Early detection of chronic wasting disease prions in urine of pre-symptomatic deer by real-time quaking-induced conversion assay

Theodore R. John, Hermann M. Schätzl, and Sabine Gilch

Chronic wasting disease (CWD) is the cervid equivalent of bovine spongiform encephalopathy (BSE), scrapie in sheep and goat or Creutzfeldt-Jakob disease (CJD) in humans. Although transmission studies of CWD prions to humanized transgenic mice or non-human primates suggest a strong species barrier, recent in vitro studies have demonstrated that human PrP can be converted by CWD prions into PrPSc upon adaptation. Therefore, a potential for zoonotic transmission, as exemplified by BSE, cannot be completely excluded.

A huge body of evidence suggests that CWD can be efficiently transmitted horizontally within and between cervid species, which may be the reason for geographical spread and increase in case numbers. Horizontal transmission is explained by the rather unusual peripheral distribution of prions in CWD affected animals and the high susceptibility to the disease by oral infection. Unlike in most other prion diseases, CWD prions can be found in a wide variety of tissues, such as skeletal and cardiac muscle or kidney, in addition to the lymphoreticular system and blood. Furthermore, they are shed in significant amounts in saliva, urine or feces, which enables oral infection of animals by foraging on contaminated pastures. In addition, it has been demonstrated that prions can persist in soil and that water in endemic areas can contain CWD-associated PrPSc.

Currently, disease surveillance is mainly based on testing hunter harvested animals. Since this testing is not obligatory, it depends on the compliance of hunters. CWD test systems are based on the detection of proteinase K resistant PrPSc, either by immunoblot, ELISA or immunohistochemistry. The main materials used for this are brain stem homogenates and tonsil or rectoanal mucosa-associated lymphoid tissue (RAMALT) biopsies, which requires anesthesia of animals. Although transmission studies of CWD prions to humanized transgenic mice or non-human primates suggest a strong species barrier, recent in vitro studies have demonstrated that human PrP can be converted by CWD prions into PrPSc upon adaptation. Therefore, a potential for zoonotic transmission, as exemplified by BSE, cannot be completely excluded.

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Since RT-QuIC offers advantages over PMCA, e.g., it does not require sonication, we have chosen this method for testing its usefulness in the detection of CWD prions in deer urine and feces. We demonstrate that CWD prions are detectable in urine...
of orally infected deer prior to the onset of clinical symptoms. Furthermore, we show that fecal extracts can be used as a seed in RT-QuIC assays. Thereby, we were able to detect CWD prions in fecal extracts collected at later stages of the disease. This study provides the first evidence that RT-QuIC can be successfully used for the preclinical diagnosis of CWD in specimens that are available by non-invasive methods.

**Results**

Detection of CWD in tissues or body fluids that are easily available and do not require invasive methods is highly desirable and would enable improved surveillance of the disease in free-ranging cervids. Therefore, our aim was to adapt the RT-QuIC assay for detection of CWD prions in deer urine and feces.

At first, we analyzed the sensitivity of detection of CWD prions using infected brain homogenate as a seed and recombinant cervid PrP as a substrate. Serial 10-fold dilutions of infected (CWD⁺) and non-infected (CWD⁻) brain homogenates (10%) were prepared and RT-QuIC reactions were seeded with these dilutions (Fig. 1A). For the lower dilutions, incorporation of Thioflavin T indicative for prion conversion was detected already after approximately 2 h. Conversion was detectable up to a brain homogenate dilution of $2 \times 10^{-7}$, whereas the $2 \times 10^{-8}$ dilution remained negative. RT-QuIC assays of serial dilutions of non-infected brain homogenates resulted in a baseline signal, except for very little spontaneous conversion for the $10^{-6}$ dilution at a late reaction time point.
In order to verify whether detection is inhibited when CWD prions are present in urine, we performed spiking experiments. CWD positive brain homogenate was serially diluted in deer urine derived from a CWD negative animal, with dilutions ranging between 10^{-3} and 10^{-4}. Each sample was used as a seed in RT-QuIC and an endpoint titration was performed. In Figure 1B, the results for the 10^{-3} and 10^{-4} brain homogenate dilutions in urine are depicted. The sensitivity of detection is similar to that in brain homogenate, with an overall detection limit of 2 \times 10^{-7} in the 10^{-3} brain homogenate dilution (upper panel) and 2 \times 10^{-6} for the 10^{-4} brain homogenate dilution (lower panel), respectively.

