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Oligomeric-Induced Activity by Thienyl Pyrimidine Compounds Traps Prion Infectivity

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Accumulation of PrP^{sc}, an abnormal form of cellular prion protein (PrP), in the brain of animals and humans leads to fatal neurodegenerative disorders known as prion diseases. Limited protease digestion of PrP^{sc} produces a truncated form called PrP(27–30) that retains prion infectivity and is the main marker of disease targeted in most diagnostic tests. In the search for new anti-prion molecules, drug-screening assays on prion-infected murine cells have been oriented toward decreasing levels of PrP(27–30). In contrast, we screened for drugs promoting multimers of PrP(27–30), illustrating a possible stabilization of mouse PrP^{sc} species, because recent studies aiming to characterize the conformational stability of various prion strains showed that stable recombinant amyloids produced more stable prion strain, leading to longest incubation time. We identified a family of thienyl pyrimidine derivatives that induce SDS-resistant dimers and trimers of PrP(27–30). Bioassays performed on mice brain homogenates treated with these compounds showed that these thienyl pyrimidine derivatives diminished prion infectivity *in vivo*. Oligomeric-induced activity by thienyl pyrimidine compounds is a promising approach not only to understanding the pathogenesis of prions but also for prion diagnostics. This approach could be extended to other neurodegenerative "prionopathies," such as Alzheimer's, Huntington, or Parkinson's diseases.

Introduction

The hallmark of prion diseases is the deposition in the brain of PrP^{Sc}, an abnormal β -sheet-rich form of the cellular prion protein (PrP^C) (Prusiner, 1982). According to the prion hypothesis, PrP^{Sc} can trigger the autocatalytic conversion of PrP^C into PrP^{Sc}, involving several PrP^{Sc} pre-amyloid intermediates, generated through a complex mechanism of oligomerization (Silveira et al., 2005; DeMarco et al., 2006; Stöhr et al., 2008). The replication

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cycle proceeds with the self-assembling of PrP^{Sc} oligomers into proto-fibrils, which in turn grow into amyloid fibrils.

These different conformational forms can be specific targets for therapeutic interventions. Targeting the PrP^C by inhibiting its synthesis using RNAi or increasing its clearance with anti-PrP antibodies is a valuable strategy for prevention of prion diseases currently being investigated through gene therapy approaches (Daude et al., 2003; Perrier et al., 2004; White et al., 2008). Another approach consists of targeting the pathological PrP^{Sc}. A fundamental biochemical property that allows differentiation of PrP^{Sc} from PrP^C is the partial resistance of PrP^{Sc} to proteinase K (PK) digestion, leading to a truncated PrP^{Sc} form called PrP(27-30), the main marker of the disease (Fig. 1.1). Many screening assays to identify new anti-prion drugs have focused on reducing the level of PrP(27-30) (Fig. 1.2). This reduction reflects decreased levels of PrP^{Sc} species in the cells, correlated in most cases with diminished infectivity. This strategy led to the identification of many potential anti-prion drugs, but minor benefits were observed in vivo (Ludewigs et al., 2007). Targeting amyloid fibrils was also explored using amyloid dyes (Adjou et al., 1999; Beringue et al., 2000), but in vivo effects were disappointing because of their low ability to cross the blood-brain barrier. Other studies showed that targeting fibrils could result in an unexpected increase in infectivity (Beringue et al., 2000; Rudyk et al., 2000), which was also observed using the protein misfolding cyclic am-

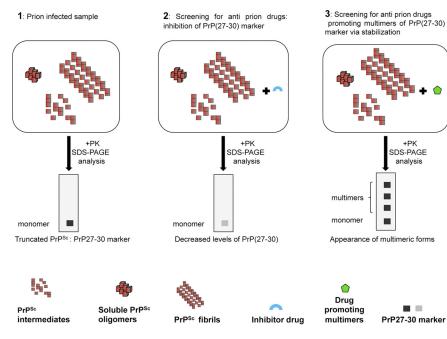


Figure 1. Schema describing classical and new drug-screening assays in prion-infected cells. **1**, The infectious agent is present under various forms in prion-infected cells (pre-amyloid intermediates, soluble oligomers of PrP ^{SC}, proto-fibrils, and amyloid fibrils of PrP ^{SC}). After a proteinase K digestion step and under SDS and reducing conditions, small and large aggregates of PrP ^{SC} undergo fragmentation, leading to a truncated PrP(27–30) marker. **2**, Classical drug-screening assays aim to search for inhibitors of prions exhibiting a decrease in the levels of PrP(27–30) marker. **3**, In our new drug-screening approach, we aimed at identifying drugs able to promote multimer formation through stabilization or cross-linking of PrP ^{SC} subunits. Because of stabilization with drugs, multimers of PrP(27–30) more resistant to denaturation can be detected.

plification (PMCA) technique, when large amyloid aggregates were broken up into smaller ones (Saborio et al., 2001; Wang et al., 2010). Altogether, these works led to the emerging concept that amyloid fibrils, rather than being a pathological entity, might be the result of a protective process to sequester more dangerous soluble oligomers (Caughey and Lansbury, 2003; Silveira et al., 2005; Simoneau et al., 2007).

Regarding this concept, we hypothesize from a therapeutic point of view that it might be better to block or trap amyloid structures rather than to solubilize or break them. Given that, after PK treatment and in SDS and reducing conditions, a complete fragmentation of PrP^{Sc} amyloid fibrils and aggregates occurs, leading to the PrP(27-30) marker, we searched for small molecules interacting with PrP^{Sc}-promoting PrP(27-30) oligomers, which would suggest a stabilization or a cross-linking step for those structures (Fig. 1.3). To accomplish this search, we developed a rational drug design approach to identify small molecules interacting with PrP, combined with a drug-screening assay using prion-infected cells. The results pinpointed a family of thienyl pyrimidine derivatives that induce SDS-resistant dimers and trimers of PrP(27-30). Bioassay experiments showed that these thienyl pyrimidine derivatives diminished prion infectivity in vivo.

Materials and Methods

Ethics statement. The Comité Régional d'Ethique de Montpellier approved the project (number CE-LR-0702).

Reagents and antibodies. Compounds selected *in silico* were purchased from Maybridge, Key Organics Limited, and Sigma. Stock solutions were prepared at 5 mM, and drugs were solubilized in DMSO (the solvent control) according to supplier recommendations.

P30 is 4-(5-bromo-2-thienyl)-2-(methylthio)pyrimidine; A6 is 2-(2'-thienyl)-5-(4"-(2"-aminopyrimidyl)thiophene); A12 is 4-(5-bromo-2- thie-

nyl)-2-pyrimidinamine; A13 is 4-(5-chloro-2thienyl)-2-pyrimidinamine; A14 is 4-(5-(thien-2-yl)-thien-2-yl)-2-

mercaptopyrimidine; A16 is 2-(methylthio)-4-(3-thienyl)pyrimidine; A17 is 5-(2-(sulfanyl-4pyrimidyl)-2-thiophenecarbonitrile; and A18 is 4-(methylthio)-6-(2-thienyl)-1,3,5-triazin-2amine. Pefabloc and PK were purchased from Roche Diagnostics. Ketamine was obtained from Mérial and xylazine from Bayer Health-Care. The protein assay kit based on bicinchoninic acid (BCA) was purchased from Pierce (Thermo Fisher Scientific). All of the anti-PrP antibodies directed against scrapie associated fibrils (SAF) used in this study were kindly provided by Dr. Jacques Grassi (Commissariat à l'Énergie Atomique et aux Énergies Alternatives, Saclay, France). Secondary antibodies were from Jackson ImmunoResearch. For immunohistology, anti-glial fibrillary acidic protein (GFAP) antibodies were purchased from AbCys, and the secondary antibody was included in the Strept ABC Complex kit (AbCys). All other chemicals were from Sigma.

