#### RESEARCH LETTERS

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#### References

- Yong D, Toleman MA, Giske CG, Cho HS, Sundman K, Lee K, et al. Characterization of a new metallo-β-lactamase gene, *bla*<sub>NDM-1</sub>, and a novel erythromycin esterase gene carried on a unique genetic structure in *Klebsiella pneumoniae* sequence type 14 from India. Antimicrob Agents Chemother. 2009;53:5046–54. http://dx.doi.org/ 10.1128/AAC.00774-09
- Peirano G, Lascols C, Hackel M, Hoban DJ, Pitout JD. Molecular epidemiology of *Enterobacteriaceae* that produce VIMs and IMPs from the SMART surveillance program. Diagn Microbiol Infect Dis. 2014;78:277–81. http://dx.doi.org/ 10.1016/j.diagmicrobio.2013.11.024
- Peirano G, Bradford PA, Kazmierczak KM, Badal RE, Hackel M, Hoban DJ, et al. Global incidence of carbapenemase-producing *Escherichia coli* ST131. Emerg Infect Dis. 2014;20:1928–31. http://dx.doi.org/10.3201/eid2011.141388
- Chou A, Roa M, Evangelista MA, Sulit AK, Lagamayo E, Torres BC, et al. Emergence of *Klebsiella pneumoniae* ST273 carrying *bla*<sub>NDM-7</sub> and ST656 carrying *bla*<sub>NDM-1</sub> in Manila, Philippines. Microb Drug Resist. 2016;22:585–8. http://dx.doi.org/10.1089/mdr.2015.0205
- Clinical and Laboratory Standards Institute. Performance standards for antimicrobial susceptibility testing: twenty-fifth informational supplement (M100–S25). Wayne (PA): The Institute, 2015.
- Nordmann P, Poirel L, Dortet L. Rapid detection of carbapenemase-producing *Enterobacteriaceae*. Emerg Infect Dis. 2012;18:1503–7. http://dx.doi.org/10.3201/eid1809.120355
- Milillo M, Kwak YI, Snesrud E, Waterman PE, Lesho E, McGann P. Rapid and simultaneous detection of *bla*<sub>KPC</sub> and *bla*<sub>NDM</sub> by use of multiplex real-time PCR. J Clin Microbiol. 2013;51:1247–9. http://dx.doi.org/10.1128/JCM.03316-12
- Milillo M, Kwak YI, Snesrud E, Waterman PE, Lesho E, McGann P. Correction for Milillo et al., Rapid and simultaneous detection of blaKPC and blaNDM by use of multiplex real-time PCR. J Clin Microbiol. 2015;53:1460. http://dx.doi.org/10.1128/ JCM.00373-15
- Schwaber MJ, Klarfeld-Lidji S, Navon-Venezia S, Schwartz D, Leavitt A, Carmeli Y. Predictors of carbapenem-resistant *Klebsiella pneumoniae* acquisition among hospitalized adults and effect of acquisition on mortality. Antimicrob Agents Chemother. 2008;52:1028–33. http://dx.doi.org/10.1128/AAC.01020-07

 Kulkova N, Babalova M, Sokolova J, Kremery V. First report of New Delhi metallo-β-lactamase-1–producing strains in Slovakia. Microb Drug Resist. 2015;21:117–20. http://dx.doi.org/10.1089/ mdr.2013.0162

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## Chronic Wasting Disease Prion Strain Emergence and Host Range Expansion

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Human and mouse prion proteins share a structural motif that regulates resistance to common chronic wasting disease (CWD) prion strains. Successful transmission of an emergent strain of CWD prion, H95<sup>+</sup>, into mice resulted in infection. Thus, emergent CWD prion strains may have higher zoonotic potential than common strains.

Chronic wasting disease (CWD) is a contagious prion disease of cervids that is spreading globally. CWD is enzootic in multiple cervid species, including deer and elk; the major foci of disease are Colorado/Wyoming (USA), Wisconsin/Illinois (USA), and Alberta/Saskatchewan (Canada). CWD is also present in captive cervids in South Korea and wild reindeer and moose in Norway (https://www.nwhc. usgs.gov/images/cwd/cwd\_map.jpg). CWD results from the conformational transformation of the host-encoded cellular prion protein (PrP<sup>c</sup>) into protease-resistant, detergent-insoluble,  $\beta$ -sheet rich, amyloidogenic conformers, termed prions (PrP<sup>CWD</sup>). Within their conformation, prion strains encipher the information that directs the templated misfolding and aggregation of PrP<sup>c</sup> molecules into additional prions (1).

Although the sequence homology of PrP among mammals is high, the ability of particular prion strains to cause disease in different species is determined by the conformational compatibility between a given strain and the host  $PrP^{C}(2)$ . We

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previously identified 2 strains of CWD prion in white-tailed deer (*3*), Wisc-1 and H95<sup>+</sup>; these strains exhibit distinct biological properties in deer and transgenic cervidized mice. To ascertain the host range of different strains from cervids, we inoculated CWD prions isolated from experimentally infected deer with different *PRNP* genotypes (Q95G96 [wild type (wt)], S96/wt, H95/wt, and H95/S96) and from elk (CWD2 strain) into hamsters and mice. All isolates have been successfully transmitted into transgenic mice expressing wt cervid PrP and contain high titers of CWD prions (*3*).

