

Isolation and Characterization of Full-Length Recombinant Cattle PrP^C Protein

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Full-length *Bos taurus* PrP^C protein was obtained in the eu- and prokaryotic expression systems. Immunoblotting and indirect enzyme immunoassay demonstrated high specificity and antigenic activity of full-length proteins in the reactions with monoclonal antibodies (anti-SAF-32 and VRQ-84). Membrane location of recombinant PrP^C protein in insect cells was shown by immunofluorescent analysis.

Key Words: prion; recombinant protein; expression

Transmitted spongiform encephalopathies (TSE) are a group of human and animal lethal neurodegenerative diseases, including Creutzfeldt—Jakob disease, Gerstmann—Straussler—Scheinker syndrome, kuru, cattle spongiform encephalopathy (CSE), sheep scrapie, mink transmitted encephalopathy, and some other diseases [1].

The concept of protein nature of TSE agent is universally acknowledged [7]. The disease is believed to be associated with conformation transition of normal cellular PrP^C protein into abnormal PrP^d form (PrP^{res}, PrP^{Sc}) with increased content of sites with β -sheet conformation [5,6]. The disease develops because of accumulation of this proteinase-resistant abnormal protein isoform, called prion.

The development of rapid and effective methods for detection of prion protein in biological fluids, essential for early life-time diagnosis, is now in progress. Immunological methods for detection of PrP^d are developed most intensely [2]. The possibility of *in vivo* blocking PrP^C conversion into PrP^d for preventing disease development is investigated [11]. Preparation of recombinant PrP^C pro-

tein opens new vistas for the development of immunological methods for its detection in biological material and for *in vitro* studies of conformation transition of PrP^C molecule into PrP^d. This latter can be used in the search for agents blocking this transition *in vivo*.

The aim of our study was to obtain a recombinant PrP^C, evaluate its immunological characteristics and prospects for diagnostic use, and determine its subcellular location.

MATERIALS AND METHODS

Genome DNA isolated from *Bos taurus* blood served as the source of gene amplification. Constructs for expression of recombinant PrP^C protein in eukaryotic (baculovirus) "Bac-to-Bac" (Invitrogen) and prokaryotic pET-system (Novagen) systems were obtained by PCR and molecular cloning. The expression in eu- and prokaryotic systems was carried out as recommended by the manufacturers. Insect Sf-21 cells were used for transfection, Sf-21 and High-Five cells for accumulation of preparative amounts of baculovirus PrP^C. A panel of 19 anti-PrP^C monoclonal antibodies (MAB) was used: 1-Surg, 2pr-Surg, 2m-Surg, 3-Surg, 4G-11, 4G-29, 4G-57, 4G-69, 5G-13, 5G-29, 5G-38, 5G-48, 5G-

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55, 5G-77, obtained to synthetic Prionics Ag peptide (NARVAC Company); anti-SAF-32 and anti-PrP^C-Hum (gracious gift from J.-F. Deslie, CEA); and VRQ-14(33), VRQ-84, and PLT-40(2), kind gift from J. Grossklaude, INRA. Recombinant proteins were purified using Ni-NTA-agarose (Qiagen) in accordance with the instruction. A 3-day monolayer of Sf-21 cells infected with a recombinant baculovirus carrying PrP^C was stained with an immunofluorescent dye.

RESULTS

We assumed that full-length recPrP^C obtained in baculovirus system of expression was glycosylated, contained maximum number of antigen epitopes, and was highly immunocompetent, because post-translation modifications in insect cells were similar to those in mammals [4]. That is why it was selected for testing with a panel of MABs. The panel consisted of 19 MABs to fragments and full-length cattle, sheep, and human PrP^C and to brain suspension of a hamster infected with adapted scrapie strain. The resultant crude full-length protein was characterized by high specificity and antigenic activity in the reactions with anti-SAF-32 and VRQ-84 MABs (Fig. 1).

The location of full-length recPrP^C in insect cells was detected by indirect immunofluorescent analysis using VRQ-84 antibodies. Full-length recPrP^C was present mainly in the cell membrane, which was seen from bright contour of specific fluorescence (Fig. 2). This location of PrP^C is characteristic of mammalian cells [3]. Therefore, the syn-

thesis, processing, and intracellular transport of recombinant protein in insect cells are similar to those in mammalian cells. Hence, this cell system for expression of PrP^C protein can be used for investigation of the subcellular location, immunological and physicochemical characteristics of the protein.

