

## Cloning, Sequencing, and Expression of cDNA Encoding Bovine Prion Protein

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**Abstract** A normal prion protein (PrP<sup>c</sup>) is converted to a protease resistant isoform (PrP<sup>sc</sup>) by an apparent self-propagating activity in bovine spongiform encephalopathies (BSE), which is a neurodegenerative disease. The cDNA encoding bovine PrP open reading frame (ORF) in Korean cattle was cloned by polymerase chain reaction (PCR). The cloned cDNA had a length of 795 base pairs which coded for a protein of 264 amino acid residues with a calculated molecular mass of 28.6 kDa. Identities of 90, 90, 79 and 78% on nucleotide and 94, 94, 84, and 84% on amino acid sequence were shown to PrP genes from sheep, goat, human, and mouse, respectively. The cloned DNA was ligated into the pQE30 expression vector and transformed into *E. coli* M15. The PrP was expressed by induction with isopropyl- $\beta$ -D-thiogalactoside (IPTG) and purified on the Ni-NTA affinity column. High specific activities of the recombinant PrP were observed in the fraction of pH 5.8 eluate and showed a molecular mass of ~29 kDa on SDS-PAGE and Western blot analysis.

**Key words:** Prion protein, cDNA, BSE, Korean cattle

Bovine spongiform encephalopathy (BSE) is thought to be the group of transmissible spongiform encephalopathy (TSE), which comprises a group of slow degenerative diseases of the central nervous system (CNS). It is characterized by neuronal vacuolation and accumulation of the abnormal isoform (PrP<sup>sc</sup>) of a host-encoded cellular membrane protein, referred to as prion protein (PrP<sup>c</sup>), in CNS [7, 26].

The epidemic that arose is considered to be due to a change in the rendering process used to prepare cattle feed from ruminant offal [8]. Since an epidemic of BSE occurred in Britain in 1986 and caused the deaths of almost

200,000 cattles it has now become a subject of general interest, because of the potential risk it poses to public health [1, 19, 25]. Even though there is no report concerning an emergency of BSE in Korea, its risk in cattle and public health still remains.

PrP<sup>c</sup> is a normal cellular protein that is expressed in the neurons and glia of the brain and spinal cord, as well as in several peripheral tissues and leukocytes [3, 4, 6, 9, 12, 13]. The normal function of PrP<sup>c</sup> still remains to be revealed. However, its localization on the cell surface would be consistent with roles in cell adhesion and recognition, ligand uptake, or transmembrane signaling [8]. Because the PrP<sup>c</sup> may fail to perform its normal function when it is converted to the PrP<sup>sc</sup> isoform, the key to understanding the pathogenesis of this disease state is the conformational conversion of a normal isoform of PrP<sup>c</sup> (a cell surface glycoprotein) into a pathogenic isoform of PrP<sup>sc</sup> [16–18]. The two isoforms, PrP<sup>c</sup> and PrP<sup>sc</sup>, differ mainly in their secondary structures, sensitivity to proteinase K (PK), and solubility [5], although these two PrP isoforms (PrP<sup>sc</sup> and PrP<sup>c</sup>) are encoded by the same chromosomal locus and have the same amino acid sequence and molecular weight [2].

The different size of PrP fragments represents a suitable tool to determine the cellular and molecular effects of PrP<sup>sc</sup> *in vitro* and the position change in the tertiary structure of PrP, which has become an important subject in the pathogenesis [5, 18]. Thus, attempts are made nowadays to produce PrP fragments using recombinant cDNA techniques.

In the present study, as the first step in PrP research, we described cloning of cDNA encoding bovine prion protein from Korean cattle, comparative sequence analysis of the gene with PrP genes from other animals, and expression of a large fragment of the recombinant PrP in *E. coli* and its purification.

Total RNA was isolated from the fetal fibroblast of Korean cattle using Trizol Reagent (Gibco, CA, U.S.A.)

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**Table 1.** Synthetic oligonucleotide sequence used for PCR amplification of bovine PrP.

Primer	Oligonucleotide Sequence	Length (bases)
Forward	5'-GCATGCATGGTGAAAAGCCACATAGGCAGTTGG-3' <i>Sph</i> I	33
Reverse	5'-AAGCTTCTATCCTACTATGAGAAAATGAGGAA-3' <i>Hind</i> III	33

and chloroform, and aqueous total RNA was precipitated by isoprophenol and centrifugation at 12,000 ×g for 10 min. The RNA pellets were washed once with 75% ethanol and dissolved in diethylpyrocarbonate (DEPC)-treated water. The RNA was treated with 2 units of RNase-free DNase at 37°C for 30 min to remove residual DNA and was purified again with Trizol Reagent.