Mice inoculated with H95<sup>+</sup> CWD prions succumbed to clinical disease at 575  $\pm$  47 or 692  $\pm$  9 days, depending on the H95<sup>+</sup> isolate (Table). Mice inoculated with Wisc-1 or elk CWD or uninfected deer homogenates were euthanized at day 708 after infection with no signs of prion disease. Clinical signs of H95<sup>+</sup> CWD in C57Bl/6 mice included ataxia, lethargy, tail rigidity, and dermatitis. Protease-resistant PrP<sup>CWD</sup> was present in all mice infected with H95<sup>+</sup> prions and was not detected in mice infected with Wisc-1 or CWD2 (online Technical Appendix, https://wwwnc.cdc. gov/EID/article/23/9/16-1474-Techapp1.pdf).

In contrast to mice, hamsters succumbed to clinical disease when inoculated with Wisc-1 CWD prions but were less susceptible to H95<sup>+</sup> CWD prions (Table). Clinical signs of CWD in hamsters began with lethargy and, upon arousal, retrocollis; as the disease progressed, lethargy declined with increased dystonic movement including ataxia and tremors. Hyperesthesia was not observed. Subclinical disease (no clinical signs but PrP-res positive by Western blot) was observed in a subset of hamsters (online Technical Appendix).

Successful interspecies prion transmission at the molecular level depends on the compatibility of the invading prion conformers and structural determinants imposed by host  $PrP^c$ . One structural motif is the loop region between  $\beta$  sheet 2 and  $\alpha$  helix 2 of  $PRP^c$  at aa 170–174 (online Technical Appendix). Host species containing  $PrP^c$  molecules with a flexible  $\beta 2-\alpha 2$  loop (mice and humans) are hypothesized to be incompatible with prions derived from species containing a rigid loop (deer and elk) (4,5). Previous attempts to transmit CWD to mice have failed (6,7). Our data show that prions from a prototypic rigid-loop species (deer) can transmit to a flexible-loop species (mice). The transmission is strain dependent. H95<sup>+</sup> overrides the conformational restriction imposed by the mouse PrP flexible loop that Wisc-1 and CWD2 cannot overcome, suggesting that the invading prion strain is a dominant contributor to the species/transmission barrier. How the N terminal amino acid polymorphism (Q95H) affects the conformation of PrP, altering the deer-to-mouse transmission barrier, is unknown. Further structural studies may clarify the effect of N terminal residues on  $\beta 2-\alpha 2$  loop rigidity.

Transmission of H95<sup>+</sup> CWD prions to mice further confirms the value of specifying strain when defining species barriers. Experimental transmission of CWD prion into macaques and transgenic mice expressing human PrP suggests a considerable transmission barrier to CWD prions (although squirrel monkeys are susceptible), and human prion protein is converted inefficiently in vitro ( $\delta$ ,9). Successful infection of a flexible-loop species (mice) with H95<sup>+</sup> CWD raises concerns for the potential pathogenicity of H95<sup>+</sup> prions to other flexible-loop species. Transmission studies with Wisc-1 and H95<sup>+</sup> in transgenic humanized and bovinized mice are ongoing.

The increasing prevalence of CWD indicates selection for cervids with resistance alleles, such as S96 and H95. Genetic resistance to a given prion strain selects for the emergence of novel prion strains with altered properties such as H95<sup>+</sup> and Nor98 (*3,10*). The iterative transmission of CWD prions to cervids with protective alleles of PrP<sup>c</sup> and the consequent emergence of new CWD prion strains highlights the dynamics of the CWD panzootic and the value of characterizing the host range of emergent CWD prion strains.

Recipient and CWD inocula	PrP-res+			
	No.	Clinical	Subclinical	Incubation period, d
Mice				
wt/wt	6	0	0	NA
S96/wt	6	0	0	NA
H95/wt	7	5	2	669, 671, 706, 706, 706
H95/S96	7	7	0	306, 593, 593, 593, 593, 673, 675
Elk	4	0	0	NA
Control	4	0	0	NA
Hamsters				
wt/wt	8	3	5	652, 653, 653
S96/wt	8	1	4	634
H95/wt	8	1	6	652
H95/S96	8	0	1	NA
Elk	8	2	2	673, 719
Control	8	0	0	NA

\*Mice infected with CWD prions were observed for up to 708 d; hamsters infected with white-tailed deer and elk CWD prions were observed for 659 and 726 d, respectively. Control mice and hamsters were inoculated with brain homogenates from CWD-negative wt/wt deer. CWD, chronic wasting disease; NA, not applicable; PrP-res+, positive for proteinase-K-resistant prion protein; wt, wild type.