At the next stage of the study crude recPrP^C proteins obtained in eu- and prokaryotic expression systems of were characterized in immunoblotting with anti-SAF-32 antibodies (Fig. 3). Immunoblotting of PrP^C of baculovirus origin showed, in addition to the main band (mol. weight 27-29 kDa), additional high molecular-weight bands, various glycoforms [8], and low molecular-weight products, emerging as a result of partial proteolytic degradation [10]. A lower molecular weight of the baculovirus product in comparison with mol. weight of recPrP^C, expressed in the prokaryotic system (30 kDa), seems to be due to removal of PrP^C signal sequence in insect cells, in contrast to *E. coli* cells where eukaryotic protein is not processed. Testing of recombinant products in indirect EIA with anti-SAF-32 and VRQ-84 gave similar results: antigens exhibited high activity and specificity of antibody binding, this indicating the absence of glycosylation effect on immunocompetence of recPrP^C protein.

Immunospecificity of fractions obtained by metal-affinity chromatography with Ni-NTA-agarose products was evaluated using anti-SAF-32 or VRQ-84 MAB by EIA and immunoblotting. The yield of the product under native conditions was lower than under denaturing (8 M urea) conditions. Further purification was carried out only under denaturing

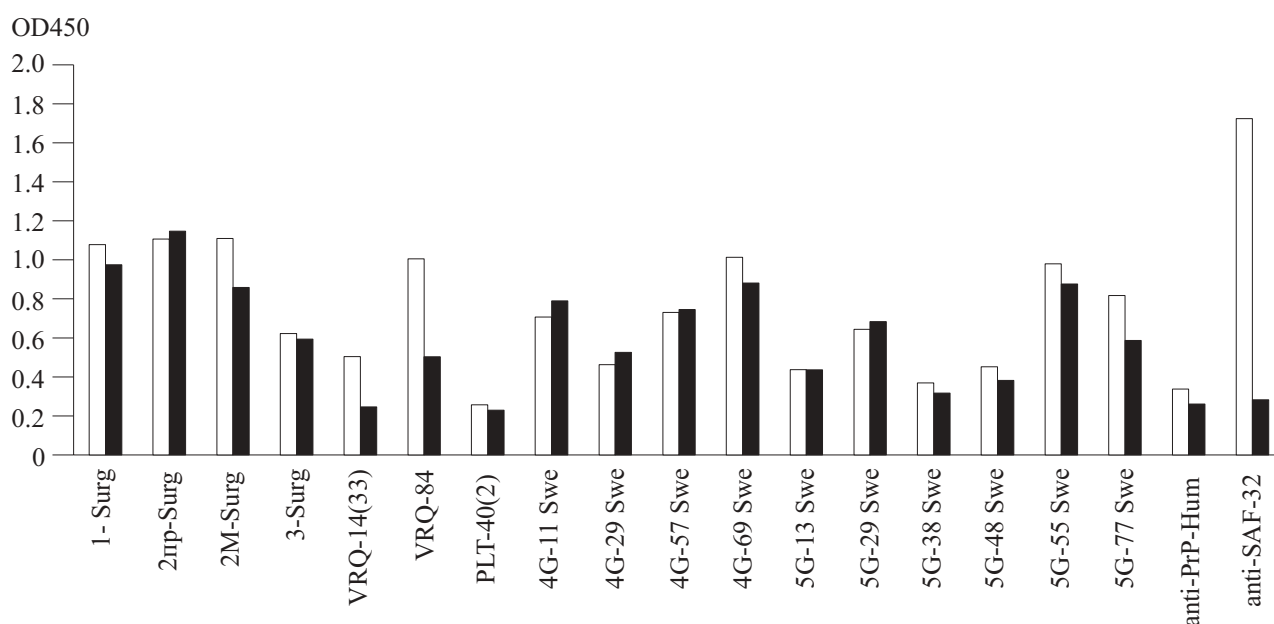


Fig. 1. Testing of a panel of monoclonal antibodies in indirect enzyme immunoassay with full-length cattle prion protein obtained in the baculovirus system of expression. Light bars: Sf-21 cells infected with recombinant baculovirus; dark bars: non-infected Sf-21 cells.

