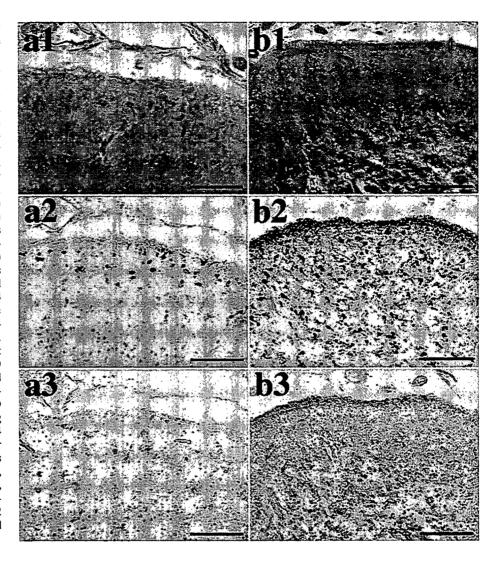
Fig. 7 Two sets of serial sections show the relation between neuropathologic findings, glial fibrillary acidic protein (GFAP) immunoreactivity aquaporin-4 (AQP-4) immunoreactivity. (a1) In subacute spongiform encephalopathy (SSE), the H&E-stained section shows gliosis and fine vacuole-type spongiform change. (a2) GFAP immunoreactivity is present in the subpial gray matter but less so in the deep cerebral neocortex. (a3) AQP-4 immunoreactivity is widespread in the cerebal neocortex. (b1) In panencephalopathic (PE)-type sporadic Creutzfeldt-Jakob (sCJD), the H&E-stained section shows severe gliosis, neuronal loss, tissue rarefaction and status spongiosus. (b2) Widespread GFAP immunoreactivity is present in the subpial gray matter and cerebral neocortex. (b3) Strong AQP-4 immunoreactivity is observed in the cerebal neocortex. The patterns of immunoreactivity of AQP-4 and GFAP differ. Analysis of serial sections showed that spongiform change and gliosis in sCJD appeared at an earlier stage than did strong AQP-4 immunoreactivity. Neuronal loss and tissue rarefaction in sCJD appeared at a later stage than did strong AQP-4 immunoreactivity. Strong AQP-4 immunoreactivity appeared at an earlier stage than did hypertrophic astrocytosis or GFAP-positive astrocytosis. (H&E, hematoxylin and eosin). Bars, 200 m. a, patient 1 (SSE case); b, patient 9 (PEtype sCJD case). a1 and b1, H&E stain; a2 and b2, GFAP immunostaining; a3 and b3, AQP-4 immunostaining.



tions, the development of sCJD pathology, particularly of neocortical lesions, occurs as follows: PrP deposition, spongiform change and gliosis, increased AQP-4 staining, hypertrophic astrocytosis, and neuronal loss and tissue rarefaction. Interestingly, increased AQP-4 staining remained in burnt-out lesions, such as those of status spongiosus.

Determining the relation between clinical data and AQP-4 immunoreactivity was an objective of the present study. We suggest a relation between AQP-4 immunoreactivity and MRI findings, particularly in diffusion-weighted images. AQP-4 immunoreactivity was intensely positive in sCJD cerebral neocortex, which usually shows high signal intensity in diffusion-weighted images in sCJD.<sup>23</sup> Because this high signal intensity may result from variable pathologic factors, including spongiform change, astrocytosis, microglial activation and PrP deposition,<sup>23</sup> evaluation of the relation between MRI findings and AQP-4 immunoreactivity may be difficult. However, we believe that

increased AQP-4 expression, particularly in the cerebral neocortex, may be involved in the high signal intensity of diffusion-weighted images in sCJD.

Western blot analysis of protease-resistant PrP showed type 1 PrP in all cases. Because most of our cases were classified as type MM1, with single cases of type MV1 and no cases of type MM2, MV2, VV1 or VV2, assessment of the influence of PRNP polymorphisms at codon 129 and PrP type relative to AQP-4 immunoreactivity in the present study cannot be made. Further studies of PRNP polymorphisms, PrP type and AQP-4 immunoreactivity in sCJD are necessary.

In conclusion, widespread AQP-4 immunoreactivity, particularly in the cerebral neocortex and cerebellar cortex in sCJD, was shown in the present study. Although this may be a secondary change of sCJD pathology associated with intense astrocytosis, it may reflect important aspects of astrocytic pathology associated with sCJD.

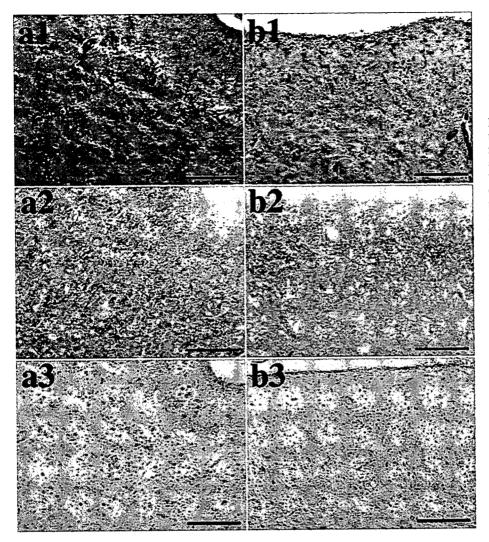


Fig. 8 Two sets of serial sections show the relation between neuropathologic findings, prion protein (PrP) deposition and aquaporin-4 (AQP-4) immunoreactivity. (a1) In subacute spongiform encephalopathy (SSE), the H&E-stained section shows hypertrophic astrocytosis; neuronal loss is not apparent. (a2) Widespread granular PrP immunoreactivity is present in the cerebal neocortex. (a3) Widespread AQP-4 immunoreactivity is present in the cerebal neocortex. (b1) In panencephalopathic (PE)-type sporadic Creutzfeldt-Jakob disease (sCJD), the H&E-stained section shows pronounced hypertrophic astrocytosis, neuronal loss and status spongiosus. (b2) PrP immunoreactivity is relatively decreased in the cerebral neocortex compared to that in SSE. (b3) Strong AQP-4 immunoreactivity is present in the cerebral neocortex. The patterns of PrP deposition and immunoreactivity of AQP-4 differ. PrP deposition appeared at an earlier stage than did spongiform change or gliosis and at an earlier stage than did strong AQP-4 immunoreactivity. Bars, 200 m. a, patient 6 (SSE case); b, patient 8 (PEtype sCJD case). a1 and b1, H&E stain; a2 and b2, PrP immunostaining; a3 and b3, AQP-4 immunostaining.

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#### Note



# Assessment of Prion Inactivation by Combined Use of *Bacillus*-Derived Protease and SDS

Miyako Yoshioka, Yuichi Murayama, Takehiro Miwa, Katsuhiro Miura, Masuhiro Таката, Takashi Yokoyama, Koji Nishizawa, and Shirou Mohri Mohri Miyako Yoshioka, Yuichi Murayama, Koji Nishizawa, Masuhiro Miura, Katsuhiro Miura, Katsuhiro Miura, Masuhiro Takata, Katsuhiro Miura, Kat

<sup>1</sup>Prion Disease Research Center, National Institute of Animal Health,

3-1-5 Kannondai, Tsukuba, Ibaraki 305-0856, Japan

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Prions, infectious agents causing transmissible spongiform encephalopathy, retain infectivity even after undergoing routine sterilization processes. We found that MSK103 protease, identified in our previous study, effectively reduces infectivity and the level of misfolded isoform of the prion protein in scrapie-infected brain homogenates in the presence of SDS. The treatment therefore can be applied to the decontamination of thermolabile instruments.