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Dr. Herbst is a research associate and Dr. Duque Velásquez is a postdoctoral fellow at the University of Alberta. Their primary research interest is the mechanism(s) of pathogenicity underlying neurodegeneration, as exemplified by prion diseases in animals and humans.

#### References

- Bessen RA, Marsh RF. Identification of two biologically distinct strains of transmissible mink encephalopathy in hamsters. J Gen Virol. 1992;73:329–34. http://dx.doi.org/10.1099/ 0022-1317-73-2-329
- Capobianco R, Casalone C, Suardi S, Mangieri M, Miccolo C, Limido L, et al. Conversion of the BASE prion strain into the BSE strain: the origin of BSE? PLoS Pathog. 2007;3:e31. http://dx.doi.org/10.1371/journal.ppat.0030031
- Duque Velásquez C, Kim C, Herbst A, Daude N, Garza MC, Wille H, et al. Deer prion proteins modulate the emergence and adaptation of chronic wasting disease strains. J Virol. 2015; 89:12362–73. http://dx.doi.org/10.1128/JVI.02010-15
- Gossert AD, Bonjour S, Lysek DA, Fiorito F, Wüthrich K. Prion protein NMR structures of elk and of mouse/elk hybrids. Proc Natl Acad Sci U S A. 2005;102:646–50. http://dx.doi.org/10.1073/ pnas.0409008102
- Sigurdson CJ, Nilsson KP, Hornemann S, Heikenwalder M, Manco G, Schwarz P, et al. De novo generation of a transmissible spongiform encephalopathy by mouse transgenesis. Proc Natl Acad Sci U S A. 2009;106:304–9. http://dx.doi.org/10.1073/ pnas.0810680105
- Raymond GJ, Raymond LD, Meade-White KD, Hughson AG, Favara C, Gardner D, et al. Transmission and adaptation of chronic wasting disease to hamsters and transgenic mice: evidence for strains. J Virol. 2007;81:4305–14. http://dx.doi.org/10.1128/ JVI.02474-06
- 7. Kurt TD, Bett C, Fernández-Borges N, Joshi-Barr S, Hornemann S, Rülicke T, et al. Prion transmission prevented by modifying the  $\beta 2-\alpha 2$  loop structure of host PrP<sup>c</sup>. J Neurosci. 2014;34:1022–7. http://dx.doi.org/10.1523/JNEUROSCI.4636-13.2014
- Race B, Meade-White KD, Phillips K, Striebel J, Race R, Chesebro B. Chronic wasting disease agents in nonhuman primates. Emerg Infect Dis. 2014;20:833–7. http://dx.doi.org/10.3201/ eid2005.130778
- Barria MA, Telling GC, Gambetti P, Mastrianni JA, Soto C. Generation of a new form of human PrP(Sc) in vitro by interspecies transmission from cervid prions. J Biol Chem. 2011;286:7490–5. http://dx.doi.org/10.1074/jbc.M110.198465
- Le Dur A, Béringue V, Andréoletti O, Reine F, Laï TL, Baron T, et al. A newly identified type of scrapie agent can naturally infect sheep with resistant PrP genotypes. Proc Natl Acad Sci U S A. 2005;102:16031–6. http://dx.doi.org/10.1073/pnas.0502296102

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# Rabies Virus Transmission in Solid Organ Transplantation, China, 2015–2016

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We report rabies virus transmission among solid organ transplantation recipients in Changsha, China, in 2016. Two recipients were confirmed to have rabies and died. Our findings suggest that more attention should be paid to the possibility of rabies virus transmission through organ transplantation for clinical and public health reasons.

In 2016, Zhou et al. reported a case of rabies virus transmission in China that was probably a result of organ transplantation (1). We report on rabies transmission that occurred among solid organ transplant recipients in Changsha, China, during December 2015–January 2016.

In November 2015, the donor, a previously healthy boy, showed development of fever, insomnia, and agitation. On day 6 of infection, these symptoms progressed, and he was sent to a healthcare center. At this time, he experienced weariness, no desire to drink water, poor appetite, and panic. One day later, he began vomiting, and was admitted to a local hospital (hospital A), where he exhibited anemophobia, convulsions, limb rigidity, and hypersalivation. The patient was moved to hospital B (days 7-14) in Changsha. At admission, some examination findings indicated a possibility of viral encephalitis (online Technical Appendix, Table 1, https://wwwnc.cdc.gov/EID/article/23/9/16-1704-Techapp1.pdf). Subhypothermia hibernation therapy and assisted ventilation were administered within 72 hours of admission, and the patient's vital signs became stable. On day 10, hyponatremia was observed, and on day 11, the patient again became febrile and tachycardic, with hypertensive abdominal distention and alimentary tract hemorrhage. On day 13, viral encephalitis was diagnosed, and rabies was suspected. However, rabies virus antibody tests performed on serum samples by using ELISA yielded negative results.

On day 14, the patient was transferred to hospital C, where he became comatose and was declared brain dead. Permission was granted for organ donation, because no specific pathogen had been detected and China's organ

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