

Cyclodextrins Inhibit Replication of Scrapie Prion Protein in Cell Culture[∇]

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Prion diseases are fatal neurodegenerative disorders that are caused by the conversion of a normal host-encoded protein, PrP^C, to an abnormal, disease-causing form, PrP^{Sc}. This paper reports that cyclodextrins have the ability to reduce the pathogenic isoform of the prion protein PrP^{Sc} to undetectable levels in scrapie-infected neuroblastoma cells. Beta-cyclodextrin removed PrP^{Sc} from the cells at a concentration of 500 μM following 2 weeks of treatment. Structure activity studies revealed that antiprion activity was dependent on the size of the cyclodextrin. The half-maximal inhibitory concentration (IC₅₀) for beta-cyclodextrin was 75 μM, whereas α-cyclodextrin, which possessed less antiprion activity, had an IC₅₀ of 750 μM. This report presents cyclodextrins as a new class of antiprion compound. For decades, the pharmaceutical industry has successfully used cyclodextrins for their complex-forming ability; this ability is due to the structural orientation of the glucopyranose units, which generate a hydrophobic cavity that can facilitate the encapsulation of hydrophobic moieties. Consequently, cyclodextrins could be ideal candidates for the treatment of prion diseases.

Transmissible spongiform encephalopathies (TSEs), also known as prion diseases, are a group of fatal neurodegenerative disorders that affect humans and animals and are unique in that they can be sporadic, inherited, or of transmissible origin (49). The TSEs affecting humans include kuru, Gerstmann-Straussler Scheinker syndrome, fatal familial insomnia, and Creutzfeldt-Jakob disease. Among the TSEs affecting other animals are natural scrapie of sheep and goats and bovine spongiform encephalopathy of cattle. Central to the control of the development of all TSEs is the prion protein, which in humans is the product of a single gene located on chromosome 20 (48). The prion protein exists in at least two conformational forms with distinct physiochemical properties. The normal cellular form of the prion protein (PrP^C) is expressed at high levels in neuronal cells (6, 59). It is a glycosyl-phosphatidylinositol (GPI)-anchored cell surface protein (59), and during TSE disease, it is converted to an abnormal, pathological form known as PrP^{Sc}. Unlike PrP^C, PrP^{Sc} is detergent insoluble and partially proteinase resistant (31). The tendency of PrP^{Sc} to aggregate in the brain leads to its characteristic neuropathological features, such as spongiform degeneration of the brain.

PrP^C and PrP^{Sc} have the same amino acid sequence but differ dramatically in conformation. PrP^C consists of 42% α-helix and 3% β-sheet, while PrP^{Sc} possesses a much higher β-sheet content of 43% and a lower α-helix content of 30% (41, 52). The conversion of PrP^C to the pathogenic isoform is a critical event. It is believed that the interaction of endogenous PrP^C substrate with the pathogenic PrP^{Sc} template causes PrP^C to unfold and refold as the β-sheet-rich isoform PrP^{Sc}. This step starts a chain reaction, where each newly converted

PrP^{Sc} molecule interacts with more PrP^C molecules, fueling the formation of PrP^{Sc} (17, 48).

Lipid rafts, which are found in the plasma membrane, are also known as detergent-resistant microdomains (DRMs) (9, 38, 63) and are enriched in cholesterol and sphingolipids. The formation of rafts is dependent on cholesterol (51), which is believed to function as a spacer between hydrocarbon chains of sphingolipids, thereby holding the raft assembly together (56). The extraction of cholesterol from membranes leads to the dissociation of proteins from rafts (15, 25, 53). These rafts are important in many cell processes, such as membrane sorting and trafficking and signal transduction (23). As has been reported for other GPI-anchored proteins, both PrP^C and PrP^{Sc} have been found to be associated with DRMs, and DRMs have been hypothesized to be involved in both the function of PrP^C (34) and the conversion of PrP^C to the pathogenic isoform (37, 38, 63). Moreover, in addition to the GPI anchor, PrP presents a sphingolipid binding domain that allows the interaction of the protein with the polar head groups of the sphingolipids (28, 64). A number of studies have provided evidence that in cell culture, conversion requires association with rafts (22, 63, 68), which may explain why cholesterol extraction modifies PrP^{Sc} formation in cell culture (63).

To date, there is no known cure for TSEs, as standard approaches to treating the diseases have proved ineffective. A range of compounds have been tested in the search for treatments for TSEs, and these compounds have been directed against targets such as preventing a rise in calcium, preventing apoptosis, directly interfering with conversion, and altering prion trafficking and dominant negative inhibition (2, 11, 12, 14, 19, 24, 29, 30, 35, 43–45, 46, 47, 54). Over the last several years, a wide variety of compounds have been tested for their effect in the treatment of TSEs, but unfortunately, none have been successful at completely eliminating the disease when given either immediately before, during, or after disease onset. Limitations on success have also related to the toxicity of antiprion compounds to humans and/or an inability to cross

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the blood-brain barrier (BBB), where most of the PrP^{Sc} accumulates.

Prion diseases share neuropathological characteristics with other neurodegenerative diseases, such as Parkinson's disease and Alzheimer's disease (AD). These characteristics include intracellular and extracellular aggregates of proteinaceous fibrils resulting from irregular protein-protein interactions (65). In a search for a new antiprion compound, β -cyclodextrin (β -CD), which reduces the toxic effects of an AD-associated protein (β -amyloid [A β] [amino acids 1 to 40]) in cell culture (10), was tested here. The formation of β -sheet fibrils is a critical event in AD, and β -CD, by attenuating fibrillization of the toxic peptide, appears to inhibit the toxic effects of A β (10). The CDs are macrocyclic, nonreducing maltooligosaccharides made from α -1,4-linked glucose units (58). α -, β -, and γ -CDs, composed of 6, 7, and 8 glucosyl units, respectively, are the parent CDs (61) and have attracted interest as natural complexing agents and as vehicles to increase drug delivery, bio-availability, and solubility (18, 58, 66).

In this report, the ability of noncytotoxic concentrations of β -CD and methyl- β -CD (M β -CD) to clear PrP^{Sc} from ScN2a cells following 2 weeks of treatment is demonstrated. Of the naturally occurring CDs, β -CD, with a half-maximal inhibitory concentration (IC₅₀) of 75 μ M, was much more efficient at clearing infection than was α -CD or γ -CD, with IC₅₀s of 750 μ M and 1,150 μ M, respectively. The antiprion activity of β -CDs was dependent not only on the ring size of the CD but also on the cyclic nature of the molecule. The antiprion activity of the CDs tested related to their ability to sequester and move molecules. β -CD modified the location of both PrP^C and PrP^{Sc}, but it did so differently, separating the isoforms into different lipid domains. Such separations have the potential to restrict the conversion process. β -CD also bound to PrP^C and interfered with abnormal prion conversion. These data indicate that the antiprion action of β -CDs is derived from a combination of mechanisms.

MATERIALS AND METHODS

Reagents and antibodies. α -, β -, and γ -CDs; M β -CD; maltohexaose (G6); maltoheptaose (G7); and water-soluble cholesterol complex were purchased from Sigma-Aldrich. Cell culture reagents, such as OptiMEM, fetal bovine serum, glutamine, and penicillin-streptomycin, were Gibco products supplied by Biosciences. Secondary antibody anti-mouse peroxidase conjugate and cholera toxin B subunit type Inaba 569B peroxidase conjugate were purchased from Calbiochem. Epoxy-activated Sepharose 6B was obtained from Amersham Pharmacia. Pefabloc was purchased from Boehringer Mannheim, and G418 was purchased from Promega; all other reagents were purchased from Sigma-Aldrich. The primary antibody 3F4 (1:10,000 dilution) used in this study is characterized as recognizing the peptide epitope comprising amino acids 109 to 112 of human PrP and was purchased from Signet Laboratories. The primary antibody 8H4 used in this study is characterized as recognizing the peptide epitope comprising amino acids 175 to 185 of PrP and was described previously (27).

Cell culture and effect of compounds on PrP^C, PrP^{Sc}, and cholesterol levels. Neuroblastoma cells (N2a) transfected with mouse PrP (N2a58) and N2a58 cells infected with the 22L scrapie strain (N2a22L), as reported previously (39), were obtained from S. Lehmann (Montpelier, France). To obtain cells producing high levels of PrP^{Sc}, the N2a22L cell line was subcloned and is referred to as N2a22L20 in this study. 3F4-tagged mouse PrP^C (MoPrP^C) cDNA in the pcDNA3 vector was obtained from S. Lehmann (Montpelier, France). N2a cells were transfected with the cDNA using FuGENE transfection reagent according to the manufacturer's instructions. Stably transfected cells were screened and maintained using G418 resistance. The N2a22L, N2a22L20, N2a58, and 3F4MoPrP^C N2a cells were cultured in OptiMEM supplemented with 10% fetal

bovine serum, penicillin-streptomycin, 2 mM glutamine, and 300 μ g/ml Geneticin (G418 sulfate) in a humidified atmosphere.

