

# Porcine Prion Protein as a Paradigm of Limited Susceptibility to Prion Strain Propagation

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Although experimental transmission of bovine spongiform encephalopathy (BSE) to pigs and transgenic mice expressing pig cellular prion protein (PrP<sup>C</sup>) (porcine PrP [PoPrP]–Tg001) has been described, no natural cases of prion diseases in pig were reported. This study analyzed pig-PrP<sup>C</sup> susceptibility to different prion strains using PoPrP-Tg001 mice either as animal bioassay or as substrate for protein misfolding cyclic amplification (PMCA). A panel of isolates representatives of different prion strains was selected, including classic and atypical/Nor98 scrapie, atypical-BSE, rodent scrapie, human Creutzfeldt-Jakob-disease and classic BSE from different species. Bioassay proved that PoPrP-Tg001-mice were susceptible only to the classic BSE agent, and PMCA results indicate that only classic BSE can convert pig-PrP<sup>C</sup> into scrapie-type PrP (PrP<sup>Sc</sup>), independently of the species origin. Therefore, conformational flexibility constraints associated with pig-PrP would limit the number of permissible PrP<sup>Sc</sup> conformations compatible with pig-PrP<sup>C</sup>, thus suggesting that pig-PrP<sup>C</sup> may constitute a paradigm of low conformational flexibility that could confer high resistance to the diversity of prion strains.

Keywords. atypical/Nor98 scrapie; BSE; classic scrapie; pig; prion conversion; prion strains; PrP; swine.

Transmissible spongiform encephalopathies (TSEs) or prion diseases are fatal neurodegenerative disorders caused by prion accumulation in the brain and lymphoreticular system [1]. Several TSEs naturally affecting animals are known [2], such as bovine spongiform encephalopathy (BSE) in cattle and scrapie in sheep and goats. Creutzfeldt-Jakob disease (CJD) is the most common human TSE. In the last century, a variant CJD associated to the dietary exposure to BSE-infected cattle was described [3]. Atypical cases of BSE are known to occur, mainly in older animals [2], classified as H-type or L-type according to their biochemical properties. Their low prevalence worldwide is consistent with a sporadic origin. The experimental transmission of atypical BSE to cattle, macaques, and mice evidenced their potentially infectious nature [4–6].

In the case of scrapie, the description of a wide variety of scrapie disease phenotypes suggests that a diversity of scrapie strains is circulating in sheep and goats [7, 8]. Moreover, in

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1998, an unusual type of scrapie was discovered in Norway (Nor98) [9]. Atypical/Nor98 scrapie has been reported in Europe, the United States, Canada, the Falkland Islands, Japan, Australia, and New Zealand and has been proposed to have a sporadic origin, because it is uniformly spread and often occurs in older animals as single cases in a flock. Transmission studies in transgenic mice and sheep demonstrated the transmissibility of atypical/Nor98 scrapie [10, 11]. Chronic wasting disease (CWD) in cervids affects both captive and wild animals [12] in North America. Cases of CWD have been detected in wild animals in Scandinavia [13], leading to rising concerns about the spreading of the disease in Europe. These animal prion diseases have become potential threats to public health and the economy.

Naturally occurring TSEs in pigs have never been reported [14, 15]. Experimental data showed that pigs can be infected after parenteral inoculation of BSE but not after oral challenge [16]. Previous studies using transgenic mice overexpressing pig protein (porcine prion protein [PoPrP]–Tg001) [17] suggested that pigs can be susceptible to BSE agent after passage in sheep or to an atypical/Nor98 scrapie isolate but not to 7 European classic scrapie isolates [18]. The susceptibility of pigs to BSE agent after experimental passage in sheep has also been demonstrated [19]. Pigs can be susceptible to US ovine scrapie isolate [20], but only a few intracranial inoculated pigs were scored positive for the infection. In a 2017 report [21], only a few pigs inoculated with an experimental inoculum of CWD were found to be TSE positive, showing low attack rates and long survival times. Moreover, adaptation of both scrapie and CWD prions to

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pigs was apparently incomplete, because further subpassage in transgenic mice expressing pig protein showed very limited attack rates. The incomplete adaptation of both scrapie and CWD to pigs could be the result of a nonadaptive prion amplification process [22].

In the present study, we use the PoPrP-Tg001 mouse model expressing pig cellular PrP (PrP<sup>C</sup>) to systematically evaluate the transmission barrier of pigs to a panel of TSE isolates from several species (cattle, sheep, goats, mice, hamsters, and humans). Additional studies have been performed using protein misfolding cyclic amplification (PMCA), an in vitro technique highly sensitive in the detection of prion propagation [23]. Brains from PoPrP-Tg001 mice were used as substrate for the PMCA reactions to evaluate the in vitro misfolding ability of pig-PrP<sup>C</sup>, using a representative collection of the isolates inoculated in PoPrP-Tg001 mice.