Key words: prion inactivation; protease; scrapie; bioassay; protein misfolding cyclic amplification

Prions are the infectious agents of transmissible spongiform encephalopathies (TSEs), including bovine spongiform encephalopathy (BSE) in cattle, scrapie in sheep and goats, and Creutzfeldt-Jakob disease (CJD) in humans.<sup>1)</sup> TSE diseases are characterized by accumulation in the central nervous system of PrPSc, a proteinase K (PK)-resistant misfolded isoform of the cellular prion protein PrPC.<sup>2)</sup> Since prions are highly resistant to conventional sterilization procedures, severe physical and chemical sterilization methods, such as high-pressure sterilization at 134 °C for 18 min or strong alkali treatment, are required to reduce infectivity.<sup>3)</sup>

The development of a secure, simple, and effective disinfection method is required as a preventive measure against prion diseases. We recently identified a *Bacillus* strain that produced protease exhibiting high-degradation activity against scrapie- and BSE-derived PrP<sup>Sc</sup>. <sup>4)</sup> The isolated protease (MSK103) was found to degrade dried PrP<sup>Sc</sup> samples more effectively than proteinase K (PK) or a keratinase. <sup>5)</sup> Since this protease can be used

under moderate conditions (optimum pH and temperature ranges were 9–10 and 60–70 °C), it can be used to decontaminate fragile, precision, and expensive instruments that are susceptible to PrPSc contamination.

In the present study, we investigated the efficacy of the MSK103 protease in scrapie prion inactivation. Since protease activity is generally enhanced in the presence of SDS, 6-8) we examined the additive effects obtained by the combined use of the protease and SDS. The hamster-adapted scrapie strain Sc237 was propagated in the Syrian golden hamster. The brains of hamsters at the terminal stage of the disease, titrating  $5 \times 10^{8.5}$  LD<sub>50</sub> per gram by bioassay, were pooled and homogenized at a concentration of 10% (w/v) in 50 mmol Tris-HCl buffer (pH 9.0). The homogenates were digested with MSK103 protease at a final concentration of 0-8 units (U)/ml at 50 °C for 20 h in the presence of 2% SDS. In a preliminary experiment, PrPSc samples containing various concentrations of SDS were digested and analyzed by western blot (WB). The addition of 2% SDS to the reaction was most effective for the degradation of PrPSc. Effective PrPSc decontamination was observed at a protease concentration of 2-3.8 U/ml against BSE-infected brain homogenate and dried Sc237 PrPSc, which had firmly attached to a plastic surface in our previous study.4) One unit of enzyme activity was defined as the amount of enzyme required to release 1 μmol of glycine per min in a 0.5% solution of human hair-derived keratin powder at 60°C for 1 h at pH 8.0.

The digested samples were centrifuged at 200,000 g for 30 min and a supernatant was carefully removed. No  $PrP^{Sc}$  signals were detected in the supernatant by WB analysis. After they were washed three times, the precipitates were resuspended in an initial volume of

proteinase-resistant prion protein; ANOVA, analysis of variance

<sup>&</sup>lt;sup>2</sup>Meiji Seika Kaisha Ltd., Kohoku-ku, Yokohama, Kanagawa 222-8567, Japan

<sup>&</sup>lt;sup>3</sup> Japan Livestock Technology Association, Tokyo 113-0034, Japan

<sup>&</sup>lt;sup>†</sup> To whom correspondence should be addressed. Tel/Fax: +81-29-838-8333; E-mail: ymura@affrc.go.jp

\*\*Abbreviations: TSE, transmissible spongiform encephalopathy; BSE, bovine spongiform encephalopathy; CJD, Creutzfeldt-Jakob disease; PrP<sup>Sc</sup>, misfolded isoform of prion protein; PrP<sup>C</sup>, cellular prion protein; PK, proteinase K; PBS, phosphate-buffered saline; PAGE, polyacrylamide gel electrophoresis; AP, alkaline phosphatase; HRP, horseradish peroxidase; WB, western blot; PMCA, protein misfolding cyclic amplification; PrP<sup>res</sup>,

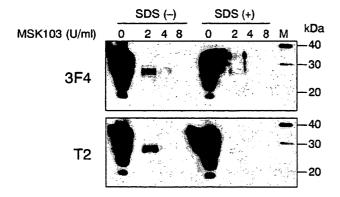


Fig. 1. Western Blot Detection of PrpSc in Sc237-Infected Hamster Brain Homogenate with the MSK103 Enzyme.

A 10% brain homogenate and 0-16 U/ml of the protease solution were mixed in equal quantities, and the mixture was then incubated at 50 °C for 20 h in the absence or presence of 2% of SDS. The positions of molecular weight standards (Magic Mark XP, Invitrogen) are indicated on the figure (the lane labeled "M").

phosphate-buffered saline (PBS). The samples were separated by SDS-polyacrylamide gel electrophoresis (PAGE) and transferred to a polyvinylidine fluoride membrane (Millipore, Bedford, MA). After blocking, the membranes were incubated for 1 h with alkaline phosphatase (AP)-conjugated 3F4 antibody (Signet Laboratories, Dedham, MA; 1/10,000) or horseradish peroxidase (HRP)-conjugated T2 antibody (1/10,000). After washing, the blotted membranes were developed with Immobilon western chemiluminescent AP or HRP substrates (Millipore). Chemiluminescence signals were analyzed using a Light Capture system (Atto, Tokyo).

Figure 1 illustrates the results of WB analysis of the digested samples in the absence or presence of SDS. Residual PrPSc was detected with 3F4 or T2 monoclonal antibodies, which recognize different epitopes of the PrPSc core fragments. As expected, residual PrPSc was significantly reduced in the presence of SDS. Although the PrPSc signal was detected in the sample digested with 2 U/ml of the protease, even in the presence of SDS there were no detectable signals in the samples treated with either 4 or 8 U/ml of the protease. When concentrated to a higher level (16 U/ml), a faint signal of residual PrPSc was detected by WB analysis, suggesting that protease activity somewhat, decreased probably due to self-digestion of the protease. Hence, the samples treated with a combination of the protease (2-8 U/ml) and SDS were analyzed in further experiments.

In addition to conventional WB analysis, protein misfolding cyclic amplification (PMCA), a highly sensitive method of detecting minute amounts of PrPSc, was performed as in our previous study. <sup>10)</sup> Briefly, the digested sample was diluted 1:10 in 10% uninfected brain homogenate, and one round of the PMCA reaction was performed by applying 40 cycles of sonication followed by incubation at 37 °C for 1 h. Next, a process diluting the PMCA product to 1:10 and its subsequent amplification was repeated twice. With regard to the

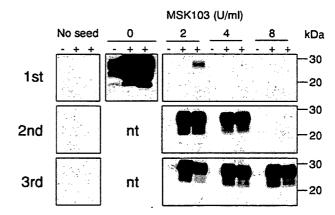


Fig. 2. Protein Misfolding Cyclic Amplification of the Samples Treated with MSK103 Protease.

The samples were analyzed before (-) and after (+), in duplicate) each round of amplification by western blotting following digestion with proteinase K  $(50\,\mu\text{g/ml})$ . The lanes labeled 'No seed' were the controls, in which  $PrP^C$  source only was treated in the same manner. No signals were detected in these samples. The positions of the molecular markers corresponding to 30 and 20 kDa, are shown in the figure. nt. not tested.

PMCA products, samples  $(10\,\mu\text{l})$  collected before and after each round of amplification were mixed with  $10\,\mu\text{l}$  PK solution  $(100\,\mu\text{g/ml})$  and incubated at 37 °C for 1 h. The digested materials were analyzed by WB using the 3F4 antibody, as described above.

Figure 2 illustrates the results of amplification of the samples treated with MSK103 protease. In the first round of amplification, weak PK-resistant PrP (PrPres) signals were detected in the sample treated with 4 U/ml of the protease in addition to the sample treated with 2 U/ml of the protease. In the second round of amplification, the signals of both these samples were enhanced, and a weak signal was observed in the sample treated with 8 U/ml of the protease. After three rounds of amplification, PrPres signals were detected in all the digested samples. These results indicate that a trace level of PrPsc, which was not detected by conventional WB, remained in the samples treated with 4 or 8 U/ml of the protease.