On passage, cells were treated with the compounds indicated in the text at the concentrations and times indicated unless otherwise stated. All compounds were solubilized in Millipore water and were sterilized by filtering through a 0.2- μ m filter prior to their being added to the medium. Cells were lysed in cold lysis buffer (0.5% [wt/vol] sodium deoxycholate, 0.5% [vol/vol] Triton X-100, 150 mM NaCl, and 50 mM Tris, pH 7.5) and analyzed for PrP^C or PrP^{Sc}. The total protein concentration was measured using the bicinchoninic acid protein assay kit (Sigma).

Total cholesterol was measured by the method of Mizoguchi et al. (32). Twenty-five microliters of sample and 150 μ l reagent solution [0.05 M MES (morpholineethanesulfonic acid) buffer, pH 6.1, 0.5 U ml⁻¹ cholesterol oxidase, 4 U ml⁻¹ cholesterol esterase, 0.2 mM 4-aminoantipyrine, 1 mM *N*-ethyl-*N*-(2-hydroxy-3-sulfoethyl)-3,5-dimethoxyaniline sodium salt, and 4 U ml⁻¹ horseradish peroxidase] were added and incubated at 37°C for 15 min, and cholesterol was measured at 600 nm.

Western blotting and densitometry. For PrP^C analysis, lysates were adjusted to 18 μ g of protein in gel loading buffer and boiled for 5 min prior to being loaded onto 12% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) gels, followed by Western blotting. For PrP^{Sc} analysis, lysates were prepared to 200 μ g for N2a22L cells and 50 μ g for N2a22L20 cells and then digested with 16 μ g of proteinase K (pK)/mg of protein at 37°C for 15 min. One millimolar Pefabloc was then added, and samples were incubated on ice for 5 min, followed by centrifugation at 14,000 rpm for 45 min. The resulting pellets were then resuspended in loading buffer, boiled for 5 min, and loaded onto 12% SDS-PAGE, followed by Western blotting using standard techniques. PrP^C and PrP^{Sc} were detected by incubating immunoblots with antibody 8H4 (3F4-tagged MoPrP^C was detected with antibody 3F4), followed by a horseradish peroxidase secondary antibody, and developed by enhanced chemiluminescence. Densitometry was performed with Bio-Rad Imager analysis software.

Sucrose density gradients. N2a22L20 cells from 3-by-60-mm dishes were lysed in 400 μ l of ice-cold sucrose gradient (SG) lysis buffer (25 mM Tris-HCl, pH 7.5, 150 mM NaCl, 5 mM EDTA and 1% Triton X-100) on ice for 30 min. Lysates were then spun at 4,850 rpm for 10 min, and the 400- μ l supernatant was adjusted to 40% (wt/vol) sucrose prepared in SG lysis buffer without detergent. One milliliter 25% (wt/vol) sucrose and 0.5 ml 5% (wt/vol) sucrose, both in SG lysis buffer without detergent, were then added. The gradient was then spun at 37,000 rpm for 18 h at 4°C. Fractions (10 \times 230 μ l) were collected from the top of the tube and processed for PrP^C, PrP^{Sc}, and ganglioside GM₁, a marker of DRM, as follows.

For PrP^C detection, 100 μ l of each fraction was diluted with 300 μ l TN (10 mM Tris, pH 7.8, and 150 mM NaCl) and then made to 1% (wt/vol) Sarkosyl (to precipitate PrP^{Sc}) and incubated on ice for 30 min. Fractions were then spun at 45,000 rpm for 2 h at 4°C, and the supernatants were methanol (MeOH) precipitated. The precipitated pellet was resuspended in gel loading buffer, and PrP^C was visualized by using 12% SDS-PAGE and Western blotting with 8H4 antibody. For PrP^{Sc} detection, 120 μ l from each protein fraction was obtained and protein was determined using the bicinchoninic acid protein assay kit and treated with 16 μ g pK/mg protein for 15 min at 37°C, followed by centrifugation at 70,000 rpm for 45 min. PrP^{Sc} was visualized by using SDS-PAGE and Western blotting with 8H4 antibody. For GM₁ detection, 10 μ l from each fraction was dotted onto a nitrocellulose membrane, which was then blocked in 3% bovine serum albumin for 3 h and incubated with cholera toxin B (1:4,000 dilution) for 1 h. Following four 10-min washes in TTBS (0.1 M Tris, 0.154 NaCl, pH 7.5, containing 0.3% [vol/vol] Tween 20), blots were developed by enhanced chemiluminescence.

Triton X-100 solubility assay. Cells were washed twice with ice-cold phosphate-buffered saline and were then left on ice for 15 min. The cells were then scraped into ice-cold buffer containing 1% Triton X-100, 150 mM NaCl, and 10 mM Tris-HCl, pH 7.8, and left on ice for 15 min. The lysate was then spun at 39,000 rpm in the ultracentrifuge Sorvall Discovery N12SE Micro Ultra at 2°C for 30 min. The pellet (insoluble protein) was resuspended in 1 \times loading buffer, and the supernatant (soluble protein) was methanol precipitated prior to centrifugation at 14,000 rpm for 10 min. The resulting pellet was resuspended in 1 \times loading buffer, and PrP^C in the soluble and insoluble fractions was analyzed by 12% SDS-PAGE and immunoblotting with antibody 8H4.

Affinity chromatography and PrP binding. Freeze-dried epoxy-activated Sepharose 6B was reconstituted in water and activated with 0.1 M NaOH according to the manufacturer's instructions (Pharmacia). The gel was incubated with or without 0.02 M β -CD in 0.1 M NaOH for 16 h at 40°C with shaking. The gel was then washed in sequence with 0.1 M NaOH, distilled water, 0.1 M NaOH, and finally 0.1 M acetate buffer, pH 4.0. Both the control gel (without β -CD) and the β -CD-complexed gel were blocked with 0.2 M glycine, pH 8.0, for 4 h at 40°C.

The gel was then washed with 0.1 M NaOH, followed by 0.1 M acetate buffer, pH 4.0.

For the binding of PrP to affinity gels containing β -CD, gel made as described above with β -CD incorporated was washed three times with 0.5 M HEPES, pH 7.5, and was resuspended in this buffer. Eight hundred micrograms of N2a cells transfected with 3F4 MoPrP^C lysed in cold lysis buffer (0.5% [wt/vol] sodium deoxycholate, 0.5% [vol/vol] Triton X-100, 150 mM NaCl, and 50 mM Tris, pH 7.5) was then added to 800 μ g of gel and mixed by inversion over a 15-min period. The mixture of gel and lysate was spun down at 1,000 rpm for 1 min, and the supernatant was removed and MeOH precipitated (S1). The gel was then washed four times with 500 μ l of 0.5 M HEPES, pH 7.5, with a 15-min mixing period between each wash. A total of 500 μ l of 0.5 M HEPES, pH 7.5, containing 500 μ M β -CD was then added, and the solution was mixed as before for 15 min and then spun at 1,000 rpm for 1 min. The supernatant was removed and MeOH precipitated (eluent 1 [E1]). A total of 500 μ l of 0.5 M HEPES, pH 7.5, containing 1 mM β -CD, was then added to the gel, and the solution was mixed as before for 15 min. It was then spun, and the supernatant was removed and MeOH precipitated (E2). The remaining 800 μ l of gel was boiled in 400 μ l of 1 \times loading buffer for 10 min. The mixture was spun at 1,000 rpm for 1 min, and the supernatant was taken and analyzed for PrP^C (E3).

Conversion reaction. Two-by-60-mm dishes of N2a22L20 and 3F4 MoPrP^C N2a cells rinsed twice with ice-cold phosphate-buffer saline were scraped into 300 μ l of ice-cold conversion buffer (0.1 M MES, pH 6.2, and 1 μ M CaCl₂) and subjected to two freeze-thaw cycles and vortexed. The suspension was then incubated at 37°C for 72 h as indicated in the text. Two volumes of ice-cold lysis buffer (0.5% [wt/vol] sodium deoxycholate, 0.5% [vol/vol] Triton X-100, 150 mM NaCl, and 50 mM Tris, pH 7.5, 1 μ g/ml pepstatin A, 1 μ g/ml leupeptin, and 2 mM EDTA) was added, and samples were MeOH precipitated. The pellet was resuspended in 50 μ l of ice-cold lysis buffer, and samples were then assayed for insolubility by centrifuging them at 70,000 rpm for 40 min at 4°C. The pellet (insoluble fraction) was resuspended in loading buffer. The supernatant (soluble fraction) was precipitated in MeOH. All samples were analyzed by SDS-12% PAGE and immunoblotting with antibody 3F4.

RESULTS

β -CD clears 22L-infected neuroblastoma cells of infection.

Prion diseases and AD are protein conformational disorders, and in both cases, cholesterol levels are thought to affect the disease process. For AD, it was identified that β -CD altered the toxic effects of the A β peptide associated with AD (10). Consequently, as β -CD is known to modify cholesterol content, its effect on PrP^{Sc} levels in cell culture models was tested here. The scrapie-infected cell line employed for this study was N2a infected with 22L (39). A subclone possessing six times the PrP^{Sc} level was also employed (N2a22L20).