In silico *screening*. The structural coordinates of human PP^{C} (Hu PP^{C}) (1QLZ, Protein Data Bank) were used for molecular modeling studies of the PrP and the search for a putative binding site conserved in both PrP^{C} and PrP^{Sc} isoforms. The *in silico* screening was performed using the Bioinfo chemical library (Kellenberger et al., 2007) and the Gold 2.0 Software (Jones and Thornton, 1997; Verdonk

et al., 2003). Candidate molecules were visualized at the conserved binding site using the Insight II Accelrys software with the Silicon Graphics SGI 02 R10000 workstation.

Cell culture. The mouse neuroblastoma cell line N2a was purchased from the American Type Culture Collection (ATCC CCL131). The N2a58 subclone, which overexpresses mouse PrP (MoPrP), was chronically infected with the mouse-adapted scrapie strain 22L (N2a58/22L cells), as described by Nishida et al. (2000). N2a58 cells and N2a58/22L cells were grown in Opti-MEM medium with L-glutamine supplemented with 10% fetal calf serum, 1% penicillin–streptomycin, and 300 μ g/ml geneticin. Cells were split twice per week, with a rotation period of 3–4 d. Cells were maintained at 37°C with 5% CO₂.

Cell screening assay and immunoblotting. N2a58/22L cells ($\sim 10^{6}/25$ cm² flasks) were incubated with various compounds at a final concentration of 20 µM (corresponding to 20 µl of drug at 5 mM, in 5 ml of medium for T25 plate) for 4 d as described previously for drug-screening test (Perrier et al., 2004). At confluence, cells were lysed in 400 µl of lysis buffer (0.5% NP-40, 0.5% deoxycholate, 10 mM Tris-HCl, pH 8, and 100 mM NaCl), and protein concentration was measured using the BCA assay. Samples of equal protein amounts and volumes were digested with 20 µg/ml PK at a ratio of 1:25 (protease/protein) at 37°C for 1 h. Digestion was stopped with 1 mM Pefabloc, and samples were centrifuged at 20,000 \times g at 4°C for 30 min. Pellets were dissolved in 20 μ l of lysis buffer and 20 μ l of 2× loading buffer (100 mM DTT, 3% SDS, 20% glycerol, 100 mM Tris-HCl, pH 6.8, and bromophenol blue) and then boiled for 3 min before loading on 12% SDS-PAGE precast Criterion gels (Bio-Rad). Western blotting was performed according to standard procedures. Mo-PrP was detected by using a mixture (SAF mix) of three anti-PrP monoclonal antibodies, SAF60, SAF69, and SAF70, as described previously (Perrier et al., 2004).

Dose–response curves. Approximately 10^{6} N2a58 or N2a58/22L cells, plated in 25 cm² flasks, were incubated with P30 or A6 at various concentrations, ranging from 5 to 40 μ M (corresponding to 5–40 μ l of drug at 5 mM, in 5 ml of medium for T25 plate) for 4 d. At confluence, cells were lysed, and the extracts were digested with PK and analyzed by West-

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ern blotting using the SAF mix for N2a58/22L samples as described previously (Nishida et al., 2000) and SAF32 antibodies for N2a58 samples (Perrier et al., 2004) as classically used.

Purification of prion protein. Full-length recombinant mouse PrP encompassing residues 23–230 (MoPrP23–230) was expressed and purified as described previously (Bocharova et al., 2005). The purified recombinant MoPrP23–230 was confirmed by SDS-PAGE and electrospray mass spectrometry to be a single species with an intact disulfide bond and correct molecular weight. After purification, MoPrP23–230 was stored in lyophilized form at –20°C until use.

Binding interaction studies by surface plasmon resonance. Surface plasmon resonance experiments were performed using a BiaCore 3000 (GE Healthcare). Full-length MoPrP23-230 was covalently immobilized on a CM5 sensor chip (GE Healthcare BiaCore AB) by the amine coupling standard procedure. A 4000-5000 RU immobilization level was obtained. For binding measurements P30, A12, A15, and the positive control quinacrine (Touil et al., 2006) were injected at 40 µM in PBS containing 5% DMSO with a flow rate of 30 μ l/min in PrP and blank flow cells. Surfaces were then washed with pulses of 10 mM NaOH until complete regeneration. The value (collected at a rate of 2.5 Hz) obtained in the reference flow cell were subtracted from those obtained in the PrP flow cell. Buffer blanks were injected before each drug injection, and binding responses were corrected for DMSO bulk differences using calibration curves (eight DMSO solutions between 4.5 and 5.8% DMSO). For K_d determination, different concentrations of P30 (from 5 to 80 μ M) were injected in PBS, 5% DMSO over the reference and PrP flow cells at a flow rate of 30 μ l/min for 60 s. Responses from the injection of drug compounds were extracted 10 s before the end of the injection. These responses were further corrected for DMSO effects by using the calibration curves, and the final response values were used for plotting the dose-response curve and for K_d determination by steady-state analysis (BiaEvaluation 4.1).

Incubation of MoPrP23–230 fibrils with thienyl pyrimidine compound. Amyloid fibrils using full-length MoPrP23–230 were formed using the manual setup protocol as described previously by Breydo et al. (2008). Fibril formation was monitored by collecting aliquots to which 10 μ M thioflavine T were added as described previously by Breydo et al. (2008). The quality of freshly made fibrils was also checked by transmission electron microscopy as described previously (El Moustaine et al., 2008). Samples of soluble MoPrP23–230 or fibrils at a final concentration of 4.4 μ M were incubated with 5 or 50 μ M P30 in MES buffer (50 mM MES, pH 5.0, 1% DMSO) at 25°C for 2 h. Ten microliters of each sample were mixed with the appropriate loading buffer (according to the protocols described previously), boiled at 90°C for 10 min before loading onto 12% SDS-PAGE gels. Silver staining was used to reveal the proteins in the gel (Chevallet et al., 2006).

Conversion of full-length MoPrP23–230 into amyloid fibrils in the presence of thienyl pyrimidine compounds using a semi-automated assay. The conversion of MoPrP23–230 into amyloid fibrils in a semi-automated setup was performed in fourfold to ensure reproducibility, according to a protocol described previously (Breydo et al., 2008). The following reagents were mixed in conical plastic tubes: 2 M GdnHCl, 50 mM MES buffer, pH 6.0, 10 μ M thioflavine T, 10 mM thiourea, 30 μ g/ml PrP in 6 M GdnHCl, and 40 μ M P30 drug, in a final volume of 500 μ l. After thorough mixing, the reaction mixture was aliquoted into four wells of a 96-well plate that was sealed. Plates were incubated at 37°C with shaking at 900 rpm (shaking diameter, 1 mm), and fluorescence was measured every 5 or 10 min with excitation at 444 nm and emission at 485 nm.