The digested samples were inoculated intracerebrally (20 µl per mouse) into five Tg52NSE transgenic mice<sup>11)</sup> that overexpressed hamster PrP<sup>C</sup> in their nerve systems. The advantage of this bioassay is its ability to confirm the onset of the disease a short period of time after inoculation. The control mice inoculated with the untreated samples developed the disease after an average period of  $45 \pm 2 \,\mathrm{d}$  (average  $\pm \,\mathrm{SD}$ , Fig. 3). The mice that were inoculated with the sample treated with SDS alone died after an average period of  $65 \pm 4 \,\mathrm{d}$ . The onset time of the disease in mice inoculated with the sample treated with 2 U/ml of the protease was significantly delayed, but the mice died after an average period of  $206 \pm 128 \,\mathrm{d}$ . On the other hand, two out of the five mice that were inoculated with samples treated with either 4 or 8 U/ml of the protease survived for more than

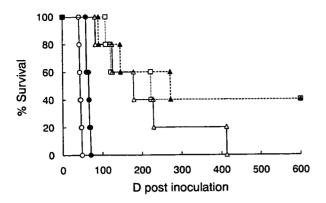


Fig. 3. Survival Curves of Tg52NSE Mice Inoculated with Infected Brain Homogenates Treated with MSK103 Protease.

The control and digested samples were injected intracerebrally into five mice. No treatment (solid line,  $\bigcirc$ ), SDS alone (solid line,  $\bigcirc$ ), 2 U/ml (solid line,  $\triangle$ ), 4 U/ml (broken line,  $\triangle$ ) and 8 U/ml (dotted line,  $\square$ ).

600 d, although the remaining mice developed the disease after an average period of  $170 \pm 73 \, d$  (4 U/ml) and  $151 \pm 62 \, d$  (8 U/ml). The survival times of the affected mice inoculated with the samples treated with the protease were statistically analyzed by one-way ANOVA. There were no significant differences among the survival times of the three experimental groups.

Although several approaches to prion inactivation by enzymatic degradation have been reported, the infectivity of the degradation products was not examined in some studies. 5.12.13) Where bioassay was performed, only a modest reduction in infectivity (approximately  $10^{-3}$ reduction) present in a low concentration (1%) of infected brain homogenate was achieved following protease digestion.<sup>14)</sup> In the present study, we found that combined treatment with MSK103 protease and SDS was effective in reducing the infectivity present in the high concentration (10%) of infected brain homogenate. This led to 40% survival of the inoculated Tg52NSE mice for more than 600 d when the protease treatment was performed under enzyme concentrations of 4 or 8 U/ml at 50 °C for 20 h. The reduction in infectivity was estimated to be less than  $10^{-6}$  on the basis of the incubation time obtained from our previous study. 10) Since a prolonged incubation time (equivalent to a  $10^{-3}$ reduction in infectivity) was also observed in the mice inoculated with the samples treated with SDS alone, the reduction in the amount of infectivity due to the protease was estimated to be less than  $10^{-3}$ .

Since PrPSc tends to aggregate, the size of the PrPSc aggregate determines the efficiency of the enzymatic degradation of PrPSc. If the PrPSc aggregate is very large, it is possible that core PrPSc surrounded by many outside PrPSc remains undigested. This is a serious problem in prion decontamination by enzymatic degradation. The variation in survival times observed in mice inoculated with samples digested with 4 or 8 U/ml of the protease may be due to the fact that core PrPSc which

remained undigested is inhomogeneous in the inoculated samples.

In conclusion, the present study indicates that the combined use of MSK103 and SDS caused effective degradation of PrPSc, resulting in a significant reduction in infectivity. Efficiency of prion decontamination is possibly improved by repeating enzymatic digestion. However, one should always keep in mind the abovementioned problem when enzymatic degradation is applied to the decontamination of instruments that are designed for reuse.

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# Urinary excretion and blood level of prions in scrapie-infected hamsters

Yuichi Murayama, Miyako Yoshioka, Hiroyuki Okada, Masuhiro Takata, Takashi Yokoyama and Shirou Mohri

Prion Disease Research Center, National Institute of Animal Health, 3-1-5 Kannondai, Tsukuba, Ibaraki 305-0856, Japan

Prions, infectious agents causing transmissible spongiform encephalopathy (TSE), are composed primarily of the pathogenic form (PrPSc) of the host-encoded prion protein. Although very low levels of infectivity have been detected in urine from scrapie-infected rodents, no reports of urinary PrPSc have been substantiated. Studies on the dynamics of urinary PrPSc during infection are needed to ensure the safety of urine-derived biopharmaceuticals and to assess the possible horizontal transmission of prion diseases. Using the protein misfolding cyclic amplification technique, a time-course study of urinary excretion and blood levels of PrPSc was performed in Sc237-infected hamsters and a high rate of PrPSc excretion was found during the terminal stage of the disease. Following oral administration, PrPSc was present in all buffy coat samples examined; it was also present in most of the plasma samples obtained from hamsters in the symptomatic stage. PrPSc was excreted in urine for a few days after oral administration; subsequently, urinary PrPSc was not detected until the terminal disease stage. These results represent the first biochemical detection of PrPSc in urine from TSE-infected animals.

Correspondence Yuichi Murayama ymura@affrc.go.jp

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### INTRODUCTION

Prion diseases, commonly referred to as transmissible spongiform encephalopathies (TSEs), are neurodegenerative disorders that have been described in humans as Creutzfeldt-Jakob disease, in sheep and goats as scrapie, and in cattle as bovine spongiform encephalopathy (Prusiner, 1991). A characteristic of TSEs is the accumulation of a protease-resistant, misfolded prion protein (PrPSc), which is the pathogenic form of the host-encoded prion protein (PrPC) (Caughey et al., 1991; Pan et al., 1993). PrPSc accumulation occurs in the central nervous system (CNS) of infected subjects through the structural conversion of PrPC. TSE infectivity has also been demonstrated in a variety of peripheral tissues (Prusiner, 1998) and body fluids such as blood (Houston et al., 2000; Taylor et al., 2000; Holada et al., 2002; Hunter et al., 2002; Cervenakova et al., 2003) of infected subjects.

Chronic wasting disease (CWD) is the only prion disease known to affect free-ranging mule and deer (Williams, 2005), and can spread horizontally (Miller et al., 1998, 2000). Therefore, it was suspected that transmission of CWD occurred through contaminated saliva, faeces and urine, and it has been demonstrated that the saliva of infected deer contain the infectious agent (Mathiason et al., 2006). With regard to excrement, a protease-resistant urinary PrP, designated uPrP<sup>Sc</sup>, has been found in the urine of infected animals and humans (Shaked et al., 2001); however, it was reported that the anti-PrP<sup>Sc</sup> reactivity was

primarily due to an immunoglobulin light chain (Serban et al., 2004; Head et al., 2005) or a contaminated bacterial component (Furukawa et al., 2004). There are several reports that demonstrate infectivity in urine during scrapie infection. Urine from scrapie-infected mice with concurrent nephritis was infectious (Seeger et al., 2005). A concentrated sample from pooled urine collected at the terminal stage of the disease exhibited a low level of infectivity in hamsters (Kariv-Inbal et al., 2006). The result obtained in hamsters is important from the viewpoint that the urinary excretion of PrPSc possibly occurred without severe complications; however, the time course and frequency of appearance of urinary PrPSc remain to be determined.

As the PrP<sup>Sc</sup> level in urine is expected to be diminutive when compared with that in the brain tissue, conventional immunoassays might not be applicable to the detection of urinary PrP<sup>Sc</sup>. Recently, it has become possible to perform *in vitro* amplification of hamster PrP<sup>Sc</sup> by protein misfolding cyclic amplification (PMCA) (Saborio *et al.*, 2001). By repeated sonication and incubation, PrP<sup>Sc</sup> converts PrP<sup>C</sup> to the protease-resistant form (PrP<sup>res</sup>); this PrP<sup>res</sup> then becomes the new nucleus, thereby facilitating PrP<sup>res</sup> formation in the cyclic incubation process. PMCA is a highly effective method for detecting minute amounts of hamster PrP<sup>Sc</sup> (Castilla *et al.*, 2005a; Saá *et al.*, 2006a) and enables the detection of PrP<sup>Sc</sup> in the blood of infected hamsters (Castilla *et al.*, 2005b; Saá *et al.*, 2006b).