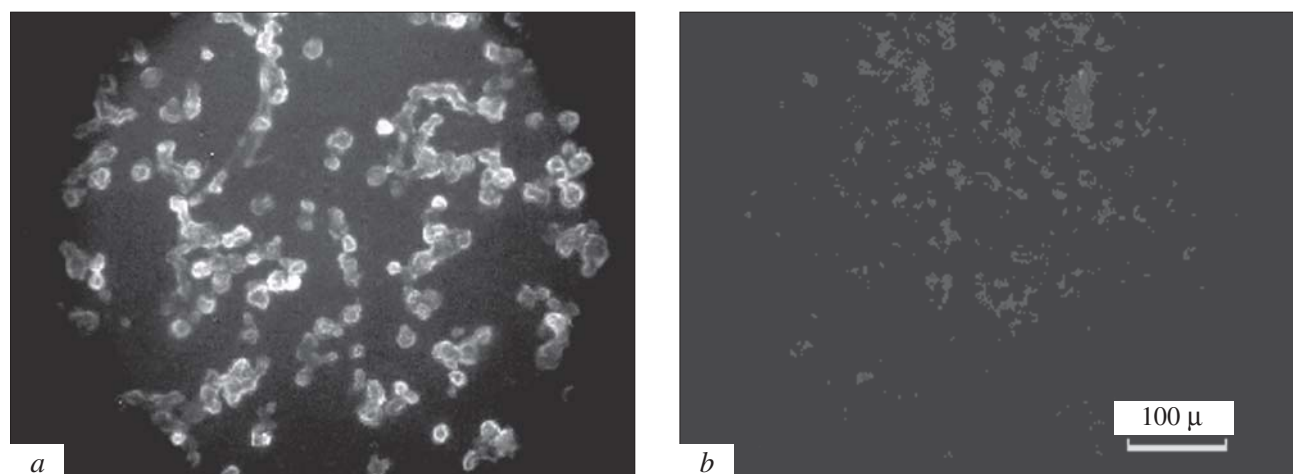


Fig. 2. Immunofluorescent staining of Sf-21 cells expressing the coding sequence of full-length cattle PrP^C protein. *a*) Sf-21 cells infected with recombinant baculovirus; *b*) negative control (Sf-21 cells infected with wild type baculovirus).

