# 医薬品 研究報告 調査報告書

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識別番号	-報告回数			報告日	第一報入手日 2005. 11. 24	<b>新医薬品</b> 該当	=	機構処理欄
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販売名(企業名)		赤十字アルブミン2 赤十字アルブミン2		研究報告の公表状況			イタリア	
○スクレイピーおよび乳腺炎に罹患した羊の乳腺におけるPrP <sup>Sc</sup> イタリアのサッサリ県で818頭の羊を調査した。7頭は脳、リンパ節、扁桃腺においてPrP <sup>Sc</sup> が検出され、臨床的に明らかなスクレイピーの症状を呈していた。4頭が乳腺炎とスクレイピーを併発していた。この4頭全ての乳腺においてPrP <sup>Sc</sup> が検出されたが、乳腺炎を併発していないスクレイピーを発症した同じ群(n=14)又は他の群(n=1)由来の羊、乳腺炎に罹患しているがスクレイピーへの感染は認められない羊(n=2)においてはPrP <sup>Sc</sup> は検出されなかった。乳腺の炎症病変部の解析では、PrP <sup>Sc</sup> のリンパ濾胞部位への集積が認められた。PrP <sup>Sc</sup> は、乳腺炎による病変部位中の主にCD68+マクロファージおよびFDCsと共局在化していた。  慢性的な炎症とスクレイピーの併発により、PrP <sup>Sc</sup> が想定外の組織まで拡大して蓄積する可能性が示された。乳房中のPrP <sup>Sc</sup> 濃度の中央値は、脾臓の0.1%、脳の0.05%と算出されたが、乳房のリンパ濾胞は確率的な分布を示しているため、局部的なPrP <sup>Sc</sup> 量には顕著なばらつきが認められた。 本研究ではMaedi-Visnaウィルス(MVV)の血清抗体陽性反応とリンパ濾胞乳腺炎の相関が示された。ヨーロッパの小型反芻動物のほとんどはMVVおよび関連レンチウィルスに感染している。ごく一般的なウィルス感染が原因となるプリオン病感染拡大の可能性が示唆された。MVVは、乳房上皮細胞やマクロファージ中に存在し、羊乳を介して子羊に伝蕾することが実験的に証明されている。PrPの乳房リンパ濾胞部位のCD68+細胞への蓄積は、乳腺炎の羊のキ乳中への大量のマクロファージの混入も併せ、プリオン感染と分泌器官の炎症の併発が分泌物のプリオン汚染を誘導し、群中におけるプリオンの水平感染の共同因子となり得るのか、という疑問を提起することとなった。								

報告企業の意見

スクレイピーおよび乳腺炎に罹患した羊の乳腺でPrPSc が検出
されたとの報告である。

これまでの疫学研究等
クレーピーを含む伝達

これまでの疫学研究等では、ヒトにおいて、血漿分画製剤を介してスクレーピーを含む伝達性海綿状脳症(TSE)が伝播するという証拠はない。また異常プリオンがアルブミン製剤の製造工程で効果的に除去されるとの報告もあるが、輸血によりvCJDに感染する可能性が示唆され

今後の対応

たことから、今後も情報の収集に努める。

(A)

# CORRESPONDENCE

Table 1 Summary of the fluorescence correlation spectroscopy measurements using 10 µM RITA

Protein	Diffusion time ± s.e.m.*	Change in diffusion time, percent
No protein	$0.063 \pm 0.011$	_
GST-p53 dN(1-63)	$0.356 \pm 0.070$	465
GST-p53 N(1-100)	$0.259 \pm 0.020$	311
GST-p53(1-393)	$0.287 \pm 0.043$	355
His-p53(1-393)	$0.198 \pm 0.007$	214
His-p53(1-312)	$0.111 \pm 0.014$	74
GST	$0.076 \pm 0.003$	20
GST-EBNA2	$0.073 \pm 0.017$	16

All experiments were performed at least three times.

subject for future research,

Accession codes. BIND identifiers (http://bind.ca):

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1. Issaeva, N. et al. Nat. Med. 10, 1321-1328 (2004).