Kinetics of PrP(27-30) dimer and trimer in the presence of P30 and A6. N2a58/22L cells were incubated with 20 μ M P30 or A6, or 20 μ l of DMSO as control for the solvent alone. After 3–4 d, confluent cells were divided into duplicate flasks, one used for Western blot analysis and the other for prolongation of the treatment with drugs for a total of 11 d. At the end of the treatment, cells were grown without drugs for 10 additional days. Then cells were lysed, and cell extracts were treated with PK and analyzed by Western blotting, as described previously (Nishida et al., 2000).

Conformational stability assay. N2a58/22L cells were incubated with 20 μ M P30 or A6, or 20 μ l of DMSO as control for the solvent alone for 4 d. At confluence, cells were lysed with 200 μ l of lysis buffer. Each sample

was normalized for equal protein amounts. Cellular extracts were incubated with 50 μ l of GdnHCl mix, at a final concentration ranging from 0 to 1.5 M, for 2 h at room temperature. Then samples were diluted by addition of 750 μ l of lysis buffer and digested with PK (8 μ g/ml) at 37°C for 1 h. Reactions were stopped by addition of 1 mM Pefabloc and subsequently centrifuged at 20,000 × g at 15°C. Pellets were dissolved in 20 μ l of lysis buffer and 20 μ l of 2× loading buffer. Western blotting was performed as described above.

Aggregation assays on cell lysates and brain homogenates. Fresh N2a58/ 22L cell extracts prepared in 400 μ l of lysis buffer were incubated with various concentrations of P30 (from 12.5 to 625 μ M, corresponding to 1 to 50 μ l of P30 at 5 mM) at 20°C for 1 h. Because cell lysates were incubated with the drug for only 1 h compared with 3 d of cellular screening assays, it was important to concentrate the reaction volume to promote interactions while keeping the same drugs/proteins ratio. Then samples were digested with PK and analyzed by Western blotting according to the protocol described above.

Brain tissues from terminally sick mice inoculated with the scrapie strain 22L were homogenized in 10% (w/v) PBS using microbead-containing tubes and a Ribolyser apparatus (Bio-Rad). Samples were shaken for 45 s, and then supernatants were passed through an insulin syringe to obtain a homogeneous suspension. Protein amounts were measured using the BCA assay. To show the oligomeric activity of P30 in brain homogenates, we incubated different quantities of 22L-infected brain extracts (50–75 μ l), diluted in PBS with 2% Sarkosyl, with 0.5–3 mM P30 (corresponding to 50–300 μ l of drug at 5 mM stock solution) or 200 μ l of DMSO as a solvent control, in a final volume of 0.5 ml for 1–3 h. Samples were then digested with 20 μ g/ml PK as described previously (Perrier et al., 2002). Fifty microliters of each sample were mixed with an equal volume of 2× loading buffer and boiled for 3 min. Thirty microliters were loaded onto 12% SDS-PAGE gels and analyzed by Western blotting.

Then we incubated 22L brain homogenate with 1.5 mM P30 or A6 during 1 h at room temperature, and samples were proteinase K digested and fractionated with loading buffer (containing SDS and DTT). Samples were loaded on non-denaturing gel (Criterion Precast Tris-HCl; Bio-Rad) and transferred on to PVDF membrane using a Tris-glycine buffer (Bio-Rad). Membrane was revealed with SAF mix antibodies as described previously.

Evaluation of proteinase K resistance. Twenty-five microliters of 22L-infected brain extract were incubated with 1.5 mM P30 or 75 μ l of DMSO in a final volume of 0.25 ml for 1 h at room temperature. Then PK was added to a final concentration ranging from 0.1 to 1.2 mg/ml for 1 h at 37°C. Reactions were stopped by addition of 1 mM Pefabloc. Fifty microliters of each sample were mixed with an equal volume of 2× loading buffer and boiled for 3 min. Thirty microliters were loaded onto 12% SDS-PAGE gels and analyzed by Western blotting as described previously.

Bioassay using N2a58/22L cells. After 38 d of treatment with 20 µM P30 or 20 µl of DMSO, confluent cells (100 mm plate) were washed twice with 5 ml PBS and scraped with 0.5 ml PBS. Cells were broken through four cycles of freezing and thawing and homogenized by repeated extrusion through a 26-gauge needle. Prion infectivity was determined by intracerebral inoculation of 15 μ l of cell homogenate (diluted 10³ with normal cell extract) in female Swiss mice (Janvier) using a stereotaxic frame (David Kopf Instruments) with coordinates as follows, using Bregma as a reference: lateral, 1.0 mm; anteroposterior, -1.0 mm; and ventral, -3.0 mm (Paxinos and Franklin, 1997). Three groups of five mice were housed in cages placed in a ventilated protective cabinet, but two mice died from intercurrent disease in DMSO and P30 groups (n =13). All experiments were performed in a biosafety level 3 laboratory, according to the local ethical committee's guidelines. Mice were scored positive for prion disease when three signs of neurologic dysfunction were observed and when progressive deterioration of the animal was apparent according to 16 diagnostic criteria, as described previously (Carlson et al., 1986; Scott et al., 1989). Once clinical signs were detected, animals were killed in extremis. Their brains were removed and either immediately frozen at -80°C or fixed in AntigenFix (Diapath) for immunohistochemical analysis.

Bioassay using 22L-infected brain homogenates. To determine the infectivity of P30-induced PrP^{Sc} oligomers in mouse brain homogenates, we incubated 50 μ l of 10% (w/v) 22L-infected brain homogenates in PBS, 2% Sarkosyl (300 μ l) with P30 at a final concentration of 1.5 mM (150 μ l), or 150 μ l of DMSO as a control, at 20°C for 2 h. Samples were then diluted 10³ times with normal brain homogenates before inoculation into animals. Prion infectivity was determined by intracerebral inoculation of 15 μ l of the P30-treated samples into female Swiss mice (n = 19), using a stereotaxic frame as described in the previous section. At the terminal stage of the disease, animals were killed, and their brains removed as described above.

Immunohistochemistry. Brain tissues were fixed in AntigenFix solution (Diapath) for 24 h. Then they were decontaminated for 30 min in a formic acid solution according to the protocol described by Andréoletti (2004) and stored in 100 mM phosphate buffer at pH 7.4 with 0.02% sodium azide. Samples were dehydrated in graded ethanol, cleared in cedar oil, and then embedded in paraffin. Frontal 6–10 μ m sections were cut using a microtome and mounted on Superfrost Plus slides (Microm). Sections were dewaxed and stained with hematoxylin and eosin. Immunolabeling with anti-GFAP antibodies was performed according to the instructions provided with the Strept ABC Complex kit. Labeling was visualized using 3–3'-diaminobenzidine chromogen solution (Sigma).