To examine the effect of β -CD on scrapie, β -CD was added to the N2a22L cell line at 100 μ M and 500 μ M at the time of passage and the cells were cultured in the presence of the compound for periods of 1 and 2 weeks. Cells were then lysed (Fig. 1A) and analyzed for PrP^{Sc} content by immunoblotting with 8H4 antibody. After 1 week of treatment, 500 μ M β -CD reduced PrP^{Sc} to 30% of the control level (Fig. 1A, lane 3, and B). Following 2 weeks of treatment, 100 μ M β -CD reduced PrP^{Sc} levels to 20% (Fig. 1A, lane 6, and B) and complete clearance was achieved with 500 μ M (Fig. 1A, lane 7, and B). A total of 500 μ M β -CD also cleared PrP^{Sc} after 2 weeks of treatment in N2a22L20, which possessed six times more PrP^{Sc} than N2a22L did (Fig. 1C, lane 4). Clearance was measured as the disappearance of the proteinase-resistant core of PrP, amino acids 27 to 30, relative to the amount in the untreated control, which was taken as 100%, and clearance was comparable to that of the known antiprion compound Congo red (Fig. 1A, lane 5, and C, lane 3).

The concentrations of β -CD used in this study were nontoxic as demonstrated by the MTT [3-(4,5-dimethylthiazol-

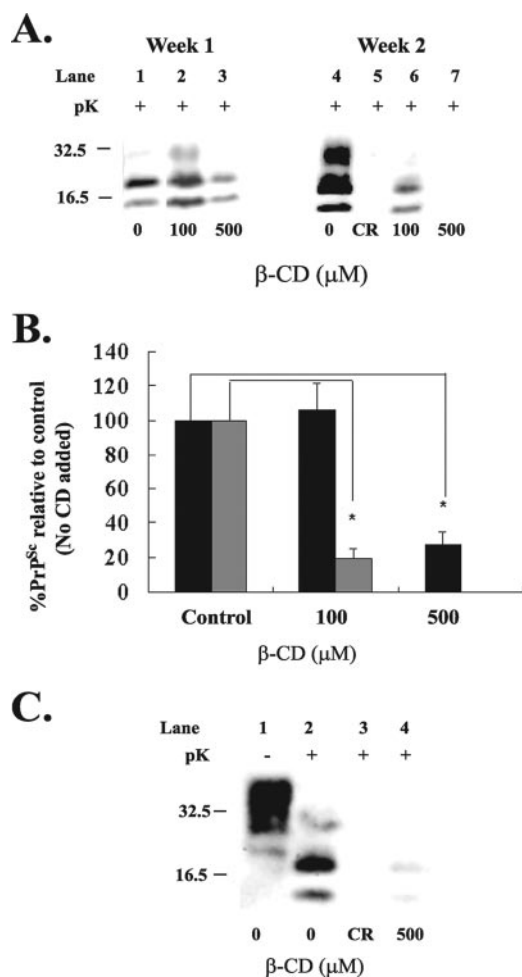


FIG. 1. β -CD reduces PrP^{Sc} to undetectable levels in the 22L-infected scrapie cell line. (A) β -CD at concentrations of 100 μ M and 500 μ M was added to N2a22L cells at the time of passage, and the cells were then passaged in the presence of the drug for 1 and 2 weeks. (B) Representative bands shown in panel A were quantified by densitometry and expressed as a percentage of the control (untreated). Black bars represent week 1, and gray bars represent week 2. *, $P < 0.005$ (Student's t test). Error bars indicate standard deviations. (C) β -CD at a concentration of 500 μ M was added to N2a22L20 cells at the time of passage, and the cells were passaged in the presence of the drug for 2 weeks. (A and C) Cells were lysed, and the lysate was then analyzed for PrP^{Sc} by 12% SDS-PAGE and immunoblotting with 8H4 antibody. Cell lysates were left untreated (-pK) or digested with pK (+pK) for the detection of PrP^{Sc}. Where indicated, as a drug control for PrP^{Sc} clearance, Congo red (CR) was added to the cells at 5 μ g/ml for 2 weeks. Results are representative of three independent experiments. Molecular mass markers in kilodaltons are shown on the left of the panels.

2-yl)-2,5-diphenyl tetrazolium bromide] cell viability assay (data not shown). To control for the possibility that clearance may have been due to a stimulation of pK activity by β -CD, β -CD was added to scrapie-infected cell lysate before the addition of pK and PrP^{Sc} detection was carried out as normal. Results from this experiment revealed that β -CD had no effect on pK activity (data not shown).

CDs have higher antiprion activity than their linear counterparts. The effect of other members of the natural CD fam-

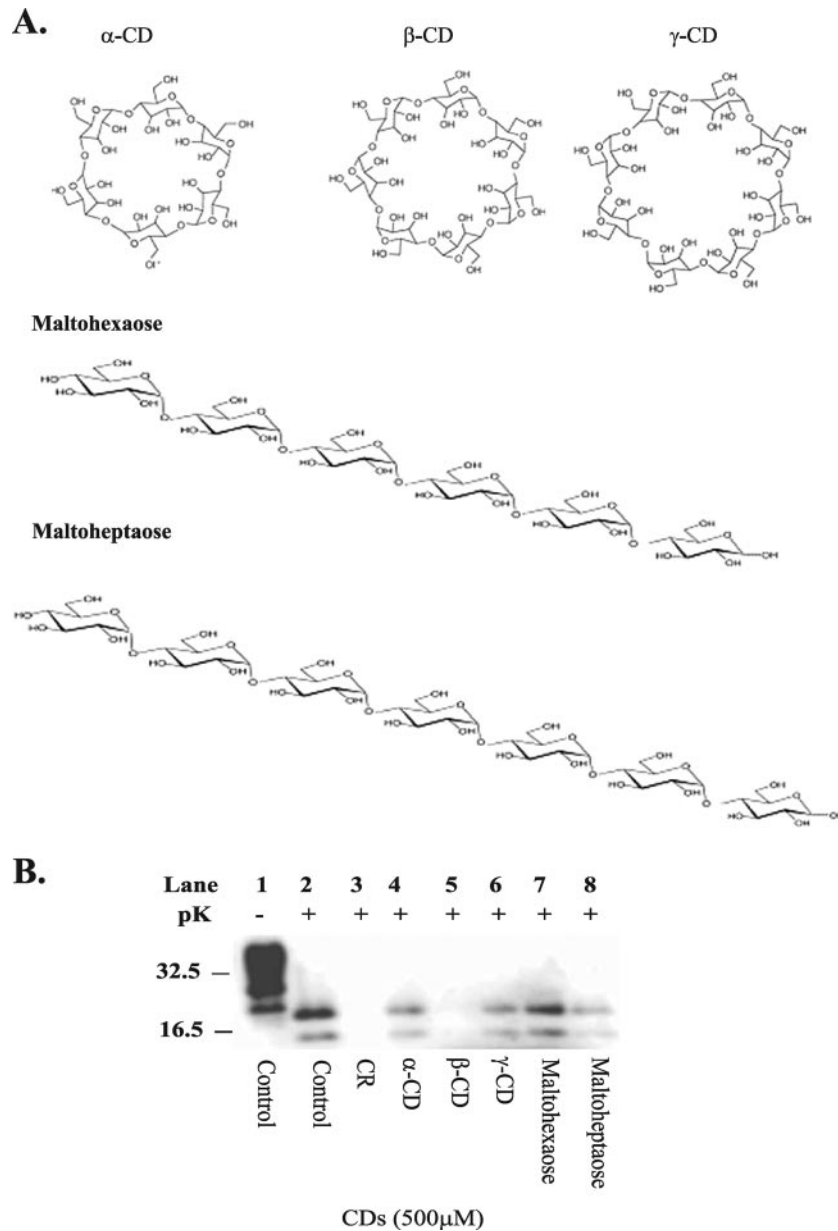


FIG. 2. Structure activity studies. (A) Structures of α -, β -, and γ -CD and their corresponding linear sugars. (B) α -, β -, and γ -CD and the linear sugars maltohexaose and maltoheptaose were added to the N2a22L cells at a concentration of 500 μ M at the time of passage, and the cells were then passaged in the presence of the drugs for 2 weeks. The cells were lysed, and the lysate was then analyzed for PrP^{Sc} by 12% SDS-PAGE and immunoblotting with 8H4 antibody. Cell lysates were left untreated ($-pK$) or were digested with pK ($+pK$) for the detection of PrP^{Sc}. Where indicated as a drug control for PrP^{Sc} clearance, Congo red (CR) was added to the cells at 5 μ g/ml for 2 weeks. Results are representative of three independent experiments. Molecular mass markers in kilodaltons are shown on the left of the panel.

ily, α -, β -, and γ -CDs (Fig. 2A), were examined for potential antiprion activity. The N2a22L cell line was treated with 500 μ M of α -, β -, and γ -CD for a period of 2 weeks, after which cells were analyzed for PrP^{Sc} levels by immunoblotting with 8H4 antibody (Fig. 2B, lanes 4 to 6). Of the CDs tested here, only β -CD (Fig. 2B, lane 5) completely cured infection, whereas α -CD and γ -CD demonstrated less antiprion activity (Fig. 2B, lanes 4 and 6). The structure-antiprion activity relationship of CDs was examined further by comparing their antiprion activity with that of their linear counterparts. The

linear counterparts for β -CD and α -CD were maltoheptaose (G7) and maltohexaose (G6), respectively (Fig. 2A). The corresponding linear sugar for γ -CD, maltooctaose, was not commercially available. Once again, the compounds were added at 500 μ M to the cells at the time of passage and the cells were cultured in this manner for a period of 2 weeks, after which the cells were lysed and analyzed for PrP^{Sc} levels. As G6 and G7 displayed less antiprion activity than α - and β -CD, respectively, the importance of the cyclic nature of the CDs for enhancing antiprion activity is demonstrated. G7 showed some ability to