# PrPSc in mammary glands of sheep affected by scrapie and mastitis

# To the editor:

Besides colonizing the central nervous system, the infectious agent of transmissible spongiform encephalopathies, termed prion, is predominantly associated with follicular dendritic cells (FDCs) of lymphoid tissues1,2. Accordingly, PrPSc, a protease-resistant isoform of the host protein PrPC representing the main prion constituent, is often detectable in spleen, tonsils, Peyer patches and lymph nodes of infected hosts.

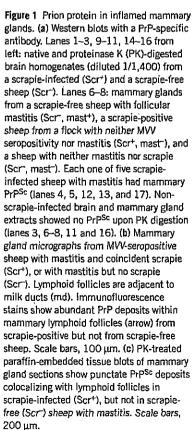
Chronic inflammatory states are accompanied by local extravasation of B cells and other inflammatory cells, which may induce lymphotoxin-dependent maturation of ectopic FDCs, Consequently, scrapie infection of mice suffering from nephritis, hepatitis or pancreatitis induces unexpected prion deposits at the sites of inflammation3. This has raised concerns that analogous phenomena might occur in farm animals.

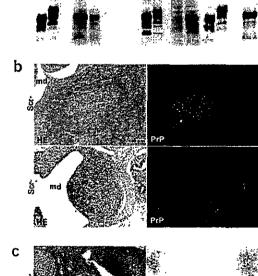
We have investigated this question in a flock of 818 Sarda sheep held in the Sassari region of Italy for production of wool and human foods. The European Surveillance Plan for Transmissible Spongiform Encephalopathies mandates the removal of all sheep of scrapie-susceptible genotypes in scrapie-infected flocks.Of the 818 sheep, 261 had Prnp alleles4 that conferred susceptibility to prion disease. Of the latter, seven had clinically overt scrapie with PrPSc in brain, lymph nodes and tonsil. All scrapie-sick sheep and 100 randomly chosen healthy sheep were killed, and mammary glands were analyzed histologically. Of these, 10 sheep had lymphocytic mastitis, and four had coincident mastitis and scrapie. Using western blots, immunohistochemistry and histoblots, we detected PrPSc in mammary glands of all

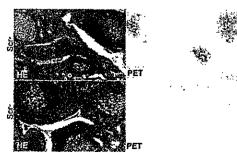
four clinically scrapie-sick sheep with mastitis (Fig. 1a,b), but not in noninflamed mammary glands from presymptomatic or scrapie-sick sheep from the same (n = 14) or a different flock (n = 1), nor in inflamed mammary glands of scrapie-uninfected sheep (n = 2). Within the inflammatory mammary lesions, PrPSc was found to be associated with lymphoid follicles

by immunofluorescent labeling and by paraffin-embedded tissue (PET) blotting (Fig. 1c). PrPSc colocalized predominantly with CD68+ macrophages and FDCs within inflamed mainmary glands (Fig. 2a).

We then surveyed a second Sarda flock (272 sheep) located 30 km away from the flock described above. One sheep was found to be