Software and statistical analysis. Survival curves were done using GraphPad Software. We used R software version 2.9.0. for statistical analysis. The nonparametric Wilcoxon's test with a probability of 0.05 defined as a significant difference was applied for comparison of mice injected with 22L-infected brain homogenates treated with P30 versus the group inoculated with 22L-infected brain homogenates treated with DMSO only.

Results

Selection of compounds that interact with PrP using a structure-based drug design approach

We developed a rational drug design approach to identify chemical molecules that could interact with PrP^{Sc} species. Crystal structure studies have revealed that, in the case of globular proteins, small molecules act directly through the plasticity of the interface or allosterically by inducing conformational changes at distant sites (Arkin and Wells, 2004); thus, we first determined by molecular modeling whether we could identify a structurally conserved region at the surfaces of both PrP^C and PrP^{Sc} that could serve as a putative binding site. According to structural studies on the scrapie PrP by Wille et al. (2002, 2009), theoretical and experimental evidence indicates that the B and C helices in PrP^C are also conserved in PrP^{Sc}. Because these segments are joined by a disulfide bridge (C179-C214) that is also retained in PrP^{Sc} and required for infectivity, we focused our studies around these two conserved structural helices in the vicinity of the disulfide bridge. Moreover, Wille and colleagues (Wille et al., 2002; Colby et al., 2009) showed that the α helices of PrP^{Sc} are located toward the outside of the oligomers, which would be expected to facilitate the docking of a small molecule.

Our molecular modeling of this region in HuPr^C (1QLZ, Protein Data Bank) revealed the existence of a putative binding site that encompasses residues Y162–C179 and C214–S222 as well as the disulfide bridge at the end of the tiny binding pocket (Fig. 2*A*). From the Bioinfo chemical library, we then selected a first group of 350,000 small aromatic and bicyclic molecules with a polar surface smaller than 100 Å that could cross the blood– brain barrier. We then screened *in silico* for ligands of these compounds using Gold 2.0 software with a 15 Å radius sphere located at the center of the disulfide bridge of HuPrP^C as the putative binding site. We retained only molecules with a binding score higher than 50 within a scale going up to 100. We visualized 37 molecules with the best scores at the putative binding site, and, among these, 32 were available for purchase.

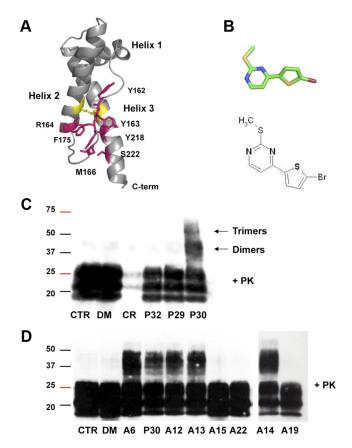


Figure 2. Identification of P30 by a structure-based drug design strategy, followed by cellular screening. **A**, Identification of a putative binding site on the HuPrP protein (1QLZ, Protein Data Bank) spanning along residues Y162–C179 (including the residue F175–C179 of α -helix 2) and C214 – S222 (α -helix 3) and the disulfide bridge at the end of the tiny binding pocket. The disulfide bridge (in yellow) and the side chains of amino acid residues Tyr163, Arg164, Met166, Phe175, and Tyr 218 (in purple) define the putative binding site for the small molecules selected in silico. The image was created by using PyMol software (www.pymol.org). B, Depiction of P30. C, Screening of small molecules selected in silico in prion-infected N2a58/22L cells. N2a58/22L cells were incubated with 20 μ M of the 32 compounds selected by virtual screening for 4 d. Then protein lysates were digested with 20 μ g/ml proteinase K (+PK), and PrP ^{Sc} accumulation was analyzed by immunoblotting (see Materials and Methods for more details) with the SAF mix (a mixture of SAF60, SAF69, and SAF70, three anti-PrP monoclonal antibodies). CTR, Negative control, untreated prion-infected N2a58/22L cells; DM, cells incubated with 20 μ l of DMSO alone; CR, cells incubated with 20 μ M Congo Red as a positive control for inhibition of PrP Sc accumulation. Among the 32 drugs tested (see Notes), only P30 triggered oligomerization of PrP(27-30) into dimers and trimers (see arrows). Molecular masses (15-75 kDa) are indicated on the left of the panel. **D**, Cellular screening of 23 derivatives of the P30 compound. Prion-infected N2a58/22L cells were incubated with 20 μM of the 23 compounds, selected for their structural analogies with P30, for 4 d and lysates processed as mentioned above. PrP ^{Sc} accumulation was analyzed by immunoblotting with the SAF mix. A6, A12, A13, A14, A15, A19, and A22 (see Materials and Methods for full names) were drug tested (see also Table 1).

Cellular screening of compounds selected in silico

We then tested whether the 32 compounds identified by our computational screening could block or promote PrP^{Sc} aggregates in prion-infected cells. Technically, control samples were submitted to a PK digestion step and to SDS denaturing and reducing conditions, including a boiling step at 100°C, leading to dissociation of the PrP^{Sc} aggregates in one main form of the PrP(27–30) marker visible on Western blot. A stabilization of PrP^{Sc} aggregates, for instance, by cross-linking PrP^{Sc} subunits with small molecules could lead under denaturing and reducing conditions to the detection of small oligomers of PrP(27–30). Our cellular test was to search for compounds with this ability. To

Compound	Structure	PrP	Compound	Structure	PrP
p • • • • • • •	H ₃ C	oligomerization	p - und	ŅH ₂	oligomerization
A1	N N S S N CH3	-	A13	N S CI	++
A2	H ₂ C ₁ S N N S H ₂ C ₁ S H ₂ C ₁ S H ₂ C ₁ S	-	A14	SH N N S S S	+ + +
A3	N N S S A	-	A15	NH2 N Br	-
A4	N N S CH3 N N S CH3 CH3 CH3 CH3 CH3 CH3	-	A16	N N N	+
A5	H ₃ C, S N N S S CH ₃ N S S S S N N CH ₃ N CH ₃	-	A17		+
A6	NH2 N N S	+++	A18ª	NH2 N N H ₂ C-S N S	+
A7	O'S D'	-	A19	N N S Br	-
A8	H ₅ C NH F F F	-	A20	N NH2 O S	-
A9	H ₃ C _{.S} N N S	-	A21		-
A10	N N N N N N N S	-	A22		-
A11	H ₂ N N N N N N N N N N N N N N N N N N N	-	A23	SH N S S Br	-
A12	NH2 NSP Br	++			

 a oligomeric activity detected with A18 at 40 μM

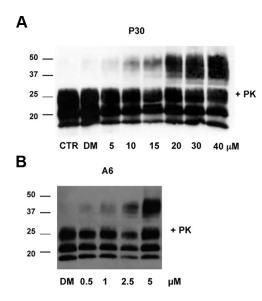


Figure 3. Oligomerization of PrP ^{5c} in prion-infected cells treated with P30 and A6 is dose dependent. *A*, Prion-infected N2a58/22L cells were incubated with 5–40 μ M P30 (corresponding to 5–40 μ l of drug at 5 mM), 40 μ l of DMSO (DM), or left untreated (CTR) for 4 d. Protein lysates were analyzed by immunoblotting after proteinase K digestion. *B*, Similarly, prion-infected N2a58/22L cells were incubated with 5–20 μ M A6 or 20 μ l of DMSO (DM) as a control, for 4 d. Protein lysates were then analyzed by immunoblotting after proteinase K digestion. Molecular masses (15–50 kDa) are indicated on the left of the panels. Immunoblots were revealed with the SAF mix.

this aim, N2a58/22L cells were incubated with 20 μ M of each of the 32 compounds for 4 d. At that time, cells reached confluence and were lysed. Negative controls were untreated N2a58/22L cells (controls) or incubated with 20 μ l DMSO (solvent control). Crude extracts were analyzed by immunoblotting after PK digestion. Only one compound, P30, increased PrP(27–30) oligomerization, especially dimer (~50 kDa) and trimer (~75 kDa) formation (Fig. 2*B*,*C*). None of the 32 compounds had an inhibitory effect on PrP^{Sc} accumulation, as observed when cells were incubated with 20 μ M Congo Red.