# **METHODS**

## **Ethic Statements**

Animal experiments were carried out in strict accordance with the recommendations in the guidelines of the Code for Methods and Welfare Considerations in Behavioural Research with Animals (directives 86/609EC and 2010/63/EU). Experiments were approved by the Committee on the Ethics of Animal Experiments of the Instituto Nacional de Investigación y Tecnología Agraria y Alimentaria (Madrid, Spain; permits CEEA 2009/004 and 2012/002).

### **Transmission Studies**

The studies used a PoPrP-Tg001 mouse line expressing porcine  $PrP^{C}$  (4-fold the level of expression in pig brain) in a background knock-out for the prion protein [17]. All inocula were prepared from brain tissues as 10% (wt/vol) homogenates in 5% glucose in distilled water. The isolates used as inocula are described in Supplementary Table 1. Individually identified 6–10-week-old mice were anesthetized with isofluorane and inoculated with 2 mg of brain homogenate in the right parietal lobe, using 25-gauge disposable hypodermic needles. Mice were observed daily, and their neurological status was assessed weekly.

When progression of a TSE disease was evident or at the end of their lifespan, animals were euthanized. Necropsy was then performed, and the brain was collected. A part of the brain was fixed by immersion in 10% formalin for histopathology and immunohistochemistry studies, and the other part was frozen at  $-20^{\circ}$ C for porcine protease-resistant PrP (PrP<sup>res</sup>) detection by means of Western blot (WB) analysis. In some cases, brain homogenates were used for second passages, either from PrP<sup>res</sup>-positive animals or for blind second passage from PrP<sup>res</sup>-negative animals. The ability of the inocula used to transmit prion disease was demonstrated using transgenic mice expressing bovine [24], ovine [10, 25], murine [26], and human-PrP<sup>C</sup> [27] or Syrian golden hamsters (*Mesocricetus auratus*).

# **WB** Analysis

A mass of 175 mg  $\pm$  20 mg of frozen brain tissue was homogenized to a concentration of 10% (wt/vol) in 5% glucose in distilled water in grinding tubes (Bio-Rad), using a TeSeE Precess 48 homogenizer (Bio-Rad). The presence of PrP<sup>res</sup> in transgenic mice brains was determined by means of WB analysis, as described elsewhere [5, 27]. Ten to 100 µL of a 10% (wt/vol) brain homogenate was digested with Proteinase K, loaded on 12% Bis-Tris Gel (Criterion XT; Bio-Rad), and detected with Sha31 monoclonal antibody [28].

## **PMCA** Procedure

PMCA was done as described elsewhere [23]. Briefly, PoPrP-Tg001 mice were perfused by cardiac puncture with 5 mmol/L ethylenediaminetetraacetic acid (Sigma-Aldrich) prepared in 1× phosphate-buffered saline without calcium and magnesium (Thermo Fisher Scientific). Brains were removed and homogenized using a Potter homogenator (Thermo Fisher Scientific) at 10% in 150 mmol/L sodium chloride (Merck) and 1% (vol/vol) Triton X-100 (Sigma-Aldrich) prepared in 1× phosphate-buffered saline without calcium and magnesium supplemented with protease inhibitors (Roche). Aliquots (500  $\mu$ L) were stored at  $-80^{\circ}$ C until use.

The different isolates were diluted 1:100 or 1:10 or undiluted into PoPrP-Tg001 substrate. Then, 7  $\mu$ L of inocula was mixed with 63  $\mu$ L of PoPrP-Tg001 brain substrate into 0.2-mL polymerase chain reaction tubes (Thermo Fisher Scientific), and 4 zirconia balls (Biospec) were added on each tube. Next, 20  $\mu$ L of the mixture was taken and immediately frozen as a nonamplification control. At least 2 tubes for each inoculumsubstrate combination were included in the experiment, and 3 independent PMCA experiments were performed. Unseeded substrate was also included in the experiment as negative amplification control. Tubes were placed into the water-filled sonicator horn (QSonica; Q700) at 37°C. Sonication-incubation cycles of 24 hours were applied to the samples. Each cycle included 20 seconds of sonication plus 30 minutes of incubation (amplitude, 40%).

# RESULTS

Isolates with differential properties representative of distinct TSE strains from several species have been used to systematically assess the potential susceptibility of pigs to different prions, using a PoPrP-Tg001 mouse model. The isolates used in this work are compiled in Supplementary Table 1. Prion infectivity of these inocula was previously tested in a homologous PrP animal model (without species barrier), as showed in Tables 1–4.