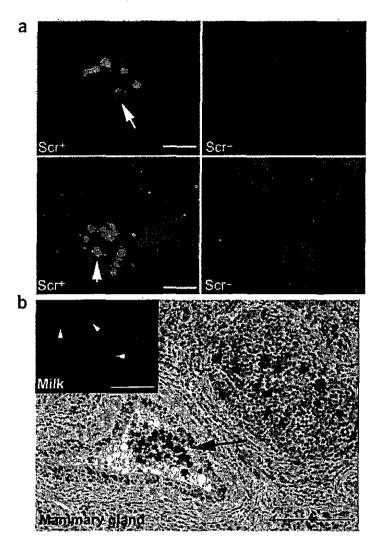


Figure 2 Mammary PrPSc localizes to macrophages and FDCs. (a) Mammary gland from a sheep with coincident mastitis, MVV seropositivity and scrapie (sheep #732). Confocal laser scanning micrographs of lymphoid follicles immunostained for PrP (green), nuclear DNA (blue) and macrophages (red, top panels) or FDC (red, bottom panels). PrPSc associates with CD68+ macrophages and FDCs in scrapie-positive (Scr\*, arrows) but not in scrapie-free sheep (Scr\*). Scale bars, 6.3 μm (top) and 7.5 μm (bottom). (b) CD68+ macrophages (arrow) and degenerating leukocytes within milk ducts and in adjacent lymphoid follicles of an inflamed mammary gland, as well as in milk sediment (inset, arrowheads). Scale bar, 100 μm (mammary gland) or 20 μm (milk cells).

scrapie-sick and was killed: necropsy showed lymphofollicular mastitis and PrPSc in the brain and tonsil. Again, PrPSc was present in the mammary gland (Fig. 1a). These results indicate that coincidence of natural chronic inflammatory conditions and natural scrapie can expand the deposition of PrPSc to unexpected tissues of sheep.

By plotting western blot signals against serially diluted scrapie-infected brain and spleen, we determined that the median mammary PrPSc concentration was 0.1% of that of spleen and 0.05% of brain. But because mammary lymphoid follicles were stochastically distributed, local PrPSc loads varied markedly. Hence these figures may underestimate PrPSc in sites

of abundant follicles, and overestimate it in sites with few or no follicles.

Common causes of lymphofollicular mastitis in sheep include Maedi-Visna virus (MVV) and inycoplasma<sup>5</sup>. We could not culture inycoplasma from mastitic glands, whereas we found that four of the five sheep with scrapie and mastitis were seropositive for MVV and that the three scrapie-sick sheep without mastitis were seronegative for MVV. In the clinically healthy group, 7 of 10 sheep with mastitis, but only 32 of 90 sheep without mastitis, were seropositive for MVV. Hence, MVV seropositivity correlated with lymphoid follicular mastitis (Fisher exact test, P = 0.01) as reported previously<sup>6,7</sup>.

MVV and related small-ruminant lentiviruses are endemic in most, if not all, European populations of small ruminants<sup>6</sup>. The above data suggest that common viral infections of small ruminants may enhance the spread of prions. MVV is found within mammary epithelial cells and macrophages8, and has been experimentally passed to lambs through milk9. Milk is believed to represent a major route of transmission for the natural spread of MVV5. The PrP deposits in CD68+ cells of mammary lymphoid follicles, in concert with the copious shedding of macrophages into milk of mastitic sheep (Fig. 2b)9,10, raises the question whether coexistence of prion infection and inflammation in secretory organs may lead to prion contamination of secretes, and may represent a cofactor for horizontal prion spread within flocks,

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# 医薬品

# 医薬部外品 研究報告 調査報告書

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識別番号・報告回数	回	<b>報告日</b> 年 月 日	第一報入手日 2005 年 10 月 25 日	新医薬品等の区分 該当なし	厚生労働省処理欄
一般的名称			Ultra-sensitive detection of prior fibrils by flow cytometry in bloo	d from	7
販売名(企業名)		   研究報告の公表状況   	cattle affected with bovine spong encephalopathy Lothar Trieschmann, Alexander Nav Santos, Katja Kaschig, Sandra Tork Maas, Hermann Schatzl and Gerald BMC Biotechnology 2005, 5:26	arrete ler, Elke	
出されることに による伝播が問 による伝播が思 た。凝集の検 た。凝集成した、 条塊の形成が仮	よってのみ下される。一方、 1題となっている。しかし PrP: :加えなかった場合のプリオン はフローサイトメトリーによ 蛍光標識したプリオン単量体 E進されることが確認された。	臨床症状を発現する前で scの血中濃度は非常に 全面単量体の重合化動 り行った。BSEを発症し を血清に添加すると、 この実験系では、発症	E(TSE)の確定診断は、死後の病理解剖でも、血液中に PrPsc が存在する可能性 低く、現在の分析法では検出することが 態の違いを利用して、高感度のプリオン た 6 頭のウシおよび 4 頭の正常ウシのご 添加しなかった場合と比較して、この単 ウシの血清 6 検体全てにおいて、10-8m 量の BSE 陽性の血清と正常血清とを識別	が示唆されており、輸血など が困難である。著者らは、「核」 蛋白凝集体の検出法を開発し 血清を使用して実験を行った。 量体が「核」となり新たな凝 M(0.24[g/mL)の低濃度でも合	その他参考事項等 BYL-2005-0199