We then used the thienyl pyrimidine scaffold of P30 to search for P30 analogs displaying the same oligomeric activity. We screened the Bioinfo chemical library again and the ZINC database (http://zinc.docking.org) and found 23 P30 derivatives (Table 1). We then performed a second cell screening in N2a58/22L cells using the same conditions as before. Among the 23 P30 derivatives, six compounds (A6, A12, A13, A14, A16, and A17) induced PrP^{Sc} oligomerization as efficiently as P30 (Table 1, Fig. 2*D*). To further characterize the properties of this new family of compounds, we focused on two, P30 and A6. To confirm the reproducibility of their oligomeric activity and to see whether this property was dose dependent, N2a58/22L cells were incubated with various concentrations of P30 and A6, from 5 to 40 μ M, for 4 d. With both compounds, PrP^{Sc} oligomerization was reproducible and dose dependent (Fig. 3*A*,*B*).

The effective dose ED₅₀ (concentration at which 50% of PrP^{Sc} oligomerization was achieved) was estimated to be 17 μ M for P30 and ~2.5 μ M for A6, indicating a more potent effect of this compound. Finally, the toxicity of P30 and A6 (at 40 μ M) in N2a58/22L cells was assessed by evaluating cell viability with Trypan Blue. We observed no difference in cell viability between P30-and A6-treated cells and control cells incubated with DMSO (data not shown).

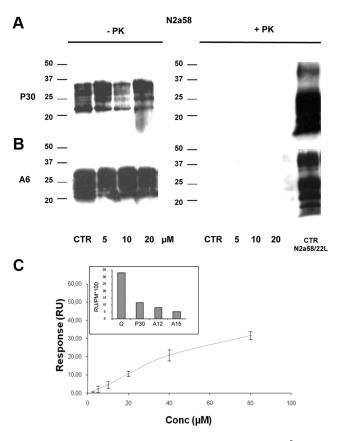


Figure 4. Thienyl pyrimidine compounds do not oligomerize the normal PrP^C isoform. *A*, N2a58 cells were incubated with 5–20 μ M P30, or not (CTR), for 4 d. Protein lysates were analyzed by immunoblotting before (–PK) or after (+PK) proteinase K digestion. *B*, N2a58 cells were incubated with 5–20 μ M A6 for 4 d and processed as in *A*. Immunoblots were probed with the monoclonal antibody SAF32. Molecular masses (20–50 kDa) are indicated on the left of the panels. *C*, Interaction of soluble MoPrP23-230 with thienyl pyrimidine compounds by surface plasmon resonance. In the inset, a histogram showing the interactions of P30, A12, A15, and quinacrine (Q) drugs at a concentration of 40 μ M, with the MoPrP23-230 immobilized on the sensor chip (4000–5000 RU). The dose–response curve illustrates the interaction between P30 drug (5–80 μ M) and the MoPrP23-230 immobilized on the sensor chip (4000–5000 RU).

Thienyl pyrimidine compounds exhibit oligomeric activity only in prion-infected cells

We then wanted to determine whether the oligomeric-induced activity of P30 and A6 was specific for PrP^C, PrP^{Sc}, or both isoforms. To this aim, N2a58 cells, which overexpress normal PrP, were incubated with various concentrations of P30 or A6 (from 5 to 20 μ M) for 4 d. Cell extracts nontreated by PK did not show oligomeric bands using SAF32 antibodies (Fig. 4A). After PK digestion of the P30/A6-treated N2a58 cell extracts, PK-resistant PrP^C forms were not detected (Fig. 4*B*). These experiments were also done using different antibodies (SAF mix), and the results were the same. Altogether, these data suggest that P30 and A6 do not allow formation of PrP^C oligomers resistant to SDS and reducing agent. We then asked whether this specificity for PrP^{Sc} could result from the fact that P30 and A6 do not bind to PrP^C, although the putative binding site is in a region that is conserved in both isoforms. To this aim, we used surface plasmon resonance (BiaCore) to assess whether P30, A12, and A15 could bind to immobilized MoPrP23-230 (Fig. 4C). Analysis of the results suggested that thienyl pyrimidine compounds could interact with PrP^C, albeit with less affinity than quinacrine (positive control). The calculated affinity for P30 was $K_{\rm d} = 147 \pm 50 \ \mu$ M. Unfortunately, we could not evaluate their binding to PrP Sc because of problems with aggregation of PrP^{Sc} to the BiaCore sensor chip.

Then we asked whether P30 could cross-link recombinant PrP fibrils to form SDS-resistant dimers/trimers. We thus incubated *in vitro* recombinant mouse PrP fibrils with 5 or 50 μ M P30 for 2 h and then evaluated dimer/trimer formation by SDS-PAGE (Fig. 5A). The absence of dimer/trimer formation suggested that P30 does not cross-link recombinant PrP fibrils. In addition, the analysis of the kinetics of recombinant mouse PrP conversion into amyloid fibrils in the presence of P30 showed that this compound did not have any effect on the kinetics of formation of recombinant amyloid fibrils (Fig. 5*B*).

PrP^{Sc} oligomers are stable after the suppression of treatment with P30 or A6

We then assessed whether prolonged incubation of N2a58/22L cells with thienyl pyrimidine compounds could increase PrP(27–30) dimer and trimer formation at the expense of monomers. N2a58/22L cells were incubated with 20 μ M P30, A6, or DMSO for 11 d. Assessment of oligomer formation at days 4, 7, and 11

showed no significant difference in the amount of dimers and trimers formed in cells incubated with A6 and P30; however, in protein extracts from cells treated with P30, less monomeric PrP(27-30) was accumulated and more trimers were detected compared with A6-treated cells (Fig. 6*A*).

After 11 d of treatment, cells were kept in culture without drugs for another 10 d (time points at days 14, 18, and 21). At day 14, PrP(27-30) dimers were still detected in extracts of P30-treated and especially A6-treated cells, in which trimers were also visible (Fig. 6A). However, by day 21, the balance tipped to the benefit of monomer formation despite the persistence of dimers. These results indicated that oligomers persist in the cells, even after the end of treatment, suggesting that these entities are quite stable. To evaluate the conformational stability possibly induced by thienyl pyrimidine compounds, prioninfected cells were first treated with 20 μ M A6 for 4 d. Then cellular lysates were denatured with guanidine hydrochloride (0.5-1.5 M) to assess conformational stability, followed by a PK digestion step. Up

to 1.25 M GdnHCl, PrP(27-30) dimers exhibited a conformational stability equivalent to those of monomers of PrP(27-30). Both DMSO- and A6-treated samples were equivalently denatured by exposure to 1.5 M GdnHCl (Fig. 6*B*).

