



# Infectious Prions in the Pregnancy Microenvironment of Chronic Wasting Disease-Infected Reeves' Muntjac Deer

Amy V. Nalls, Erin McNulty, Clare E. Hoover, Laura A. Pulscher, Edward A. Hoover, Candace K. Mathiason

Department of Microbiology, Immunology and Pathology, Colorado State University, Fort Collins, Colorado, USA

**ABSTRACT** Ample evidence exists for the presence of infectious agents at the maternal-fetal interface, often with grave outcomes to the developing fetus (i.e., Zika virus, brucella, cytomegalovirus, and toxoplasma). While less studied, pregnancy-related transmissible spongiform encephalopathies (TSEs) have been implicated in several species, including humans. Our previous work has shown that prions can be transferred from mother to offspring, resulting in the development of clinical TSE disease in offspring born to muntjac dams infected with chronic wasting disease (CWD) (1). We further demonstrated protein misfolding cyclic amplification (PMCA)-competent prions within the female reproductive tract and in fetal tissues harvested from CWD experimentally and naturally exposed cervids (1, 2). To assess whether the PMCA-competent prions residing at the maternal-fetal interface were infectious and to determine if the real-time quaking-induced conversion (RT-QuIC) methodology may enhance our ability to detect amyloid fibrils within the pregnancy microenvironment, we employed a mouse bioassay and RT-QuIC. In this study, we have demonstrated RT-QuIC seeding activity in uterus, placentome, ovary, and amniotic fluid but not in allantoic fluids harvested from CWD-infected Reeves' muntjac dams showing clinical signs of infection (clinically CWD-infected) and in some placentomes from pre-clinically CWD-infected dams. Prion infectivity was confirmed within the uterus, amniotic fluid, and the placentome, the semipermeable interface that sustains the developing fetus, of CWD-infected dams. This is the first report of prion infectivity within the cervid pregnancy microenvironment, revealing a source of fetal CWD exposure prior to the birthing process, maternal grooming, or encounters with contaminated environments.

**IMPORTANCE** The facile dissemination of chronic wasting disease within captive and free-range cervid populations has led to questions regarding the transmission dynamics of this disease. Direct contact with infected animals and indirect contact with infectious prions in bodily fluids and contaminated environments are suspected to explain the majority of this transmission. A third mode of transmission, from mother to offspring, may be underappreciated. The presence of pregnancy-related prion infectivity within the uterus, amniotic fluid, and the placental structure reveals that the developing fetus is exposed to a source of prions long before exposure to the infectious agent during and after the birthing process or via contact with contaminated environments. These findings have impact on our current concept of CWD disease transmission.

**KEYWORDS** CWD, maternal, pregnancy, prions

**T**ransmissible spongiform encephalopathies (TSEs), or prion diseases, are proteinaceous infectious neurodegenerative diseases affecting animals, including humans (3, 4). The disease is caused by the accumulation of an aberrant misfolded form

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Address correspondence to Candace K. Mathiason, [candace.mathiason@colostate.edu](mailto:candace.mathiason@colostate.edu).

A.V.N. and E.M. contributed equally to this article.

(protease-resistant, Pr<sup>PRE5</sup>) of a normal cellular protein, the prion protein (Pr<sup>PC</sup>) (5). A hallmark of all TSE-infected hosts is their ability to harbor prion infectivity in tissues and bodily fluids throughout the protracted asymptomatic, or silent-carrier, phase of disease, which can last years, decades, or perhaps an entire life span (6, 7).

The transmission of prions, while remaining poorly understood, has obvious animal and human public health significance. It is clear that TSEs are spread by horizontal means of transmission via animal to animal and by environmental contact with prions, as well as by medical means, including blood transfusion or iatrogenic exposure (reviewed in references 3 and 4). Less studied is the potential for prion transmission from mother to offspring. Although it is rare for pathogens to cross the placental barrier and infect the fetus, certain viruses, bacteria, and protists have demonstrated this ability (8). Therefore, it is not surprising that there are several reports of prions in the female reproductive tract (9, 10) and pregnancy-associated milieu (11–14). These reports, in conjunction with the recent demonstration of a silent variant Creutzfeldt-Jakob disease (vCJD) carrier state (1 in 2,000 people) in the United Kingdom (15) and with the first report of chronic wasting disease (CWD) in Europe (16), mandate a thorough investigation of the role maternal transmission plays in prion transmission and disease pathogenesis.

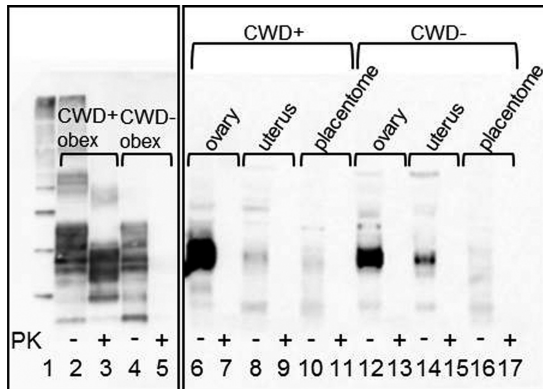
Few studies have been conducted to assess the presence of prions in female reproductive and pregnancy-related tissues and in fluids from women displaying clinical CJD. The studies that have been published provide conflicting results. Uterine and gestational tissues from a 41-year-old woman with clinical CJD did not reveal deposition of the abnormal prion protein (Pr<sup>PRE5</sup>) (17). Nor has there been a report of the manifestation of clinical TSE symptoms in children born to women with vCJD (18). Yet other studies demonstrate prions in tissues of the human reproductive system. Proteinase K (PK)-resistant prions were demonstrated in the uterus and ovaries of a nonpregnant 25-year-old woman diagnosed with vCJD (9, 19). Infectious prions were detected by mouse bioassay in blood, colostrum, and placental tissues from a 38-year-old woman diagnosed with CJD between her 20th and 30th weeks of pregnancy (14). Considerable evidence exists for maternal scrapie transmission resulting in TSE disease and perinatal trafficking of prions (10, 12, 13, 20–29). In addition, maternal transmission of feline spongiform encephalopathy (FSE) has been suspected (30). Vertical transmission of bovine spongiform encephalopathy (BSE) has not been realized (31–33), yet enhanced risk factors for maternal/vertical transmission have been reported (32, 34); vertical transmission has been evidenced in transgenic (Tg) mouse studies (35). Thus, the biological significance associated with gestational prion exposure remains unanswered.