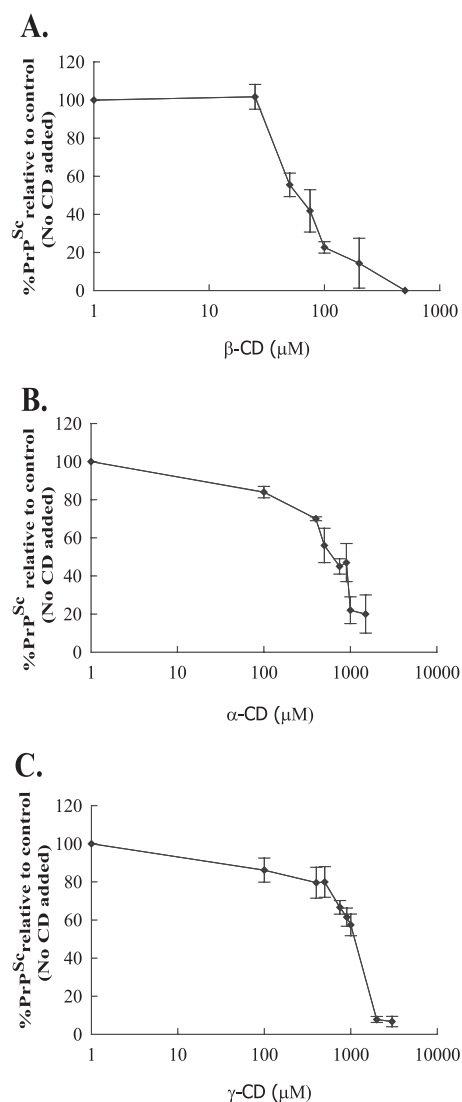


FIG. 3. IC_{50} determination for α -, β -, and γ -CD. Various concentrations of α -, β -, and γ -CD were added to the N2a22L cells at the time of passage, and the cells were then passaged in the presence of the compounds for 2 weeks. Cells were then lysed, and the lysate was then analyzed by 12% SDS-PAGE and immunoblotting with 8H4 antibody for the effect of β -CD (A), α -CD (B), and γ -CD (C) on PrP^{Sc} levels. Representative bands from immunoblots were quantified by densitometry and expressed as a percentage of the control (untreated). Results are representative of three independent experiments. Error bars indicate standard deviations.

reduce PrP^{Sc} levels, but it was not able to reduce infection like β -CD. CDs differ in size with α -CD possessing 6, β -CD possessing 7, and γ -CD possessing 8 glucopyranose units (Fig. 2A), leading to differences in the cavity sizes of the compounds (8 Å for γ -CD, 6 Å for β -CD, and 4.5 Å for α -CD) (66). These data indicate that the 6-Å size of β -CD is important for antiprion activity, and this indication is further emphasized by comparing the IC_{50} values of β -CD, α -CD, and γ -CD for clearing infection in N2a22L.

IC_{50} s were determined for β -, α -, and γ -CD by adding a range of concentrations of the compounds to the cells at the time of passage over a period of 2 weeks, after which the cells

were lysed and analyzed for PrP^{Sc}. β -CD with an IC_{50} of 75 μ M (Fig. 3A) was 10 times more efficient at clearing infection than was α -CD, which had an IC_{50} of 750 μ M (Fig. 3B). γ -CD had an IC_{50} of 1,150 μ M (Fig. 3C).

β -CD does not act through altering PrP^C levels or through an acidic compartment. The conversion of PrP^C to pathogenic PrP^{Sc} is believed to be a central event leading to the disease state in TSEs, and based on this belief, we reasoned that β -CD could be working by reducing substrate levels (PrP^C) for conversion. To test this hypothesis, noninfected N2a cells (N2a58 cell line) were treated with both β - and α -CD for a period of 2 weeks to investigate whether the compounds had an effect on PrP^C levels (Fig. 4A and B). Cells were then lysed and analyzed for PrP^C levels by immunoblotting with antibody 8H4. β -CD did not have a significant effect on PrP^C levels compared to the control (Fig. 4A, lane 2, and B). On the other hand, α -CD, which is 10 times less efficient than β -CD as an antiprion compound, lowered PrP^C levels by 30% (Fig. 4A, lane 3, and B).

The possibility that all known antiprion compounds are trafficked through endosomes or lysosomes has been suggested previously (20, 60, 62). Consequently, the effect of NH_4Cl on the antiprion activity of β -CD was investigated. NH_4Cl is a lysomotropic agent that alkalinizes endosomes and lysosomes without influencing PrP^{Sc} levels (60). N2a22L cells were treated as before, with 500 μ M β -CD for 2 weeks. The cells were also treated with NH_4Cl (30 mM) and a combination of NH_4Cl and β -CD for the 2-week period, and the cells were then lysed and analyzed for PrP^{Sc} content. On its own, NH_4Cl did not clear PrP^{Sc} (Fig. 4C, lane 3), and it was unable to block the antiprion activity of β -CD (Fig. 4C, compare lanes 4 and 5), indicating that β -CD may not be acting through lysosomes.

Effect of β -CD on PrP^{Sc}, PrP^C and GM_1 association with DRMs. CDs are powerful extractors of cholesterol, but of the CDs, β -CD had the highest capacity to sequester lipids (16, 40, 57). The accumulation of proteins in rafts is altered or abolished upon cholesterol depletion (46). This holds true for PrP^C, and altering cholesterol levels interferes with PrP^{Sc} production in cell culture (63). So, to investigate how β -CD acts against PrP^{Sc}, its effect on the floatation properties of PrP^C and PrP^{Sc} was determined. Like most GPI-anchored proteins, PrP^C and PrP^{Sc} are insoluble in cold Triton X-100 (9, 38, 63). This is due to their association with membrane domains or rafts that are enriched in cholesterol, sphingolipids, and glycosphingolipids; these domains are also referred to as DRMs (4, 8). The association of PrP^C with such domains is linked with the conversion to PrP^{Sc} (37, 38, 63). The solubility of DRMs is thought to be associated with the way their lipid content affects their rigidity and, consequently, the inability of cold Triton X-100 to penetrate and solubilize (9). Consequently, PrP^C and PrP^{Sc} float in buoyant fractions in density gradients and float in similar fractions to the raft resident, cell surface ganglioside GM_1 (38).

To examine for the effect of β -CD on floatation properties, floatation gradients were carried out with the N2a22L20 cell line. Cells were lysed after 3 days of growth, as for SGs, and the gradients were applied for 18 h at 37,000 rpm. Fractions were then processed for PrP^C, PrP^{Sc}, and GM_1 content as described in Materials and Methods and analyzed by immunoblotting with 8H4 antibody for PrP^C and PrP^{Sc} and by dot blot analysis with CTXB for GM_1 . PrP^C was separated from PrP^{Sc} by Sar-