報	告企	業0.	)意見
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プリオン病の発症前診断の手がかりとなるかもしれない。

# 今後の対応

弊社の血漿分画製剤の製造工程におけるプリオン除去能は 4 log を上回ることが確認されている。本論文の実験結果が実用化されれば、プリオンの理論的伝播リスクがさらに低減することが期待される。

現時点で弊社において新たな安全対策上の措置を講じる必要はないと考える。引き続き本方法の実用化および PrPss の検出・除去技術に関する関連情報の収集に努める。



# **BMC Biotechnology**



Research article

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# Ultra-sensitive detection of prion protein fibrils by flow cytometry in blood from cattle affected with bovine spongiform encephalopathy

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### **Abstract**

**Background:** The definite diagnosis of prion diseases such as Creutzfeldt-Jakob disease (CJD) in humans or bovine spongiform encephalopathy (BSE) in cattle currently relies on the post mortem detection of the pathological form of the prion protein (PrPSc) in brain tissue. Infectivity studies indicate that PrPSc may also be present in body fluids, even at presymptomatic stages of the disease, albeit at concentrations well below the detection limits of currently available analytical methods.

**Results:** We developed a highly sensitive method for detecting prion protein aggregates that takes advantage of kinetic differences between seeded and unseeded polymerization of prion protein monomers. Detection of the aggregates was carried out by flow cytometry. In the presence of prion seeds, the association of labelled recombinant PrP monomers in plasma and serum proceeds much more efficiently than in the absence of seeds. In a diagnostic model system, synthetic PrP aggregates were detected down to a concentration of approximately 10-8 nM [0.24 fg/ml]. A specific signal was detected in six out of six available serum samples from BSE-positive cattle.

**Conclusion:** We have developed a method based on seed-dependent PrP fibril formation that shows promising results in differentiating a small number of BSE-positive serum samples from healthy controls. This method may provide the basis for an *ante mortem* diagnostic test for prion diseases.

# **Background**

A group of fatal transmissible neurodegenerative diseases, including Creutzfeld-Jakob disease (CJD), bovine spongiform encephalopathy (BSE), chronic wasting disease (CWD) and scrapie, is caused by an unusual infectious agent that has been termed prion [1]. Prions consist of an

aberrant isoform (PrPSc) of the normal cellular prion protein (PrPC). PrPC is a cell surface glycoprotein expressed in neurons [2] and other cell types [3,4]. The precise physiological function of the cellular prion protein is not known yet. PrPSc differs from PrPC in its higher content of  $\beta$ -sheet structure [5,6], its partial resistance to protease digestion

[7], and its tendency to form large aggregates [8]. PrPSc propagates by converting the cellular prion protein to the PrPSc conformation [9]. PrPSc aggregates accumulate predominantly in the central nervous system (CNS), and definitive diagnosis of prion diseases currently relies on the post mortem detection of PrPSc in CNS tissue by immunohistochemistry, Western blotting, or ELISA [10]. Transmission studies indicate that prions may also be present in blood, potentially allowing for ante mortem diagnosis, but the sensitivity of the currently available analytical methods is insufficient for the detection of the extremely low prion titers that can be expected in body fluids [11].

Here, we report the development of a method based on kinetic differences between seeded and unseeded aggregation of prion protein that allows the detection of PrP aggregates in blood down to attomolar concentrations by flow cytometry.