In the present study, using the PMCA technique, we examined the urinary excretion of PrPSc in hamsters that were inoculated intracerebrally or orally with the scrapie prion strain Sc237. We also investigated PrPSc levels in the plasma, buffy coat and urine during the period from latent to terminal stages of the disease and compared PrPSc dynamics among the above-mentioned samples. This paper is the first demonstration of urinary PrPSc in any disease model.

### **METHODS**

**Inoculation of the hamster prion.** The hamster-adapted scrapie prion strain Sc237 was propagated in hamsters. The brains of hamsters at the terminal stage of the disease, with a titre of  $5 \times 10^{8.5}$  LD<sub>50</sub> g<sup>-1</sup> by bioassay (Murayama *et al.*, 2006), were pooled and homogenized at a 10 % concentration (w/v) in PBS. The homogenate was injected intracerebrally (20  $\mu$ l per hamster) or administered orally (100  $\mu$ l per hamster) using a gastric tube. All animal experiments were performed in accordance with National Institute of Animal Health guidelines.

**Preparation of urine and blood samples.** Urine samples were collected at several time points after inoculation. From noon to about 3 p.m., the hamsters were held with both hands for spontaneous urination in a sterilized Petri dish. To separate urinary exudate cells such as leukocytes and epithelial cells of renal tubules from urine, the individual or pooled urine sample (20  $\mu$ l) was diluted 1:10 with PBS and passed through a 0.45  $\mu$ m membrane filter unit (Ultrafree-MC; Millipore). The flow-through in the collection tube was removed and 40  $\mu$ l of 2% Triton X-100, 8 mM EDTA in PBS was added to the membrane. The entire surface of the membrane was rinsed with the detergent solution by pipetting, and the membrane unit was incubated for 30 min at room temperature. After centrifugation, the flow-through was recovered and stored at -80 °C until use.

For blood collection, hamsters were sacrificed at several time points after inoculation and blood (1.5 ml) was collected from the heart and heparinized. Blood samples were centrifuged at 830 g for 15 min, and the plasma and buffy coat fractions were recovered. The plasma fraction was centrifuged again at 9170 g for 5 min to remove precipitates and the supernatants were stored at -80 °C until further use. Erythrocytes contained in the buffy coat fraction were haemolysed in 0.75 % NH<sub>4</sub>Cl, 17 mM Tris/HCl (pH 7.2). The white blood cells and platelets were then washed three times in PBS and the pellets stored at -80 °C until analysis.

PMCA and Western blotting. The PMCA and Western blotting procedures have been described in our previous study (Murayama et al., 2006). Briefly, normal hamster brains were homogenized at a 10% concentration (w/v) in PBS containing complete protease inhibitors (Roche Diagnostics), 1 % Triton X-100, and 4 mM EDTA. To avoid contamination, the PrPC source was prepared in a laboratory in which infected materials had never been handled. The urine and plasma samples were diluted 1:10 and 1:100, respectively, in the  $PrP^{\hat{C}}$  source (total 100  $\mu$ l). Buffy coat samples were prepared by three freeze-thaw cycles and mixed with 100 µl of the PrP<sup>C</sup> source. Amplification was performed using a fully automatic cross-ultrasonic protein activating apparatus (ELESTEIN 070-GOT; Elekon Science Corp.), which has a capacity to generate high ultrasonic power (700 W). PMCA was performed by 40 cycles of sonication (a pulse oscillation for 3 s was repeated five times at 1 s intervals), followed by incubation at 37 °C for 1 h with gentle agitation. The amplified product obtained after the first round of amplification was diluted 1:10 in the PrPC source and a second round of amplification was

performed. The process was repeated to obtain the amplified products. Samples were treated with proteinase K ( $50 \mu g ml^{-1}$ ), separated by SDS-PAGE and transferred to a PVDF membrane (Millipore). After blocking, the membrane was incubated for 1 h with 1:10000-diluted alkaline phosphatase-conjugated 3F4 monoclonal antibody (mAb; Signet Laboratories). After washing, the blotted membrane was developed using Immobilon Western Chemiluminescent AP substrate (Millipore) according to the manufacturer's instructions. After exposure for 5 min, chemiluminescence signals were analysed using a Light Capture System (ATTO).

Histological analysis. The urinary PMCA product obtained by 160 amplification cycles (see Fig. 1b, lane f) was diluted 1:10 with PBS and injected intracerebrally into Tg52NSE mice (20 µl per mouse), which overexpress hamster PrP<sup>C</sup> in their nervous system (Race et al., 1995). Densitometric analysis of Western blots revealed that the PrP<sup>res</sup> signal intensity in the urinary PMCA product was approximately one-eighth of that in the 1% homogenate of Sc237-infected brain. Therefore, the 10% homogenate of Sc237-infected brain was diluted 1:800 and injected into Tg52NSE mice as a control. Tg52NSE mice exposed to the urinary PMCA product and Sc237-infected brain homogenate were sacrificed at 148 and 62 days post-inoculation (p.i.), respectively.

The left hemisphere of the brain was fixed in 10% buffered formal saline (pH 7.4). Coronal slices of 2 mm thickness were immersed in 98% formic acid to diminish prion infectivity (Taylor et al., 1997) and embedded in paraffin. Serial sections were mounted on new silane II-coated glass slides (Muto Pure Chemicals Co. Ltd) and routinely stained with haematoxylin and eosin. For the detection of immunoreactive PrPSc, dewaxed sections were immersed in 98% formic acid for 5 min and then treated with 3 % hydrogen peroxide in methanol to block endogenous peroxidase activity. Sections were treated by autoclaving for 3 min at 121 °C in a low-ionic-strength buffer [2.1 mM Tris/HCl (pH 7.8), 1.3 mM EDTA, 1.1 mM sodium citrate; Asante et al., 2006]. After washing in distilled water, the sections were placed on an automated immunohistochemical stainer (Autostainer Universal Staining System; DakoCytomation). mAb T1, which was generated in PrP-deficient mice by immunization with recombinant mouse PrP<sub>121-231</sub> and which recognizes mouse PrP<sub>138-144</sub>, was used as the primary antibody. Anti-mouse universal immunoperoxidase polymer (Histofine Simple Stain MAX-PO (M); Nichirei) was used as the secondary antibody, and 3'-3diaminobenzidine tetrachloride was used as the chromogen. Astrocytes were immunostained with a mouse anti-human glial fibrillary acid protein antibody (GFAP; DakoCytomation) after heatmediated antigen retrieval. All sections were counterstained with haematoxylin.