	Survival Time, Mean (SD), d [Diseased, PrP <sup>res</sup> -Positive/Inoculated Mice, No.] <sup>a</sup>							
	PoPr							
Inocula	1st Passage	2nd Passage	(1st Passage)					
Ca-BSE 2	498 (9) [2/12] <sup>b</sup>	198 (6) [15/15] <sup>b</sup>	308 (5) [5/5] <sup>c</sup>					
Ca-BSE 2/Tg110 Bo6OR	372 [1/6] <sup>d</sup>	208 (12) [6/6] <sup>d</sup>	265 (35) [6/6] <sup>d</sup>					
Ca-BSE 2/Tg008 Bo5OR	461 (100) [3/18]	ND	331 (73) [7/7]					
Ca-BSE-H 02.2695	>650 [0/6]	>650 [0/6]	328 (15) [12/12] <sup>e</sup>					
Ca-BSE-H 02.2695/Tg110 H	>650 [0/6]	ND	292 (12) [6/6] <sup>e</sup>					
Ca-BSE-H 07-644	>650 [0/6]	ND	274 (3) [6/6] <sup>e</sup>					
Ca-BSE-H 07-644/Tg110 H	>650 [0/6]	>650 [0/6]	298 (10) [7/7] <sup>e</sup>					
Ca-BSE-L 02.2528	>650 [0/6]	>650 [0/6]	207 (7) [6/6]					
Ca-BSE-L 02.2528/Tg110 L	>650 [0/6]	>650 [0/6]	198 (1) [6/6]					
Ca-BSE-L 43	>650 [0/6]	ND	188 (1) [6/6]					
Sheep-Sc Langlade/Tg110	>650 [0/5]	ND	321 (8) [5/5]					
Sheep-Sc PS48/Tg110	>650 [0/6]	ND	197 (1) [5/5]					
Sheep-Sc198-9/Tg110	>650 [0/5]	ND	165 (7) [7/7]					
Sheep-ScPS21/Tg110	>650 [0/6]	ND	187 (2) [5/5]					
Sheep-Sc pool pre-75 cattle P75-7	>650 [0/6] <sup>f</sup>	>650 [0/6] <sup>f</sup>	203 (5) [6/6] <sup>f</sup>					
Sheep-Sc pool post-90 cattle P90-1	>650 [0/6] <sup>f</sup>	>650 [0/6] <sup>f</sup>	173 (3) [6/6] <sup>f</sup>					

Abbreviations: BoPrP, bovine prion protein (PrP).BSE, bovine spongiform encephalopathy; ND, not done.; PoPrP, porcine PrP; PrP<sup>res</sup>, protease-resistant PrP.

<sup>a</sup>The mean survival time is indicated for all mice scored positive for PrP<sup>res</sup>.

<sup>b</sup>Published elsewhere [18].

<sup>c</sup>Published elsewhere [17].

<sup>d</sup>Published elsewhere [29].

<sup>e</sup>Published elsewhere [5].

<sup>f</sup>Published elsewhere [30].

#### Susceptibility of PoPrP-Tg001 Mice to TSE Isolates

As described elsewhere [17, 18], PoPrP-Tg001 mice can be infected with classic BSE prions but show a high transmission barrier (Table 1). A similar high transmission barrier was also observed in PoPrP-Tg001 mice inoculated with classic BSE passaged in bovine-PrP<sup>C</sup> transgenic mice with either 5 or 6 octarepeats (Table 1). Considering that porcine-PrP<sup>C</sup> harbors 5 octarepeats, this result suggests that the identity in the number of octarepeats does not affect the bovine-porcine transmission barrier for classic BSE.

When atypical BSE-L or atypical BSE-H isolates were inoculated, no transmission was detected in PoPrP-Tg001 mice even after 2 iterative passages. Overall, classic BSE was the only bovine prion able to infect mice expressing pig-PrP<sup>C</sup>. This observation can be extended to classic BSE after passage in other species. The outcome of the inoculations with a wide panel of prions from other species (sheep, goats, mice, hamsters, and humans) indicates that only those isolates derived from classic BSE strain were able to infect PoPrP-Tg001 mice (Tables 2–4) showing similar survival times after second passage, as reported elsewhere [29]. Furthermore, brains from PoPrP-Tg001 mice exhibited similar neuropathological features for all the classic BSE-derived prions (data not shown), regardless of the species of origin of the inoculum, as reported elsewhere [29].

When other prion isolates representative of different strains, such as classic scrapie, atypical/Nor98 scrapie, or sporadic CJD (sCJD), were used as inocula, no transmission to PoPrP-Tg001 mice was detected (Tables 2–4), neither by PrP<sup>res</sup> detection with WB analysis nor by histopathological analysis. This occurs independently of the PrP expressed in the donor used as inoculum (cattle, sheep, goats, mice, hamsters, or humans), suggesting that differences in amino acid sequences between pig-PrP<sup>C</sup> and donor scrapie-type PrP (PrP<sup>Sc</sup>) are not mainly responsible for this transmission barrier.

Strikingly, none of the 9 atypical/Nor98 scrapie isolates from either sheep or goat cases were able to infect PoPrP-Tg001 mice (Table 2). These results contrast with those previously published by our group [18], in which sheep-AtSc 152 isolate was able to infect 2 of 12 inoculated mice. In the present work, sheep-AtSc 152 isolated from the same sheep brain was inoculated in another 9 mice but there was no evidence of transmission to any of them. We also attempted, without success, to infect PoPrP-Tg001 mice with the inoculum previously amplified in VRQ-ovine-PrP-Tg338 mice (sheep-AtSc 152/Tg338 in Table 2).