Altogether, these data demonstrate that CWD prions can be detected in both brain homogenate and urine. The sensitivity does not differ significantly from that of detection in brain homogenate, indicating that urine does not or not significantly inhibit RT-QuIC reactions.

Next we were interested in whether it is possible to detect CWD prions in urine samples of orally infected mule deer or white-tailed deer. Depending on the genotype at codon 96, white-tailed deer in this study were tested positive for CWD prions in tonsil biopsies as early as 8.4 mo post infection (96GG and 96GS), whereas animals with the genotype 96SS remained negative until > 11.4 mo post infection. Clinical symptoms were observed at the earliest at 17.2 mo post infection in the 96GG animals, both 96GS and 96SS animals exhibited longer incubation times of the disease. Mule deer showed clinical signs between 16.2 and 25.9 mo post inoculation. Samples of two animals (W804; mule deer and W1004; white-tailed deer) collected 5, 13 and 16 mo post oral inoculation, respectively, were chosen for analysis. Notably, all samples were collected at a pre-symptomatic stage of the disease. A 2 \times 10^{-3} dilution was used to seed RT-QuIC reactions, recombinant cervid PrP was employed as a substrate. Urine of three different non-infected deer served as a negative control, and these samples did not induce conversion (Fig. 2, nos. 1 and 2 and nos. 2 and 3, respectively). However, samples of both infected animals taken at 13 and 16 mo post inoculation were positive, whereas urine collected five months post inoculation did not contain detectable amounts of seed. In animal W804 (Fig. 2, upper panel), the conversion reaction started after approximately 18 h for the 16 mo sample and after approximately 30 h for the 13 mo sample, indicating that the seeding activity in the 16 mo sample was higher than in that collected 13 mo post inoculation. In the samples of animal W1004 (Fig. 2, lower panel), conversion started after 10 h in the 13 mo sample and after approximately 28 h in the 16 mo sample, respectively.

**Figure 2.** Detection of CWD prions in urine of pre-clinical deer. Urine of orally infected animals (W804 = mule deer; W1004 = white-tailed deer) collected 5, 13 and 16 mo post infection, respectively, was diluted 20-fold in RT-QuIC buffer and was then used to seed RT-QuIC reactions as described using cervid PrP as a substrate. Urine of non-infected animals served as a negative control.
In summary, we demonstrate that CWD prions can be detected by RT-QuIC in urine of orally infected white-tailed deer and mule deer at a pre-symptomatic stage of the disease.

Since urine might be difficult to collect from free-ranging animals, we were interested in whether it is possible to use feces extracts as a seed in RT-QuIC reactions. Fecal samples can be easily collected, both from captive and free-ranging animals, and testing of such samples, even if they cannot be assigned to a certain individual animal, might add valuable information about the distribution of CWD in certain areas or the occurrence of new cases in areas that were assumed to be free of CWD. To this end, fecal extracts of animal W1004, collected 20 or 30 mo post inoculation were prepared. RT-QuIC reactions were seeded with either undiluted extract or a $2 \times 10^{-3}$ dilution thereof. Feces extracts of non-infected animals were used as a negative control (Fig. 3).

Although no seeding activity was detectable in the 20 mo sample, both dilutions of the 30 mo sample were positive. However, conversion started late even in the undiluted extracts compared with the results of the urine or brain homogenate samples, which is indicative for low sensitivity.

Nevertheless, we provide strong evidence that it is possible to detect CWD prions in feces of orally inoculated mule deer by RT-QuIC assay.