Single-stranded cDNA was synthesized using Superscript Preamplification System for First Strand cDNA synthesis Kit (Gibco, CA, U.S.A.). Five µg of purified total RNA was incubated with 50 units of Superscript II reverse transcriptase at 42°C for 50 min in the presence of 10 × RT buffer, 25 mM MgCl<sub>2</sub>, 0.1 M DTT, 10 mM dNTP, and 0.5 µg/µl oligo (DT). Samples were treated with 2 units of RNase H at 37°C for 20 min and the synthesized single-stranded cDNA was used as a template in the polymerase chain reaction (PCR).

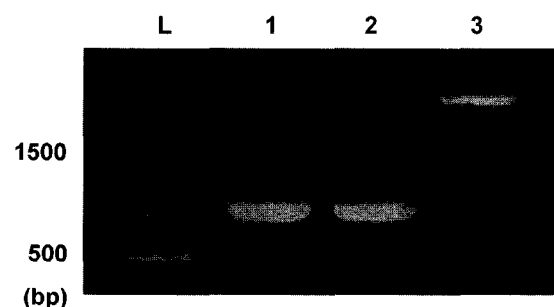
Primers were designed to obtain *Sph*I and *Hind*III enzyme sites by modification of the flanks representing nucleotides of the GenBank nucleotide entry (S55629) for bovine PrP (Table 1). The PCR solution was composed of 2 µl of 10 × buffer, 0.4 µl of dNTPs (2.5 mM), 0.5 µl each of 5'CS/3'CS (10 pmol/µl), 0.2 µl of *Taq* DNA polymerase (5 U/µl, Promega, Wisconsin, U.S.A.), 15.4 µl of distilled water, and 1 µl of template DNA (50 ng/µl). The PCR amplification consisted of an initial denaturation at 94°C for 3 min, followed by 30 cycles of denaturation at 94°C for 30 sec, annealing at 60°C for 30 sec, and extension at 72°C for 1 min and 30 sec, and then a final extension at 72°C for 15 min. PCR products were analyzed through electrophoresis on 1.0% agarose gel. The PCR product identical with bovine PrP in size (795 bp) was extracted by QIAquick Gel Extraction Kit (Qiagen, Venlo, Netherlands), and the purified amplicon was ligated with TA cloning vector (Invitrogen, California, U.S.A.) and transformed into One Shot Cell (Invitrogen, California, U.S.A.) for cloning. The cloned DNA fragment was sequenced using an automated DNA sequences (ABI PRISM 377 × L, perkin Elmer, U.S.A.) and compared with the GenBank database using the software DNassist.

PCR product was digested by *Sph*I and *Hind*III to obtain an insert DNA fragment with a staggered ends. This fragment was ligated with pQE30 prokaryote expression vector (Qiagen, Venlo, Netherlands), digested by the same enzymes, and transformed into *E. coli* M15 strain. Twenty ml of overnight culture of the transformed M15 were inoculated to 1 l of LB medium containing both 100 µg/ml of ampicillin and 25 µg/ml of kanamycin, and then grown

at 37°C with vigorous shaking until an OD<sub>600</sub> of 0.6 was reached. The expression of the mass cultured M15 was induced by 1 mM final concentration of isopropyl-β-D-thiogalactoside (IPTG), and they were incubated for additional 4 h. The M15 was harvested, lysed by lysis buffer (100 mM NaH<sub>2</sub>PO<sub>4</sub>, 10 mM Tris-Cl, 8 M urea, pH 8.0), and sonicated for 10 min. The lysate was applied onto Ni-NTA agarose column (Qiagen, Venlo, Netherlands) and eluted with elution buffers having a series of pH from 6.3 to 4.5. To confirm the expression of recombinant PrP, each eluat was loaded, resolved on 15% polyacrylamide gel (SDS-PAGE), and then transferred to a nitrocellulose membrane. The membrane was incubated overnight at 4°C with mouse anti-PrP monoclonal antibody (F89/160.1.5) [15]. Bound mAbs were detected with an alkaline phosphate (AP) conjugated goat anti-mouse IgG (Bio-Rad, CA, U.S.A.), and the blots were developed using AP-Substrate Kit (Bio-Rad, CA, U.S.A.).

Prion is a proteinaceous infectious particle that is thought to be the causative agent of transmissible spongiform encephalopathies (TSEs), comprising a group of fatal neurodegenerative disorders, and these disorders can arise sporadically or can have an infectious or genetic etiology [10, 18, 24]. PrP encoded by bovine PrP cDNA comprised 264 amino acids and has a predicted unglycosylated molecular weight of 28,614 Da [20].