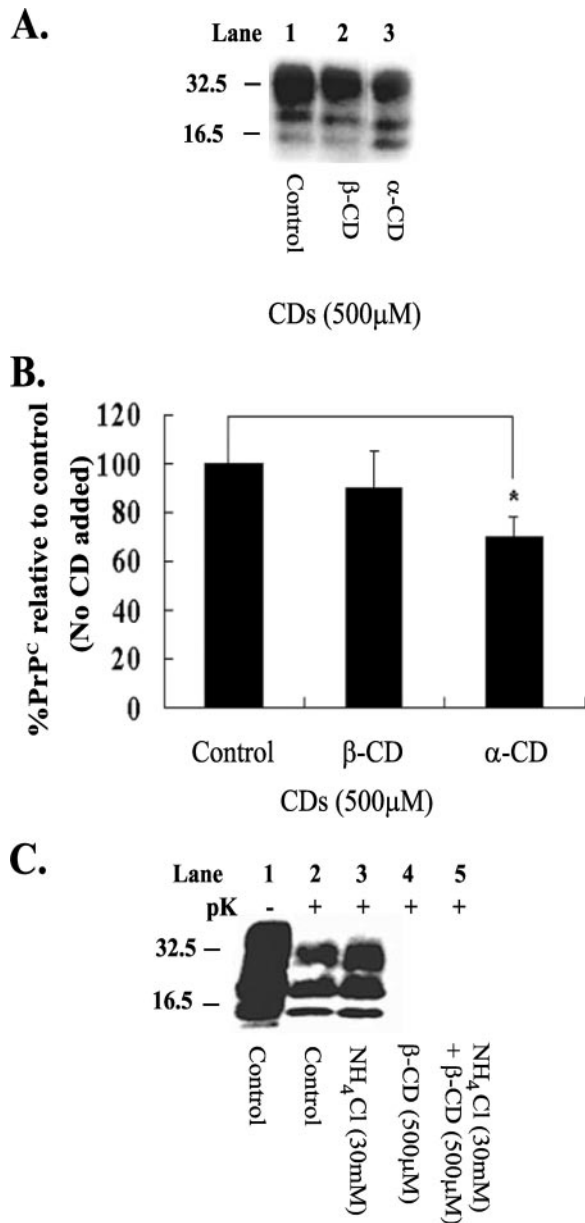


FIG. 4. β -CD does not act through altering PrP^C levels or through an acidic compartment. (A) β - and α -CD were added to noninfected N2a cells at 500 μ M each time the cells were passaged, and the cells were then lysed at 2 weeks. The lysate was prepared to 18 μ g of protein, as described in Materials and Methods, before SDS-PAGE and Western blotting, and PrP^C was detected with 8H4 antibody. (B) Representative bands shown in panel A were quantified by densitometry and expressed as a percentage of the control (untreated). Error bars indicate standard deviations. (C) The compounds NH₄Cl (30 mM) and β -CD (500 μ M) were added separately at the time of passage to N2a22L cells for 2 weeks (lanes 3 and 4, respectively). In addition, 30 mM NH₄Cl was added to the N2a22L cells at the time of passage and β -CD (500 μ M) was added an hour later. The cells were passaged in this manner for a period of 2 weeks (lane 5). The cells were lysed, and the lysate was then analyzed for PrP^{Sc} by 12% SDS-PAGE and immunoblotting with 8H4 antibody. Cell lysates were left untreated ($-pK$) or were digested with pK ($+pK$) for the detection of PrP^{Sc}. *, $P < 0.05$ (Student's t test). Results are representative of three independent experiments. Molecular mass markers in kilodaltons are shown on the left of the panels.

kosyl treatment. Naslavsky et al. (38) reported that this methodology does not affect PrP^C floatation properties. In the absence of β -CD, as expected, PrP^C, PrP^{Sc} and the raft marker, GM₁, floated in light fractions (Fig. 5). To examine for the effect of β -CD on the floatation characteristics, N2a22L20 cells were treated with β -CD (500 μ M) for 3 days, and floatation gradients were carried out as before. In keeping with the hypothesis that the removal of cholesterol increases the solubility of raft domains and consequently decreases the floatation of the associated GPI-anchored proteins, treatment with β -CD caused PrP^{Sc} and the GM₁ marker to float in heavier fractions. PrP^{Sc} peaked in fraction 4 prior to the addition of β -CD and between fractions 5 to 8 afterwards (Fig. 5B, compare upper panel, lane 4, and lower panel, lanes 5 to 8, and E). The GM₁ marker similarly shifted from fraction 5 to 6 under β -CD treatment (Fig. 5C, compare lane 5, upper panel, and lane 6, lower panel, and F). Unexpectedly, however, upon treatment with β -CD, PrP^C floated in lighter fractions (fractions 3 and 4) (Fig. 5A, lower panel, lanes 3 and 4, and D) to the control (fraction 5) (Fig. 5A, upper panel, lane 5, and D). It would appear that β -CD acts through modification of the lipid raft domain, but surprisingly, the two isoforms reacted differently, with PrP^C floating in more insoluble fractions than PrP^{Sc}. This separation of the localization of the isoforms could have the potential to restrict conversion. However, although the major portions of PrP^C and PrP^{Sc} have been separated, an overlap in gradients that could allow for a low level of conversion to occur does exist.

Reintroduction of cholesterol encapsulated in M β -CD does not reverse PrP^{Sc} clearance but facilitates it. Cholesterol depletion by M β -CD has previously been shown to inhibit A β formation in neurons, when it was combined with lovastatin (57) and when, on its own, it redistributed the *N*-methyl-D-aspartic acid receptor subunit, postsynaptic density protein, from Triton X-100 insoluble to soluble fractions (1). In both reports, the CD effect was reversed by reintroducing cholesterol in a cholesterol-M β -CD complex (chol-M β -CD). As β -CD altered the floatation properties of PrP^C and PrP^{Sc}, the ability of chol-M β -CD to reverse the antiprion activity of β -CD was investigated. In addition, to control for an effect of the M β -CD, which is the cholesterol vector, the effect of M β -CD on PrP^{Sc} levels was determined. N2a22L20 cells were cultured with both β -CD and M β -CD, each at 500 μ M, for 1 week. Cells were then lysed and analyzed for PrP^{Sc} content by immunoblotting with antibody 8H4 (Fig. 6A). As expected, after 1 week, β -CD lowered PrP^{Sc} levels to 50% of the control (Fig. 6A, lane 2, and B). M β -CD presented with marginally higher antiprion activity, lowering PrP^{Sc} levels to 40% (Fig. 6A, lane 3, and B). The IC₅₀ for M β -CD was determined to be 60 μ M after 2 weeks of treatment of N2a22L20 cells (data not shown). To investigate the ability of chol-M β -CD to interfere with the antiprion activity of M β -CD, N2a22L20 cells were grown for 1 week in the presence of M β -CD (500 μ M) in combination with chol-M β -CD (46 μ M, a concentration chosen from toxicity assays, but also used by others for this purpose [1]) and cells were analyzed for PrP^{Sc} (Fig. 6A, lane 4, and B). The presence of cholesterol enhanced, rather than inhibited, the clearance of PrP^{Sc}, as levels fell to 20%, indicating that the replenishment with cholesterol could not inhibit the antiprion activity of M β -CD.

To further demonstrate that chol-M β -CD could not inhibit

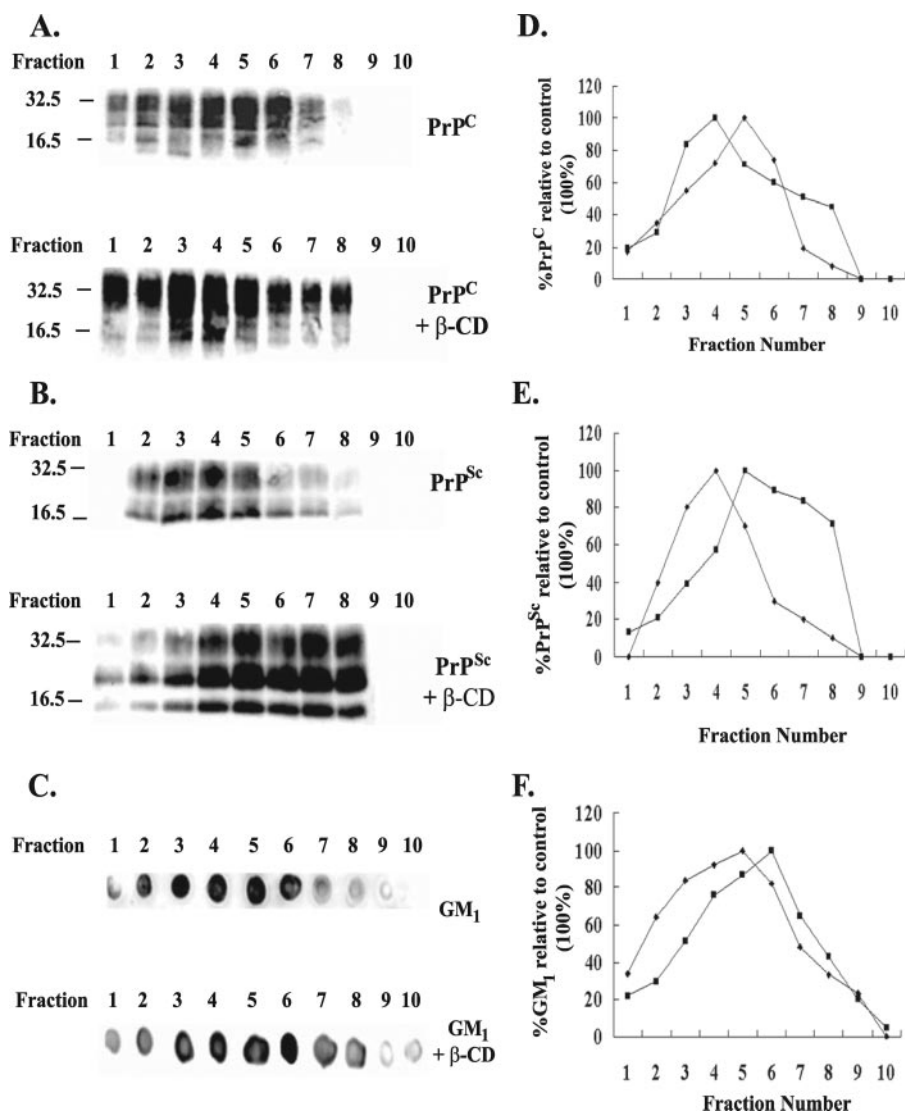


FIG. 5. Effect of β -CD on PrP^C, PrP^{Sc}, and GM₁ association with DRMs. The N2a22L20 cell line was either treated with 500 μ M β -CD at the time of passage (PrP^C, PrP^{Sc}, and GM₁, each with β -CD) or left untreated (PrP^C, PrP^{Sc}, and GM₁), and the cells were grown for 72 h. The cells were then lysed as for density gradients and adjusted to 40% sucrose, and a density gradient was applied and spun at 37,000 rpm for 18 h at 4°C. Fractions (10 \times 230 μ l) were collected from the top of the tube and processed for PrP^C, PrP^{Sc}, and GM₁ detection as described in Materials and Methods. Fraction 1 is the top of the gradient and fraction 10 is the bottom. Fractions for PrP^C (A) and PrP^{Sc} (B) detection were prepared as described in Materials and Methods and analyzed by 12% SDS-PAGE and immunoblotting with 8H4 antibody. Fractions for GM₁ detection (C) were blotted onto a nitrocellulose membrane and visualized using cholera toxin B conjugated to horseradish peroxidase. (D, E, and F) Representative bands from panels A (D) and B (E) and dots from panel C (F) were quantified by densitometry and were expressed as a percentage of the control (the fraction presenting with the highest signal [100%]). Results are representative of three independent experiments. Molecular mass markers in kilodaltons are shown on the left of the panels. Control results are plotted as closed diamonds, and β -CD results are plotted as closed squares.