**Discussion**

With this study, we provide proof of concept that RT-QuIC is a highly useful method for the detection of CWD prions in urine and feces of orally infected deer. Of note, we were able to seed conversion reactions using urine collected at an early presymptomatic stage of the disease. Fecal extracts that were tested positive were taken at stages of the disease when clinical symptoms already might have occurred.

It has been shown previously that RT-QuIC can be used for the detection of CWD prions using full-length cervid PrP as a substrate. In this study, a $10^{-6.3}$ dilution of a 10% brain homogenate was used to optimize the NaCl concentration in the reaction buffer. In our endpoint titration experiment (Fig. 1A) the detection limit was reached at a dilution of $2 \times 10^{-7}$ of a 10% brain homogenate, with a NaCl concentration of 300 mM. A similar sensitivity of detection was observed in our spiking experiments. Here, the detection limit was an overall dilution of $2 \times 10^{-7}$, which is the combination of a $10^{-3}$ brain homogenate dilution in urine, and the further $2 \times 10^{-4}$ dilution in RT-QuIC buffer, indicating that urine does not negatively influence the detection of seeding activity. Spontaneous conversion was only observed in the $2 \times 10^{-6}$ dilution after approximately 42 h when reactions were seeded with negative brain homogenates and in the spiking experiment (Fig. 1B, lower panel) in an overall brain homogenate dilution of $2 \times 10^{-5}$ upon a reaction time of approximately 36 h. These results indicate that at dilutions higher than $2 \times 10^{-5}$ positive results have to be carefully evaluated since spontaneous conversion might occur.

Conversion efficiency of the cervid PrP substrate might be negatively influenced by differences in the primary structure of the seed due to polymorphisms. In order to confirm that differences between the primary structures of seed and substrate are tolerated we used recombinant mouse PrP as a substrate and obtained similar sensitivity and reaction kinetics as with cervid PrP (data not shown).

When using deer urine for seeding RT-QuIC, samples from both infected animals collected at 13 and 16 mo post oral inoculation resulted in positive signals, whereas the 5 mo sample and urine from non-infected deer were negative. However, collection time points of 13 and 16 mo are still in the pre-symptomatic stage of disease, since mule deer developed clinical signs between 16.2 and 25.9 mo post oral inoculation and white-tailed deer even later with the earliest signs 17.2 mo post inoculation. Until now, infectivity in urine of CWD infected animals has been determined by PMCA or bioassays using transgenic cervidized mice, however, in this study only urine from terminally sick white-tailed deer has been analyzed. When PMCA was combined with a highly sensitive immunoassay (surround optical fiber immunoassay; SOFIA), prion-disease associated seeding activity was found in white-tailed deer urine collected approximately 30 mo (891 d) post inoculation, which is still later in the course of the disease.

**Figure 3.** CWD prions are detectable in feces by RT-QuIC. Extracts of white-tailed deer (W1004) feces collected at 20 or 30 mo post oral inoculation were seeded either undiluted or in a $2 \times 10^{-1}$ dilution into quadruplicate reactions. Fecal extracts from non-infected deer served as a negative control.
Prion 257

Brain homogenate preparation. A section of brain stem from a CWD+ or CWD− mule deer (Odocoileus hemionus) was excised, weighed, then transferred to a Dounce homogenizer. PBS [20 mM sodium phosphate (pH 7.4), 130 mM NaCl] was added to 10% (w/v) and the tissue was homogenized first with the loose, then the tight plunger. Homogenates were stored at −80°C.

Source of urine and feces. Deer urine and feces of known CWD status were collected from captive animals involved in a study in which white-tailed (Odocoileus virginianus) and mule deer (Odocoileus hemionus) were orally inoculated with CWD brain homogenate followed by monthly collection of blood, saliva, urine and feces. A herd of captive mule deer in Pullman, WA (a non-CWD endemic area) served as the source of CWD negative samples. After collection, samples were stored frozen.