In the present study, the PrP ORF of Korean cattle amplified by PCR using cDNA of fetal fibroblast of healthy cattle was cloned into TA cloning vector. The plasmid vector was treated with *Sph*I and *Hind*III, and the insert DNA was determined to have 795 bp through electrophoresis on 1.0% agarose (Fig. 1). To avoid



**Fig. 1.** Agarose gel (1.0%) electrophoresis pattern of the insert DNA and pQE30 expression vector digested with *Sph*I and *Hind*III.

Lane L, 100 bp DNA ladder (Promega); lanes 1 and 2, insert DNA; lane 3, pQE30 expression vector.

M V K S	H I G	S W I	L V L F	V A M	W S D
10	20	30	40	50	60
atgggtgaaa	gccacatagg	cagttgggac	ctgggttctct	ttgtggccat	gtggagtgcac
taccactttt	cggtgtatcc	gtcaacctag	gaccaagaga	aacaccggta	cacctcactg
V G L C	K K R	P K P	G G G W	N T G	G S R
70	80	90	100	110	120
gtgggctctt	gcaagaagcg	accaaaactt	ggaggaggat	ggaacactgg	ggggagccga
caccgggaga	cgttcttcgc	tggttttgga	cctcttccta	ccttgtagcc	ccccctggct
Y P G Q	G S P	G G N	R Y P P	Q G G	G G W
130	140	150	160	170	180
taccaggagc	agggcagtc	tggaggcaac	cgttatccac	ctcaggaggg	gggtggctgg
atgggtctct	tcccctcagg	accctcgttg	gcaataggtg	gagtcctctc	cccaccgacc
Q P H	G G G	W G Q	P H G G	G W G	Q P H
190	200	210	220	230	240
gttcagcccc	atggaggtgg	ctggggccag	cctcatggag	gtggctgggg	ccagcctcat
cagctcgggg	taccctcacc	gaccctggtc	ggagtaccct	caccgacccc	ggtcggagta
G G W	G Q P	H G G	G W G Q	P H G	G G G
250	260	270	280	290	300
ggaggtggct	ggggtcagcc	ccaatggtag	ggctggggac	agccacatgg	tggtaggggc
cctccaccga	ccccagtcgg	ggtaccacca	ccgacccttg	tcgggtgacc	accacctccg
* G Q G	G T H	G Q W	N K P S	K P K	T N M
310	320	330	340	350	360
tggggctcaag	gtggtagcca	cggtcaatgg	aacaaacca	gtaagccaaa	aaccacatg
atccagcttc	caccatgggt	gccaagtacc	ttgtttgggt	cattcggttt	ttggtgtac
K H V A	G A A	A A G	A V V G	G L G	G Y M
370	380	390	400	410	420
aagcatgtgg	caggagctgc	tgcagctgga	gcagtggtag	ggggccttgg	tggctacatg
ttcgtacacc	gtcttcgacg	acgtcgcact	cgtcacatc	ccccggaacc	accgatgtac
I G S A	M S R	P L I	H F G S	D Y E	D R Y
430	440	450	460	470	480
ctgggaagtg	ccatgagcag	gcctcttata	cattttggca	gtgactatga	ggaccgttac
gacccttcac	ggtactcgtc	cggaagaat	gtaaaaccgt	cactgtaact	cctggcaatg
Y R E N	M H R	Y P N	Q V Y Y	R P V	D Q Y
490	500	510	520	530	540
tatcgtgaaa	acatgcaccg	ttaccccaac	caagtgtact	acaggccagt	ggatcagtat
atagcacttt	tgtacgtggc	aatggggttg	gttcacatga	tgctccggtca	cctagtcata
S N Q N	N F V	H D C	V N I T	V K E	H T V
550	560	570	580	590	600
agtaaccaga	acaactttgt	gcattgaact	gtcaacatca	cagtcaagga	acacacagtc
ttatgtgtct	tgttgaaa	cgactgaca	cagttgtagt	gtcagttcct	tgtgtgtcag
T T T T	K G E	N F T	E T D I	K M M	E R V
610	620	630	640	650	660
aacaccacga	ccaaggggga	gaacttcacc	gaaactgaca	tcaagatgat	ggagcagatg
tgtgggtggc	ggttccccct	cttgaagtgg	ccttgaactgt	agttctacta	ctctcgtcac
V E Q M	C I T	Q Y Q	R E S Q	A Y Y	Q R G
670	680	690	700	710	720
gtggagcaaa	tgtgcattac	ccagtaccag	agagaatccc	aggcttatta	ccaacgaggg
caccctgttt	acacgtaatg	ggtcatggtc	ttctcttaggg	tccgaataat	ggttgctccc
A S V I	L F S	S P P	V I L L	I S F	L I F
730	740	750	760	770	780
gaaagtgtga	tcctcttctc	ttccccctct	gtgatcctcc	tcatctcttt	cctcattttt
ctttcacact	aggagaagag	aaggggagga	cactagaggg	agtagagaaa	ggagtaaaaa
L I V G	*				
790	795				
ctcatagtag	gatag				
gagtatac	ctatc				