the antiprion activity, N2a22L20 cells were treated with M β -CD or β -CD (each at 500 μ M) for 1 week. The compounds were then removed, and the cells were allowed to grow in their absence for 72 h or they were replaced by chol-M β -CD (46 μ M) for the 72-h period. Cells were then lysed and analyzed for PrP^{Sc} levels by immunoblotting with antibody 8H4 (Fig. 6C). Passage of the cells for 1 week in the presence of M β -CD and then in its absence for 3 days lowered PrP^{Sc} levels to 35% (Fig. 6C, lane 3), which was equivalent to the level of PrP^{Sc} reduction observed after 7 days of treatment (compare Fig. 6A, lane 3, with C, lane 3). This result indicates that PrP^{Sc}

clearance does not continue on drug removal. However, when the cells were grown for 3 days with chol-M β -CD (46 μ M), following the 1-week treatment with M β -CD, PrP^{Sc} clearance continued to 20% of the control (Fig. 6C, lane 4), reaffirming that the reintroduction of cholesterol for 3 days did not reverse the antiprion activity. The same pattern was seen when cells were grown in the presence of β -CD for 1 week, followed by 3 days in the absence or presence of chol-M β -CD (Fig. 6C, lanes 5 and 6, respectively).

CDs affect cholesterol levels but do not move PrP^C to soluble Triton X-100 domains. CDs are known for their ability to

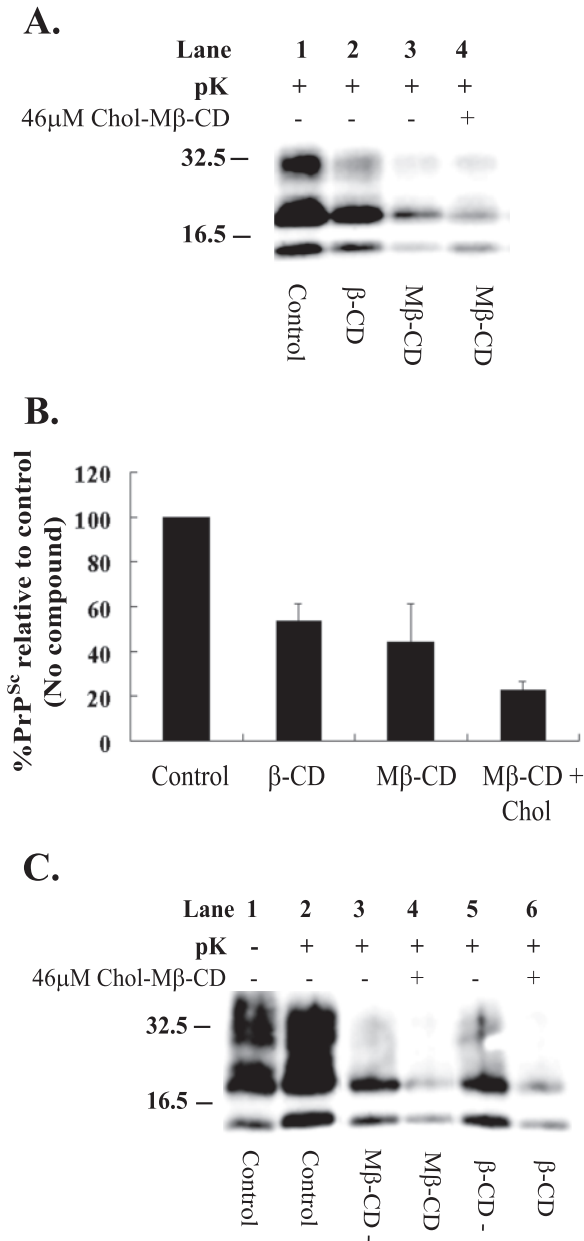


FIG. 6. Reintroduction of cholesterol encapsulated in M β -CD does not reverse PrP^{Sc} clearance, but rather facilitates it. (A) β -CD and M β -CD at a concentration of 500 μ M were added to N2a22L20 cells at the time of passage, and the cells were then passaged in the presence of the drugs for 1 week (lanes 2 and 3, respectively). N2a22L20 cells were also treated with a combination of M β -CD (500 μ M) and chol-M β -CD (46 μ M), and the cells were then passaged in their presence for 1 week (lane 4). (B) Representative bands shown in panel A were quantified by densitometry and were expressed as a percentage of the control (untreated). Error bars indicate standard deviations. (C) N2a22L20 cells were treated with M β -CD or β -CD at a concentration of 500 μ M at the time of passage, and the cells were passaged in the presence of the drugs for 1 week and then passaged in the absence of the drugs for 3 days (lanes 3 and 5) or passaged in the presence of chol-M β -CD (46 μ M) alone for 3 days (lanes 4 and 6). (A and C) Cells were lysed, and the lysate was then analyzed for PrP^{Sc} by 12% SDS-PAGE and immunoblotting with 8H4 antibody. Cell lysates were left untreated (-pK) or digested with pK (+pK) for the detection of PrP^{Sc}. Results are representative of three independent experiments. Molecular mass markers in kilodaltons are shown on the left of the panels.

complex with and sequester cholesterol. To examine this ability in N2a22L20 cells, cells were incubated for 1 h in the presence of increasing concentrations of either β -CD or M β -CD and their effects on total cholesterol levels were determined. The ability of chol-M β -CD to reintroduce cholesterol was also examined. After the hour of treatment with the CDs, the medium was replaced, and cells were incubated for a further hour in the presence of chol-M β -CD (Fig. 7A). Both CDs lowered cholesterol levels; however, M β -CD was more efficient. At 0.5, 1, and 10 mM, β -CD lowered levels to 97, 91, and 23% of those of the control, whereas M β -CD lowered levels to 79, 63, and 25% (Fig. 7A). Results are in line with those reported previously (1). Chol-M β -CD introduced 40% extra cholesterol into control cells and reintroduced cholesterol into CD-treated cells. Cells that had been treated with 0.5, 1, and 10 mM β -CD had levels increased by chol-M β -CD to 110, 128, and 81.8%, respectively, of those of the untreated control. For cells that had been treated with 0.5, 1, and 10 mM M β -CD, the levels were brought to 118, 100, and 53%, respectively, relative to levels of the untreated control (Fig. 7A).

β -CD and M β -CD are known to complex cholesterol and have been employed by others to modify cholesterol levels (16). In light of the ability of chol-M β -CD to introduce cholesterol into the cell, if the CDs were working solely by removing cholesterol, it was surprising that chol-M β -CD could not reverse the antiprion effect in the experiment for Fig. 6. In the past, lovastatin (63) and squalastatin (5), compounds working by reducing cholesterol levels, have moved PrP^C from cholesterol-rich lipid rafts, which are insoluble in cold Triton X-100, to domains that are soluble. To reconfirm our density gradient data that PrP^C was present in more insoluble domains after β -CD treatment, the effect of M β -CD on PrP^C location was examined (Fig. 7B and C). As for β -CD, M β -CD moved PrP^C to more insoluble domains (data not shown). This effect of CDs was reconfirmed by a simple solubility assay. Confluent N2a cells were grown in the presence and absence of 500 μ M M β -CD for 1 h, and the solubility of PrP^C in cold Triton X-100 was determined. The majority of PrP^C was insoluble, and treatment with M β -CD did not redistribute PrP^C to soluble domains (Fig. 7B, lanes 3 and 4, and C). Cells were also grown from the time of passage in the presence and absence of M β -CD for 72 h, and again PrP^C was maintained in the insoluble fraction (Fig. 7B, lanes 7 and 8, and C) and became slightly more insoluble.

β -CD binding to PrP^C. With CDs moving PrP^C to domains of lower density in sucrose density experiments, an effect that chol-M β -CD could not inhibit, it is possible that CDs (through cholesterol movement) could facilitate the generation of some altered density compartments to which PrP^C, and not PrP^{Sc}, is directed. Alternatively, PrP^C may be directed by these compounds to naturally existing high-lipid domains as a consequence of drug presence. Due to their hydrophobic cores, CDs possess the ability to encapsulate hydrophobic moieties. If CDs bound PrP, this binding could aid in the antiprion mechanism. Consequently, the ability of β -CD to bind PrP^C was determined.