#### In Vitro Conversion of Pig-PrP by TSE Isolates

Some isolates tested on the PoPrP-Tg001 mice were selected for testing of their ability to propagate in vitro, using PMCA with PoPrP-Tg001 brain as substrate (Table 5). These isolates were subjected to 15 rounds of PMCA and further analyzed to detect

#### Table 2. Transmission of Sheep and Goat Inocula to PoPrP-Tg001 and Ovine Transgenic Mice

	Survival Time	Survival Time, Mean (SD), d [Diseased, PrP <sup>res</sup> -Positive/Inoculated Mice, No.] <sup>a</sup>								
Inocula	PoPrP-	PoPrP-Tg001								
	1st Passage	2nd Passage	Ovine PrP-1g Mice <sup>o</sup> (1st Passage)							
Sheep-Sc pool pre-75	>650 [0/6]	>650 [0/6]	69 (1) [6/6] <sup>c</sup> (Tg338)							
Sheep-Sc pool post-90	>650 [0/6]	>650 [0/6]	480 (19) [6/6] <sup>c</sup> (Tg338)							
Sheep-Sc PS09	>650 [0/6]	>650 [0/5]	431 (18) [6/6] (Tg338)							
Sheep-Sc PS21	>650 [0/6]	>650 [0/6]	182 (17) [6/6] (Tg338)							
Sheep-Sc PS42	>650 [0/7]	>650 [0/5]	67 (7) [6/6] (Tg338)							
Sheep-Sc198-9	>650 [0/5]	ND	629 (27) [5/5]							
Sheep-ScO100	>650 [0/6]	>650 [0/5]	364 (61) [12/12] <sup>d</sup> (TgOvPrP4)							
Sheep-ScO104	>650 [0/5]	>650 [0/5]	248 (50) [10/10] <sup>d</sup> (TgOvPrP4							
Goat-Sc S2	>650 [0/6]	>650 [0/5]	449 (62) [9/9] <sup>e</sup>							
Goat-F10	>650 [0/5]	ND	465 (17) [7/7] <sup>e</sup>							
Goat-Sc Goujon	>650 [0/6]	>650 [0/5]	253 (7) [5/5]							
Goat-AtSc I15	>650 [0/6]	ND	552 (78) [6/6] <sup>f</sup>							
Sheep-AtSc Engavagen	>650 [0/10]	>650 [0/5]	227 (15) [11/11] (Tg338)							
Sheep-AtSc Leknes	>650 [0/9]	ND	ND							
Sheep-AtSc Kjerringøy	>650 [0/5]	>650 [0/5]	285 (43) [6/6] (Tg338)							
Sheep-AtSc Suldalsosen	>650 [0/6]	>650 [0/5]	ND							
Sheep-AtSc Tennevoll	>650 [0/5]	ND	245 (15) (Tg338)							
Sheep-AtSc Vinje	>650 [0/5]	ND	ND							
Sheep-AtSc 152	300–600 [2/12] <sup>g</sup>	162 (13) [9/9] <sup>g</sup>	418 (6) [6/6]							
Sheep-AtSc 152	>650 [0/9]	ND	ND							
Sheep-AtSc 152/Tg338	>650 [0/6]	>650 [0/5]	ND							
BSE in sheep ARQ/ARQ	458 (11) [15/15] <sup>9</sup>	162 (4) [13/13] <sup>g</sup>	485 (62) [7/7] <sup>e</sup>							
BSE in sheep ARR/ARR	471 (58) [9/9]	185 (5) [5/5]	358 (20) [6/6]							
BSE in TgOV ARQ	389 (37) [5/5]	ND	ND							
BSE in TgOV ARR	417 (18) [5/5]	183 (14) [6/6]	ND							
Experimental BSE in goat	459 (27) [5/5]	176 (5) [5/5]	497 (31) [5/5] <sup>e</sup>							
BSE in goat CH636	505 (60) [7/7]	182 (9) [7/7]	ND							

Abbreviations: BSE, bovine spongiform encephalopathy; ND, not done; PoPrP, porcine prion protein; PrPres, protease-resistant prion protein.

<sup>a</sup>The mean survival time is indicated for all mice scored positive for PrP<sup>res</sup>

<sup>b</sup>Data from ovine PrP-Tg501 mice [25] except for those indicated as either Tg338 [10] or TgOvPrP4 [31].

<sup>f</sup>Published elsewhere [32].

positive propagation by means of PrP<sup>res</sup> detection with WB analysis (Supplementary Figure 1 shows a representative WB). The results presented here (Table 1) were obtained using 1:100 dilutions of the isolates as inocula. When a higher prion seed concentration was used (1:10 or undiluted), isolates negative at 1:100 dilution remained negative at lower dilutions, whereas for positive isolates, in some cases, additional amplification rounds were needed for positive detection. This may be owing to the presence of PMCA inhibitors in the inocula. Alternatively, the dilution of the inocula may result in a concentration-dependent dissociation of the aggregates, thereby releasing and increasing the concentration of available seeds for PMCA, as suggested elsewhere [34].