P30 traps prion infectivity in N2a58/22L cells

We asked whether this oligomeric activity induced by thienyl pyrimidine drugs could diminish prion infectivity. Thus, N2a58/22L cells were supplemented with 20 μ M P30 or 20 μ l of DMSO for 38 d (Supattapone et al., 1999). Western blot analysis showed that oligomeric activity was still strong after 34 d of treatment (Fig. 7*A*). Then homogenates of P30- or DMSO-treated N2a58/22L cells were inoculated into the brains of Swiss mice. All inoculation experiments were performed using a stereotaxic frame to obtain accurate and reproducible injected volumes and sites of injections, because these variables can cause variations in incubation time (Kim et al., 1987). Mice inoculated with P30-treated cell

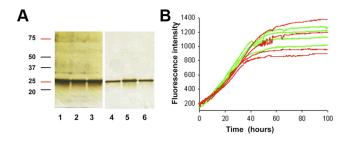


Figure 5. A, P30 does not cross-link full-length MoPrP23–230 or recombinant fibrils. Mo-PrP23–230 was solubilized in 50 mM MES buffer, pH 5.0, 1% DMSO and incubated during 2 h at 25°C, either without drug (lane 1) or 5 or 50 μ M P30 drug (lanes 2 and 3, respectively). Already formed MoPrP fibrils by a manual process were incubated during 2 h at 25°C either without drug (lane 4) or 5 or 50 μ M P30 drug (lanes 5 and 6, respectively). **B**, P30 does not modify the kinetics of conversion of MoPrP23–230 into amyloid fibrils. The kinetic of conversion of MoPrP23–230 into amyloid fibrils in a semi-automated setup was performed in presence of 40 μ M P30 drug (red curves) or DMSO alone (green curves) at 37°C.

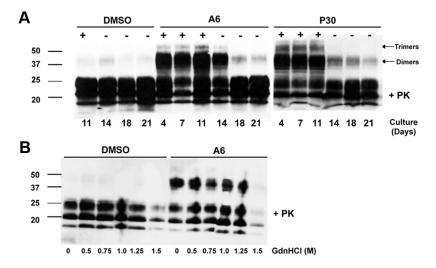


Figure 6. *A*, Oligomerization of PrP(27–30) by thienyl pyrimidine compounds persists after suppression of treatment. Prioninfected N2a58/22L cells were incubated (+) with 20 μ M P30 or A6 or 20 μ l of DMSO alone as control for the solvent for 11 d (time points at days 4, 7, and 11). At the end of the treatment, cells were grown without drugs (-) for 10 d (time points at days 14, 18, and 21). At the different time points, cells were lysed, total proteins quantified, and cell extracts digested with proteinase K (+ PK) before Western blotting. Blots were probed with the SAF mix. **B**, Thienyl pyrimidine compounds do not induce a conformational stability. Prion-infected N2a58/22L cells were incubated with 20 μ M A6 or 20 μ l of DMSO alone as control for the solvent for 4 d. At confluence, cells were lysed and treated with a final concentration of GdnHCl: 0, 0.5, 0.75, 1.0, 1.25, or 1.5 M. Samples were digested with proteinase K before Western blotting. Blots were probed with the SAF mix.

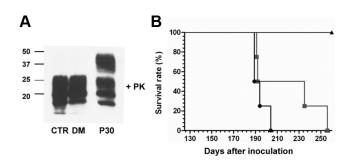


Figure 7. Decrease in prion infectivity in prion-infected N2a58/22L cells treated with P30. Prion-infected N2a58/22L cells were treated in duplicate with 20 μ M P30, 20 μ l of DMS0 (DM), or untreated (CTR) for 38 d. **A**, Samples were analyzed by immunoblotting, as described in Figure 2, to check for PrP(27–30) oligomerization before injection into brains of healthy Swiss mice. **B**, Kaplan–Meier survival plots of Swiss mice inoculated with cell homogenates prepared from prion-infected N2a58/22L cells treated with P30 (n = 4) (squares) or DMS0 (n = 4) (circles). Swiss mice were inoculated with PBS (triangles).

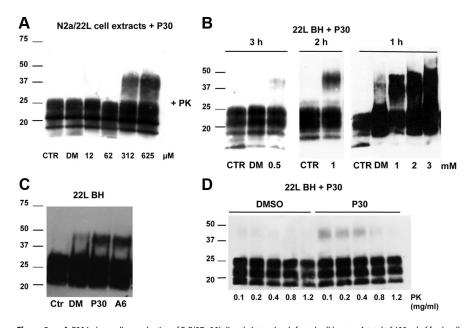


Figure 8. A, P30 induces oligomerization of PrP(27–30) directly into prion-infected cell lysates. A total of 400 μ l of fresh cell lysates from prion-infected N2a58/22L cells (each containing equivalent amounts of proteins) was incubated with various quantities of P30 (from 12.5 to 625 µm, corresponding to 1 to 50 µl of P30 at 5 mm) or 50 µl of DMS0 (DM) at room temperature for 1 h. Then samples were digested by proteinase K before analysis by Western blotting and probed with the SAF mix. CTR, Negative control, untreated N2a58/22L cell lysate. B, P30 oligomeric activity is attributable to direct aggregation in prion-infected brain homogenates. The experimental parameters leading to oligomerization of PrP ^{Sc} in prion-infected brain homogenates were investigated: (i) 75 µl of 1% mouse brain homogenate infected with the 22L scrapie strain diluted in PBS with 2% Sarkosyl (22Linfected brain homogenate) was incubated with 0.5 mm P30 (50 μ l) on ice for 3 h; (2) 50 μ l of 22L-infected brain homogenate was incubated with 1 mm P30 at room temperature for 2 h; (3) 50 μ l of 22L-infected brain homogenate was incubated with 1–3 mm P30 (corresponding to 100–300 μ l of P30 at 5 mm) at room temperature for 1 h. At the end of the incubation time, samples were digested with proteinase K before analysis of PrP sc accumulation by Western blotting according to the protocol described in Figure 2. CTR, Negative control, untreated 22L-infected brain homogenate; DM, 22L-infected brain homogenate incubated with 200 μ L of DMSO alone. C, Native gel experiments did not show any P30-induced oligomers above dimers. Fifty microliters of 22L-infected brain homogenate were incubated with 1.5 mm P30 or A6 (150 μ l at 5 mm) at room temperature for 1 h. At the end of the incubation time, samples were digested with proteinase K for 1 h and reaction was stopped with Complete (Roche). Fifty microliters of the mixture were mixed with an equal volume of denaturating loading buffer for fractionation. Then samples were charged on a native gel (Criterion Precast Tris-HCI; Bio-Rad), and the migration was done using Tris-glycine buffer (Bio-Rad). Proteins were transferred onto PVDF membrane and revealed using SAF mix antibodies. CTR, Untreated 22L-infected brain homogenate; DM, 22L-infected brain homogenate incubated with 150 μ l of DMSO. *D*, PrP(27–30) oligomers induced by thienyl pyrimidine compounds are less resistant to increasing concentrations of proteinase K. Twenty-five microliters of 22L-infected brain extract were incubated with 1.5 mM P30 or 75 μ l of DMSO for 1 h at room temperature. Resistance of PrP(27–30) oligomers to proteinase K was evaluated using various concentrations of proteinase K: 0.1, 0.2, 0.4, 0.8, or 1.2 mg/ml. Samples were digested with proteinase K before Western blotting. Blots were probed with the SAF mix.