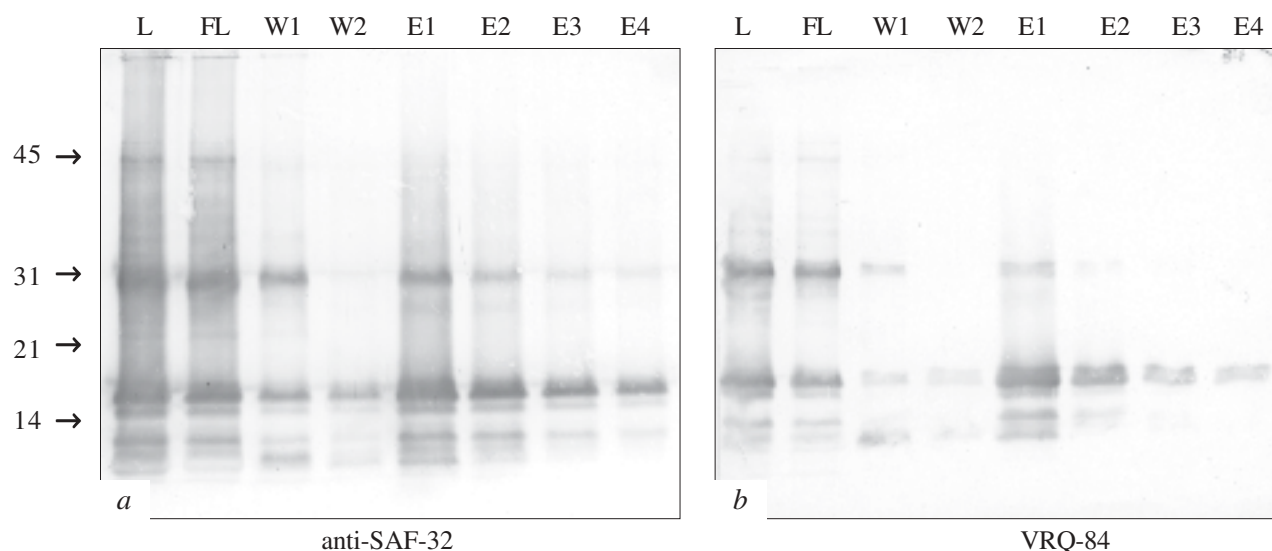


Fig. 3. Detection of recPrP of baculovirus origin by immunoblotting with anti-SAF-32 (*a*) and VRQ-84 monoclonal antibodies (*b*) in protein fractions purified by metal affinity chromatography on Ni-NTA-agarose. L: cell lysate; FL: free protein; W1-W2: washing buffer (pH 6.3); E1-E4: elution buffer (pH 4.5). Arrows show reference mol. weights.

conditions by the Qiagen protocol. The bulk of immunocompetent PrP was eluted from the column at pH 5.9-4.5, the eluates contained, along with full-length PrP with a mol. weight of 27-29 kDa, appreciable amounts of immunocompetent product with a mol. weight of about 17-19 kDa. The appearance of a low-molecular-weight protein seems to be due to partial degradation of full-length protein. Further use of proteinase inhibitor mixtures (Sigma) prevented proteolytic degradation of full-length baculovirus product and increased its output. Distribution of purified prokaryotic PrP^C protein by fractions was similar. The resultant fractions contained no proteolytic degradation products (even without proteinase inhibitor treatment). It seems

that amino acid sequence of PrP^C protein carries no recognition sites for *E. coli* proteinases. The expression of recombinant protein in the prokaryotic system was significantly higher than in baculovirus system.

Presumably, glycosylation has no appreciable effect on immunological recognition of recombinant PrP^C by specific MAB. The process of PrP^C to PrP^d conversion depends on the presence of sugars [9], and therefore conditions of *in vitro* conformation transition can be different for the resultant polypeptides. Higher level of expression in *E. coli* and absence of differences in antigenic activities of recombinant products are advantages suggesting the use of prokaryotic antigen in experiments re-

quiring appreciable amounts of recPrP^C. Our present findings and published data [1,7,8] indicate the possibility of using variants of *B. taurus* recombinant PrP^C as antigens for obtaining specific anti-PrP^C/PrP^d monoclonal antibodies, which can be useful in the development of approaches to the diagnosis of CSE. In addition, creation of a model for *in vitro* studies of PrP^C/PrP^d conformation transitions and understanding of the mechanism of this process are essential for further studies of therapeutic agents blocking the formation of infectious isoform of the prion in the body [11].

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REFERENCES

1. V. B. Grigor'ev, *Vopr. Virusol.*, **6**, 4-12 (2004).
2. L. Cervenakova and P. Brown, *Exp. Rev. Anti. Infect. Ther.*, **2**, 873-880 (2004).
3. D. A. Harris, *Clin. Microbiol. Rev.*, **12**, 429-444 (1999).
4. V. A. Luckow, *Curr. Opin. Biotechnol.*, **4**, 564-572 (1993).
5. K.-M. Pan, M. Baldwin, J. Nguyen, et al., *Proc. Natl. Acad. Sci. USA*, **90**, 10,962-10,966 (1993).
6. P. Pergami, H. Jaffe, and J. Safar, *Anal. Biochem.*, **236**, 63-73 (1996).
7. S. B. Prusiner, *Proc. Natl. Acad. Sci. USA*, **95**, 13,363-13,383 (1998).
8. P. M. Rudd, T. Endo, C. Colominas, et al., *Ibid.*, **96**, 13,044-13,049 (1999).
9. P. M. Rudd, M. R. Wormald, D. R. Wing, et al., *Biochemistry*, **40**, 3761-3766 (2001).
10. A. Sakudo, M. Hamaishi, T. Hosokawa-Kanai, et al., *Biochem. Biophys. Res. Commun.*, **307**, 678-683 (2003).
11. C. Weissman and A. Aguzzi, *Annu. Rev. Med.*, **56**, 321-344 (2005).