Preparation of deer feces extracts. Single pellets of deer feces were weighed, then placed into 8 ml polypropylene bottles (Nalgene). Buffer [20 mM sodium phosphate, pH 7.1, 130 mM NaCl, 0.05% (v/v) Tween 20, 1 mM PMSF, 1× Roche Complete Protease Inhibitors (11-697-498-001)] was added to 20% (w/v) final concentration. Bottles were placed onto a platform rotary shaker at −30° angle and were shaken gently for 1 h at room temperature. About 1.8 ml of solution was then transferred to a 2.0 ml screw cap microcentrifuge tube and insoluble debris was pelleted by centrifugation at 18,000 × g for 5 min at room temperature. Supernatants were transferred to fresh 2.0 ml screw cap microcentrifuge tubes and the samples were stored frozen.

Purification of recombinant prion protein. A plasmid containing a DNA sequence coding for a cervid prion protein (residues 24–234, accession AF156185; described in ref. 27) was a generous gift from Dr Byron Caughey (NINDS, NIH, Rocky Mountain Laboratories). Recombinant PrP was expressed in E. coli Rosetta DE3 and purified as described. In brief, cells were grown for 22–24 h in LB media supplemented with Overnight Express Autoinduction System 1 (EMD, 71300-4). Harvested cells were subjected to 2 freeze/thaw cycles in liquid nitrogen and the cell paste was stored at −20°C. Inclusion bodies containing rPrP were prepared from the cell paste using BugBuster Master Mix (EMD, 71456-3) to lyse the cells. Inclusion bodies were then washed twice with 0.1× BugBuster Master Mix, pelleted by centrifugation and stored at −20°C. Proteins from the inclusion bodies were denatured in 8 M guanidine-HCl, then batch-bound to Ni-NTA Superflow resin (Qiagen, 30430) that had been pre-equilibrated in denaturing buffer [100 mM sodium phosphate, 10 mM Tris, 6 M guanidine-HCl (pH 8.0)]. This was poured into an AKTA XK-16 column and attached to an AKTA Explorer chromatography system for protein purification at room temperature. Recombinant PrP was refolded on the column using a linear gradient of 100 mM sodium phosphate, 10 mM Tris (pH 8.0), then eluted using another linear gradient of 100 mM sodium phosphate, 10 mM Tris, 500 mM imidazole (pH 5.8). Eluted protein was collected in 2 ml fractions; 2 ml dialysis buffer [10 mM sodium phosphate (pH 5.8)] was previously added to the collection tubes to dilute the product. Fractions containing rPrP were pooled, filtered through 0.22 µm, then added to SnakeSkin dialysis tubing (7000 MWCO, Thermo Scientific, 68700) for overnight dialysis at 4°C. Following dialysis, the sample was again filtered through 0.22 µm and protein concentration was determined using a Pierce BCA Protein Assay Kit (Thermo Scientific, 23227). Aliquots of purified rPrP were stored at −80°C.

RT-QuIC assay. Real time QuIC was performed as described. Briefly, reactions were set up in a buffer containing 20 mM sodium phosphate (pH 6.9), 300 mM NaCl, 1 mM EDTA, 10 µM Thioflavin T, 0.1 mg/ml rPrP substrate. A reaction cocktail was prepared and 98 µl aliquots were added to the wells of a 96-well optical bottom plate (Nalge Nunc International, 265301). Quadruplicate reactions were seeded with 2 µl of brain homogenate, urine or feces extract that were diluted in 20 mM sodium phosphate (pH 6.9), 130 mM NaCl, 0.1% (w/v) SDS, 1× N2 Supplement (Invitrogen, 17502048); note that the final detergent concentration in each reaction was 0.002%. The plate was sealed with with Nunc Amplification Tape (Nalge Nunc International, 232702) and placed in a BMG Labtech FLUOstar Omega fluorescence plate reader that was pre-heated to 42°C. A program of 1 min rest followed by 1 min shaking (700 rpm, double orbital) with fluorescence readings (450 nm excitation, 480 nm emission) every 15 min was run. This cycling was allowed to continue for 50 h (200 readings). Data was plotted as the average of quadruplicate reactions using every third time point (45 min) with GraphPad Prism software.

Disclosure of Potential Conflicts of Interest
No potential conflict of interest was disclosed.
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