Our development and use of a small polyestrous cervid host, the Reeves' muntjac deer, has permitted investigation of questions related to CWD mother-to-offspring prion transmission in the native cervid host (1). We have validated our muntjac system by demonstrating the presence of protein misfolding cyclic amplification (PMCA)-competent prions in *in utero*-harvested tissues from free-range naturally exposed elk (2). Building on these studies we have further assessed CWD prions (Pr<sup>PCWD</sup>) at the maternal-fetal interface.

Using bioassay and real-time quaking-induced conversion (RT-QuIC), we undertook this study to determine the infectious nature and RT-QuIC seeding activity of (i) female reproductive tissues and fluids associated with the pregnancy microenvironment and (ii) the placentome, which is the semipermeable interface between mother and fetus throughout pregnancy.

## RESULTS

**Maternal reproductive tissue.** Our previous studies (1, 2) led to the discovery that Pr<sup>PRE5</sup> deposition in maternal tissues of the reproductive system was too low to detect by immunohistochemistry. This led us to the use of PMCA (1, 2) and now RT-QuIC and bioassay to investigate the seemingly low concentration of prions within these tissues and the pregnancy microenvironment. Here, we further investigated these tissues for



**FIG 1** Western blot demonstrating the presence of PrP<sup>C</sup> in muntjac reproductive tissues. Lane 3 demonstrates that PrP<sup>RES</sup> is present in a CWD-infected muntjac obex. Lane 5 shows complete proteinase K (PK) digestion of PrP<sup>C</sup> in a negative-control muntjac obex. Lanes 6, 8, 10, 12, 14, and 16 demonstrate PrP<sup>C</sup> detection in ovary, uterus, and placentome tissues from CWD-infected and negative-control muntjac. Lane 1, molecular weight marker. No PrP<sup>RES</sup> was detected in these reproductive tissues after PK digestion (2-mg tissue equivalents/lane).

the presence of the normal cellular form of the prion protein, PrP<sup>C</sup>. The detection of the normal cellular form of the prion protein in uterus, ovary, and placentome, i.e., pregnancy-related tissues (Fig. 1), suggests the presence of a substrate capable of fueling PrP<sup>C</sup> conversion. To this end, we detected RT-QuIC PrP<sup>C</sup> seeding activity (amyloid formation) in uterine (7/7 samples) and ovary (5/7 samples) tissue harvested from muntjac dams showing clinical signs of infection with CWD agent (clinically CWD-infected dams) ( $P < 0.0001$ ). No amyloid formation was detected in the uterine (0/4) or ovary (0/3) tissue from pre-clinically CWD-infected muntjac dams (Table 1 and Fig. 2). We have further demonstrated the presence of infectious prions in uterine tissues harvested from a clinically ill CWD-infected muntjac dam (animal 45) by mouse bioassay: 8/8 mice inoculated with uterine tissue from dam 45 developed clinical signs consistent with TSE disease (ataxia, weight loss, and stiff tail) between 237 and 343 days postinfection (dpi). All 8 clinically ill mice revealed demonstrable PrP<sup>RES</sup> deposition in brain tissue as determined by Western blotting (Fig. 3). RT-QuIC seeding activity was demonstrated in brain tissue of all 8 mice and in spleen of 7/8. Note that 1/9 mice suffered intercurrent death (Table 2; Fig. 3). All mice inoculated with uterine tissue from a naive muntjac dam (animal 64) remained healthy for the same period of time and negative for PrP<sup>RES</sup> deposition and amyloid formation (Table 2 and Fig. 3).

**Maternal-fetal interface at the placentome.** We were not able to detect PrP<sup>RES</sup> deposition in placentomes by Western blot analysis or immunohistochemistry (data not shown), similar to other maternal reproductive tissues, yet we were able to demonstrate PMCA seeding activity in tissues at this intimate interface between mother and fetus (1, 2). Thus, RT-QuIC and bioassay studies were initiated. Significant RT-QuIC seeding activity was evident in 9/11 placentomes harvested from clinically CWD-infected dams ( $P \leq 0.0018$ ) (Table 1 and Fig. 2). Mice intracranially (i.c.) inoculated with homogenate of placentome sample 1 from clinically CWD-infected dam 45 (Fig. 2, 45-P1) remained healthy for 500 dpi and lacked PrP<sup>CWD</sup> deposition as determined by Western blotting or RT-QuIC seeding activity in brain or spleen tissues (Table 2). RT-QuIC analysis of dam 45-P1 suggested a lack of amyloid fibrils in this placentome (Fig. 2). Two placentomes with the highest RT-QuIC seeding activity rates (Fig. 2, 47-P1 and 47-P5) were assessed by bioassay. Prion infectivity was revealed, and 7/8 mice developed terminal TSE disease between 180 and 270 dpi (Table 2). PK-resistant prions were evident in brain tissue of 6/8 mice by Western blotting (Table 2 and Fig. 3), and upon RT-QuIC analysis all 8 mice were demonstrated to have RT-QuIC seeding activity in brain and spleen tissues. Note that 1/9 mice suffered intercurrent death (Table 2 and Fig. 3). RT-QuIC analysis of placentome tissue from pre-clinically CWD-infected dams

**TABLE 1** Muntjac dam and fetal study animals, their disease and gestational stages, and reproductive tissue and fluid RT-QuIC results