# Results and discussion

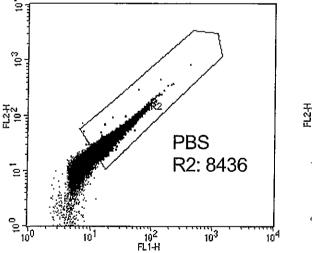
# Detection of synthetic prion protein aggregates in serum or plasma

Kinetic differences between seeded and spontaneous polymerization of peptide monomers can be used for the detection of amyloid  $\beta$ -protein aggregates in the cerebrospinal fluid of Alzheimer's disease patients [15]. Here, we

extend the principle of seeded polymerization to the detection of prion protein aggregates.

While trying to establish conditions for the labeling of synthetic prion protein aggregates with a fluorescently labeled prion protein probe, we observed that the formation of prion protein aggregates proceeds much less efficiently in serum or plasma (not shown) than in PBS (Fig. 1). This inhibition is probably caused by interactions of the prion protein probe with serum proteins.

Next, we found that the addition of preformed prion protein aggregates to plasma can partially overcome this inhibition (Fig. 2). The preformed aggregates presumably function as seeds that facilitate the formation of new aggregates in the inhibitory environment of plasma. The seeds stimulated the formation of prion protein aggregates at all concentrations tested, from 5 nM [120 ng/ml] to 10-8 nM [0.24 fg/ml] (Fig 2C). The average ratio of event counts in seeded samples to those in samples without seeds was 6.4. The number of events, however, was not proportional to the seed concentration, but remained relatively constant over the whole concentration range. Thus, the seed-dependent formation of prion protein aggregates can be used to detect extremely low amounts (down to the attomolar range) of spiked prion protein aggregates in blood.



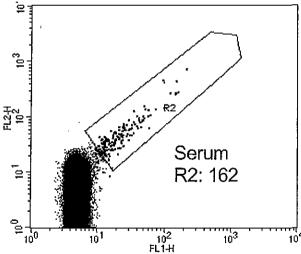
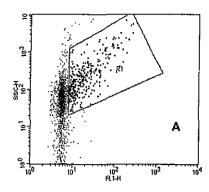
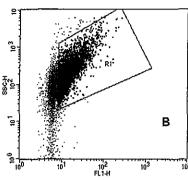


Figure I Inhibition of PrP aggregation in serum. FITC-labeled recombinant bovine prion protein (concentration 10 nM) was incubated at 37°C for 20 h with continuous shaking, either in 150 µl PBS (left panel) or in the same volume of serum (right panel), followed by flow cytometry. The measurements are depicted in a Fluorescence I (FLI-H) vs. Fluorescence 2 (FL2-H) dot-plot. The number of counts in the area containing specific signals (R2) is given in the figures. Aggregate formation in serum is strongly inhibited.





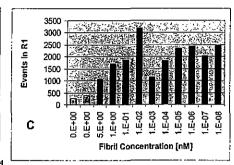


Figure 2
Seed-dependent PrP aggregate formation in plasma. FITC-labeled recombinant prion protein (5 nM) was incubated in plasma as described in the methods section for 20 h either in the absence (panel A) or presence (panel B) of 10-8 nM PrP aggregates. Panel C: quantification of measurements shown in A and B, and of measurements (not shown) with different seed concentrations. The measurements are depicted in a Fluorescence 1 (FL1-H) vs. Side-Scatter (SSC) dot-plot. Aggregate formation (signal in region R1) was strongly enhanced by all seed concentrations tested, from 5 nM to 10-8 nM.

### Analysis of serum from clinical-stage, BSE-positive cattle

Studies demonstrating the transmission of prion diseases by blood transfusion suggest that prions are present in the blood of afflicted animals and people, even at pre-symptomatic stages of the disease [16-18]. We used the method of seed-dependent fibril formation to analyze serum from six confirmed cases of clinical-stage, BSE-positive cattle and four controls. Based on the spiking experiments described above, our hypothesis was that any PrPSc aggregates present in serum may act as seeds for the formation of easily detectable amounts of labeled PrP aggregates, whereas in the absence of seeds the formation of PrP aggregates would be inhibited. The serum samples from BSE-positive cattle and controls from healthy cattle were incubated with 10 nM of a FITC-labeled bovine PrP probe at 37°C for 20 h with continuous shaking, followed by analysis in a flow cytometer. All six BSE-samples could be clearly distinguished by a population of events that was absent in the controls (Fig. 3A-J, green dots in region R3; quantification in fig. 3K).