β -CD was bound to epoxy-activated Sepharose 6B as described in Materials and Methods. N2a cells expressing 3F4-tagged MoPrP^C were lysed in cold lysis buffer (0.5% [wt/vol] sodium deoxycholate, 0.5% [vol/vol] Triton X-100, 150 mM

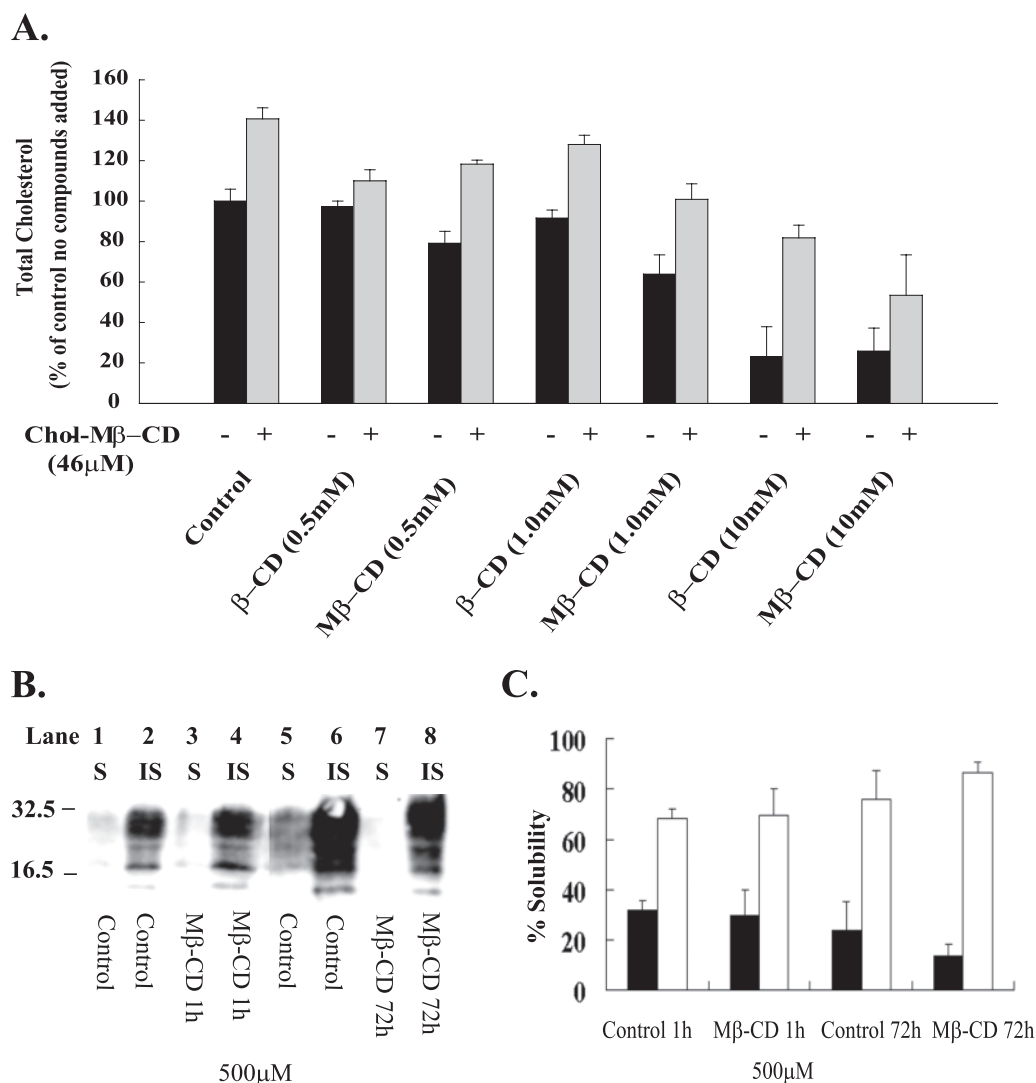


FIG. 7. Effect of CDs on cholesterol levels and Triton X-100 insolubility of PrP^C. (A) N2a22L20 cells were incubated with 0, 0.5, 1, and 10 mM β -CD or M β -CD for 1 h, the medium was removed, and cells were treated (+) or not treated (-) with chol-M β -CD (46 μ M) for 1 h. Cholesterol levels were then determined. Black bars represent cells treated with CDs as indicated, and gray bars represent cells treated with CDs, followed by treatment with chol-M β -CD. Error bars indicate standard deviations. (B) N2a cells were grown to confluence, and the medium was then replaced with fresh medium containing 500 μ M M β -CD for a period of 1 h or cells were left untreated (lanes 1 to 4). N2a cells were also treated with 500 μ M M β -CD at the time of passage for a period of 72 h or cells were left untreated (lanes 5 to 8). The cells were then lysed, the lysate was prepared for the TX-100 solubility assay, and then soluble (S) and insoluble (IS) fractions were separated. Samples were then analyzed for PrP^C by 12% SDS-PAGE and immunoblotting with 8H4 antibody. (C) Representative bands shown in panel B were quantified by densitometry and were expressed as percent soluble and percent insoluble. Black bars represent the percentage of soluble PrP^C, and white bars represent the percentage of insoluble PrP^C. Results are representative of at least three independent experiments. Molecular mass markers in kilodaltons are shown on the left of the panels.

NaCl and 50 mM Tris, pH 7.5). The lysate was then added to gel containing β -CD that had been equilibrated to pH 7.5 with 0.5 M HEPES buffer, and the binding process was carried out as described in Materials and Methods. Samples were analyzed by 12% SDS-PAGE and immunoblotting with 3F4 antibody. When binding was carried out with β -CD incorporated into the gel, approximately 50% of the 3F4-tagged PrP^C bound (Fig. 8A, lane 2, and B) and no PrP^C was eluted off in subsequent washes with 0.5 M HEPES buffer, pH 7.5, alone (data not shown). Depending on the strength of binding, it is sometimes possible to elute a bound protein with the complexing agent

(β -CD) in solution. However, neither 500 μ M β -CD nor 1 mM β -CD in solution could elute PrP^C from the gel (Fig. 8A, lanes 3 and 4, and B). Some protein-to-ligand complexes require more stringent methods for elution; therefore, the gel with PrP^C bound was boiled in loading buffer, and 12% of the eluate was analyzed for PrP^C content. A 100% elution of bound PrP^C occurred under these conditions (Fig. 8A, lane 5, and B). On the other hand, when binding was carried out without β -CD incorporated into gel, PrP^C did not bind (data not shown).

CDs inhibit an in vitro conversion of PrP. As CDs have been reported to modify protein conformation (50) and as PrP^C

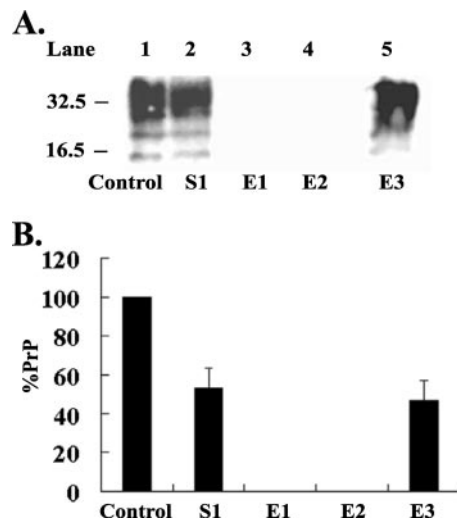


FIG. 8. PrP^C binding to Sepharose gel containing β -CD. (A) Eight hundred micrograms of lysate from N2a cells expressing 3F4-tagged MoPrP^C was loaded onto epoxy-activated Sepharose 6B gels prepared with β -CD. The gels were processed as described previously. The following were analyzed for PrP^C: 2.25% (18 μ g) of cell lysate (control), equivalent level (2.25%) of protein in unbound material (S1), 100% β -CD eluents (E1 and E2), and 12% of final eluent (E3). PrP^C in these fractions was analyzed by 12% SDS-PAGE and immunoblotting with 3F4 antibody. (B) Representative bands shown in panel A were quantified by densitometry, and PrP levels were adjusted to the level that would be present if 100% samples were loaded. The percentage of PrP was then plotted relative to the control (the level of PrP^C prior to binding to the gel). Results are representative of three independent experiments. Molecular mass markers in kilodaltons are shown on the left of the panels. Error bars indicate standard deviations.

bound to β -CD, the ability of β -CD to interfere with an in vitro conversion assay, based on the abnormal insolubility property of PrP^{Sc}, was determined. Cell lysate from N2a22L20 cells and N2a cells expressing 3F4-tagged PrP^C were mixed and incubated for up to 72 h as described above, and the conversion of 3F4-tagged PrP^C to insoluble 3F4-tagged PrP^C was determined. After 72 h, PrP^C was converted into insoluble 3F4-tagged PrP (Fig. 9A, lane 4, and B, lane 4) (this was not seen in the absence of N2a22L20 lysate [data not shown]). When the assay was carried out in the presence of Congo red (which is known to interfere with in vitro conversion) and β -CD (500 μ M), the in vitro conversion was inhibited (Fig. 9A, lane 6, and B, lane 6, respectively).

DISCUSSION

CDs, a family of cyclic maltooligosaccharides, have been used for decades as complexing agents for humans. The glucose units of these compounds are linked by α -1-4 bonds, and due to the structural orientation of the glucopyranose units, a hydrophobic cavity, which can facilitate the encapsulation of hydrophobic moieties, is generated (58, 61). This report demonstrates that CDs also possess antiprion activity, which appeared to be due to their ability to sequester compounds.

