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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, H951, 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 (PrPc) into protease-resistant, detergent-insoluble, β-sheet rich, amyloidogenic conformers, termed prions (PrPSc). Within their conformation, prion strains encipher the information that directs the templated misfolding and aggregation of PrPSc 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 PrPc (2). We

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previously identified 2 strains of CWD prion in white-tailed deer (3), Wisc-1 and H95\(^+\); 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\(^+\) CWD prions succumbed to clinical disease at 575 ± 47 or 692 ± 9 days, depending on the H95\(^+\) 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\(^+\) CWD in C57Bl/6 mice included ataxia, lethargy, tail rigidity, and dermatitis. Protease-resistant \(PrP^{\text{res+}}\) was present in all mice infected with H95\(^+\) prions and was not detected in mice infected with Wisc-1 or CWD2 (online Technical Appendix, https://wwwnc.cdc.gov/EID/article/23/9/14-1747-Techapp1.pdf).

In contrast to mice, hamsters succumbed to clinical disease when inoculated with Wisc-1 CWD prions but were less susceptible to H95\(^+\) 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 \(PrP\). One structural motif is the loop region between \(\beta\) sheet 2 and \(\alpha\) helix 2 of \(PRP\) at aa 170–174 (online Technical Appendix). Host species containing \(PrP\) molecules with a flexible \(\beta\)-2-\(\alpha\) 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\(^+\) 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-\(\alpha\) loop rigidity.

Transmission of H95\(^+\) CWD prions to mice further confirms the value of specifying strain when defining species barriers. Experimental transmission of CWD prions to 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 (8, 9). Successful infection of a flexible-loop species (mice) with H95\(^+\) CWD raises concerns for the potential pathogenicity of H95\(^+\) prions to other flexible-loop species. Transmission studies with Wisc-1 and H95\(^+\) 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\(^+\) and Nor98 (3,10). The iterative transmission of CWD prions to cervids with protective alleles of \(PrP\) 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.

### Table. Results of CWD prion inoculation into rodents\(^*\)

<table>
<thead>
<tr>
<th>Recipient and CWD inocula</th>
<th>(PrP)-res+</th>
<th>Incubation period, d</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Mice</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>wt/wt</td>
<td>6</td>
<td>0</td>
</tr>
<tr>
<td>S96/wt</td>
<td>6</td>
<td>0</td>
</tr>
<tr>
<td>H95/wt</td>
<td>7</td>
<td>5</td>
</tr>
<tr>
<td>H95/S96</td>
<td>7</td>
<td>7</td>
</tr>
<tr>
<td>Elk</td>
<td>4</td>
<td>0</td>
</tr>
<tr>
<td>Control</td>
<td>4</td>
<td>0</td>
</tr>
<tr>
<td><strong>Hamsters</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>wt/wt</td>
<td>8</td>
<td>3</td>
</tr>
<tr>
<td>S96/wt</td>
<td>8</td>
<td>1</td>
</tr>
<tr>
<td>H95/wt</td>
<td>8</td>
<td>1</td>
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<tr>
<td>H95/S96</td>
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<td>0</td>
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<tr>
<td>Elk</td>
<td>8</td>
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<tr>
<td>Control</td>
<td>8</td>
<td>0</td>
</tr>
</tbody>
</table>

\(^*\) 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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References

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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 weakness, 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 anemophoria, 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...