homogenates survived longer (245 \pm 14.8 d) than animals injected with DMSO-treated cell homogenates (199 \pm 7.1 d) (Fig. 7*B*). Statistical analyses were not possible because of the low number of animals per group; however, the experiment was repeated with brain homogenates and validated (see Fig. 9). This result shows that P30 drug reduces the infectious titer of prion-infected cells. These experiments were performed only with P30 because A6 exhibited poor solubility that could have been detrimental for injection into the brain.

Thienyl pyrimidine compounds trap PrP^{Sc} in cell lysates and brain homogenates infected with prions

To test whether the oligomeric activity of the thienyl pyrimidine compounds required the *in vivo* cell machinery or relied on direct interaction, fresh N2a58/22L cell lysates were incubated with quantities of P30 that were equivalent to those used to treat cells with $20-40 \ \mu\text{M}$ P30 for 4 d (see Fig. 3, dose–response curve). After a 1 h incubation, cellular lysates were digested with PK, and oligomeric activity was assessed by Western blotting. P30 directly

induced formation of PrP(27-30) dimers and trimers in cell lysates in a dosedependent and reproducible manner (Fig. 8A). This result suggests that the mechanism of action of P30 does not require the in vivo cell machinery but rather is a direct effect. In addition, if the mechanism of action is free from the cell machinery, we should also be able to observe PrP(27-30) oligomers in brain homogenates of mice infected with the scrapie strain 22L after treatment with the P30 drug. To this aim, several experimental parameters were tested (i.e., quantity of 22L-infected brain extracts, volume of P30, and incubation times) to finally determine that a short incubation time (1 h), small quantities of brain homogenates (50 μ g), and at least 1 mM P30 induced PrP(27-30) oligomers in 22L-infected brain homogenates (Fig. 8B).

To test whether non-denaturating conditions could allow us to see oligomers of higher molecular weight, we performed experiments on native gels. A first assay was done using native loading and migration buffers without SDS and DTT but samples stacked in the wells of the native gel (data not shown). A second assay was done using a denaturating loading buffer to fractionate proteins, and the migration was performed on native gel using a native migration buffer without SDS and DTT (Fig. 8C). We did not observe differences with SDS-PAGE gel in the migration pattern, because higher-molecular-weight oligomers did not appear. This last experiment suggests that the two compounds did not increase the total aggregated state of the PrP Sc.

Furthermore, we evaluated the resistance to PK by using various concentrations of PK on brain homogenates previously treated with P30 drug or

DMSO for 1 h at room temperature. A first experiment using a range of concentrations from 10 to 100 μ g/ml showed no differences in the levels of monomers and dimers of PrP(27–30) (data not shown). A second experiment was performed with higher concentrations of PK, from 0.1 to 1.2 mg/ml (Fig. 8 *D*). When we used an increasing concentration of PK, no difference was observed in the quantity of PrP(27–30) monomers between the samples pretreated with P30 and those treated with DMSO (Fig. 8 *D*, immunoblot). However, the signal of PrP(27–30) dimers disappeared in the presence of 1.2 mg/ml PK, indicating that these species are more sensitive to high concentrations of PK.

Survival time is increased in mice infected with prions treated with P30

Previously, we showed that the oligomeric activity induced by thienyl pyrimidine drugs could diminish the infectivity of prioninfected cells; thus, we checked whether the P30 drug could also diminish PrP^{Sc} infectivity in brain homogenates. To test this hypothesis, 22L-infected brain homogenates were incubated with

1.5 mM P30 or 150 µl of DMSO for 2 h and then injected into the brains of healthy Swiss mice using a stereotaxic frame. These experiments were performed only with P30 because of the poor solubility of A6. Immunoblot analysis of P30-treated inoculi was performed to check for oligomeric activity before intracerebral injection (Fig. 9A). Mice inoculated with 22Linfected brain homogenates treated with P30 lived significantly longer than those inoculated with 22L-infected homogenates treated with DMSO alone (Fig. 9B). The mean \pm SD survival time of mice injected with P30-treated inoculi was 167 \pm 2.3 d (n = 10) compared with 154 ± 1.3 d (n = 9) for control mice (DMSO-treated inoculum), which was statistically significant according to the nonparametric Wilcoxon's test (W = 7, p = 0.0019). Kaplan– Meier survival curves showed that half of the mice infected with P30-treated inoculi survived for up to 172 d. This result shows that oligomers induced by P30 reduces the infectious titer of prion-infected inoculi. Analysis of brain homogenates from mice inoculated with P30-treated homogenates did not show the presence of oligomeric bands as in the inoculum, suggesting that we did not create a new prion strain (Fig. 9C). Histopathological analysis of brain tissues at the terminal stage of disease revealed no difference in spongiosis and astrocytic gliosis in mice that were inoculated with homogenates treated with P30 or with DMSO (Fig. 9D).

Discussion

PrP^{Sc} is a key element in the pathogenesis of prion diseases and thus represents privileged targets for the development of therapeutic strategies to combat these fatal transmissible agents. Our objective was to target PrP^{Sc} species with small molecules with the idea of blocking PrP^{Sc} species to prevent their recycling in the pathological aggregation pathway. Our combined approach of virtual and drug screenings showed that thienyl pyrimidine compounds trap the pathological isoform PrP^{Sc} (either small PrP^{Sc} entities or large aggregates present in cells or brain homogenates), which then become SDS-resistant PrP(27-30) dimers and trimers after PK digestion and denaturation. The resulting effect is a small decrease in prion infectivity. Additional experiments to gain insights into the mode of action of two of these thienyl pyrimidine compounds (P30 and A6) showed the following: (1) their oligomer-inducing activity is observed only in the presence of infectious material and appears to be PrP^{Sc} specific; (2) the oligomer-inducing effect on cells treated with thienyl pyrimidine compounds remains stable even after termination of the treatment with P30 or A6; (3) brain homogenates treated with these compounds also exhibit dimers of PrP27-30; and (4) this oligomeric-induced activity is free from the in vivo cell machinery, suggesting a direct effect.