## Conclusion

We have developed a method based on seed-dependent PrP fibril formation that shows promising results in differentiating a small number of BSE-positive serum samples from healthy controls. More samples need to be tested in order to validate its potential as an *ante mortem* diagnostic test for BSE and other prion diseases.

# Methods

### Biological fluids

Serum samples from six confirmed cases of BSE in cattle and four control animals were obtained from BFAV, Insel Riems, Germany. Control plasma was obtained from a blood bank.

### Labeling of prion protein

Recombinant full-length bovine PrP was produced as described previously [12,13]. The purified protein was labeled with a FITC-labeling kit (Roche) according to the manufacturer's instructions.

### Preparation of fibrils from recombinant prion protein

25 µM of unlabeled bovine prion protein in PBS containing 0.2 % SDS was incubated for 10 min at room temperature, followed by a twentyfold dilution with PBS. For fibril formation, the diluted reaction mixture was incubated for 48 h at room temperature [14].

### PrP fibril formation in serum or plasma

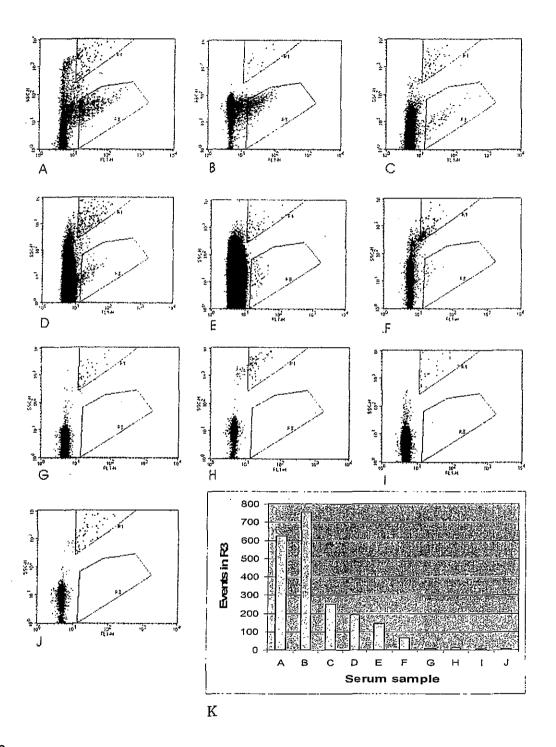
Recombinant FITC-labeled bovine prion protein was incubated in 150  $\mu$ l serum or plasma at a concentration of 5 or 10 nM for 5–10 min. at 20 °C, shaking at 550 rpm in an Eppendorf thermomixer, followed by an increase of the temperature to 37 °C h at constant shaking speed. The incubation was continued for 20 h. Samples were then analyzed by flow cytometry.

### Flow cytometry

Analysis of the samples was carried out on a FACSVantage flow cytometer (BD Biosciences) at room temperature, measurement time was 30 sec per sample.

### **Authors' contributions**

LT participated in the design of the study, carried out the measurements and drafted the manuscript. ANS participated in the analysis of the data. EM prepared the recom-



Analysis of serum from BSE-positive cattle. FITC-labeled recombinant prion protein (10 nM) was incubated in 150 µl of the serum samples as described in the methods section and analyzed by flow cytometry. The measurements are shown in a Fluorescence 1 (FLI-H) vs. Side-Scatter (SSC) dot-plot. All six BSE-samples (A-F) can be differentiated from the controls (G-J) by a population of events in region R3 (green dots). Panel K: Quantification of measurements shown in panels A-J.

binant protein. KK and ST were also involved in protein expression and purification. HS participated in the design and coordination of the study. GB conceived of the study and helped to draft the manuscript. All authors read and approved the final manuscript.

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