Dam disease status and identification no. (fetus)	Inoculum (amt and route) <sup>a</sup>	Dam disease stage (no. of mos. p.i.) <sup>b</sup>	Fetal gestational stage (no. of mos.) <sup>c</sup>	Tissue or fluid <sup>f</sup>	QuIC median rate <sup>e,g</sup>
CWD positive					
11	Brain (1 g p.o.)	25 <sup>c</sup>	NA	<b>Uterus</b>	<b>0.0470****</b>
				Ovary	0.0107
15	Brain (0.5 g s.c and 2 g p.o.)	26 <sup>c</sup>		<b>Uterus</b>	<b>0.0878****</b>
				Ovary	0
43	Brain (0.35 g s.c. and 0.5 g p.o.)	23 <sup>c</sup>		<b>Uterus</b>	<b>0.1060****</b>
				<b>Ovary</b>	<b>0.0991****</b>
45 (58)		22 <sup>d</sup>	4.8	<b>Uterus</b>	<b>0.0700****</b>
				<b>Ovary</b>	<b>0.0799****</b>
				Placentome 1	0
				<b>Placentome 2</b>	<b>0.0598****</b>
				<b>Placentome 3</b>	<b>0.0660****</b>
				<b>Placentome 4</b>	<b>0.0980****</b>
				<b>Placentome 5</b>	<b>0.0880****</b>
				Allantoic fluid	0
46		24 <sup>c</sup>	NA	<b>Uterus</b>	<b>0.0793****</b>
				<b>Ovary</b>	<b>0.0714****</b>
47 (57)		20 <sup>d</sup>	>4.5	<b>Uterus</b>	<b>0.0611****</b>
				<b>Ovary</b>	<b>0.0676****</b>
				<b>Placentome 1</b>	<b>0.1200****</b>
				<b>Placentome 2</b>	<b>0.1010****</b>
				<b>Placentome 3</b>	<b>0.0176**</b>
				<b>Placentome 4</b>	<b>0.1092****</b>
				<b>Placentome 5</b>	<b>0.1084****</b>
				Placentome 6	0
				<b>Amniotic fluid</b>	<b>0.0527****</b>
				Allantoic fluid	0
50 (59)		23 <sup>c</sup>	2	<b>Amniotic fluid</b>	<b>0.0185**</b>
78		22	NA	<b>Uterus</b>	<b>0.0592****</b>
				<b>Ovary</b>	<b>0.0392****</b>
16 (115)		7	3.5	Uterus	0
				Ovary	0
				Placentome 1	0
				Placentome 2	0
				Placentome 3	0
				Placentome 4	0
				Placentome 5	0
				Amniotic fluid	0
				Allantoic fluid	0
19 (116)		7	3.5	Uterus	0
				Placentome 1	0
				<b>Placentome 2</b>	<b>0.0194***</b>
				Placentome 3	0
				<b>Placentome 4</b>	<b>0.0172**</b>
				Placentome 5	0
				Amniotic fluid	0
				Allantoic fluid	0
73 (89)		12	3	Uterus	0
				Ovary	0
				<b>Placentome 1</b>	<b>0.0086*</b>
				<b>Placentome 2</b>	<b>0.0283**</b>
				Placentome 3	0
				<b>Placentome 4</b>	<b>0.0168*</b>
				Placentome 5	0
				<b>Placentome 6</b>	<b>0.0086*</b>
77 (90)		12	3	Uterus	0
				Ovary	0
				Placentome 2	0
				Placentome 3	0
				Placentome 4	0
				Placentome 5	0
				<b>Placentome 6</b>	<b>0.0245****</b>
				Amniotic fluid	0
84 (98)		5	7	<b>Placentome 1</b>	<b>0.0106*</b>
				Placentome 2	0
				Placentome 3	0
				<b>Placentome 4</b>	<b>0.0081*</b>
				Amniotic fluid	0

(Continued on next page)

**TABLE 1** (Continued)

Dam disease status and identification no. (fetus)	Inoculum (amt and route) <sup>a</sup>	Dam disease stage (no. of mos. p.i.) <sup>b</sup>	Fetal gestational stage (no. of mos.) <sup>c</sup>	Tissue or fluid <sup>f</sup>	QuIC median rate <sup>e,g</sup>
CWD negative					
62	Naive	None	NA	Uterus Ovary	0
64	Naive		5	Ovary Placentome 1 Placentome 2 Amniotic fluid	
75	Brain (0.5 g s.c. and 0.5 g p.o.)		NA	Uterus Ovary	
86	Naive		NA	Uterus	
107 (119)	Naive		2	Amniotic fluid	

<sup>a</sup>s.c., subcutaneously; p.o., *per os*.

<sup>b</sup>Terminal CWD clinical disease in experimentally inoculated muntjac dams manifests between 23 and 26 months postinfection (p.i.).

<sup>c</sup>Euthanized due to terminal clinical CWD disease, with symptoms of hypersalivation, weight loss, and ataxia.

<sup>d</sup>Early signs of clinical CWD (weight loss).

<sup>e</sup>The full-term gestational period for Reeves' muntjac deer is 7 months. NA, not available (no fetus at necropsy).

<sup>f</sup>Boldface indicates statistical significance (\*,  $P \leq 0.05$ ; \*\*,  $P \leq 0.01$ ; \*\*\*,  $P \leq 0.001$ ; \*\*\*\*,  $P < 0.0001$ ).

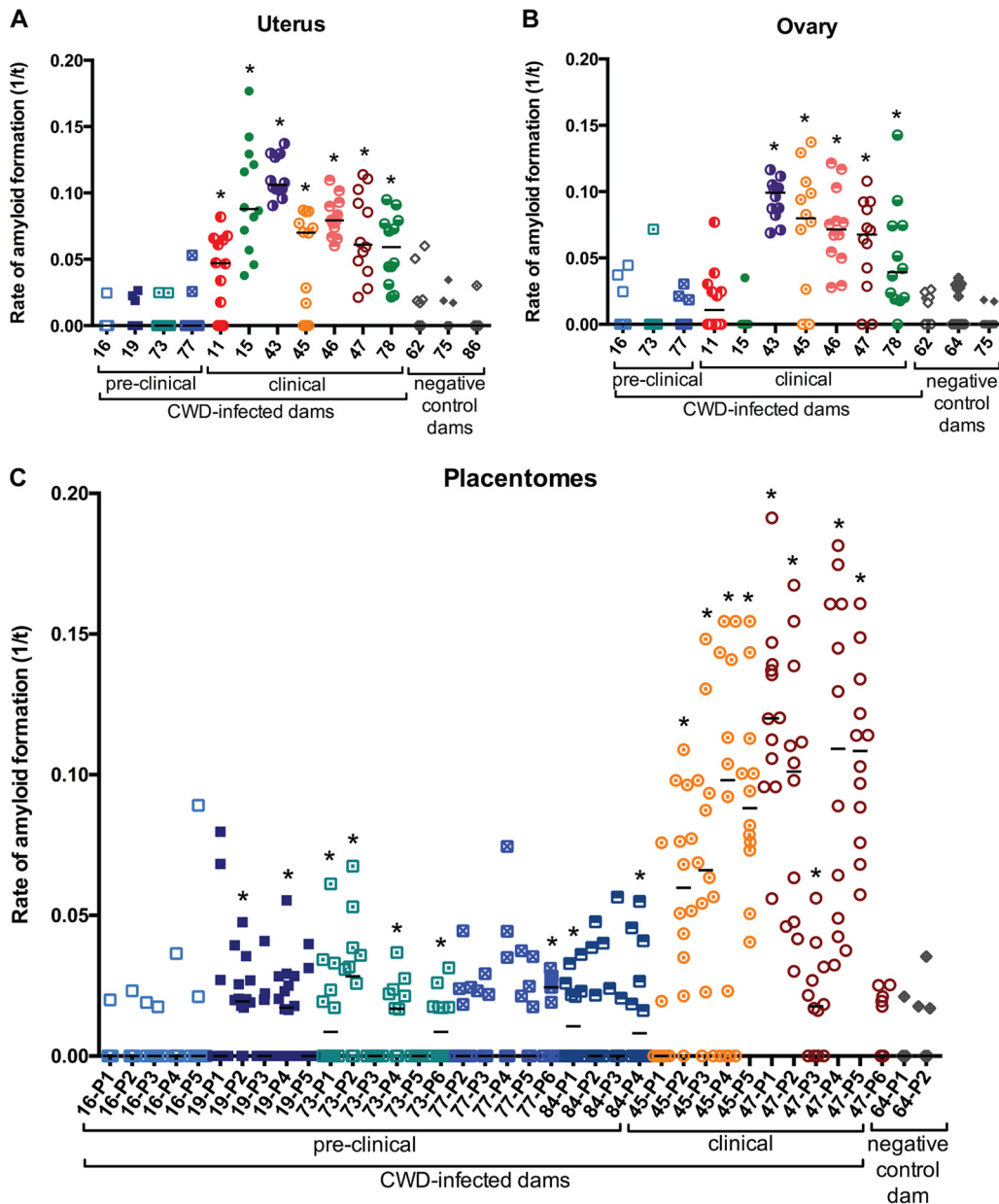
<sup>g</sup>Values represent 1/time to the ThT fluorescence threshold.

