstrated that four had a point mutation of V180I and one had MV at codon 129, and the postmortem examination revealed that two had VV2 and one had MM2-thalamic (see table 1). Generally, these types of CJD patients do not show PSWCs.[3]

Brain-specific proteins in CSF. The 14-3-3 protein was examined in 25 patients and NSE in 30. Twenty-one of those 25 patients (84.0%) were positive for 14-3-3 protein

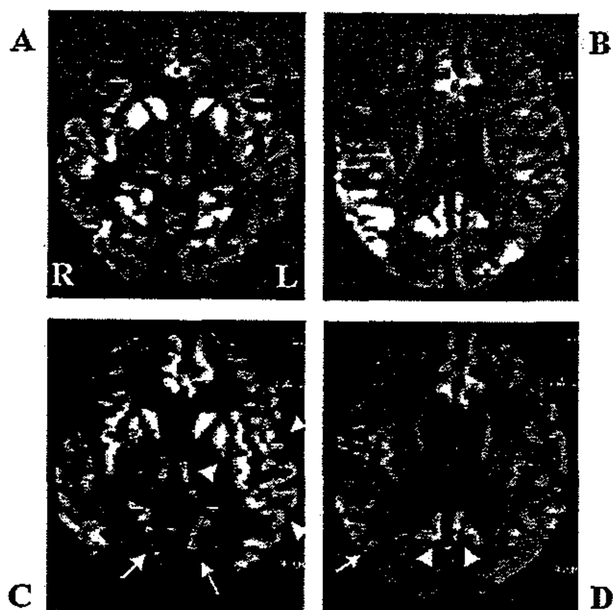

Figure 3. Chronologic change of the cortical lesions in sporadic Creutzfeldt–Jakob disease (sCJD). The cortical high intensity seen in a case of sCJD changed with time, with both increases and decreases in signal intensity in different areas. The high intensity in the bilateral occipital cortices (A) decreased (C, arrows), whereas the signal intensity in the left insular and temporal cortices (A) apparently increased (C, arrowheads). The interval between (A) and (C) was 1 month. The high intensity in the right temporal cortex and bilateral occipital cortices (B) decreased (D, arrow for the left temporal cortex and arrowheads for the bilateral medial occipital cortices). The interval between (B) and (D) was 1 month. Note that the high-intensity lesions depicted in diffusion-weighted imaging did not simply expand with the advance of the disease.

and 22 of those 30 patients (73.3%) were positive for NSE. In 24 patients examined for both brain-specific proteins, 16 were positive for both, 4 were negative for both, and 4 NSE-negative patients were positive for 14-3-3 protein (see table 1).

Comparison of sensitivity of DWI, PSWCs on EEG, and brain-specific CSF protein assay in diagnosing CJD. The sensitivity of DWI examined for the differential diagnosis was 92.3%, of PSWCs 50.0%, of 14-3-3 protein 84.0%, and of NSE 73.3%. DWI was more sensitive than PSWCs ($p < 0.0005$). 14-3-3 protein was more sensitive than PSWCs ($p < 0.01$). DWI tended to be more sensitive than 14-3-3 protein and NSE, but the differences were not significant ($p = 0.36$ and $p = 0.06$) (figure 5). In all 10 patients who had PSWCs in the sequential EEG recording, the lesions had already been detected earlier by DWI. DWI was positive in three of four 14-3-3 protein-negative patients and in six of seven NSE-negative patients. In only one patient who was classified as a rare variant of MM2-thalamic,[3] DWI, PSWC, 14-3-3 protein, and NSE were all negative.

Discussion. MRI had not been thought to be a sensitive noninvasive diagnostic test of CJD[21]; it was previously thought that EEG was the most reliable diagnostic test.[5] Increased signal intensity in the basal ganglia on T2I was first described in 1988,[22] and it was demonstrated that MRI was useful in depicting the lesions of CJD.[23] Although the usefulness of DWI for the early diagnosis of CJD has been suggested,[11,13] no one has compared the ability to depict the lesions among MR sequences such as T2I, FLAIR, and DWI or the accuracy of DWI in diagnosing CJD, especially for the early clinical diagnosis of CJD, with other noninvasive tests including EEG. In this study, we found that the sensitivity of DWI was 92.3% and that DWI was significantly more sensitive than conventional T2I and FLAIR in detecting the CJD-related lesions. T2I and FLAIR, whose sensitivities were 40 to 50%, are inadequate as a test for the first-line differential diagnosis. Further, the CJD-related lesions that we demonstrated on DWI were not detected in a small number of controls with AD, dementia with Lewy bodies, and cerebrovascular dementia, which are the major differential diagnoses of CJD.[18] We have demonstrated the superiority of DWI over the other noninvasive diagnostic tests by comparing its sensitivity with that of PSWCs on EEG, 14-3-3 protein, and NSE examined for the differential diagnosis. Unexpectedly, the interobserver agreement rate of DWI was 100%, and it was significantly higher than that of T2I and FLAIR. This indicates that DWI, which can depict the CJD-related lesions clearly and reliably, may represent a very important diagnostic test for the differential diagnosis. As we demonstrated previously,[13] DWI is more tolerant of motion artifacts than T2I and FLAIR. This advantage is especially important in CJD patients with the involuntary movement of myoclonic jerk. This tolerance may be one of the reasons for the higher sensitivity and the higher interobserver agreement rate of DWI compared with T2I and FLAIR.

The positive rates of T2I by our two observers were 36.4 and 50.0%, and these were significantly lower than a previously reported positive rate for T2I of 79.3%.[23] We think that this discrepancy can be accounted for by the difference in the time when the MRI was examined; the mean duration for our patients from the onset to MRI examination was 10.7 weeks (2.6 months), whereas that of the previous report was 8.1 months.[23] A positive rate for T2I of 67.3% was also reported; however, the positive rate of the first T2I in that report was 43.2%.[24] This is almost the same as in our results.