### RESULTS

### **Detection of urinary PrP<sup>Sc</sup> in Sc237-infected** hamsters

In the Western blot analysis,  $PrP^{Sc}$  propagation and accumulation in hamster brains became apparent 38 days after intracerebral inoculation (data not shown). The mean incubation time of the inoculated hamsters was  $70\pm2$  days (mean  $\pm$  SD, n=58) and the onset of clinical signs appeared 4–7 days prior to death. Fig. 1(a) illustrates the results of the amplification of the urine samples. No  $PrP^{res}$  was amplified in the uninfected control samples and pooled samples obtained during the period 26–46 days p.i. after 280 amplification cycles. In contrast,  $PrP^{res}$  was detected

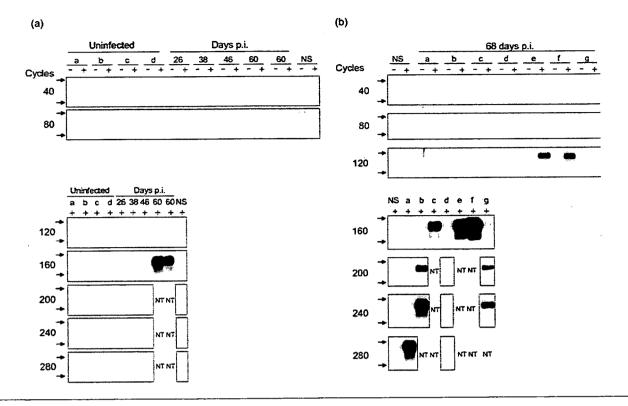
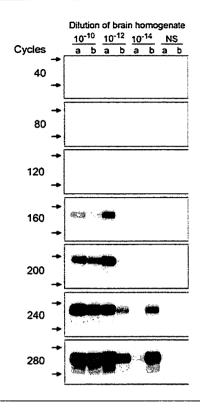


Fig. 1. (a) PMCA results from urine samples obtained from hamsters inoculated intracerebrally with Sc237. The PrP<sup>C</sup> source was mixed with 0.1 vols of the pooled urine samples prepared as described in Methods, and one round (40 cycles) of the PMCA reaction was performed. This process of 1:10 dilution of the PMCA product and subsequent amplification was repeated six times. Samples before (-) and after (+) amplification were analysed by Western blotting following digestion with proteinase K. Lanes: a-d, negative controls in which the urine sample from an uninfected hamster was used for PMCA. No signals were detected in these samples. Arrows indicate the positions of molecular mass markers corresponding to 30 and 20 kDa. (b) PMCA results from urine samples obtained at the terminal stage of disease. Urine samples were prepared from seven individual hamsters (labelled 'a'-'g') and used for sequential PMCA. NS, No seed: negative control reaction in which uninfected brain homogenate (PrP<sup>C</sup> source only) was treated in the same manner; NT, not tested.

after 160 cycles in two different pools of urine collected from five hamsters at 60 days p.i. To examine the incidence of urinary PrPSc excretion at the terminal stage of the disease, urine was collected from seven individual hamsters at 68 days p.i. and amplified (Fig. 1b). PrPres signals were detected in two samples (Fig. 1b, hamsters 'e' and 'f') after 120 amplification cycles. Based on previous data (Murayama et al., 2006), we estimated that the level of PrPSc present in these samples corresponded to that present in a 10<sup>-9</sup>-diluted infected brain homogenate; such a dilution is well below the level detectable by any other method. PrPres signals were detected in one sample (Fig. 1b, hamster 'c') after 160 cycles, in two samples (Fig. 1b, hamsters 'b' and 'g') after 200 cycles, and in one sample (Fig. 1b, hamster 'a') after 280 cycles of amplification. In contrast, no signals were detected in hamster 'd' after 280 amplification cycles.

We also examined the detection limit of our amplification system (Fig. 2) and confirmed that PrP<sup>Sc</sup> present in an infected brain homogenate diluted 1:10<sup>10</sup> could be

detected after 160 cycles, and that PrPres signals were enhanced in a similar manner during the further amplification process in the duplicated samples. However, in the more extreme dilution range, the number of amplification cycles required to amplify PrPSc to a detectable level varied in the duplicated samples. After 240 amplification cycles, both samples diluted 1:1012 became positive for PrPres; however, the PrPres signal remained very weak in one of the samples that was diluted 1:1014, even after 280 cycles. As PrPSc tends to aggregate, the weak detection of PrPres in the reaction may have been due to the near-absence of PrPSc seed, which would have been almost completely diluted out. Therefore, it was assumed that PMCA with 280 cycles could reproducibly detect PrPSc present in the samples at a level equivalent to that present in an infected brain homogenate diluted 1:1012. Taken together, the PMCA analysis results indicated that PrPSc was excreted in the urine in a higher rate (86 %, six out of seven hamsters in Fig. 1b) at the terminal stage of the disease, although PrPSc concentrations differed considerably among the urine samples.



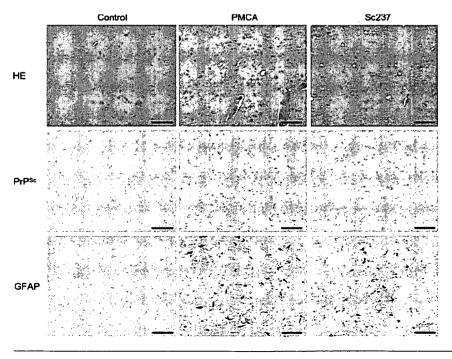
**Fig. 2.** Sensitivity of detection of hamster scrapie Sc237 using sequential PMCA. The PrPSc seed was diluted from 10<sup>-10</sup> to 10<sup>-14</sup> and used as the PrPC source. Lanes a and b represent the results of duplicate samples. Arrows indicate 30 and 20 kDa molecular mass markers. NS, No seed.

## Histological analysis of brain tissue infected with urine-derived PMCA product

Neuropathologically, vacuolar degeneration was detected in the thalamus, mesencephalon and medulla oblongata of Tg52NSE mice inoculated with Sc237 and the PMCA product (Fig. 3). This vacuolation was present mainly in the neuropil but was rare in the neurons. No vacuolation was observed in control mice brains. Granular-type PrPSc accumulation principally associated with the presence of vacuolation was observed in the neuropil and neurons of the thalamus, mesencephalon and medulla oblongata stained with mAb T1. Swollen or hypertrophic astrocytes proliferated in the same lesion that showed PrPSc accumulation. No significant difference was observed in vacuolar change, PrPSc accumulation and astrogliosis between Tg52NSE mice inoculated with Sc237 and those inoculated with the PMCA product.

### PrPSc levels in the blood

As it has been demonstrated that PrPSc binds to a blood component (Fischer et al., 2000), we examined PrPSc levels in both the plasma and buffy coat samples collected during the period of detection of urinary PrPSc. Fig. 4(a) shows the results of the amplification of buffy coats obtained at 60 and 68 days p.i. All samples became positive for PrPres signals after 160 (60 days p.i.) or 120 (68 days p.i.) amplification cycles; therefore, we confirmed that PrPSc could be detected in the buffy coat samples by PMCA, as reported previously (Castilla et al., 2005b). With regard to



**Fig. 3.** Pathology of Tg52NSE mice exposed to urine-derived PMCA product from hamster 'f' in Fig. 1(b) (PMCA) and Sc237-infected brain homogenate (Sc237). Thalamus sections of mice that succumbed to scrapie after intracerebral inoculation with the PMCA product or the infected brain homogenate showed similar levels of vacuolation,  $PrP^{Sc}$  accumulation (labelled with mAb T1) and astrogliosis (labelled with anti-GFAP antibody). Bars, 50 μm.

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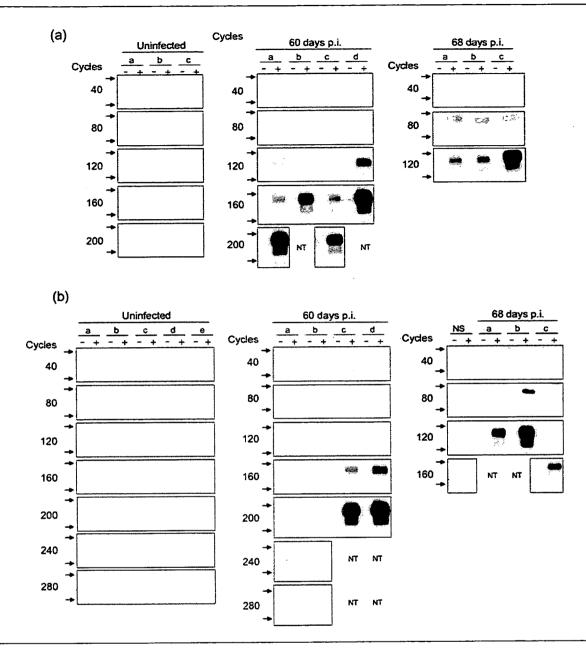
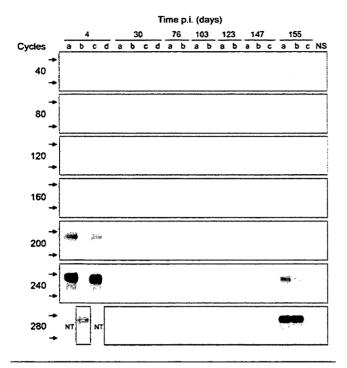