The modulation of cholesterol levels with M β -CD on its own has been shown to alter the susceptibility of cells to A β toxicity in PC12 cells (3) and, when M β -CD is combined with lova-

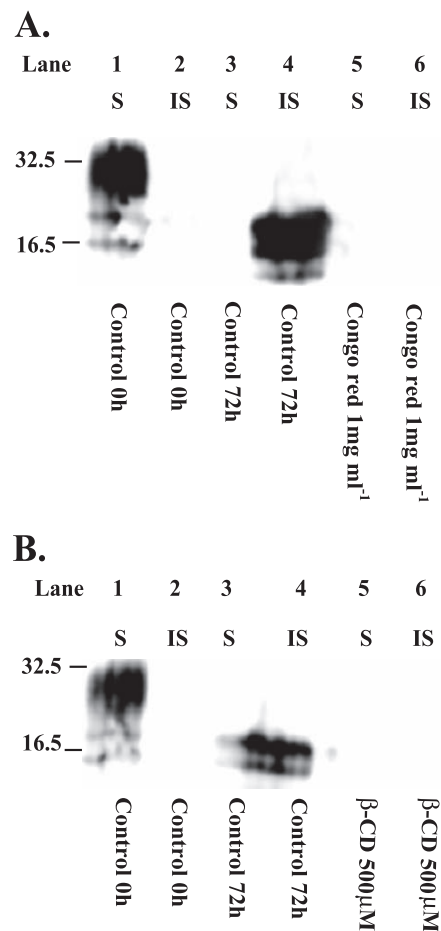


FIG. 9. β -CD blocks the conversion of PrP^C to PrP^{Sc}. (A and B) Samples were incubated as described in Materials and Methods for 0 h (lanes 1 and 2) or 72 h at 37°C (lanes 3 and 4). Samples were also incubated in the presence of Congo red (1 mg/ml) for 72 h (panel A, lanes 5 and 6) or β -CD (500 μ M) (panel B, lanes 5 and 6). After incubation, soluble (S) and insoluble (IS) PrP fractions were separated and levels were then analyzed by 12% SDS-PAGE and immunoblotting with 3F4 antibody.

statin, to modify A β production in hippocampal cells (57). In this report, the capacity of β -CD to clear PrP^{Sc} from infected cell culture is demonstrated. The antiprion activity of β -CD was time dependent, with PrP^{Sc} clearance in N2a22L cells occurring in 2 weeks with 500 μ M β -CD. Structural studies revealed the cyclic nature of the CDs to be important to the antiprion activity, and this is reflected in the IC₅₀ for β -CD (75 μ M), which was 10 times lower than that for α -CD (750 μ M). The linear form of β -CD also reduced PrP^{Sc} levels, but as it could not clear PrP^{Sc}, we infer that cyclicization of the molecule is significant for activity.

The antiprion action of β -CD did not appear to relate to a reduction in the substrate PrP^C within the cell for conversion. However, M β -CD (which is also reported here, for the first time, to have antiprion activity) has recently been reported to facilitate the shedding of PrP^C from neuronal cells (42). This shedding of PrP^C could have the potential to prevent conversion by reducing the access of PrP^{Sc} to PrP^C. This possibility would be in line with the report stating that the release of PrP

from the plasma membrane by phosphatidylinositol-specific phospholipase C (13) prevents PrP^{Sc} formation. Additionally, although it has been reported that known antiprion compounds pass through endosomes and lysosomes and act at these compartments to interfere with PrP^{Sc} production (60, 62), β -CD does not appear to act through these compartments, as the lysototropic agent NH₄Cl could not inhibit the action of β -CD.

It is more likely that the antiprion activity of CDs relates to their ability to sequester and move molecules. CDs are known for their ability to remove cholesterol from cell membranes (40, 57), and as cholesterol levels affect PrP^{Sc} production (63), it is possible that their action may be through an effect on the lipid raft environment where both PrP^C and PrP^{Sc} reside. Rafts are composed of sphingolipids and cholesterol, and the latter is thought to act as a spacer between the sphingolipids, keeping the lipid raft domain together (55). These domains house a number of protein networks, including an array of GPI-anchored proteins, and the ability of the domains to concentrate proteins/molecules may facilitate protein-to-protein interactions, including the conversion of PrP^C to PrP^{Sc} (38, 63). Cholesterol extraction with M β -CD has been reported to dissociate proteins from rafts and to move proteins from Triton X-100 insoluble to soluble compartments (46). The modulation of the raft environment has also been shown by others to interfere with the PrP conversion. However, reports on this are conflicting. Lovastatin, an inhibitor of 3-hydroxy-3-methylglutaryl-coenzyme A reductase, which reduces cholesterol levels, reduced prion infection in cell culture (63). Likewise, squalenstatin, an inhibitor of cholesterol production, also cured prion-infected neurons (5). Both squalenstatin and lovastatin moved PrP^C from Triton X-100 insoluble fractions to soluble fractions on extraction at 4°C (5, 63). M β -CD did not do this. However, the depletion of sphingolipid with fumonisins B actually increased the formation of PrP^{Sc} in neuroblastoma cells and had no effect on the floatation properties of either PrP^C or PrP^{Sc} (37), suggesting that the modulation of cholesterol and sphingolipids have different effects on the conversion.

Of the natural CDs, β -CD presented with the highest antiprion activity. This result was not surprising, as the ability of CDs to extract cholesterol is highly dependent on CD size, with β -CD being more effective than either α - or γ -CD (33). However, in light of the report of fumonisins B, β -CD's antiprion action, rather than proprion action, was unexpected. CDs are powerful cholesterol effluxors, but whereas α -CD preferentially sequesters phospholipids, β -CD preferentially extracts shingomyelin (33). Of course β -CD and fumonisins B are most probably acting very differently; however, Naslavsky et al. (37) could not rule out whether the effect of fumonisins B on PrP^{Sc} related to the level of sphingolipids in rafts or the head group identity.

Unlike fumonisins B, β -CD affected the floatation properties of both PrP^C and PrP^{Sc}. Prior to treatment with β -CD, PrP^C, PrP^{Sc}, and the DRM marker GM₁ all floated in buoyant fractions (Fig. 5A to F), which is in agreement with the results of other reports (38). However, on β -CD treatment, PrP^C and PrP^{Sc} were resolved into separate fractions (Fig. 5). PrP^{Sc} moved to the less buoyant fractions along with the GM₁ marker. This result could be expected as an effect of cholesterol extraction on the location of GPI-anchored proteins, but

PrP^C moved to more buoyant fractions. This physical separation of PrP^C and PrP^{Sc} could be sufficient to hinder the conversion process. However, a slight overlap in gradients still existed after treatment, and this overlap could still have allowed some conversion. In keeping with these data, M β -CD, which had higher antiprion activity than that of β -CD and had the highest ability to remove cellular cholesterol, failed to move PrP^C to domains that were Triton X-100 soluble, but a higher percentage of PrP^C was located in domains that were Triton X-100 insoluble after treatment. On extraction of neuroblastoma cells at 37°C prior to density floatation gradients, Naslavsky et al. (38) identified that in certain cell lines and in scrapie brain homogenate, PrP^C and PrP^{Sc} could be resolved into different fractions, indicating that the two isoforms reside in rafts of different contents. This separation was seen only in certain clones, and the researchers proposed that the lipid content of cellular membranes differed between cell types. It is possible that β -CD affects the lipid content of the rafts hosting PrP^C differently than those hosting PrP^{Sc}, with PrP^C residing in domains of higher lipid content after β -CD treatment. Cholesterol can transfer between different lipid vesicles, and CDs have been reported, by binding cholesterol in water-soluble complexes, to donate cholesterol to cellular membranes (16, 26) and in these locations PrP^C would reside.

As CDs have the capacity to sequester (33) and move hydrophobic molecules, the possibility that CDs act via a combination of mechanisms cannot be ruled out. β -CD binds PrP^C in affinity chromatography, and although this binding would not be specific to PrP^C, as CDs can bind to a range of proteins (21), this binding may play a part in its action. CDs have also been reported to modify protein conformation (18). CDs interfere with insulin self-association (67) and the toxicity of A β (10). Qin et al. (50) demonstrated the ability of β -CD to inhibit the A β peptide 12-28 β -sheet conformational change. Interestingly, in keeping with our antiprion data that are specific for β -CD, only β -CD, and not α -CD or γ -CD, possessed the ability to inhibit the structural conversion of A β . It could be hypothesized that β -CD may interact with the hydrophobic central core of PrP; such an interaction could lead to the ability of β -CD to interfere with the abnormal conversion.

The successfulness of antiprion compounds is dependent on a number of factors, including their ability to pass the BBB. CDs have been used for a long time to increase the solubility of compounds and to aid drug delivery. They are biocompatible, show resistance to human enzymes, have low toxicity in humans, and do not elicit an immune response (18). One millimolar of β -CD has been reported to pass the BBB without affecting the tight junctions; a tight junction breakdown has only been observed to start at 5 mM β -CD (33). Consequently, β -CD's antiprion IC₅₀ of 75 μ M falls well below the toxic level reported for this compound. CDs are a new class of antiprion compounds, and their usage has the potential to open up a number of avenues in the search for novel therapeutics in TSEs.

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