Currently, PSWCs play a central role in the diagnosis of CJD.[12] However, in the case of PSWC-negative patients, the diagnosis of CJD is sometimes difficult, and such cases are classified as "possible CJD."[12] We have demonstrated in this study not only that DWI was positive earlier than the presence of PSWCs but also that DWI was positive in CJD subjects without PSWCs throughout their disease. In the suspected CJD patients who are diagnosed as "possible CJD," the accuracy of the diagnosis is different between DWI-positive patients and DWI-negative patients. The likelihood of CJD is higher in

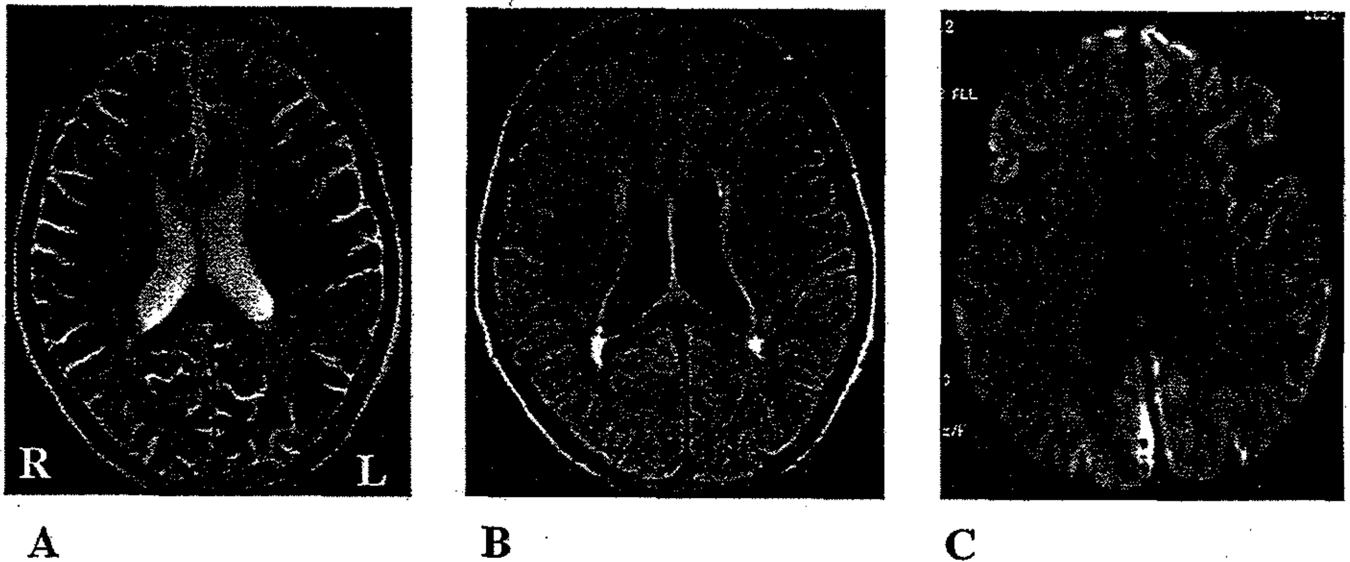

Figure 4. Comparison of conspicuity of Creutzfeldt–Jakob disease–related changes of the same patient on different MRI sequences. T2-weighted imaging (A) and fluid-attenuated inversion recovery imaging (B) show normal findings, and diffusion-weighted MRI (C) demonstrates high-intensity lesions in the cerebral cortex.

DWI-positive patients and lower in DWI-negative patients.

Each observer judged as false positive 2 of 32 disease controls. However, two observers did not agree on the result: one observer judged DWI of cryptococcal meningoencephalitis and interval form of CO poisoning as false positive, and another observer judged DWI of herpes simplex encephalitis and alcoholic encephalopathy as false positive. However, a careful history taking and the presence of pleocytosis in the CSF study significantly reduced the possibility of CJD. DWI was very useful to distinguish CJD from AD, vascular dementia, and dementia with Lewy bodies, which are the major differential diagnoses of CJD[18] and account for the vast majority of dementia in elderly patients.[25] DWI of mitochondrial encephalopathy, lactic acidosis, and stroke-like episodes (MELAS), Wilson disease, and Wernicke encephalopathy can demonstrate similar abnormalities. However, the clinical course and laboratory findings easily distinguish them from CJD. DWI in hypoglycemia,[26] anoxia,[27] and reversible posterior leukoencephalopathy syndrome[28] has also been reported to demonstrate high-intensity lesions similar to those of CJD. However, these have peculiar episodes, and the onset is apparently different from that of CJD. We must mention a 15-year-old boy with a final diagnosis of CNS lupus who was referred to the Japanese CJD Surveillance Committee. His consciousness disturbance developed subacutely, and his DWI showed scattered high-intensity lesions in the cerebral cortex and basal ganglia. His neurologic symptoms improved after the administration of prednisolone. CNS vasculitic disease also needs to be excluded in the differential diagnosis.

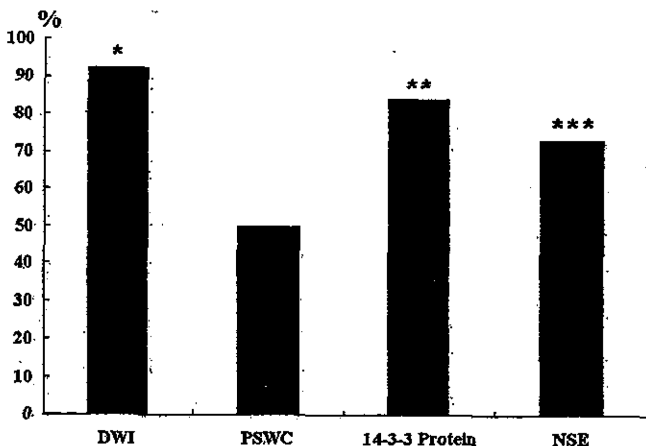

*Figure 5. Percentage of Creutzfeldt–Jakob disease cases with positive test. The positive rates of diffusion-weighted MRI (DWI), periodic sharp wave complexes (PSWCs), 14-3-3 protein, and neuron-specific enolase (NSE) examined for the differential diagnosis were 92.3, 50.0, 84.0, and 73.3%. *DWI was more sensitive than PSWCs ($p < 0.0005$). **14-3-3 protein was more sensitive than PSWCs ($p < 0.01$). *DWI tended to be more sensitive than 14-3-3 protein ($p = 0.36$) and NSE ($p = 0.06$). ***NSE tended to be more sensitive than PSWCs ($p = 0.053$). However, these were not significant.*

The kinds of pathologic findings that correlate with the CJD-related high-intensity lesions demonstrated in DWI are still controversial. Spongiform changes[29] and prion protein deposits[30] are candidates. The time lag from DWI examination to postmortem pathologic examination impedes an accurate understanding. The postmortem examination in one case of familial CJD with V180I whose DWI showed prominent high intensity in the cerebral cortex revealed severe spongiform changes and rather weak prion protein staining there immunohistochemically. The postmortem examination in a patient with spo-

radic CJD with MM2-thalamic whose DWI demonstrated negative findings throughout the disease course revealed no spongiform changes and rather weak prion protein staining. Based on our limited experience, we speculate that the high-intensity lesions depicted by DWI are related to spongiform changes rather than to prion protein deposition; however, more animal and postmortem studies are required to confirm this. It remains unclear why some high-intensity lesions become less prominent with the advance of the disease. We need to accumulate radiopathologic studies for several types of CJD.