Fig. 4. PMCA of buffy coat (a) and plasma (b) samples obtained at 60 and 68 days p.i. from individual hamsters inoculated intracerebrally with Sc237. The PrP<sup>C</sup> source was mixed with the buffy coat sample prepared as described in Methods or with a 1:100 dilution of the plasma sample. Sequential PMCA was performed on three to five hamsters (labelled 'a'-'e'). Arrows indicate 30 and 20 kDa molecular mass markers. NS, no seed; NT, not tested.

the plasma samples, PrP<sup>res</sup> signals were detected after 160 amplification cycles in two samples ('c' and 'd') collected at 60 days p.i., whereas no signal was detected, even after 280 cycles, in other samples (Fig. 4b). In the three samples collected at 68 days p.i., 80–160 amplification cycles were required to amplify PrP<sup>res</sup> to a detectable level. These results indicated that PrP<sup>Sc</sup> was also present in the plasma at symptomatic and terminal stages of the disease. However, the detection sensitivity of PrP<sup>Sc</sup> in plasma was low compared with that in buffy coats.

## $PrP^{Sc}$ levels in urine and blood after oral administration

The mean survival time of orally administered hamsters was  $171\pm15$  days (mean  $\pm$  sp, n=18). Fig. 5 illustrates the results of the amplification of urine samples collected between 4 and 155 days p.i. PrP<sup>res</sup> was detected after 200–240 amplification cycles in two of four samples collected at 4 days p.i., and an additional sample became positive for PrP<sup>res</sup> after 280 cycles. In contrast, PrP<sup>res</sup> signals could not



**Fig. 5.** PMCA results from urine samples obtained from individual hamsters after oral administration. Urine was collected from two to four hamsters (labelled 'a'-'d') at each time point. Arrows indicate 30 and 20 kDa molecular mass markers. NS, no seed; NT, not tested.

be detected in urine obtained between 30 and 147 days p.i. At 155 days after administration, urinary PrP<sup>Sc</sup> was detected in two of three samples after 240–280 amplification cycles. This observation was in agreement with the results that urinary PrP<sup>Sc</sup> was detected in the symptomatic and terminal stage of disease after intracerebral inoculation.

Fig. 6 (a) and (b) illustrate the results of amplification of plasma and buffy coats, respectively, obtained from animals sacrificed during the period 30-166 days p.i. PrPSc was not detected in plasma collected at the asymptomatic stage (<75 days p.i.) (Fig. 6a). At 103 days p.i., the propagation and accumulation of PrPSc became apparent in brains (Fig. 6c); at this stage, PrPres signals were detected in the plasma after 160-200 amplification cycles (Fig. 6a). However, no PrPres signal could be detected in one plasma sample (sample b) collected at 103 days, even after 280 amplification cycles. No remarkable PrPSc accumulation was observed in the brain of the plasma donor (Fig. 6c, 103 days p.i., sample b) when compared with the other donor (Fig. 6c, 103 days p.i., sample a). With regard to the buffy coats, PrPres signals were detected after 120-160 cycles in all hamsters examined during the period 75-166 days p.i (Fig. 6b). We could not examine the buffy coat samples at 30 days p.i. because of a technical failure and could not verify the previous observation that PrPSc disappeared in the buffy coat

fraction during the period of asymptomatic stage of the disease (Saá et al., 2006b).

### DISCUSSION

In the present study, using PMCA, we have revealed for the first time the presence of PrPSc in the urine of hamsters infected intracerebrally or orally with scrapie, without any dialysis and concentration procedures. A characteristic property of the urinary PrPSc was its detectability only at the terminal stage of the disease, except for a few days immediately after oral administration (Fig. 7). We also examined PrPSc levels in blood and clarified that PrPSc could be detected in plasma as well as in buffy coat fractions. There was an obvious time lag between PrPSc excretion in the urine and the appearance of PrPSc in the blood, and urinary PrPSc did not increase in proportion to PrPSc accumulation in the CNS. Furthermore, in some cases, urinary PrPSc could not be detected, even during the terminal stage of the disease, using the highly sensitive PMCA technique. Hence, urinary PrPSc appears to be an unsuitable marker for the early diagnosis of prion infection.

With regard to urinary PrPSc amplification, 120 or more cycles were required to amplify PrPSc to a detectable level, indicating that the amount of PrPSc present may be insufficient to induce infection. Although an extremely small amount of PrPSc is excreted in the urine, PrPSc can survive in the natural environment for a long time. In fact, prions remain infectious after incubation for several years in soil (Brown & Gajdusek, 1991), and recent research has demonstrated that PrPSc adsorbs firmly to soil components (Leita et al., 2006) and that the soil minerals-PrPSc complex remains infectious (Johnson et al., 2006). Therefore, it is conceivable that PrPSc is captured and absorbed effectively on soil and is then concentrated. Furthermore, it has been indicated that soil ingestion behaviours have been observed frequently in wild animals including mule deer (Arthur & Alldredge, 1979). A recent study indicated that excrement-borne transmission was not proved in an experimental CWD infection (Mathiason et al., 2006); however, the possibility exists that some prion diseases may spread via contaminated soil in nature.

Where is the urinary PrP<sup>Sc</sup> derived from? PrP<sup>res</sup> was amplified successfully from urinary material trapped on a filter membrane. The urinary material probably included leukocytes and epithelial cells from the renal tubules. In preliminary experiments, PrP<sup>res</sup> could not be amplified when the flow-through fraction was concentrated and used as a PMCA seed (data not shown); this suggests that PrP<sup>Sc</sup> was primarily present in the insoluble and solid fraction in the urine samples. Kidney dysfunction or bacterial urinary tract infection might occur following disease progression, and lymphocyturia might arise at the terminal stage of the disease. However, the above elucidation does not provide a reasonable explanation for amplification of PrP<sup>res</sup> in urine

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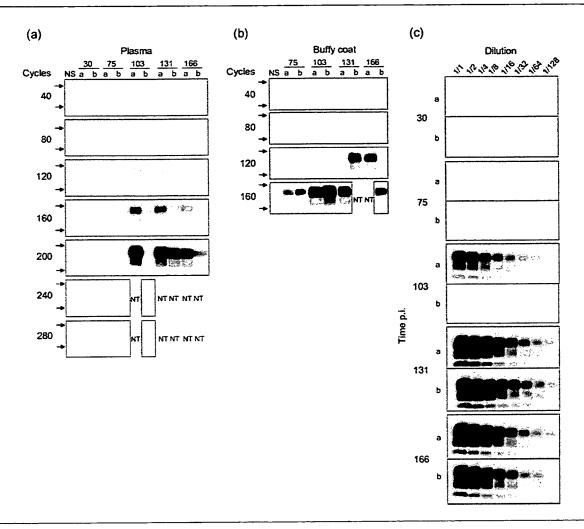
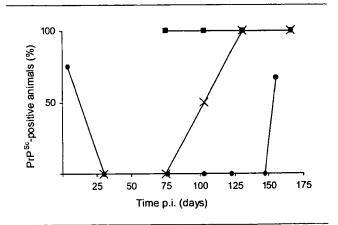


Fig. 6. (a, b) PMCA results from plasma (a) and buffy coat (b) samples obtained at 30–166 days p.i. from hamsters inoculated orally with Sc237. Two hamsters (labelled 'a' and 'b') were sacrificed at each time point, and blood and brain samples were prepared. (c) PrPSc accumulation in the brains of hamsters after oral administration of Sc237. Serial twofold dilutions of 10% brain homogenate were prepared and the results analysed by Western blotting. NS, no seed; NT, not tested.