PMCA results were comparable to those obtained in animal bioassays: classic BSE was the only strain able to amplify in

PoPrP-Tg001 substrate independently of the species-PrP in the isolate (Table 5). In the absence of species barrier, as the case of classic BSE from pigs, the amplification was detected in the first round, whereas several PMCA rounds were required to detect positive amplification when a species barrier existed (ie, classic BSE from cattle, sheep, goats, or humans). Atypical BSE, classic scrapie, and sCJD prion strains were unable to propagate in PoPrP-Tg001 substrate, as also found with the animal bioassay (Table 5).

## **Biochemical Characterization of Pig-PrP**<sup>res</sup>

Comparison of the brain PrP<sup>res</sup> collected from PoPrP-Tg001 mice inoculated with classic BSE–derived prions revealed the same profile in WB analysis, irrespective of the species-PrP in the inoculum (Figure 1A). In all cases, a PrP<sup>res</sup> glycosylation

<sup>&</sup>lt;sup>c</sup>Published elsewhere [30].

<sup>&</sup>lt;sup>d</sup>Published elsewhere [31].

<sup>&</sup>lt;sup>e</sup>Published elsewhere [25].

<sup>&</sup>lt;sup>g</sup>Published elsewhere [18].

#### Table 3. Transmission of Mouse and Hamster Inocula in PoPrP-Tg001, Tga20 transgenic Mice or Hamsters

Inocula	S	Survival Time, Mean (SD), d [Diseased, PrP <sup>res</sup> -Positive/Inoculated Mice, No.] <sup>a</sup>								
	PoP	rP-Tg001	7 00							
	1st Passage	2nd Passage	I ga20 (1st Passage)	Hamsters (1st Passage)						
BSE in Tga20	506 [1/6] <sup>b</sup>	ND	154 (21) [5/5]	ND						
BSE in wt mouse	650 [1/6] <sup>b</sup>	201 (12 <sup>b</sup> ) [6/6]	158 (39) [5/5]	ND						
22L	>650 [0/6]	>650 [0/6]	112 (13) [4/4]	ND						
RML	>650 [0/6] <sup>b</sup>	>650 [0/6] <sup>b</sup>	75 (7) [5/5]	ND						
263K	>650 [0/7]	>650 [0/5]	ND	87 (3) [5/5]						

Abbreviations: BSE, bovine spongiform encephalopathy; ND, not done; PoPrP, porcine prion protein; PrP<sup>res</sup>, protease-resistant prion protein; wt, wild-type.

<sup>a</sup>The mean survival time is indicated for all mice scored positive for PrP<sup>res</sup>

<sup>b</sup>Published elsewhere [29].

pattern with a predominant monoglyglcosylated band was observed. This profile was similar to that reported elsewhere in pigs inoculated with classic BSE [18, 21, 29, 35]. Comparison of PrP<sup>res</sup> obtained from both inoculated mice and PMCA with classic BSE–derived isolates demonstrated that the biochemical strain properties (glycoform proportion and molecular weight of the unglycosylated band) of classic BSE were maintained (Figure 1B). This suggests that the classic BSE prion's conformation is reliably transmitted to the pig-PrP in vitro, as described elsewhere for PrP from other species [36].

# DISCUSSION

In this work, the transmissibility of a panel of TSE isolates representing diverse prion strains from cattle, sheep, goats, mice, hamsters, and humans was assayed both in vivo using mice overexpressing pig- $PrP^{C}$  and in vitro using the PMCA technique. Although different combinations of prion strains and  $PrP^{C}$  expressing donors have been used, only the classic BSE strain was able to propagate in mice expressing pig- $PrP^{C}$  independently of the donor PrP amino acid sequence. It is interesting that classic BSE after passage in other species, such as sheep, goats, or humans, propagate in mice expressing pig- $PrP^{C}$ 

with better transmission efficiency than cattle BSE, resembling previous observations made in human and bovine PrP transgenic mice [27, 37]. Moreover, all classic BSE–derived prions, regardless of the originating species, exhibited similar strain features, such as survival time and a PrP<sup>res</sup> glycosylation pattern characterized by a predominant monoglycosylated band, matching that reported elsewhere in pigs infected with BSE [18, 21, 29, 35].

The strain-dependent transmission barrier observed in pig-PrP mice is in accordance with previous observations evidencing that prion strain properties, probably associated with different PrP<sup>Sc</sup> conformers, have a determinant impact on the ability of prions to cross the species barrier [29, 38, 39]. The PMCA results reinforce those obtained in the animal bioassay, confirming that only the classic BSE strain seems able to propagate in a pig-PrP context. In addition, the results indicates that for the isolates analyzed in this study, PMCA is a valuable tool as a complementary method to animal bioassays to assess more quickly the susceptibility or resistance to TSEs in PoPrP-Tg001, because the results obtained using both techniques were equal in terms of isolate propagation and PrP<sup>res</sup> WB profile.