The weaknesses of this study are that the number of ideal controls, CJD suspects with a final alternative diagnosis, was too small, because it was difficult to obtain a large number of such patients, and also the lack of pathologic diagnosis in the majority of CJD patients, with the result that only 9 of 36 patients were definite because of the difficulty in obtaining a postmortem examination in many cases. To overcome these weak points, a multicenter analysis of pathologically verified CJD patients and ideal controls is needed.

Last, in interpreting the results of the diagnostic tests, we must understand that such laboratory tests as DWI, brain-specific protein, and EEG reflect different aspects of the disease. Brain-specific proteins reflect the ongoing rapid and massive destruction of the neurons, and EEG reflects the current state of the injured brain.

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Accumulation of prion protein in muscle fibers of experimental chloroquine myopathy: *in vivo* model for deposition of prion protein in non-neuronal tissues

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Prion protein (PrP) is known to accumulate in some non-neuronal tissues under conditions unrelated to prion diseases. The biochemical and biological nature of such accumulated PrP molecules, however, has not been fully evaluated. In this study, we established experimental myopathy in hamsters by long-term administration of chloroquine, and we examined the nature of the PrP molecules that accumulated. PrP accumulation was immunohistochemically demonstrated in autophagic vacuoles in degenerated muscle fibers, and this was accompanied by the accumulation of other molecules related to the neuropathogenesis of prion diseases such as clathrin, cathepsin B, heparan sulfate, and apolipoprotein J. Accumulated PrP molecules were partially insoluble in detergent solution and were slightly less sensitive to proteinase K digestion than normal cellular PrP. Muscle homogenates containing these PrP molecules did not cause disease in inoculated hamsters. The findings indicate that the PrP molecules that accumulated in muscle fibers have distinct biochemical and biological properties. Therefore, experimental chloroquine myopathy is a novel and useful model to investigate the mechanism of deposition of PrP in non-neuronal tissues and might provide new insights in the pathogenesis of prion diseases.
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Keywords: detergent-solubility; experimental chloroquine myopathy; lysosome; non-neuronal tissues; prion protein; protease sensitivity

Prion diseases such as Creutzfeldt–Jakob disease in humans, and scrapie and bovine spongiform encephalopathy in animals are neurodegenerative disorders characterized by the accumulation in the brain of a protease-resistant, detergent-insoluble abnormal isoform of prion protein (PrP). This abnormal isoform of PrP (PrP^{Sc}) is pathogenic itself and replicates by altering the conformation of a protease-sensitive, detergent-soluble normal cellular isoform of prion protein (PrP^{C}).[1] In addition to the central nervous system, PrP^{Sc} deposition is observed in non-neuronal tissue such as tonsils and skeletal muscles in human prion diseases[2,3] and experimental animals.[4]

PrP is also known to accumulate in non-neuronal tissues under certain pathological conditions unrelated to prion diseases. Frederiske *et al*[5] recently revealed increased PrP immunoreactivity in the regions of fiber-cell degeneration in cataractous lenses in humans. Askanas *et al*[6] reported the accumulation of PrP in vacuolated muscle fibers, in angulated and round atrophic fibers with sarcolemmal enhancement, and in the perivascular inflammatory cells of sporadic inclusion-body myositis in humans.[6,7] It was also reported that the accumulated PrP molecules were sensitive to protease treatment.[7] However, the biochemical and biological characteristics of these PrP molecules have not been fully evaluated.

Chloroquine, a widely used antimalarial agent, is known to be concentrated in lysosomes and to cause elevation of intralysosomal pH.[8] Long-term

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administration of chloroquine sometimes causes myopathy, termed chloroquine myopathy (CM), which is characterized by degenerated muscle fibers with numerous autophagic, rimmed vacuoles.[9] Tsuzuki *et al*[10] established experimental CM in the rat to investigate the mechanism of accumulation of the proteins related to Alzheimer's disease in rimmed vacuoles, because of its histopathological similarity to human myopathies where amyloid β deposition is observed in rimmed vacuoles.

To explore the biochemical and biological properties of PrP molecules that accumulate under pathological conditions unrelated to prion diseases, we established experimental CM in hamsters and characterized the PrP molecules (PrP^{CQ}) that accumulated in affected muscle fibers.

Materials and methods

Animals and Reagents

Female Syrian hamsters, 3–8-week old, were purchased from SLC (Hamamatsu, Japan). Chloroquine diphosphate and Nonidet P-40 (NP-40) were purchased from Sigma Chemical (MO, USA). Proteinase K (PK) and complete mini protease inhibitor cocktail were obtained from Roche Molecular Biochemicals (Germany). Monoclonal antibody 3F4 recognizing hamster PrP109-112 was from Senetek (St Louis, MO, USA). Anti-prion protein polyclonal antibody PrP2B was raised by immunization of rabbits with a hamster PrP89-103 fragment. Polyclonal antibodies for apolipoprotein J (clusterin) and for cathepsin B, and monoclonal antibodies CHC5.9 for clathrin and HepSS-1 for heparan sulfate were purchased from Chemicon (Temecula, CA, USA), Calbiochem (Cambridge, MA, USA), PROGEN Biotechnik GmbH (Germany), and Seikagaku Corporation (Japan), respectively.

Experimental CM in Hamsters

Hamsters received 50 mg/kg chloroquine diphosphate (10 mg/ml in sterile saline, pH 7.6) as daily intraperitoneal injections for 60 days. Then the hamsters were killed by decapitation under deep anesthesia. Bilateral soleus, tibialis anterior, and quadriceps muscles were removed and immediately snap-frozen in isopentane cooled with liquid nitrogen. Frozen muscles were kept at −80°C until analysis.

Immunohistochemical Studies

After the blockage of endogenous peroxidase with 0.3% hydrogen peroxide in methanol, serial 10-μm thick sections of frozen muscle were incubated overnight at 4°C with the primary antibodies diluted with 10 mM phosphate-buffered saline (PBS) containing 1% normal hamster serum. The sections were then incubated with horseradish peroxidase-conjugated secondary antibodies for 1 h followed by reactions with 3,3′-diaminobenzidine/H_2O_2 and counterstained with hematoxylin. The serial sections were stained with hematoxylin and eosin (HE), or stained for acid phosphatase or by a modified Gomori-trichrome method.