( $n = 5$ ) indicated low to moderate levels of RT-QuIC seeding activity in 9/25 placentomes tested ( $P = 0.0004$  to  $0.0476$ ); 4/5 pre-clinically infected dams had at least one RT-QuIC-positive placentome (Table 1 and Fig. 2). We initiated a mouse bioassay of these tissues to assess infectivity. RT-QuIC analysis of two placentomes and bioassay of one placentome from a naive muntjac dam (dam 64) showed neither amyloid formation nor infectivity (Tables 1 and 2; Fig. 2 and 3).

**Fetal microenvironment.** Fluids collected from the fetal amniotic and allantoic sacs of the aforementioned muntjac pregnancies were concentrated through lyophilization, dialyzed, and assessed by RT-QuIC and/or bioassay (Table 2; Fig. 3 and 4). RT-QuIC seeding activity was present in amniotic fluid harvested from clinically ill CWD-infected dams (two, dams 59 and 47) in 79% of RT-QuIC replicates ( $n = 12$ /each) but was absent in allantoic fluid tested from the same pregnancies (Table 1; Fig. 4) and in amniotic and allantoic fluids harvested from pre-clinically CWD-infected dams ( $n = 4$ ) (Table 1; Fig. 4). Mouse bioassay of amniotic fluid harvested from clinically ill CWD-infected dam 47 confirmed the presence of very low concentrations of infectious prions. One of nine (1/9) mice presented with signs consistent with TSE disease at 320 dpi and was euthanized. The remaining mice remained asymptomatic and were euthanized at 500 dpi. Although Pr<sup>PCWD</sup> was not demonstrated in brain tissue of any of these mice by Western blotting, further analysis by RT-QuIC revealed prion seeding activity in brain (1/9 mice; the clinically ill mouse) and spleen (2/9 mice, including the clinically ill mouse) (Table 2; Fig. 3). None (0/9) of the mice inoculated with allantoic fluid from CWD-positive dams developed TSE disease or demonstrated Pr<sup>PRE5</sup> deposition or RT-QuIC seeding activity in brain or spleen tissues (Table 2; Fig. 3). RT-QuIC analysis and bioassay of amniotic and allantoic fluids harvested from naive muntjac dams showed neither amyloid formation nor infectivity (Table 2; Fig. 3).

## DISCUSSION

Our previous work demonstrates that prions can be transferred from mother to offspring, resulting in the development of clinical TSE disease in offspring born to CWD-infected muntjac dams (1). Because the offspring were born to and raised by their infected mothers, we were unable to parse the point source of infectivity, i.e., whether infection originated from contact with maternal prions during parturition or shortly thereafter or via *in utero* exposure. We addressed this question by revealing amplification-competent prions in *in utero*-harvested fetal tissues from CWD-infected muntjac dams regardless of disease or gestational stage (1) and in reproductive and fetal tissues harvested from free-range cervid dams naturally exposed to CWD but not presenting clinical signs of disease (preclinical dams) (2). What remained unanswered