Table 4. Transmission of Human I	nocula in l	PoPrP-Ta001	and HuPrP-To	1340 Transc	penic Mice
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	Survival Time, Mean (SD), d [Diseased, PrP <sup>res</sup> -Positive/Inoculated Mice, No.] <sup>a</sup>							
	PoPri							
Inocula	1st Passage	2nd Passage	(1st Passage)					
sCJD 129 M/MT1 NHBX0/0001	>650 [0/6] <sup>b</sup>	>650 [0/6] <sup>b</sup>	214 (6) [5/5] <sup>b</sup>					
sCJD 129 M/MT1 0.08.02523_001	>650 [0/6]	ND	187 (11) [6/6]					
sCJD 129 V/VT2 0.08.02497_001	>650 [0/5]	ND	522 (36) [6/6]					
vCJD 129 M/M NHBY0/0014	556 (81) [6/6] <sup>b</sup>	212 (6) [6/6] <sup>b</sup>	626 (29) [6/6] <sup>b</sup>					
vCJD 129 M/M BC1458	530 (48) [6/6]	197 (9) [5/5]	545 (146) [5/5] <sup>c</sup>					
BSE in HuPrP-Tg340	486 (31) [5/6]	ND	614 (87) [6/6] <sup>c</sup>					

Abbreviations: BSE, bovine spongiform encephalopathy; HuPrP, human prion protein (PrP); ND, not done; PoPrP, porcine PrP; PrP<sup>res</sup>, protease-resistant PrP; sCJD, sporadic Creutzfeldt-Jakob disease; vCJD, variant Creutzfeldt-Jakob disease.

<sup>a</sup>The mean survival time is indicated for all mice scored positive for PrP<sup>res</sup>.

<sup>b</sup>Published elsewhere [29].

<sup>c</sup>Published elsewhere [33].

#### Table 5. Protein Misfolding Cyclic Amplification of Selected Inocula Using PoPrP-Tg001 as Substrate

	Amplification by Serial PMCA Round, % <sup>a</sup>														
Seed	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
Ca-BSE/Pig <sup>b</sup>	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
Ca-BSE 2	0	0	0	0	0	0	100	100	100	100	100	100	100	100	100
Ca-BSE-H 07-644	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Ca-BSE-L 43	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sheep-Sc pool pre-75 cattle P75-7	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sheep-Sc pool post-90 cattle P90-1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sheep-Sc PS48°	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sheep-Sc PS13°	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sheep-Sc PS21	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sheep-Sc PS42	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Sheep-Sc198-9	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Goat-Sc F10	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
BSE in sheep ARQ/ARQ	0	0	100	100	100	100	100	100	100	100	100	100	100	100	100
BSE in TgOV ARQ	0	0	100	100	100	100	100	100	100	100	100	100	100	100	100
Experimental BSE in goat	0	0	0	0	0	0	100	100	100	100	100	100	100	100	100
22L	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
RML	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
263K	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
sCJD 129 M/M T1 0.08.02523_001	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
sCJD 129 V/V T2 0.08.02497_001	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
vCJD129 M/M BC1458	0	0	0	0	0	0	50	50	100	100	100	100	100	100	100
BSE in HuPrP-Tg340	0	0	0	0	0	0	100	100	100	100	100	100	100	100	100
Unseeded	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0

Abbreviations: BSE, bovine spongiform encephalopathy; HuPrP, human prion protein (PrP); PMCA, protein misfolding cyclic amplification; PoPrP, porcine PrP; sCJD, sporadic Creutzfeldt-Jakob disease; vCJD, variant Creutzfeldt-Jakob disease.

<sup>a</sup>Percentage of positive tubes (showing PrP<sup>res</sup>) of the total number of tubes sonicated (n = 6).

<sup>b</sup>Described elsewhere [29].

<sup>c</sup>Described elsewhere [18].

None of the several atypical/Nor98 scrapie isolates used in this work, including a new inoculation of sheep-AtSc 152 isolate, was transmitted, supporting the contention that porcine species is highly resistant to atypical/Nor98 scrapie prions. This is in contrast with previous results obtained with sheep-AtSc 152 [18], where material from the same infected sheep brain was able to be transmitted, although with a very low attack rate. The ability of sheep-AtSc 152 to infect mice may be due to particular properties distinguishing this isolate from other atypical/Nor98 scrapie isolates. Other possibilities cannot be excluded, such as the coexistence of BSE agent as a minor component present in the donor sheep brain. Moreover, contamination with classic BSE agent in any of the different steps (sample harvesting, homogenization, or inoculation) related with the preparation of the inoculum used in the experiment reported elsewhere [18] cannot be excluded. In any case, the transmission barrier for atypical/Nor98 scrapie infection in pigs can be considered very high as assayed in the mouse model expressing pig-PrP<sup>C</sup>.