For augmentation of PrP(27–30) signal and diminution of prion infectivity, how should the apparent contradiction be reconciled? For two decades, the search for new anti-prions drugs

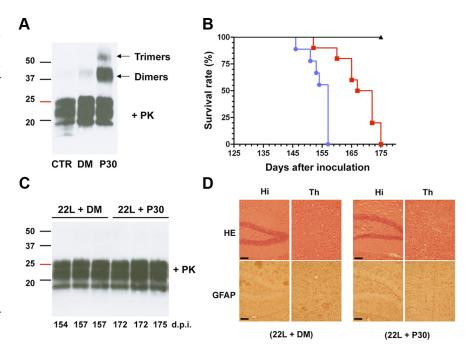


Figure 9. P30 increased survival time in mice injected with brain homogenates infected with the 22L strain. *A*, Fifty microliters of 1% brain homogenate infected with the 22L strain were incubated with 1.5 mM (volume, 150 μ J) of P30, 150 μ J of DMS0 (DM), or nothing (CTR) at room temperature for 2 h. This experiment was done in duplicate for analysis by both immunoblotting (to confirm P30-induced oligomerization of PrP^{Sc}; this panel) and injection into mice brains. *B*, Kaplan–Meier survival plots of Swiss mice (n = 19) inoculated with 22L-infected brain homogenates incubated with P30 (22L + P30) (squares) or DMS0 (22L + DM) (circles). Swiss mice were inoculated with PBS (triangles). *C*, Brain homogenates from mice inoculated with 22L-infected brain homogenate incubated with P30 (22L + P30) or DMS0 (22L + DM) were analyzed by immunoblotting after proteinase K digestion. The numbers below the panel indicate the day when the mice were killed (days post-inoculation, d.p.i.). *D*, Histological analysis of hippocampus (Hi) and thalamus (Th) tissue sections from mice inoculated with 22L-infected brain homogenates incubated with P30 (22L + P30) or DMS0 (22L + DM). Tissue sections were stained with hematoxylin and eosin (HE) and probed with anti-GFAP antibodies as a marker of astrocytic gliosis. Scale bars, 10 μ m.

using prion-infected cells or in vitro assays focused on decreasing levels of PrP(27–30), the main marker of the prion diseases (Fig. 1). In contrast, one can speculate that compounds favoring the appearance of PrP(27-30) species would increase infectivity and/or toxicity, which is probably the main reason that screening for "anti-prion" drugs promoting oligomers of the PrP(27-30) signal has not yet been explored. In one way, an increase in the level of the monomeric PrP(27-30) marker would strongly suggest an amplification of prion and infectivity. Indeed, Murayama et al. (2010) succeeded in amplifying the monomeric PrP(27-30) from bovine prions when they added sulfated dextran during the PMCA method used for diagnostic purposes (Murayama et al., 2010). In contrast, in the present study, we searched for drugs promoting the formation of small oligomers of PrP(27-30) because of a suggested stabilization of PrP Sc associated with the idea of trapping prion infectivity. Indeed, many recent studies aiming to characterize the conformational stability of various prion strains to elucidate the molecular aspects of PrP conformation showed a direct relationship between stability and incubation time. Colby et al. (2009) demonstrated that stable recombinant amyloids produced a more stable prion strain, leading to the longest incubation time, whereas more labile amyloids generated less stable strains and shorter incubation times (Colby et al., 2009). Guanidine hydrochloride experiments on cells preincubated with thienyl pyrimidine compounds have shown that these drugs do not ostensibly induce a conformational stability of the inoculum. They could interact and stabilize a small subpopulation of infectious PrP^{Sc} molecules, but this effect is not representative of the overall population, and the conformational stability thus cannot be observed (Colby et al., 2009). This hypothesis could explain why we obtained a minor decrease in infectivity. It is likely that some analogs of P30 that exhibit a more potent activity could completely block the infectivity of the inoculum, and we will focus our efforts in this direction.

Although we could determine that the thienyl pyrimidine compounds do not need the *in vivo* cell machinery to form PrP(27-30) dimers and trimers, their mechanism of action remains to be clarified. In addition, we do not know yet which PrP^{Sc} molecular species are targeted. We thus propose two possible hypotheses.

(1) Thienyl pyrimidine compounds could bind to preexisting PrP^{Sc} dimers and trimers. PrP^{Sc} dimers and trimers could be transitory and unstable species (DeMarco et al., 2006; Silveira et al., 2005), possibly referring to pre-amyloid states (Stöhr et al., 2008). Indeed, from electron crystallographic studies established using PrP^{Sc} purified from infected brain homogenates, two models of prion proto-fibril propagation are currently emerging: the β -helix model and the spiral model. Both models contain oligomeric discs composed of PrP^{Sc} trimers that can assemble into proto-fibrils (Wille et al., 2002; DeMarco and Daggett, 2004; Govaerts et al., 2004; DeMarco et al., 2006). The true existence of such small oligomeric species is a matter of much debate, but we cannot exclude that possibility. We hypothesize that, during binding of thienyl pyrimidine drugs, dimers and trimers of PrP^{Sc} are diverted from the pathological aggregation pathway and trapped into a thermodynamic landscape that is not competent for additional propagation of infection (El Moustaine et al., 2008). After the PK digestion assay, those putative P30-stabilized dimers and trimers of PrP^{Sc} become SDS-resistant (PrP27-30) dimers and trimers under denaturation conditions. In addition, dimers and trimers of PrP^{Sc} might represent a minority of the heterogeneous infectious PrP^{5c} species present in the inoculum. Thus, trapping these small oligomers with drugs can block only a fraction of the prion infectivity in the inoculum. This hypothesis correlates well with the results of our in vivo experiments in which we observed a partial reduction in prion infectivity.

(2) The thienyl pyrimidine compounds could cross-link PrP^{Sc} subunits belonging to amyloidogenic structures. To check this hypothesis, we performed *in vitro* experiments using recombinant mouse PrP fibrils incubated with thienyl pyrimidine compounds and showed that P30 does not cross-link amyloid fibrils or influence the kinetics of amyloid fibril formation (Fig. 5A, B). Despite this result, we cannot completely rule out this hypothesis because recombinant fibrils substantially differ from natural fibrils (Wille et al., 2009). Wille et al. have observed some repeated units along the natural fiber axis purified from brain-derived prions that were barely observed in recPrP fibrils, suggesting a key role in infectivity. Our compounds could conceivably target these specific structures.

Because of their specific effect on PrP^{sc}, the thienyl pyrimidine compounds have interesting potential applications in the field of detection assays and/or decontamination procedure. In addition, PrP(27–30) dimers or trimers observed with thienyl pyrimidine compounds could also represent a new marker of prion disease in biological fluids. In the therapeutic field, trapping the main marker of prion diseases, the PrP(27–30), into dimeric/trimeric forms represents a breakthrough in the identification of new anti-prion drugs. Furthermore, this approach opens new avenues for the development of new drug-screening assays. We believe that the combination of several anti-prion drugs that target different intermediates of the prion replication cycle would be a better way to prevent prion propagation and to possibly identify a cure for prion-related diseases. The strategy, which consists of stabilizing the main marker of prion diseases into small oligomers, could also be envisaged for use in the treatment of other proteinopathies that lead to neurodegenerative disorders, such as Alzheimer's and Huntington diseases.

Notes

For results of the first screening, see Table S1. Supplemental material for this article is available by e-mailing Dr. Perrier at veronique.perrier@univ-montp2.fr, or at http://mmdn.um2.fr start-ing December 2011. This material has not been peer reviewed.

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