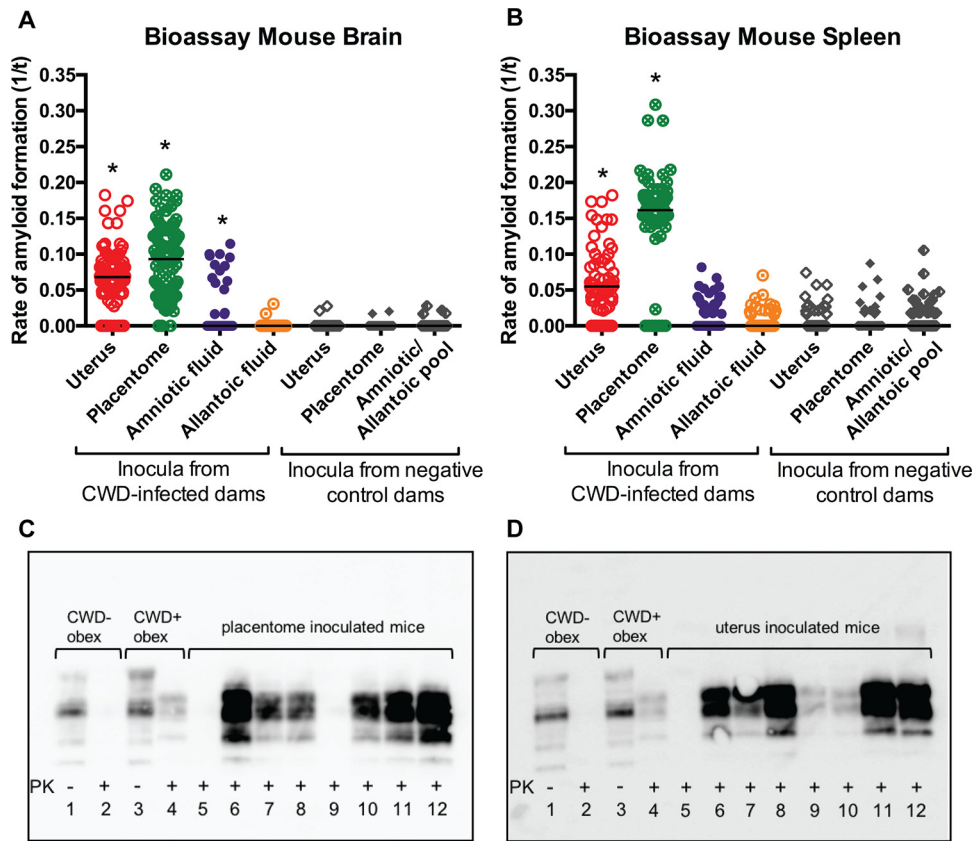


**FIG 2** RT-QuIC detection of amyloid formation in muntjac dam uterus, ovary, and placentomes. Detection of amyloid formation is displayed as reaction rates (1/time [t] to ThT fluorescence threshold). Square, pre-clinically CWD-infected dams; circles clinically CWD-infected dams; diamonds negative-control dams. Horizontal lines indicate the medians ( $n = 12$  replicates for CWD-infected dams and  $n = 12$  to 16 replicates for CWD-negative dams). Statistical significance is indicated above the sample replicates with an asterisk ( $P < 0.0001$  for uterus and ovary;  $P \leq 0.0476$  for preclinical placentome;  $P \leq 0.0018$  for clinical placentome).

was whether the prions detected in *in utero*-harvested tissues were infectious. Here, we demonstrate, for the first time, prion infectivity within the maternal reproductive and pregnancy-related tissues and milieu of clinically CWD-infected dams. We further assessed tissues by an *in vitro* conversion assay, RT-QuIC, to determine if correlates could be drawn between the prion seeding activity detected by this assay and infectivity realized by animal bioassay. We found a direct correlation between RT-QuIC seeding activity and bioassay infectivity in samples harvested from clinically CWD-infected dams.

In our experience, CWD-infected muntjac dams showing clinical signs of disease maintain fecundity, successfully breeding and producing full-term viable and nonviable





**FIG 3** Detection of amyloid formation in cervid transgenic mice [Tg(CerPrP-E226)5037<sup>+/-</sup>] inoculated with placentome and uterine tissues and birthing fluids from CWD-infected muntjac. (A and B) RT-QuIC detection of amyloid formation displayed as reaction rates (1/time to ThT fluorescence threshold). Replicates are shown as circles for mice inoculated with material from CWD-infected muntjac (uterus and placentome, *n* = 8 mice per cohort; amniotic and allantoic fluids, *n* = 9 mice per cohort; 12 replicates each) and as diamonds for mice inoculated with material from negative muntjac (uterus, *n* = 9 mice; placentome, *n* = 8 mice; amniotic and allantoic fluids, *n* = 7 mice per cohort; 12 replicates each). Horizontal lines indicate the medians. Statistical significance is indicated above the sample replicates with an asterisk. For panel A, *P* < 0.0001 for values from mouse brains [not NaPTA treated] tested from uterus and placentome bioassay and *P* = 0.0277 for values in the concentrated amniotic fluid bioassay compared to control values (values for the allantoic fluid were not significantly different). For panel B, *P* < 0.0001 for values from mouse spleens tested from the uterus and placentome bioassay compared to control values (values from the concentrated amniotic and allantoic fluid bioassay were not significantly different from control values). (C and D) Lanes 2 demonstrate complete proteinase K (PK) digestion of PrP<sup>C</sup> in a negative deer obex, and lanes 4 demonstrate PrP<sup>RES</sup> in a CWD-positive deer obex. In panel C, lanes 6 to 8 and 10 to 12 demonstrate PrP<sup>RES</sup> detection in the brains of 6/8 mice inoculated with placentome tissue from a CWD-infected muntjac. In panel D, lanes 6 to 12 demonstrate PrP<sup>RES</sup> detection in the brains of 7/8 mice inoculated with uterine tissue from a CWD-infected muntjac.

offspring. Our findings (i) provide further evidence that CWD prions are present during gestation, placing the developing fetus at risk of TSE infection long before exposure to the infectious agent during the birthing process or maternal grooming behaviors shortly after parturition or via exposure to CWD-contaminated environments and (ii) suggest a correlation between RT-QuIC seeding activity and infectivity.

**Maternal tissues: infectious prions are present in CWD-infected deer reproductive tissues.** We found PrP<sup>C</sup>, RT-QuIC seeding activity, and infectious CWD prions in the uterus of CWD-infected dams. Infectious agents can access reproductive tissues (i) via transfer from the abdominal cavity to fallopian tubes, (ii) from the oral cavity by means of the circulatory system, (iii) by direct hematogenous infection, and (iv) by ascending from the vagina to the female reproductive tract (36). Substantial evidence has been established for the presence of pathological prions in tissues of the oral (37–40) and abdominal cavities (41–43), as well as the blood of TSE-infected hosts (44–50), providing opportunity for reproductive tissue exposure to prions by the aforementioned

**TABLE 2** Cervid transgenic mouse bioassay: inocula and Western blotting and RT-QuIC analysis

Muntjac dam disease status and identification no(s).	Inoculum		Clinical TSE disease <sup>a</sup>			RT-QuIC assay <sup>c</sup>		
	Tissue or fluid source (n)	Amt and route	No. of positive mice/total no. of mice	Time to disease (avg dpi ± SD)	Western blotting of brain <sup>c</sup>	Brain	Spleen	
CWD positive 45	Uterus (8)	30 μl i.c. of 10% homogenate	8/8	302.6 ± 45.8	8/8	8/8	7/8	
	Placentome 1 (6)	30 μl i.c. of 10% homogenate	0/6	500	0/6	0/6	1/6	
	47	Placentomes 1 and 5 (8)	30 μl i.c. of 10% homogenate	7/8	198.3 ± 45.4	6/8	8/8	8/8
		Amniotic fluid (9)	30 μl i.c.	1/9	320	0/9	1/9	2/9
		Allantoic fluid (9)	30 μl i.c.	0/9		0/9	0/9	0/9
CWD negative 64	Uterus (9)	30 μl i.c. of 10% homogenate	0/24 <sup>b</sup>		0/24	0/9	0/9	
	Placentome 1 (8)	30 μl i.c. of 10% homogenate				0/8	0/8	
	64 and 107	Amniotic and allantoic fluid (7)				0/7	0/7	

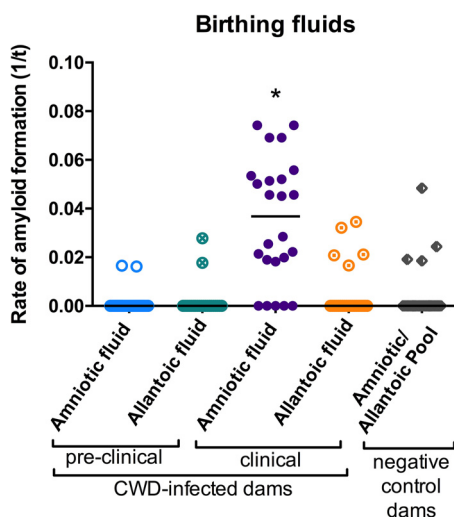
<sup>a</sup>Clinical disease, demonstrated by lethargy, stiff tail, circling, and hind limb ataxia.