Brain specimens obtained in some experiments were immersion-fixed in 10% buffered formalin for 24 h at 4°C and embedded in paraffin for immunohistochemical examination. For detection of abnormal PrP deposition, deparaffinized 8-μm thick sections were treated with hydrolytic autoclaving prior to incubation with 3F4 monoclonal antibody.[11]

Protease Sensitivity Assay of PrP^{CQ}

After confirmation of the histological findings, the remaining frozen muscle was homogenized using Tissue-Tearor (Biospec Products, Oklahoma) in 10 volumes of lysis buffer A (0.5% NP-40, 0.5% sodium deoxycholate in PBS pH 7.4). Homogenates were centrifuged at 3300 $\times g$ for 15 min to remove the nuclear fraction and debris. The supernatant was then treated with the indicated amount of PK at 37°C for 20 min. After stopping the digestion with 4 mM 4-[2-aminoethyl]-benzenesulfonyl fluoride (Pefabloc, Roche, Germany), an aliquot corresponding to 4 mg of muscle tissue was analyzed by Western blotting using polyclonal antibody PrP2B. Labeled PrP was visualized by using CDP-star detection reagent (Amersham, UK).

Detergent Solubility Assay of PrP^{CQ}

The detergent solubility of PrP^{CQ} was determined as described by Lehmann and Harris[12] with minor modification. Briefly, soleus muscle samples from either control or CM hamsters were homogenized in lysis buffer B (15 mM NaCl, 50 mM Tris-HCl pH 7.5, complete-mini protease inhibitor cocktail) containing the designated concentration of NP-40. Homogenate was centrifuged for 5 min at 1600 *g* to remove debris and the nuclear fraction. The supernatant was ultracentrifuged at 265 000 *g* for 40 min at 25°C. Proteins in the supernatant and in the pellet were separately recovered and analyzed by Western blotting using monoclonal antibody 3F4.

Intracerebral Inoculation of PrP^{CQ}

Inoculum was prepared by homogenizing muscular tissue from CM hamsters or the control in 10 volumes of sterile saline, and 20 μl of the inoculum was injected into the brain of 19 3-week-old female hamsters under deep anesthesia. The hamsters were observed for over 2 years and killed to examine PrP molecules in the brain immunohistochemically.

Densitometry and Statistical Analysis

Blots of the gels were scanned with a CanoScan D2400UF (Canon, Japan). Densities of bands were quantified using NIH Image software. Statistical significance of densitometric data was analyzed by repeated measure ANOVA, and statistical comparison at each dose point between groups was made by Student's t-test or Welch's t-test.

Ethics

Animal handling and killing were in accordance with the nationally prescribed guidelines, with ethical approval for the study granted by the Animal Experiment Committee of Kyushu University.

Results

Histochemical Findings of CM

All the muscle specimens taken from chloroquine-treated hamsters showed various degrees of myopathic changes accompanied with rimmed vacuoles (Figure 1a), and this was consistent with histopathological findings of experimental CM in the rat.[9,10] In addition to the rimmed vacuoles, many muscle fibers contained coarse granular structures that were strongly stained by hematoxylin (Figure 1c) and modified Gomori trichrome stain (Figure 1g), and they showed enhanced acid phosphatase activity (Figure 1h). These abnormal structures were observed in about 10% of muscle fibers in the soleus muscle which was the most affected in all the muscles of chloroquine-treated hamsters. In contrast, muscles of control hamsters did not contain any rimmed vacuoles or coarse granular structures.

All of the rimmed vacuoles and the coarsely granular structures in degenerated muscle fibers of CM were positively stained by an anti-PrP monoclonal antibody, 3F4, recognizing hamster PrP109-112 (Figure 1b, d). In control hamster muscles, the 3F4 monoclonal antibody reacted with sarcolemmal membranes (data not shown).

To investigate the possible involvement of some molecules that relate to the metabolism of PrP^{C} or the deposition of PrP^{Sc}, serial sections were immunostained for apolipoprotein J (Figure 1e), clathrin (Figure 1f), cathepsin B (Figure 1i), and heparan sulfate (Figure 1j). Immunoreactivities for these molecules were enhanced in the rimmed vacuoles and coarse granular structures in the affected muscle fibers of CM.

