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 (PrPc) (porcine PrP [PoPrP]–Tg001) has been described, no natural cases of prion diseases in pig were reported. This study analyzed pig-PrPc 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-PrPc into scrapie-type PrP (PrPSc), independently of the species origin. Therefore, conformational flexibility constraints associated with pig-PrP would limit the number of permissible PrPSc conformations compatible with pig-PrPc, thus suggesting that pig-PrPc 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.
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\textsuperscript{C}) 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\textsuperscript{C}, 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 86/609/EC 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\textsuperscript{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 into PoPrP-Tg001 substrate. Then, 7 µL of inocula was diluted into PoPrP-Tg001 substrate. Then, 7 µL of inocula was diluted 1:100 or 1:10 or undiluted into PoPrP-Tg001 substrate. The different isolates were diluted 1:100 or 1:10 or undiluted into PoPrP-Tg001 substrate. Then, 7 µL of inocula was mixed with 63 µL 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 µL of the mixture was taken and immediately frozen as a nonamplification control. At least 2 tubes for each inoculum-substrate 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.

**WB Analysis**

A mass of 175 mg ± 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\textsuperscript{res} 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].
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-PrPC transgenic mice with either 5 or 6 octarepeats (Table 1). Considering that porcine-PrPC 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 (Table 1). 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 1).

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
positive propagation by means of PrP\textsuperscript{res} 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\textsuperscript{res}**

Comparison of the brain PrP\textsuperscript{res} 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\textsuperscript{res} glycosylation...
pattern with a predominant monoglycosylated band was observed. This profile was similar to that reported elsewhere in pigs inoculated with classic BSE [18, 21, 29, 35]. Comparison of PrPres 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 and in vitro using the PMCA technique. Although different combinations of prion strains and PrP<sup>C</sup> expressing donors have been used, only the classic BSE strain was able to propagate in mice expressing pig-PrP<sup>C</sup> 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<sup>C</sup> 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 3. Transmission of Mouse and Hamster Inocula in PoPrP-Tg001, Tga20 transgenic Mice or Hamsters**

| Inocula                  | Survival Time, Mean (SD), d [Diseased, PrP<sup>res</sup>-Positive/Inoculated Mice, No.]<sup>a</sup> |
|--------------------------|----------------------------------------------------------------------------------------------------|
|                           | PoPrP-Tg001                                                                                      |
|                           | 1st Passage                                                                                      |
|                           | 2nd Passage                                                                                      |
| BSE in Tga20              | 506 [1/6]<sup>b</sup>                                                                           |
| BSE in wt mouse           | 650 [1/6]<sup>b</sup>                                                                           |
| 22L                      | >650 [0/6]                                                                                       |
| RML                      | >650 [0/6]<sup>b</sup>                                                                           |
| 263K                     | >650 [0/7]                                                                                       |

| Inocula                  | Tga20 (1st Passage)                                                                              |
|--------------------------|--------------------------------------------------------------------------------------------------|
|                           | 154 (21) [5/5]                                                                                   |
|                           | 185 (39) [5/5]                                                                                   |
|                           | 112 (13) [4/4]                                                                                   |
|                           | 75 (7) [5/5]                                                                                     |
|                           | 87 (3) [5/5]                                                                                     |

| Inocula                  | Hamsters (1st Passage)                                                                            |
|--------------------------|--------------------------------------------------------------------------------------------------|
| sCJD 129 M/M T1           | >650 [0/6]<sup>b</sup>                                                                           |
| sCJD 129 M/M T1 0.08 02523,001 | ND                                                                                                 |
| sCJD 129 V/V T2           | >650 [0/6]<sup>b</sup>                                                                           |
| vCJD 129 M/M NHBY0/0014   | 556 (81) [6/6]<sup>b</sup>                                                                         |
| vCJD 129 M/M BC1458       | 530 (48) [6/6]                                                                                     |
| BSE in HuPrP-Tg340       | 488 (31) [5/6]                                                                                     |

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].