Because positive transmission was not detected in animals inoculated with any of the prion strains used in this work, other than classic BSE, our results indicate a high resistance of the

mouse model expressing pig-PrP<sup>C</sup> to all of them. PMCA is an extremely sensitive technique used to detect prion propagation [23]. Thus, the absence of positive amplification after 15 rounds of PMCA for strains different from classic BSE strongly supports the low susceptibility of PoPrP-Tg001 to prions other than classic BSE. In a recent work, pigs were inoculated with a pool of brains of white tailed deer intracranially inoculated with CWDaffected elk, white-tailed deer, and mule deer [21]. Although RT-QuIC (real-time quaking-induced conversion) enabled detection of PrPSc in both orally and intracranially CWDinoculated pigs as early as 6 months after inoculation, brain PrP<sup>res</sup> was detectable with WB analysis a long time (45 months) after inoculation in only a few animals, and in pigs inoculated intracranially but not those inoculated orally. However, 1 orally inoculated pig was positive at immunohistochemistry and enzyme-linked immunosorbent assay 45 months after inoculation. Furthermore, second passages in a pig-PrP<sup>C</sup> transgenic mouse model showed reduced attacks rates, suggesting that pig-PrP<sup>C</sup> can support low-level propagation of CWD prions, though with a high species barrier.

Although pig- $PrP^{C}$  could sustain replication of some prion strains, the transmission barrier of pig- $PrP^{C}$  for the analyzed



Figure 1. Immunoblot of porcine protease-resistant prion protein (PrPres). A, Brain PrPres Western blot profile in porcine PrP (PoPrP)-Tg001 mice inoculated with the different classic bovine spongiform encephalopathy (BSE)-derived inocula: Ca-BSE 2 (lane 1), Ca-BSE 2/Tg110 Bo6OR (lane 2), Ca-BSE 2/Tg008 Bo5OR (lane 3), BSE in sheep ARQ/ARQ (lane 4), BSE in sheep ARR/ARR (lane 5), BSE in TgOV ARQ (lane 6), BSE in TgOV ARR (lane 7), experimental BSE in goat (lane 8), BSE in goat CH636 (lane 9), BSE in Tga20 (lane 10), BSE in wild-type mouse (lane 11), variant Creutzfeldt-Jakob disease (vCJD) 129 M/M NHBY0/0014 (lane 12), vCJD 129 M/M BC1458 (lane 13), and BSE in HuPrP-Tg340 (lane 14). B, Brain PrPres Western blot profile in the different inocula and in the PoPrP-Tg001 brain homogenates after in vivo or in vitro experiments. Original inocula used included Ca-BSE/Pig (lane 1), Ca-BSE 2 (lane 4), BSE in Sheep ARQ/ARQ (lane 7), and vCJD 129 M/M BC1458 (lane 10). PoPrP-Tg001 mouse infected with: Ca-BSE/Pig (lane 2), Ca-BSE 2 (lane 5), BSE in Sheep ARQ/ARQ (lane 8) and vCJD 129 M/M BC1458 (lane 11). Brain homogenates from PoPrP-Tg001 were used in protein misfolding cyclic amplification seeded with Ca-BSE/Pig (Iane 3), Ca-BSE 2 (Iane 6), BSE in sheep ARQ/ARQ (Iane 9), and vCJD 129 M/M BC1458 (lane 12). PrPres was detected using Sha31 monoclonal antibody. (Molecular weight in kilodaltons is shown at the right of each panel.)

strains is, at least, higher than the already known strong transmission barrier for classic BSE, a prion difficult to transmit in both transgenic mice expressing pig-PrP<sup>C</sup> and pigs [16, 17]. Interestingly, the ability of pig-PrP to sustain replication of the classic BSE strain regardless of the donor PrP<sup>Sc</sup> amino acid sequence used as inoculum suggests that only a very restricted number of PrP<sup>Sc</sup> conformers, such as classic BSE, present molecular compatibility with pig-PrP for prion propagation. Thus, strain-specific PrP<sup>Sc</sup> conformers seem to play a determinant role in prion strain transmission barrier, even more decisively than amino acid sequence differences (species barrier) [18, 39]. The control of prion host range is thus dictated by selective constraints imposed by the PrP<sup>Sc</sup> rather than the PrP<sup>C</sup> encoded by the host.

On the other hand, the high resistance of pig-PrP<sup>C</sup> to replicating a broad diversity of prion strains is in contrast to the prion susceptibility observed in other related species showing minor differences at the PrP amino acid sequence. The pig-PRNP gene is considered very homogeneous, because no relevant polymorphisms have been described [40]. The amino acid sequence of pig-PrP shows ~94% identity with either cattle or sheep-PrP amino acid sequences (see Figure 2A). This supports the notion that the pig-PrP amino acid sequence has a limited proficiency for recognizing and/or adopting the different PrP<sup>Sc</sup> conformations associated with the diversity of prion strains.

This can be due to limitations in the conformational flexibility of the pig-PrP amino acid sequence.