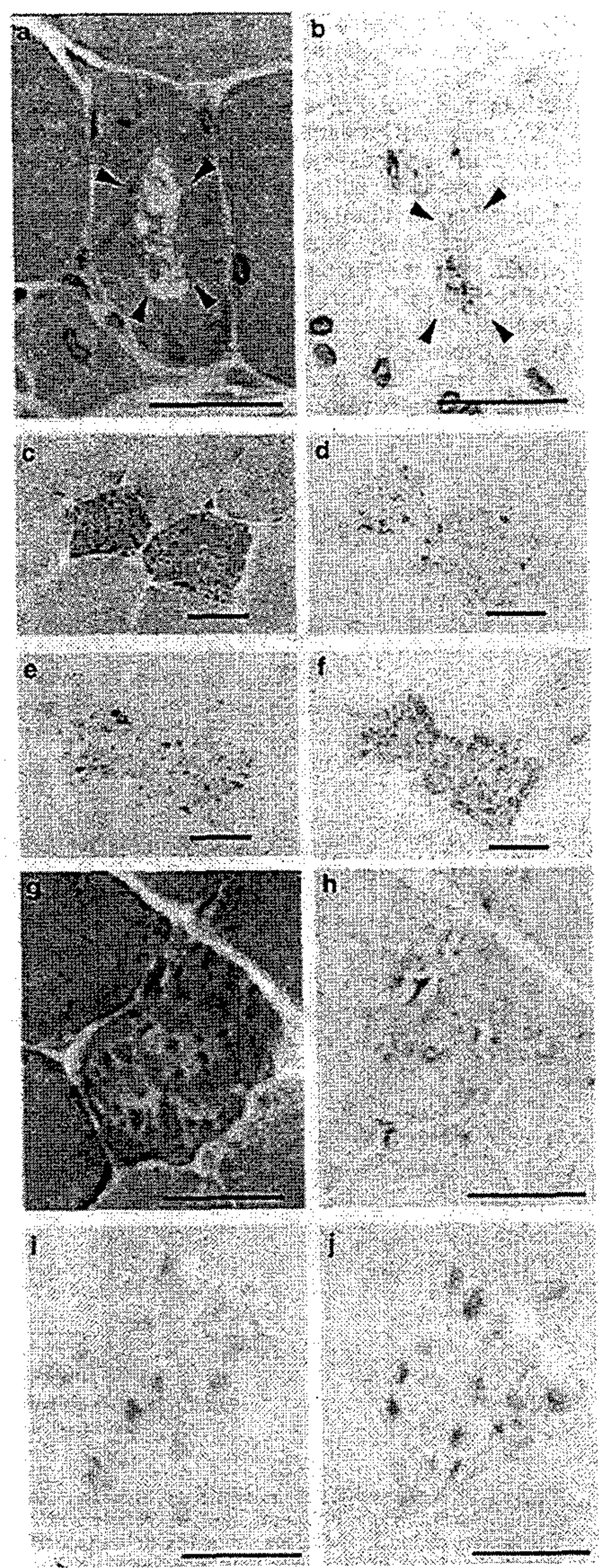

Figure 1 Immunohistochemical findings in CM muscles. Transverse sections of chloroquine-treated (50 mg/kg/day for 60 days) hamster soleus muscles are shown. (a) HE stain shows rimmed vacuole formation (arrow heads). (b) Immunoreactivity for PrP is detected in a rimmed vacuole (arrow heads) shown in (a). (c–f), (g–j) Serial sections stained with HE (c), modified Gomori trichrome (g), acid phosphatase (h), and immunostained for PrP (d), apolipoprotein J (e), clathrin (f), cathepsin B (i), and heparan sulfate (j). Scale bars = 20 μm.

PrP^{CQ} is Slightly Less Sensitive to PK Digestion

Western blot analysis using a polyclonal antibody PrP2B, raised against hamster PrP89-103, revealed prominent bands of 27 and 30 kDa in either CM muscular homogenates or the control, before digestion with PK. There was no significant difference in the intensity of PrP signals between CM muscles and controls (first lanes, Figure 2a, b). The specificity of these bands was confirmed by absorbing the PrP2B antibody with synthetic peptide PrP89-103 (Figure 2c). Although both of the PrP molecules were completely digested with 50 μg/ml of PK, which was the stringent condition to distinguish PrP^{Sc} from PrP^{C} (data not shown), digestion with a smaller amount of PK revealed different PK sensitivity between the PrP molecules from CM muscles (PrP^{CQ}) and PrP^{C} from the control muscles. PrP^{C} derived from control hamster muscle was apparently digested with 0.375 μg/ml of PK, whereas a considerable amount of PrP^{CQ} of 27 kDa still remained after treatment with 1.0 μg/ml of PK (Figure 2a and b). Statistical analysis of the relative density of the bands revealed a significant difference between PrP^{C} and PrP^{CQ} after treatment with 0.5, 0.75, or 1.0 μg/ml of PK (Figure 2d).

PrP^{CQ} is Partially Insoluble in Detergent

Western blot analysis using a monoclonal antibody 3F4 revealed prominent signals at 35 kDa and additional signals at about 30 kDa in the supernatants from either CM muscle homogenate or the control (first lanes, Figure 3a, b). While PrP^{C} from control muscle was completely solubilized in the lysis buffer containing 0.5% NP-40 (Figure 3a), a considerable amount of PrP^{CQ} remained in the

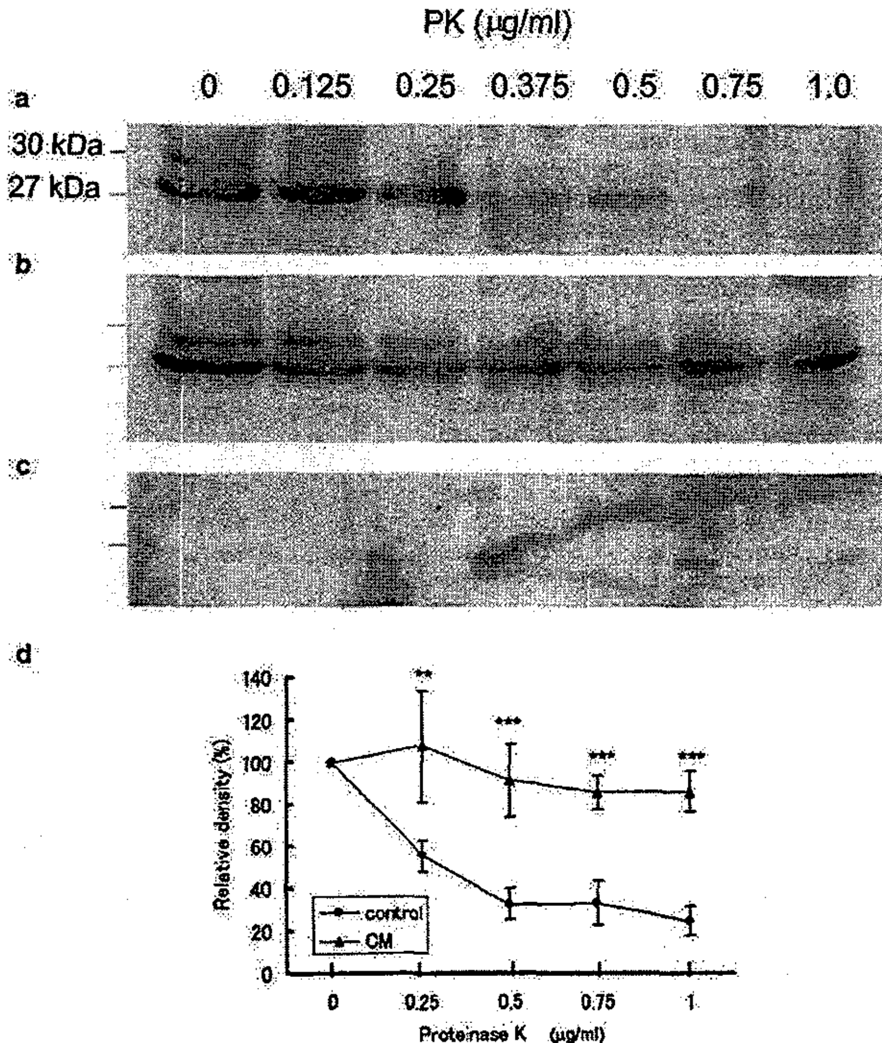

Figure 2 PK sensitivity of PrP molecules in CM muscles. (a–c) PrP molecules in the homogenate of control hamsters (a) or CM hamsters (b) were detected with PrP2B antibody after digestion with a designated amounts of PK at 37°C for 20 min. Prominent bands of 27 kDa and 30 kDa were diminished after the antibody had been absorbed by a synthetic polypeptide used for immunization (c). (d) Densitometric analysis of 27 and 30 kDa bands. Data from three independent experiments are indicated. $^{**}P<0.05$, $^{***}P<0.01$.