**Table 4. Transmission of Human Inocula in PoPrP-Tg001 and HuPrP-Tg340 Transgenic Mice**

| Inocula                  | Survival Time, Mean (SD), d [Diseased, PrP<sup>res</sup>-Positive/Inoculated Mice, No.]<sup>a</sup> |
|--------------------------|----------------------------------------------------------------------------------------------------|
|                           | PoPrP-Tg001                                                                                      |
|                           | 1st Passage                                                                                      |
|                           | 2nd Passage                                                                                      |
| sCJD 129 M/M T1 NHBX0/0001 | >650 [0/6]<sup>b</sup>                                                                           |
| sCJD 129 M/M T1 0.08 02523,001 | ND                                                                                                 |
| sCJD 129 V/V T2           | >650 [0/5]                                                                                       |
| vCJD 129 M/M NHBY0/0014   | 556 (81) [6/6]<sup>b</sup>                                                                         |
| vCJD 129 M/M BC1458       | 530 (48) [6/6]                                                                                     |
| BSE in HuPrP-Tg340       | 488 (31) [5/6]                                                                                     |

| Inocula                  | HuPrP-Tg340 (1st Passage)                                                                        |
|--------------------------|--------------------------------------------------------------------------------------------------|
|                           | 214 (6) [5/5]<sup>b</sup>                                                                           |
|                           | 187 (11) [6/6]                                                                                     |
|                           | 522 (38) [6/6]<sup>b</sup>                                                                         |
|                           | 626 (29) [6/6]<sup>b</sup>                                                                         |
|                           | 545 (148) [5/5]<sup>b</sup>                                                                         |
|                           | 614 (87) [6/6]<sup>b</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].
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 CWD-affected elk, white-tailed deer, and mule deer [21]. Although RT-QuIC (real-time quaking-induced conversion) enabled detection of PrP<sub>Sc</sub> in both orally and intracranially CWD-inoculated pigs as early as 6 months after inoculation, brain PrPres 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<sup>C</sup> could sustain replication of some prion strains, the transmission barrier of pig-PrP<sup>C</sup> for the analyzed

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

| Seed | Amplification by Serial PMCA Round, %<sup>a</sup> |
|------|-----------------------------------|
|      | 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  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  |
| Ca-BSE-H 07-644 | 0  | 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  | 0  |
| Sheep-Sc pool pre-75 cattle P75-7 | 0  | 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  | 0  |
| Sheep-Sc PS48<sup>c</sup> | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  |
| Sheep-Sc PS13<sup>c</sup> | 0  | 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  | 0  |
| Sheep-Sc PS42 | 0  | 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  | 0  |
| Goat-Sc F10 | 0  | 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  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  | 0  |
| 22L | 0  | 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  | 0  |
| 263K | 0  | 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  | 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  | 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  | 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.

*aPercentage of positive tubes (showing PrPres) of the total number of tubes sonicated (n = 6).*

*bDescribed elsewhere [29].*

*cDescribed elsewhere [18].*
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-PrPSc and pigs [16, 17]. Interestingly, the ability of pig-PrP to sustain replication of the classic BSE strain regardless of the donor PrPSc amino acid sequence used as inoculum suggests that only a very restricted number of PrPSc conformers, such as classic BSE, present molecular compatibility with pig-PrP for prion propagation. Thus, strain-specific PrPSc 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 PrPSc rather than the PrPc encoded by the host.

On the other hand, the high resistance of pig-PrPc 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 PrPSc 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-PrPc. 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 β2-α2 loop of PrPSc protein (residues 169–179 of porcine sequence) may considerably affect the transmission barrier [41–43]. As a paradigm, the Q171R polymorphism present in the β2-α2 loop of sheep-PrPSc 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-PrPSc is present only in species with low susceptibility to prion infection, such as rabbits (Figure 2B).

The amino acid change found in the β2-α2 loop in pig-PrPSc would alter the flexibility of the β2-α2 loop, strengthening the transmission barrier for diverse prion strains other than classic BSE. Other amino acid changes in the pig-PrPSc would also participate in the limited capacity of this protein to sustain replication of different prion strains and hence to adopt the PrPSc conformations associated with those strains. From these changes, 226Y227 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 226Y227 amino acid changes are present in PrPSc from other species alleged to be reluctant to conformational conversion to PrPSc, such as horses [22]. E226 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 CWD-susceptible cervids. CWD prion strain propagation is stable in transgenic mice expressing E226 cervid-PrPSc, whereas mice expressing Q226 cervid-PrPSc unsteadily generate CWD mixed strains [45]. Furthermore, these 226Y227 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 226Y227 amino acids can be relevant in the restricted ability of pig-PrP to sustain prion replication.

Overall, our results demonstrate that pig-PrPc can be converted in pig-PrPSc 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-PrPc into pig-PrPSc.
Therefore, conformational flexibility constraints associated with pig-PrP would limit the number of permissible PrPSc conformations compatible with pig-PrPC, thus suggesting that pig-PrPC 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-PrPC 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 PrPSc 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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**Potential conflicts of interest.** All authors: No reported conflicts. All authors have submitted the ICMJE Form for Disclosure of Potential Conflicts of Interest. Conflicts that the editors consider relevant to the content of the manuscript have been disclosed.

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