**Fig. 7.** Time course for the appearance of PrP<sup>Sc</sup> in urine (●), plasma (×) and buffy coat (■) of orally infected hamsters.

samples collected at 4 days after peroral inoculation. It has been demonstrated that PrPSc can be detected on migrating intestinal dendritic cells (DCs) immediately after oral administration (Huang et al., 2002), and that infected DCs can transmit the PrPSc prion to the CNS in recombination activating gene (RAG) knockout mice (Aucouturier et al., 2001). These observations indicated that DCs could transport PrPSc from the intestinal tract to other tissues and body fluids without any propagation of the prion in the peripheral tissues. DCs also exist in the kidney, and it has been demonstrated that renal DCs form a contiguous network that continuously surveys the tubulointerstitium (Soos et al., 2006). Renal DCs would capture PrPSc from the tubular lumen. Therefore, migrating DCmediated transport may also be involved in urinary excretion of PrPSc, particularly in the early stage of infection.

The results of PrPres amplification in the plasma samples after oral administration provide a possible explanation for the origin of plasma PrPSc. PrPres could not be amplified from the plasma samples collected at 30 and 75 days p.i. and from one sample at 103 days p.i.; PrPSc accumulation could not be detected in the brains of these hamsters (Fig. 6). In contrast, PrPres could be amplified from all plasma samples that were obtained when PrPSc propagation was apparent in the brains. These observations imply that PrPSc might leak into the plasma from the infected brain, probably due to destruction of the blood-brain barrier. The source of PrPSc in the buffy coats during the asymptomatic stage appeared to be the spleen and other lymphoid organs, as suggested previously (Saá et al., 2006b). A more detailed analysis needs to be performed to confirm whether plasma PrPSc can serve as an indicator for the propagation of PrPSc in the CNS.

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### Note



### Assessment of Prion Inactivation by Fenton Reaction Using Protein Misfolding Cyclic Amplification and Bioassay

Kyozo Suyama,<sup>1</sup> Miyako Yoshioka,<sup>2</sup> Mitsugu Akagawa,<sup>3</sup> Yuichi Murayama,<sup>2,†</sup> Hiroko Horii,<sup>2</sup> Masuhiro Таката,<sup>2</sup> Takashi Yokoyama,<sup>2</sup> and Shirou Mohri<sup>2</sup>

<sup>1</sup>Sports Nutrition Department, Sendai University, 2-2-18 Funaoka Minami, Shibata 989-1693, Japan

Graduate School of Life and Environmental Science, Osaka Prefecture University,

1-1 Gakuen-cho, Naka-ku, Sakai 599-8531, Japan

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An abnormal isoform of the prion protein, associated with transmissible spongiform encephalopathies, retains infectivity even after undergoing routine sterilization processes. We found that a formulation of iron ions combined with hydrogen peroxide effectively reduced infectivity and the level of abnormal isoforms of the prion protein in scrapie-infected brain homogenates. Therefore, the Fenton reaction has potential for prion decontamination.

**Key words:** prion inactivation; Fenton reaction; scrapie; bioassay; protein misfolding cyclic amplification

Prions, infectious agents causing transmissible spongiform encephalopathy (TSE), primarily comprise the proteinase K (PK)-resistant pathogenic isoform (PrPSc) of the host-encoded prion protein (PrPC).1) Prion diseases include Creutzfeldt-Jakob disease (CJD) in humans, bovine spongiform encephalopathy (BSE) in cattle, and scrapie in sheep and goats.2) PrPSc accumulates in the central nervous system of the infected subjects via structural conversion of PrPC.3.4) PrPSc deposition or TSE infectivity has also been found in a variety of peripheral tissues1) and body fluids, such as the blood<sup>5)</sup> of scrapie-infected subjects. Since PrP<sup>Sc</sup> has also been detected in the peripheral tissues of CJD patients,6) there is a serious risk of iatrogenic transmission through human-derived products, medical equipment, and instruments such as endoscopes.<sup>7)</sup>

PrPSc retains infectivity even after undergoing routine

sterilization processes, and complete inactivation necessitates very harsh treatment such as complete incineration, strong alkali treatment, or high-temperature autoclaving.<sup>8)</sup> Hence, the livestock and healthcare industries desire the development of practical techniques for PrPSc inactivation. Hence, novel sterilization methods involving chemical<sup>9,10)</sup> or enzymatic<sup>11,12)</sup> treatment have been devised to inactivate prions effectively.

In the present study, we investigated the efficacy of the Fenton reaction in scrapie prion inactivation. The Fenton reaction consists of the reduction of Fe (III) by an electron donor and the generation of a hydroxyl radical *via* the reduction of hydrogen peroxide  $(H_2O_2)$  by the reduced metal. In Iron is an essential redox transition element that is capable of inducing ion-mediated damage in proteins. Here, we present evidence justifying interest in an Fe (II)/ $H_2O_2$  formulation for use in prion inactivation.

The hamster-adapted scrapie prion strain Sc237 was propagated in hamsters. The brains of hamsters at the terminal stage of the disease, titrating  $5 \times 10^{8.5}$  LD<sub>50</sub> per gram by bioassay, <sup>16)</sup> were pooled together and homogenized at a concentration of 20% (w/v) in phosphate-buffered saline (PBS). The homogenate was mixed with an equal volume of 0.2% (w/v) FeCl<sub>2</sub>-PBS solution in the presence of 5% (v/v) H<sub>2</sub>O<sub>2</sub>. The samples were heated to 50 °C and kept for 22 h. The Fenton reaction was performed on the basis of experimental results obtained in our previous study. <sup>15)</sup> A high concentration of H<sub>2</sub>O<sub>2</sub> was used in the present study, but H<sub>2</sub>O<sub>2</sub> itself does not disinfect prions. <sup>17)</sup> After treatment, the samples

<sup>&</sup>lt;sup>2</sup>Prion Disease Research Center, National Institute of Animal Health,

<sup>3-1-5</sup> Kannondai, Tsukuba 305-0856, Japan

<sup>&</sup>lt;sup>3</sup>Department of Biological Chemistry, Division of Applied Life Science,

<sup>&</sup>lt;sup>†</sup> To whom correspondence should be addressed. Tel/Fax: +81-29-838-8333; E-mail: ymura@affrc.go.jp

Abbreviations: TSE, transmissible spongiform encephalopathy; PK, proteinase K; PrPSc, pathogenic isoform of prion protein; PrPC, host-encoded prion protein; CJD. Creutzfeldt-Jakob disease; BSE, bovine spongiform encephalopathy; PBS, phosphate-buffered saline; HRP, horseradish peroxidase; WB, western blot; PMCA, protein misfolding cyclic amplification; PrPres, protease-resistant prion protein

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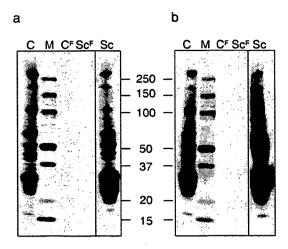


Fig. 1. Western Blotting Analysis of the Prion Protein in Hamster Brain Homogenate Following the Fenton Reaction Using 3F4 (a) or SAF32 (b) Antibodies.

Uninfected brain homogenate before (C) and after ( $C^F$ ) the Fenton reaction; infected homogenate before (Sc) and after (Sc<sup>F</sup>) the Fenton reaction. The positions of molecular weight standards (15–250 kDa) are also shown in the figure (the lanes labeled M).

were washed three times by ultracentrifugation to remove the  $H_2O_2$ , and the pellets were resuspended in an initial volume of PBS.

The treated samples were separated by SDS-polyacrylamide gel electrophoresis and transferred to a polyvinylidene fluoride membrane (Millipore, Bedford, MA). After blocking, the membranes were incubated for 1 h with horseradish peroxidase (HRP)-conjugated 3F4 (Signet Laboratories, Dedham, MA) or SAF32 (Cayman Chemical, Ann Arbor, MI) antibodies, which recognize different epitopes of the prion protein. After it was washed, the blotted membrane was developed by the ECL+Plus Western Blotting Detection System (Amersham Biosciences, Buckinghamshire, England), according to the manufacturer's instructions. Chemiluminescence signals were analyzed by the Light Capture System (Atto, Tokyo).