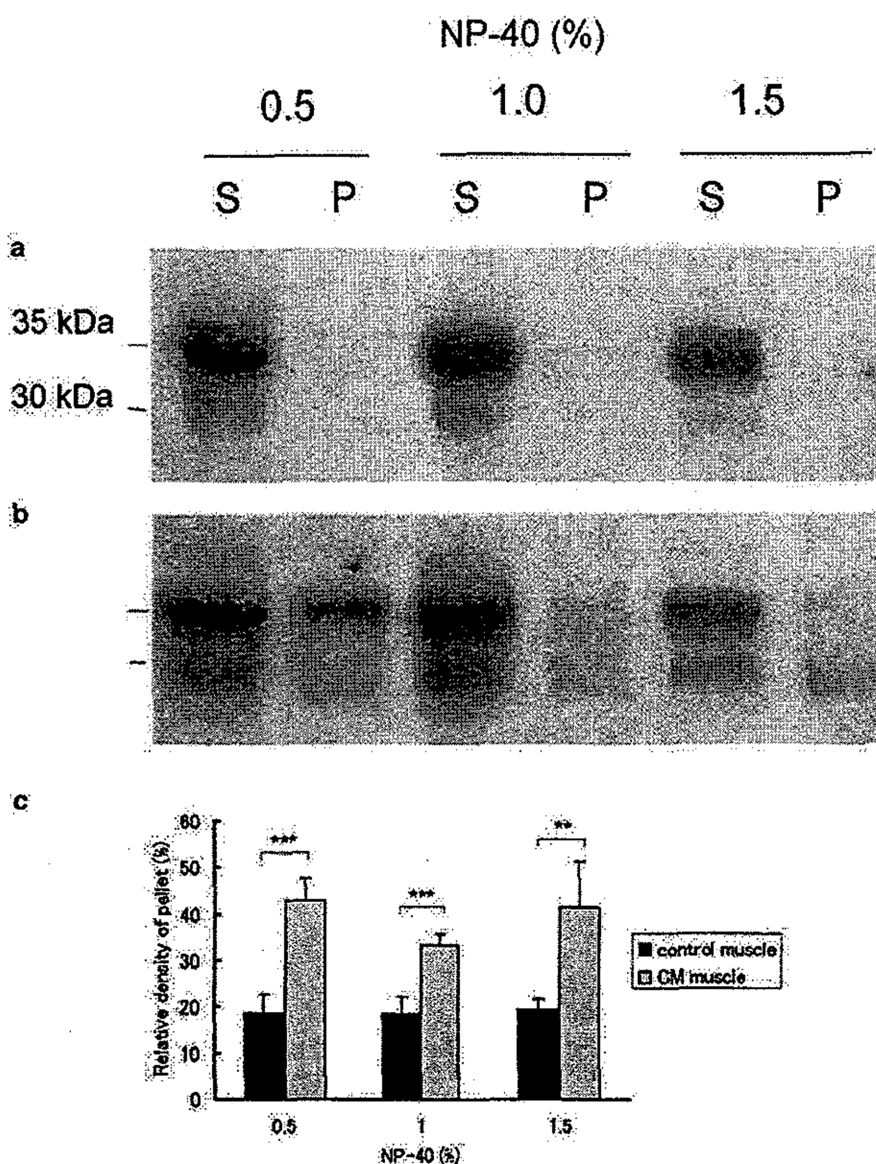

Figure 3 Solubility in NP-40 of PrP molecules in CM muscles. (a, b) Muscles from control hamsters (a) or CM hamsters (b) were homogenized in the lysis buffer containing designated amount of NP-40 and subsequently ultracentrifuged at 265 000 *g* to separate detergent-soluble PrP molecules in the supernatant (S) and detergent-insoluble molecules in the pellet (P). PrP molecules were labeled with 3F4 monoclonal antibody. (c) Relative PrP amount in the pellet fraction. Percentage of PrP signals (30 and 35 kDa) of the pellet fraction in the sum of those of the pellet and the supernatant is shown. Data from three independent experiments. $^{**}P<0.05$, $^{***}P<0.01$.

insoluble fraction in the presence of 0.5, 1.0, or 1.5% NP-40 (Figure 3b, c).

PrP^{CQ} is not Pathogenic

To investigate whether PrP^{CQ} is able to cause pathological changes characteristic of prion diseases *in vivo*, 10% muscular homogenates containing PrP^{CQ} were injected into the brain of Syrian hamsters. The hamsters were observed over 2 years after the inoculation had been given, and none of them developed any signs of prion diseases nor muscle disorders (data not shown). Histological examination of the brain revealed no significant pathological findings or abnormal PrP deposition (Figure 4).

Discussion

In this study, we have demonstrated that slightly less PK-sensitive and partially detergent-insoluble PrP^{CQ} accumulated in affected muscle fibers of experimental CM in hamsters. While PrP molecules from control muscle were sensitive to PK digestion at 0.375 μg/ml and fully soluble in buffer containing 0.5% NP-40, PrP^{CQ} molecules were less sensitive to PK digestion up to 1.0 μg/ml, and a considerable portion of them was insoluble in buffer containing