<sup>b</sup>An age-matched negative-control mouse was harvested with each mouse euthanized with clinical TSE disease or at study termination at 343 days postinfection (dpi).

<sup>c</sup>Values are the number of positive mice/total number of mice tested.

routes of infection. Prions have been demonstrated in uterine tissues of vCJD-infected women (9) and scrapie-infected sheep (39). The normal cellular prion protein (PrP<sup>C</sup>), necessary for the conversion and maintenance of prion infection (51), has been found in high concentrations in reproductive tissues, including uterus (17, 52–55). It is therefore feasible that high levels of PrP<sup>C</sup> substrate within female reproductive tissues fuel amplification of infectious uterine prions, leading to sustained gestational exposure and subsequent *in utero* transmission to the developing embryo.

The role of the ovary in *in utero* prion exposure remains unclear. Here, we detected ample PrP<sup>C</sup> and RT-QuIC seeding activity in ovarian tissue harvested from clinically CWD-infected muntjac deer. The expression of PrP<sup>C</sup> in ovary tissues has been demon-



**FIG 4** RT-QuIC detection of amyloid formation in birthing fluids from clinically CWD-infected muntjac dams. For concentrated amniotic fluid from CWD-infected dams, values from preclinical specimens (4 dams) were not significantly different from control values, and values from clinical samples (2 dams) were significant at a *P* value of <0.0001. For allantoic fluid, values for both preclinical and clinical samples (2 dams each) from CWD-infected dams were not significantly different from values for negative dams (*n* = 2 dams). Number of replicates, 12 per dam per treatment group.



strated in cervids, humans, sheep, and cattle (9, 10, 55, 56) and in ovarian follicles of cattle and mice (52, 57), providing a substrate to sustain prion infection. The prion protein may have a role in regulating embryonic stem cell pluripotency and differentiation in early embryonic developmental stages (58). PK-resistant prion protein has been detected in human ovary tissue (9). Embryo transfers of 6-day-old embryos recovered from scrapie-infected ewes and implanted into scrapie-free ewes resulted in the transmission of scrapie to the offspring (23). These data imply a role for the transmission of prions during the very early stages of embryogenesis or with the germ line. Studies to determine the infectious nature of ovaries harvested from CWD-infected cervids are ongoing.

**Maternal-fetal interface: infectious prions are found within placentomes of CWD-infected dams.** Our bioassay study indicates that placentomes, i.e., the interface for metabolic exchange between dam and fetus during pregnancy, harbor infectious prions. The potential for prion trafficking across the placental structure has been evidenced by the identification of scrapie deposition within sheep placentomes (10, 20, 29, 59, 60) and in highly motile and phagocytic fetus-derived trophoblast cells (25) and by transmission of scrapie infectivity with placental tissues fed to sheep (12, 61). Prion infectivity has been demonstrated in the placenta of a woman showing clinical signs of CJD (14). A number of studies have established that infectious prions are present in the blood of symptomatic and asymptomatic TSE-infected hosts (45, 47–50, 62–64), and blood cells have been localized within trophoblast cells (65). Therefore, it is possible that infectious prions are trafficked via maternal blood to the fetal trophoctoderm at the placentome.

Exposure of the fetal trophoctoderm to maternal blood varies among species, dependent upon placental structure (66). A more intimate connection between mother and baby exists in human hemochorial placentation where the fetal trophoctoderm is directly bathed in maternal blood (67). This placentation contrasts with that of the ruminant placentome, epitheliochorial placentation, where three to five tissue layers separate the maternal circulatory system from the offspring (67–69). Increased angiogenesis to ensure maintenance of the developing fetus is a hallmark of all pregnancies (70). Subchorionic hematoma, separation of the chorion from the endometrium, permits maternal blood leakage and pooling within the placenta/placentome during pregnancy (71, 72). In addition, the normal events associated with blastocyst implantation in hemochorial placentation, i.e., remodeling of the maternal decidua (67), directly expose the fetal trophoctoderm to maternal blood. Thus, a breach in the uterine/placental membranes at implantation or during gestation may present an opportunity for infectious blood-borne or uterine-derived prions to enter the fetal environment. In this study, the detection of prion infectivity within uterine tissues of CWD-infected dams provides a local and perhaps persistent *in utero* prion source.

The muntjac placentome, densely interdigitated with both maternal and fetal tissues, is technically challenging to separate, ruling out contamination of one region with the other. Our bioassay inocula were therefore generated from tissue containing both maternal (caruncle) and fetal (cotyledon) portions of the placentome. Thus, we are unable to determine the degree to which infectivity is associated with fetal versus maternal placentome tissues. Nevertheless, our findings add to the body of evidence that maternal prions, whether maintained within the female reproductive microenvironment or hematogenously sourced, are present at the placental barrier.

**Fetal microenvironment: infectious prions are detected in the amniotic fluid of CWD-infected deer.** The fetal microenvironment of a pregnant, clinically CWD-infected dam contained prion infectivity. Here, we have demonstrated low levels of prion infectivity in amniotic fluid but not in allantoic fluid harvested from a clinically CWD-infected dam. PrP<sup>RES</sup> has been detected in amniotic fluid of scrapie-infected sheep (60), yet the infectivity of this material has not been determined. The fetus is bathed in amniotic fluid throughout gestation. During pregnancy, to further the development of fetal respiratory and digestive systems, the fetus recirculates amniotic fluid (up to 500 ml/day) via oronasal ingestion (73–76). Amniotic fluid consists of maternal plasma and

fetal urine, depending upon gestational stage (77–79). It is possible that blood-borne prions are trafficked with maternal plasma to amniotic fluid or that infectious prions are transported across the maternal-fetal interface via trophoblast cells from the uterine environment. In either case, susceptible fetal mucosal surfaces may be continuously bathed in, and thus exposed to, infectious prions throughout gestation.