Because only 1 amino acid substitution may drastically alter prion resistance or susceptibility, it is difficult to determine the particular effect of any of the amino acid changes with the high resistance to different prion strains revealed by pig-PrP<sup>C</sup>. However, we can speculate about the potential effect of some of the amino acid changes observed in the pig-PrP sequence when compared with either bovine or sheep PrP, because both bovine and sheep PrP can adopt the PrPSc conformations associated with the different prion strains used in this work. It is known that minor changes in the  $\beta 2-\alpha 2$  loop of  $PrP^{C}$  protein (residues 169-179 of porcine sequence) may considerably affect the transmission barrier [41-43]. As a paradigm, the Q171R polymorphism present in the  $\beta 2-\alpha 2$  loop of sheep-PrP<sup>C</sup> is strongly linked to resistance to classic scrapie, but not to BSE [44]. In this sense, the N-to-S amino acid change in the 178 position of the pig-PrP<sup>C</sup> is present only in species with low susceptibility to prion infection, such as rabbits (Figure 2B).

The amino acid change found in the  $\beta 2-\alpha 2$  loop in pig-PrP<sup>C</sup> would alter the flexibility of the  $\beta 2-\alpha 2$  loop, strengthening the transmission barrier for diverse prion strains other than classic BSE. Other amino acid changes in the pig-PrP<sup>C</sup> would also participate in the limited capacity of this protein to sustain replication of different prion strains and hence to adopt the PrP<sup>Sc</sup> conformations associated with those strains. From these changes,  $_{226}YE_{227}$  amino acids present in the pig-PrP primary sequence are absent in PrP from other species susceptible to prions showing SQ amino acids at the equivalent position (see Figure 2C).

These <sub>226</sub>YE<sub>227</sub> amino acid changes are present in PrP<sup>C</sup> from other species alleged to be reluctant to conformational conversion to PrP<sup>Sc</sup>, such as horses [22]. E<sub>227</sub> amino acid in pig-PrP is the equivalent of the Q226E polymorphism observed in cervids, which is E226 in Rocky Mountain elk and Q226 in other CWDsusceptible cervids. CWD prion strain propagation is stable in transgenic mice expressing E<sub>226</sub> cervid-PrP<sup>C</sup>, whereas mice expressing Q226 cervid-PrP<sup>C</sup> unstably generate CWD mixed strains [45]. Furthermore, these 226 YE,227 amino acid changes are close to the equivalent position of the Q222K polymorphic variant described in goat populations, considered to confer resistance to classic scrapie prions and reduce susceptibility to the classic BSE strain [25], and the human E219K polymorphism that has been linked to protecting humans against sCJD in epidemiological studies in Asiatic populations [46]. Together, these data suggest that  $_{226}YE_{227}$  amino acids can be relevant in the restricted ability of pig-PrP to sustain prion replication.

Overall, our results demonstrate that pig-PrP<sup>C</sup> can be converted in pig-PrP<sup>Sc</sup> only by the classic BSE prion strain, irrespective of the donor species, but not by any other strain used in this work, though other strains not used in this work, such as CWD, may also be able to convert pig-PrP<sup>C</sup> into pig-PrP<sup>Sc</sup>.



**Figure 2.** Amino acid sequence alignment. The amino acid sequences of different species' prion protein (PrP) were compared with pig-PrP (*top rows*). Points indicate identical residues; dashes, deletions. Amino acid numbering is indicated on the right, and species on the left. *A*, Amino acid alignment of pig, cattle, sheep, and mouse prion proteins. Amino acid changes in the pig-PrP amino acid sequence conserved in cattle, sheep, and mice are in bold and underlined. *B*, Pig-, cattle-, sheep-, mouse-, rabbit-, and horse-PrP alignment of the β2-α2 loop region and the surrounding amino acids. *C*, Pig-, cattle-, sheep-, mouse-, rabbit-, and horse-PrP alignment of the region around <sup>206</sup> YE<sub>202</sub> of pig-PrP.

Therefore, conformational flexibility constraints associated with pig-PrP would limit the number of permissible PrP<sup>Sc</sup> conformations compatible with pig-PrP<sup>C</sup>, thus suggesting that pig-PrP<sup>C</sup> amino acid sequence may constitute a paradigm of low conformational flexibility that could confer high resistance to a wide diversity of prion strains. Amino acid changes in pig-PrP<sup>C</sup> would be responsible for its limited conformational flexibility compared with other, more susceptible species. This strengthens the transmission barrier for prion strains other than classic BSE, which may represent a thermodynamically favored PrP<sup>Sc</sup> conformation that is readily imprinted on PrP from a range of different species, accounting for the high promiscuity of the BSE strain in mammals.

Finally, the susceptibility of pigs to the classic BSE prion agent and their potential susceptibility to other prion strains not

tested here, such as CWD, should not be neglected and underlines the importance of continued monitoring of classic BSE cases and the prohibition of meat and bone meals to reduce the risk of prion transmission to pigs.

# **Supplementary Data**

Supplementary materials are available at *The Journal of Infectious Diseases* online. Consisting of data provided by the authors to benefit the reader, the posted materials are not copyedited and are the sole responsibility of the authors, so questions or comments should be addressed to the corresponding author.

## Notes

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