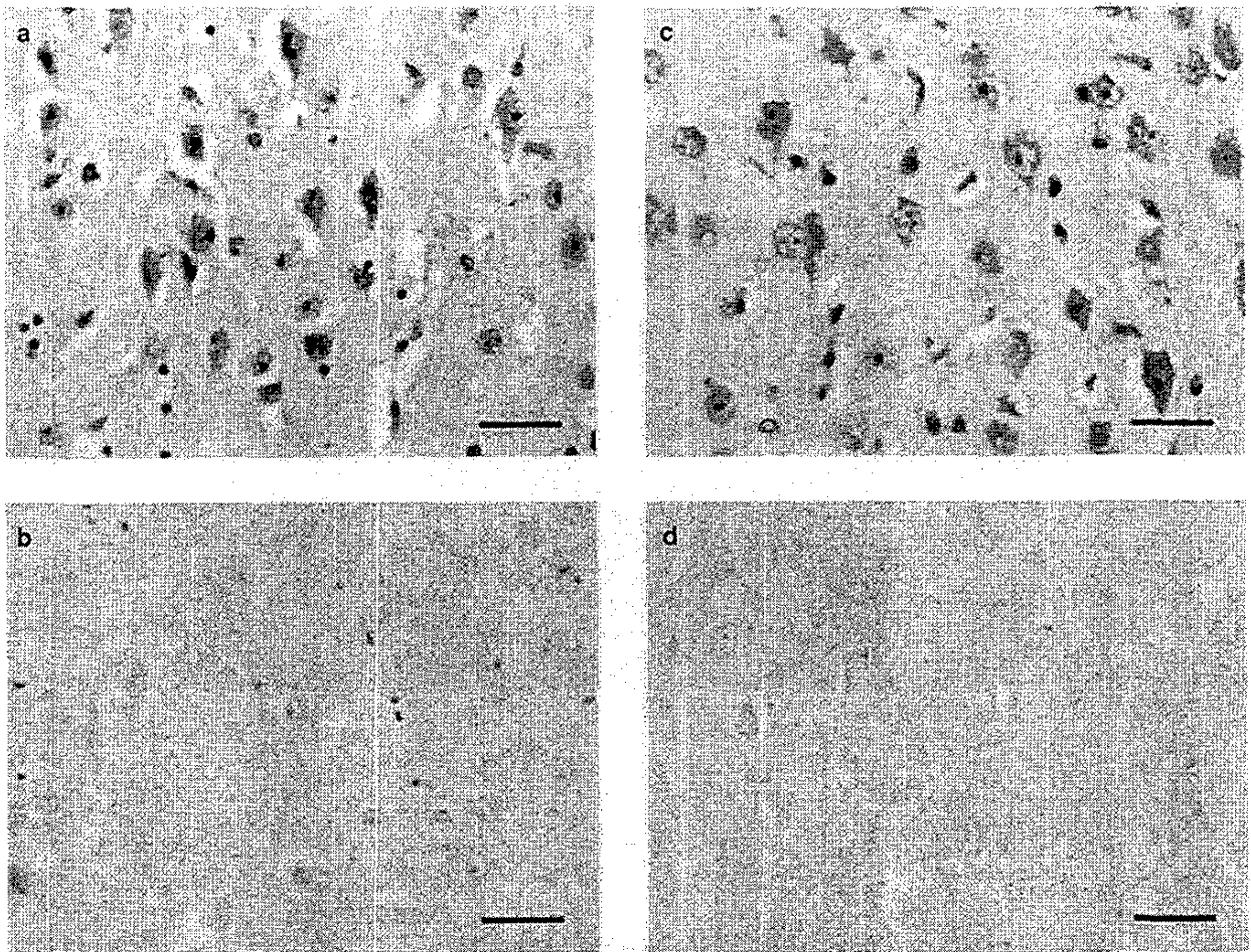

Figure 4 Histological analysis of hamster brain inoculated with CM muscle homogenate. Sections of the brain inoculated with PrP^C-containing muscle homogenate (**a, b**) or PrP^{CQ}-containing muscle homogenate (**c, d**) were stained with HE (**a, c**) or immunostained using 3F4 monoclonal antibody (**b, d**). Neither neurodegenerative change nor abnormal PrP deposition was revealed. Scale bars = 30 μm.

1.5% NP-40. These biochemical properties of PrP^{CQ} molecules are distinct from PrP^C.

There have been several attempts to create *de novo* PrP^{Sc}-like molecules *in vitro*. Using a cell-free conversion system, Horiuchi *et al*[13] were able to convert PrP^C into a protease-resistant isoform by the addition of PrP^{Sc} under physiological conditions. Even in the absence of PrP^{Sc}, an acidic buffer can give a β-sheet-dominant conformation and PK resistance to PrP^C *in vitro*.[14,15] Nevertheless, no *de novo* PrP^{Sc}-like molecules succeeded to reproduce the disease *in vivo*.[16] In the present study, PrP^{CQ} molecules were not capable of causing any pathological changes in the central nervous system. PrP^{CQ} molecules seem to be distinct from both of these *de novo* PrP^{Sc}-like molecules and PrP^{Sc}, because PrP^{CQ} molecules possess neither marked resistance to PK and detergent insolubility nor transmissibility of the disease conditions.

The CM model reported in this study is different from other PrP-conversion models previously reported, in that PrP^{CQ} molecule was generated *in vivo* in the absence of exogenous input of PrP^{Sc} or PrP^{Sc}-like molecules. Instead, the microenvironment in the lysosome was altered in hamsters by the injection of chloroquine to produce this distinct PrP molecule. Lysosomes are acidic compartments that have been reported to play an important role in the conformational conversion of PrP in prion diseases.[17,18] Chloroquine raises intralysosomal pH to as high as 6.0–6.5, causing marked changes in intracellular protein processing and trafficking.[8] As a consequence of the long-term administration of chloroquine, skeletal muscle fibers degenerate, with numerous autophagic vacuoles.[9] In the process of forming autophagic vacuoles, endogenous muscular PrP^C could acquire the properties of PrP^{CQ}.

In the present study, it remains unclear whether chloroquine modifies PrP^C molecules directly or indirectly. A previous study revealed that chloroquine does not directly interact with PrP molecules in scrapie-infected neuroblastoma cells.[19]

Furthermore, it has been reported that unfolded recombinant human prion protein PrP90-231 forms a stable protein folding intermediate rich in β-sheet at pH lower than 4.[14] Matsunaga *et al*[20] reported that pH is a crucial factor in determining the conformational state of some amyloidgenic proteins. They found that synthetic Aβ42 and stefin B peptides, showing similar amino-acid alignment to PrP90-144, tend to form amyloid fibrils at acidic pH. Considering these observations, it is unlikely that chloroquine directly interacts with PrP^{C} molecules. An increase in lysosomal pH due to chloroquine, and subsequent metabolic changes in lysosomal systems might be responsible for the biosynthesis of PrP^{CQ} molecules.

Besides experimental CM, there are a few experimental models in which PrP molecules of skeletal muscle are rendered partially PK-resistant and detergent-insoluble. Chiesa *et al*[21] established transgenic mice expressing PrP molecules with nine-octapeptide insertional mutation. Mutated PrP molecules obtained PrP^{Sc}-like properties in the brain and the periphery, producing neurodegeneration similar to an inherited prion disease in humans. In their model, the primary structure of PrP molecules was changed and the mutated PrP was overexpressed not only in the brain but also in the skeletal muscle and heart. This model is quite different from our CM model, in that the primary structure of PrP molecules was not manipulated.

The other experimental model is a transgenic mouse harboring high copy numbers of wild-type PrP transgenes, which spontaneously exhibited necrotizing myopathy, demyelinating polyneuropathy, and focal vacuolation of the central nervous system without apparent deposition of PrP^{Sc}.[22] In spite of severe neurodegeneration and neuromyopathy, only small amount of PK-resistant PrP was detected in affected muscles and brains. They concluded that low level of PK-resistant PrP might reflect aggregation of PrP^{C} and was not correlated with neuropathological changes in these transgenic mice. In our study, PrP^{CQ} after PK digestion did not show molecular characteristics of PrP^{Sc} in prion diseases, suggesting that PrP^{CQ} acquires less PK sensitivity through a different mechanism from that of PrP^{Sc}. Although the expression level of PrP was not increased in CM (data not shown), it is possible that distinct biochemical properties of PrP^{CQ} might simply be due to protein aggregation or alteration in PK–protein ratio, not to the conformational change of monomeric PrP^{C} molecules.