Another potential reservoir for prion infectivity is the allantoic fluid, which consists of metabolic wastes produced by the fetus. Both PrP<sup>C</sup> (10) and PrP<sup>RES</sup> (29) have been detected in allantoic fluid collected from scrapie-infected ewes. This is not surprising because it is known that infectious prions are excreted in the shed by-products (urine and feces) of CWD- and scrapie-infected deer and sheep (80–83). However, in other studies, infectious prions were not found in fetal allantoic fluid (79, 84). If prion infectivity is present in fetal metabolic waste shed by the fetus (i.e., within allantoic fluid), the prions may be present at a concentration too low for detection, masked, or inactivated by pH or inhibiting properties present in these waste products, or the process of prion shedding may differ during gestation.

We are further investigating the ability of prions to traffic from the pregnancy and fetal microenvironment across the placental structure to establish infection in the developing fetus. To this end, bioassay studies to determine the infectivity of *in utero*-derived fetal tissues harvested from early- and late-stage experimental CWD-infected muntjac dams and free-range naturally infected elk dams are ongoing.

**RT-QuIC and infectious prions.** RT-QuIC provides a rapid methodology for analysis of tissues and fluids containing minute quantities of prions (85, 86) and may have utility in quantitative and correlative investigations of early and carrier states of prion diseases and potentially other protein misfolding disorders (86–89). There is little to no evidence that RT-QuIC generates, or represents, the presence of prion infectivity. Yet, similar to PMCA (1, 2), RT-QuIC consistently detected the presence of prion seeding activity in the reproductive milieu of clinically CWD-infected dams that, in this study, correlated with bioassay infectivity. We further demonstrated the presence of RT-QuIC seeding activity in some placentomes from pre-clinically CWD-infected dams lacking prion seeding activity in uterine and ovary tissues. These results are intriguing in that they suggest the accumulation of amyloid fibrils at the maternal-fetal interface prior to that in maternal reproductive organs, suggesting the potential for hematogenous- rather than uterine-sourced prions at this semipermeable membrane between mother and baby. It is well established that prion peripheral distribution and accumulation increase as TSE disease progresses (90–93). Our ability to demonstrate RT-QuIC seeding activity and prion infectivity within the female reproductive and pregnancy-related milieu of clinically CWD-infected dams may reflect these findings. The demonstration of RT-QuIC seeding activity in tissues harvested from preclinical dams was much lower than that demonstrated in the same tissues harvested from clinically ill dams, and infectivity of these tissues has not yet been demonstrated. We have initiated bioassays of these tissues to further assess the biological significance of our RT-QuIC results.

Our previous studies have compared RT-QuIC seeding activity to CWD cervid brain in a 50% lethal dose (LD<sub>50</sub>) bioassay (88, 94). Because we used the same conditions and controls for this study, we applied this principle to our RT-QuIC data. We estimated an LD<sub>50</sub> range from 18.7-fold less to 13.5-fold greater than that determined for CWD cervid brain ( $3.33 \times 10^6$  LD<sub>50</sub>s/g tissue) (88), suggesting a wide and variable range of brain equivalents in the reproductive milieu. More work is certainly needed to understand the significance of prion seeding activity and infectivity, as well as to discern a clearer picture of the isoforms detected by each methodology.

**Implications associated with maternal CWD infections.** The facile dissemination of CWD within captive and free-range cervid populations has led to questions regarding the transmission dynamics of this TSE. Direct contact with infected animals (95, 96) and indirect contact with infectious prions in bodily fluids and contaminated environments (97, 98) are suspected to explain the majority of this transmission. The effects of maternal CWD infections and a gestational source of prions may be underappreciated.

The presence of prion infectivity at the maternal-fetal interface, i.e., within the uterus, placentome, and fetal microenvironments, would help explain the high rates of CWD among cervids. In our experience, CWD-infected muntjac dams successfully breed and produce viable and nonviable offspring throughout the course of CWD infection even when exhibiting clinical signs of TSE disease. In a natural setting, however, the day-to-day challenges encountered by cervid dams to nurse, care for, and protect their young are much greater than those in a laboratory setting. This would surely be compounded by CWD infection, particularly in the later stages of disease. This, in combination with the high nonviable birth rates observed in offspring born to CWD-infected muntjac dams (60%) (1), may result in increased offspring mortality leading to lower annual recruitment rates (99). Yet a portion of offspring born to CWD-infected dams that survive to weaning and reproductive age would enter the population as potential asymptomatic carriers.

Infectious CWD prions in the reproductive milieu of cervid dams may spread disease by the following: (i) placing each developing fetus in direct contact with infectious prions throughout gestation; (ii) passage of CWD prions with ova to each offspring, increasing the likelihood of multigenerational mother-to-offspring transmission; (iii) postpartum fetal and placental membranes contributing to environmental contamination; and (iv) availability of postpartum tissues for consumption by scavenger species, increasing the potential for cross-species transmission.

Here, using a native CWD-susceptible host, the Reeves' muntjac deer, we share the first report of prion infectivity within the pregnancy microenvironments and placental structure of CWD-infected cervid. These findings reveal a source of infectious gestational prions that expose the developing fetus to CWD and are likely shed during parturition, contributing to CWD environmental contamination, and may contribute to CWD asymptomatic carriers in cervid populations. These findings have impact on our current concept of CWD disease transmission.

## MATERIALS AND METHODS

**Reproductive tissue source and processing. (i) Tissue source.** Uterus, ovary, and placentome tissues and amniotic and allantoic fluids were harvested from muntjac dams as part of a previous CWD maternal transmission study performed at Colorado State University and approved by the International Animal and Care Use Committee (IACUC) (1). Dams received CWD-positive brain homogenate via oral and subcutaneous inoculation and were humanely euthanized for maternal and fetal tissue collections at time points prior to (preclinical) or after (clinical) presentation of clinical signs consistent with TSE disease. Reeves' muntjac dams show terminal clinical signs of TSE disease (hypersalivation, weight loss, and ataxia) at approximately 24 months postexposure (1). All dams were confirmed to be CWD positive (1). Negative-control dams were dosed in the same manner with naive brain, remained healthy throughout the observation period, and were confirmed to be free of CWD. Genotyping of the dams was performed and previously reported (1). Sample details are provided in Table 1.

**(ii) Fluid and tissue collection and processing.** Great care was taken to minimize cross contamination between tissues and fluids as per previously published protocols (48, 49, 97). Amniotic and allantoic fluids were collected using single-use animal- and fluid-specific syringes and needles directly after each dam was euthanized and the uterus was removed for necropsy. New single-use, animal- and tissue-specific blades and forceps were used to harvest each tissue and fluid. Each ovary and placentome was frozen whole and bisected sagittally, and a midline section was taken for the homogenate. The uterus was bisected at necropsy, and the tissue was divided, with a portion stored at  $-80^{\circ}\text{C}$  for bioassay, Western blotting, and RT-QuIC homogenates. A portion of each tissue was fixed in McLean's paraformaldehyde-lysine-periodate (PLP) solution for immunohistochemistry.