Figure 1 illustrates the results of western blot (WB) analysis performed on the samples treated by the Fenton reaction. Following the Fenton reaction, no detectable PrP signals were observed in the uninfected or infected samples. In addition to conventional WB analysis, protein misfolding cyclic amplification (PMCA), a highly sensitive method of detecting minute amounts of PrPSc, 18.19) was performed to detect residual PrPSc. The PMCA procedure followed was based on that reported in our previous study. 16) Briefly, the treated sample was diluted 1:10 in 10% uninfected brain homogenate, and one round of the PMCA reaction was performed by applying 40 cycles of sonication, followed by incubation at 37 °C for 1 h. Next, the process diluting the PMCA product to 1:10 and its subsequent amplification was repeated two times. With regard to the PMCA products, samples (10 µl) collected before and after each round of amplification were mixed with 10 μl

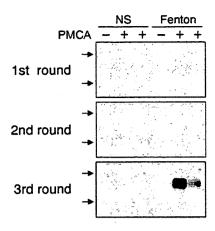


Fig. 2. Protein Misfolding Cyclic Amplification of Samples Treated by the Fenton Reaction.

The samples were analyzed by western blotting before (-) and after (+, in duplicate) each round of amplification following digestion with proteinase K. The lanes labeled NS denote the controls, in which the uninfected brain homogenate (PrP<sup>C</sup> source only) was treated in an identical manner. No signals were detected in these samples. The arrows indicate the positions of the molecular weight markers corresponding to 30 and 20 kDa respectively.

PK solution ( $100 \,\mu\text{g/ml}$ ) and incubated at 37 °C for 1 h. The digested materials were analyzed by WB using the 3F4 antibody, as described above.

Figure 2 illustrates the results of amplification of the samples treated by the Fenton reaction. In the first and second rounds of amplification, no signals from the protease-resistant prion protein (PrPres) were detected in the samples. After three rounds of amplification, PrPres signals were detected in both the duplicated samples. These results suggest that only a trace amount of PrPSc remained in the sample treated by the Fenton reaction.

The sample treated by the Fenton reaction was inoculated intracerebrally  $(20\,\mu l)$  per mouse) into 3–5 Tg52NSE transgenic mice that overexpressed hamster PrP<sup>C</sup>.<sup>20)</sup> The control mice inoculated with the untreated samples developed the disease after an average period of  $45\pm2\,d$  (average  $\pm$  SD, n=5). The mice that were inoculated with the sample subjected only to heat treatment died after an average period of  $46\pm0\,d$  (n=3). The disease onset time significantly decreased in the mice inoculated with the sample treated by the Fenton reaction, but the mice died after an average period of  $163\pm28\,d$  (n=3). Based on the incubation time obtained in a previous study, <sup>16)</sup> the reduction in infectivity was estimated to be approximately  $10^{-6}$ .

According to the Fenton reaction, Fe (II) generates a hydroxyl radical *via* the breakdown of H<sub>2</sub>O<sub>2</sub>. This radical is a highly reactive oxygen species that causes direct damage to nucleic acids and proteins, a property that has been already used in the inactivation of conventional pathogens.<sup>21)</sup> A copper-H<sub>2</sub>O<sub>2</sub> formulation has been proved an effective method for prion decontamination.<sup>10)</sup> We also confirmed that the most effective oxidative deamination by H<sub>2</sub>O<sub>2</sub> was obtained in the presence of Cu<sup>2+</sup>,<sup>15)</sup> but we adopted the iron-H<sub>2</sub>O<sub>2</sub>

formulation for prion decontamination considering all possible risks relating to chemical toxicity against a living body, corrosive impact against medical apparatus, and environmental pollution. The present study indicated that iron ions cause effective oxidative damage to PrPSc in the presence of high concentrations of H<sub>2</sub>O<sub>2</sub>, resulting eventually in an effective reduction in infectivity and a significant decrease in PrPSc levels. Hence, the Fenton reaction using iron ions can be applied in decontamination of fragile, precision, or expensive instruments that are susceptible to PrPSc contamination.

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### Prion inactivation by the Maillard reaction

Kyozo Suyama <sup>a</sup>, Miyako Yoshioka <sup>b</sup>, Mitsugu Akagawa <sup>c</sup>, Yuichi Murayama <sup>b,\*</sup>, Hiroko Horii <sup>b</sup>, Masuhiro Takata <sup>b</sup>, Takashi Yokoyama <sup>b</sup>, Shirou Mohri <sup>b</sup>

Sports Nutrition Department, Sendai University, Miyagi, Japan
 Prion Disease Research Center, National Institute of Animal Health, 3-1-5 Kannondai, Tsukuba, Ibaraki 305-0856, Japan
 Department of Biological Chemistry, Graduate School of Life and Environmental Science, Osaka Prefecture University, Osaka, Japan

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#### Abstract

Since variant Creutzfeldt-Jakob disease (vCJD) has been suspected to be attributable to the infectious agents associated with bovine spongiform encephalopathy (BSE), it is important to prevent the transmission of pathogenic forms of prion protein (PrPSc) through contaminated feeding materials such as meat and bone meal (MBM). Here, we demonstrate that the Maillard reaction employing a formulation of glucose in combination with sodium hydrogen carbonates effectively reduced the infectivity (approximately 5.9-log reduction) of a scrapie-infected hamster brain homogenate. In addition to a bioassay, a protein misfolding cyclic amplification (PMCA) technique, in which PrPSc can be amplified in vitro, was used as a rapid test for assessing PrPSc inactivation. The PMCA analysis also indicated that the PrPSc level in the infected material significantly decreased following the Maillard reaction. Therefore, the Maillard reaction can be employed for the decontamination of large amounts of byproducts such as MBM.

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Keywords: Prion inactivation; Meat and bone meal; Maillard reaction; Bioassay; Protein misfolding cyclic amplification

Prion diseases or transmissible spongiform encephalopathies (TSEs) are neurodegenerative disorders that have been described in humans as Creutzfeldt-Jakob disease (CJD), in sheep and goats as scrapie, and in cattle as bovine spongiform encephalopathy (BSE) [1]. A characteristic of TSE is the accumulation of a protease-resistant, misfolded prion protein (PrPSc), which is a pathogenic variant of the host-encoded prion protein (PrPC) [2,3]. Since PrPSc retains its infectivity after undergoing routine sterilizing processes, harsh physical or chemical sterilization procedures are required for its inactivation [4,5].

BSE is an emerging disease that first appeared in the United Kingdom after 1986 [6]. It appears that the cause of the BSE outbreak was feeding PrPSc-contaminated meat and bone meal (MBM) acquired from rendered carcasses of BSE- or scrapie-infected ruminants to healthy cattle

In the present study, we report a novel method for prion inactivation by chemical modification of proteins based on the principle of the Maillard reaction [10–13]. The Maillard reaction is a very complicated reaction between reducible carbohydrates and reactive amino acid residues of proteins such as lysine, arginine, and tryptophan [14–16]. The reaction products consist of a large variety, for example, brown protein polymers called "melanoidins." Since the Maillard reaction irreversibly modifies the proteins, the structural changes in PrPSc induced by the Maillard reaction are expected to be effective for reducing the infectivity. We therefore investigated the efficacy of the Maillard reaction in scrapie prion inactivation by a bioassay that examined

<sup>[7,8].</sup> It is thought that the recycling of BSE-infected bovine tissues augmented the concentration of PrPSc in commercial MBM, thus causing the subsequent BSE epidemics [9]. Thus, in order to ensure that byproducts such as MBM can be safely used in the future, the development of a technology that can be used for treating large amounts of byproducts at low costs is required.

<sup>\*</sup> Corresponding author. Fax: +81 29 838 8333. E-mail address: ymura@affrc.go.jp (Y. Murayama).