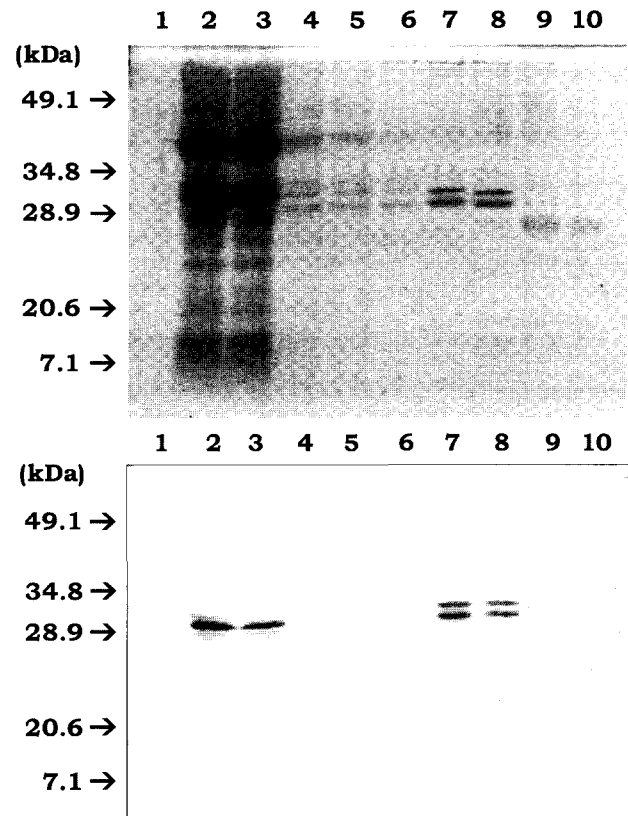
**Fig. 2.** Nucleotide and predicted amino acid sequence of bovine PrP.

The deduced amino acid sequence is shown by the single-letter amino acid codes over the nucleotide sequence. Stop codon is indicated by asterisk.

potential artifacts due to PCR, five positive clones from different trials were isolated, sequenced, and compared with each other. The confirmed plasmid vector with insert DNA was sequenced and reported to GenBank database

**Table 2.** Comparison of nucleotide and amino acid sequences of the clones with PrP genes originated from different animals.

	Bovine <i>Bos taurus</i>	
	Identities of nucleotide (%)	Identities of amino acid (%)
Sheep <i>Ovis aries</i>	90	94
Goat <i>Capra hircus</i>	90	94
Human <i>Homo sapiens</i>	79	84
Mouse <i>Mus musculus</i>	78	84



**Fig. 3.** SDS-PAGE (top) and Western blot (bottom) analysis. Fractions were obtained in each purification step of Ni-NTA column chromatography. Lane 1, LB broth used for culturing M15; lane 2, lysate; lane 3, non-bound fraction; lane 4, 4th washing; lanes 5 and 6, pH 6.3 eluates; lanes 7 and 8, pH 5.8 eluates; lanes 9 and 10, pH 4.5 eluates.

with Accession No. AF517842 (Fig. 2). Comparison of the nucleotide sequences of the present bovine PrP ORF with that reported by Prusiner [20] (GenBank Accession No. S55629) revealed that both were identical. The nucleotide sequence of bovine PrP ORF in the cDNA was approximately 90, 90, 79 and 78% homologous and the predicted amino acid sequence of the cDNA was approximately 94, 94, 84, and 84% homologous to those from sheep, goat, human, and mouse [11, 14, 21, 22], respectively (Table 2).

To express the bovine PrP in *E. coli* M15, the cloned vector plasmid was digested with *SphI* and *HindIII* to obtain an insert DNA fragment with staggered ends and ligated into pQE30 prokaryote expression vector system. Although positive transformants were screened by antibiotics, both ampicillin and kanamycin in growth media, a false positive could not be avoided. Therefore, additional analysis with restriction enzymes and sequencing should be performed to confirm the presence of insert DNA. The recombinant PrP was expressed in *E. coli* M15 by IPTG induction and purified by applying it to an affinity chromatography. The expressed bovine PrP was found by

Western blot using mouse anti-PrP monoclonal antibody to be identical with that previously reported [15], and it was purified by immobilized-metal affinity chromatography column, Ni-NTA resin, with pH 5.8 elution buffer (Fig. 3). However, the protein concentration was found to be extremely low. Significant difficulties have been reported in the expression of recombinant PrP, probably due probably to their insolubility, aggregativity, sensitivity towards proteolytic digestion and instability, as well as environmental conditions [11, 12, 17]. In some cases, this was caused in mammalian cells by misfolding and accumulating into inclusion bodies [5, 23].

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