**Western blotting.** Tissue homogenates were prepared from the obex region of the medulla oblongata at 10% (wt/vol) and from reproductive tissues (ovary, uterus, and placentome) at 20% (wt/vol) in Igepal buffer (10 mM Tris-HCl buffer, pH 7.5, 0.5% Igepal, 0.5% sodium deoxycholate) in an Omni Bead Ruptor 24 (Omni International, Inc.). Homogenates (2-mg tissue equivalent) were mixed with proteinase K (PK; Invitrogen) at a final concentration of 70  $\mu\text{g}/\text{ml}$  and incubated at  $37^{\circ}\text{C}$  for 30 min with shaking. Samples were mixed with reducing agent (10 $\times$ )-lithium dodecyl sulfate (LDS) sample buffer (4 $\times$ ) (Invitrogen) at a final concentration of 1 $\times$ , heated to  $95^{\circ}\text{C}$  for 5 min, and then run through a NuPAGE 12% Bis-Tris gel at 125 V for 1.5 h. Proteins were transferred to a polyvinylidene fluoride (PVDF) membrane at 80 V for 1 h in transfer buffer (0.025 M Trizma base, 0.2 M glycine, 20% methanol, pH 8.3). The membrane was incubated with blocking buffer (5% nonfat dried milk in 1 $\times$  Tris-buffered saline [TBS] with 0.1% Tween 20 [TBST]) for 20 min and then for 1 h with the primary antibody BAR224 (0.2- $\mu\text{g}/\text{ml}$  final concentration; Cayman Chemical) diluted in TBST, followed by a 30-min wash with TBST. The membrane was incubated for 30 min with the secondary antibody, peroxidase-labeled goat anti-mouse IgG (0.05- $\mu\text{g}/\text{ml}$  final concentration; KPL) diluted in TBST, followed by a 30-min wash with TBST. The

membrane was developed with ECL Plus Western blotting detection reagents (Pierce) and viewed on a Luminescent Image Analyzer LAS-4000 (Fujifilm).

**NaPTA precipitation.** Ten percent (10%; wt/vol) tissue homogenates made in 1× phosphate-buffered saline (PBS) (130 mM NaCl, 7 mM Na<sub>2</sub>HPO<sub>4</sub>·7H<sub>2</sub>O, 3 mM NaH<sub>2</sub>PO<sub>4</sub>·1H<sub>2</sub>O) in an Omni Bead Ruptor 24 (Omni International, Inc.) were diluted 1:10 in 0.1% SDS in 1× PBS for a total of 100 μl (0.1-mg tissue equivalent). This volume (100 μl) was subjected to protein precipitation (adapted from reference 100) by adding 7 μl of a freshly made solution containing 5% sodium phosphotungstate (NaPTA) octadecahydrate and 112 mM magnesium chloride hexahydrate prepared in sterile-filtered water (Sigma-Aldrich), followed by a 1-h incubation at 37°C and centrifugation for 30 min at 15,000 rpm. Supernatants were decanted, and the remaining pellet was resuspended in 10 μl of 0.1% SDS in 1× PBS for subsequent analysis by RT-QuIC.

**RT-QuIC assay.** All real-time quaking-induced conversion (RT-QuIC) assays (adapted from references 85, 87, and 101) were performed blinded, whereby sample identities were not revealed until after the analysis was completed. CWD-positive and -negative spleen samples were treated alongside the test samples and were used as plate controls in every experiment (spleen was used in ovary and uterus experiments; uterus was used in placentome experiments). Reaction mixtures consisted of 2 μl of NaPTA-precipitated sample into 98 μl of RT-QuIC buffer (final concentrations in 1× PBS: 130 mM NaCl, 1 mM EDTA, 10 mM thioflavin T [ThT], 0.1 mg/ml truncated recombinant hamster PrP<sup>C</sup> substrate) loaded into wells of a 96-well black-bottom optic plate (Nalgene Nunc). Prepared plates containing quadruplicate wells per sample were placed in a BMG FLUOstar Omega microplate reader and subjected to 700-rpm double-orbital shaking for 1 min every other minute for 15 min. After each shaking cycle, ThT fluorescence was read at an excitation of 450 nm and emission of 480 nm. Gain was set at 1,700 and read using orbital averaging with 20 flashes per well with a 4-mm setting. Fluorescent readings were recorded for all sample reactions for a total time of 62 h at a temperature of 42°C. Three or four separate experiments were run for each sample in quadruplicate, totaling 12 to 16 replicates per sample. Sample replicates were considered positive if they crossed the plate threshold (5 standard deviations [SD] above the mean of the initial five readings). Amyloid formation rates for positive replicates were determined using the inverse of the time when each positive reaction exceeded a threshold value paraformaldehyde-lysine-periodate. Statistical analyses were run in GraphPad Prism. The QuIC median rates of replicates for a given tissue are listed in Table 1. A Mann-Whitney test was used to generate *P* values (Table 1) by comparing the sample rates to the rates of known negative-control tissues. The confidence interval considered for statistical significance is 95%.

**Bioassay of cervid transgenic mice: inoculations and monitoring.** To further assess the presence of prion infectivity at the maternal-fetal interface, uterine tissue, placentomes, and amniotic and allantoic fluids harvested from CWD-infected and -negative muntjac dams were inoculated into Tg(CerPrP-E226)5037<sup>+/-</sup> mice for bioassay (102). Cohorts of 9 mice each were intracranially (i.c.) inoculated with 30 μl of a 10% (wt/vol) homogenate from clinically CWD-infected dam tissues (uterine, dam 45; placentome, dams 45 and 47 [3-mg tissue equivalent]) or 30 μl of concentrated amniotic (dam 47) and allantoic (dam 47) fluids (birthing fluids; 150-μl birthing fluid equivalent). The birthing fluids were concentrated by lyophilization to 5-fold less than their original volume and were dialyzed to remove the high salt concentrations. Biweekly weights and observations were taken until the mice were terminated due to disease or at 500 dpi, whichever occurred first. The negative-control mice received identical volumes of negative tissues and fluids and were followed similarly. Brain and spleen tissues harvested from these mice were assessed for the presence of prions by RT-QuIC and/or Western blotting.

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