PrP^{CQ} in the affected muscles of the present model was accompanied by the accumulation not only of lysosomal markers but also of those molecules known to be involved in prion disease pathogenesis, such as clathrin, heparan sulfate proteoglycan, and apolipoprotein J.[23–25] It is known that certain sulfated glycans, such as heparan sulfate and pentosan polysulfate, stimulate PrP conversion *in vitro*.[26] Then, it might be possible that accumulated heparan sulfate in the CM muscles contribute to the acquisition of altered PK sensitivity and partial detergent insolubility of the PrP molecules.

Experimental CM in the rat has been established previously as a model of myopathies with rimmed vacuoles, including distal myopathy with rimmed vacuole formation and inclusion-body myositis. Owing to of the deposition of amyloid β in inclusion-body myositis,[27] experimental CM has been utilized by several groups as a peripheral model to investigate the pathogenesis of Alzheimer's disease.[10,28] The precise mechanism of rimmed vacuole formation in CM is still unknown; however, it has been reported that chloroquine causes an increase in endogenous autophagosomes in mammalian cells.[29] Similar mechanism(s) might be shared between amyloid β deposition in the CM rat model and PrP^{CQ} accumulation in our CM hamster model.

The PrP2B polyclonal antibody revealed prominent 27 kDa signals with additional 30 kDa signals (Figure 2), while the 3F4 monoclonal antibody reacted with dominant 35 kDa signals and additional 30 kDa signals (Figure 3), which were similar to Cp33-37 signal in skeletal muscle of hamster.[30] The common signals of 30 kDa were detected by both of the two antibodies. Minor epitope differences between the two antibodies might account for such a diversity of PrP signals, but it remains to be elucidated.

Finally, together with the biochemical and biological properties of PrP^{CQ}, the immunohistochemical findings in CM muscles of the molecules known to be involved in prion disease pathogenesis indicate that experimental CM in hamsters is a useful *in vivo* model to investigate the mechanism of PrP accumulation in the pathogenesis of PrP-related diseases.

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Treatment of Transmissible Spongiform Encephalopathy by Intraventricular Drug Infusion in Animal Models

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The therapeutic efficacy of direct drug infusion into the brain, the target organ of transmissible spongiform encephalopathies, was assessed in transgenic mice intracerebrally infected with 263K scrapie agent. Pentosan polysulfate (PPS) gave the most dramatic prolongation of the incubation period, and amphotericin B had intermediate effects, but antimalarial drugs such as quinacrine gave no significant prolongation. Treatment with the highest dose of PPS at an early or late stage of the infection prolonged the incubation time by 2.4 or 1.7 times that of the control mice, respectively. PPS infusion decreased not only abnormal prion protein deposition but also neurodegenerative changes and infectivity. These alterations were observed within the brain hemisphere fitted with an intraventricular infusion cannula but not within the contralateral hemisphere, even at the terminal disease stage long after the infusion had ended. Therapeutic effects of PPS were also demonstrated in mice infected with either RML agent or Fukuoka-1 agent. However, at doses higher than that providing the maximal effects, intraventricular PPS infusion caused adverse effects such as hematoma formation in the experimental animals. These findings indicate that intraventricular PPS infusion might be useful for the treatment of transmissible spongiform encephalopathies in humans, providing that the therapeutic dosage is carefully evaluated.

Transmissible spongiform encephalopathies (TSEs), or prion diseases, are fatal neurodegenerative disorders that include Creuzfeldt-Jakob disease (CJD) and Gerstmann-Sträussler-Scheinker disease in humans, in addition to scrapie and bovine spongiform encephalopathy in animals. These disorders are characterized by deposition in the brain of a protease-resistant isoform of prion protein (PrP), which is thought to be the main pathogenic component responsible for the pathogenesis (18). Outbreaks of acquired forms of human TSEs, such as variant CJD (21) and iatrogenic CJD with cadaveric growth hormone or dura grafts (3) in younger people, are prompting the development of prophylactic and therapeutic interventions.

There are some chemicals that are effective in inhibiting the accumulation or conformational change of PrP molecules in vitro and/or in prolonging the incubation period when they are administered around the time of infection in TSE animal models (2, 17). However, no chemicals, except for amphotericin B and one of its derivatives, have been reported to improve prognosis when they are administered late in the disease course or after the infectious agent has already invaded the brain (8). This implies that it may be difficult to improve the prognosis in human patients by using chemicals, because patients first come to medical attention after the onset of neurological symptoms.

Most of the previously reported chemicals have been large and hydrophilic, and whether their ineffectiveness was actually due to their poor accessibility to the brain has not been evaluated. We have developed a more sensitive drug evaluation system, which does not depend on drug accessibility to the brain, by implanting a continuous intraventricular drug infusion device into an intracerebrally TSE-infected animal model. Using this model, we have examined clinically applicable chemicals that have previously been reported to be effective either in vitro or in vivo, including antimalarial drugs such as quinacrine and chloroquine, the E-64d cysteine protease inhibitor, amphotericin B, and pentosan polysulfate (PPS). We report that intraventricular administration of PPS through the infusion device inhibited not only abnormal PrP accumulation but also neurodegenerative changes and infectivity in the brain, thereby giving rise to a dramatic prolongation of the life spans of intracerebrally infected animals.

MATERIALS AND METHODS

Experimental animals and in vivo evaluation. Tg7 mice, which are derived from Tg10 mice (19) and express hamster PrP but not endogenous mouse PrP, were inoculated with 20 μl of 1% 263K agent hamster homogenate in the right parietal portion of the brain. An Alzet osmotic pump (Durect, Cupertino, Calif.) filled with a chemical was placed in a subcutaneous area of the back. An intraventricular infusion cannula connected to the osmotic pump through a catheter was implanted in the left frontal portion of the brain in order to place the cannula tip into the left ventricle at either day 10 or 35 postinoculation or at another designated time. The pump worked stably from 40 h after implantation and continuously for 4 weeks. Five to 10 male mice per group, each weighing about 35 g, were used. Each mouse was kept in an individual cage under the same feeding and watering conditions in an air-conditioned, light time-controlled, specific-pathogen-free room. The mice began to exhibit ambiguous signs of reduced activity about 2 days prior to death, followed by obvious signs of ruffled

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