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Molecular Cloning and Sequence Analysis of Prion Protein Gene of Dezhou Donkey in China

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Abstract: Transmissible Spongiform Encephalopathy (TSE) or prion disease has been reported in many herbivores but not in *equus* and the species barrier might be playing a role in resistance of these species to the disease. Therefore, analysis of genotype of prion protein (PrP) in these species may help understand the transmission of the disease. Dezhou donkey is a species of *equus* reared widely in China for service, food and medicine but its PrP gene has not been studied. Based on the reported PrP sequence in GenBank we designed primers and amplified, cloned and sequenced the PrP gene of Dezhou donkey. The sequence analysis showed that the Dezhou donkey PrP gene was consisted of an open reading frame of 767 nucleotides encoding 256 amino acids. Sequence comparison showed that the PrP gene of Dezhou donkey and horse shared 91.53% homology in nucleic acids and 94.03% homology in amino acids. Amino acid residues unique to donkey as compared with horse, mink, bovine, goat, human, elk and hamster were identified. The results provided the PrP gene data for an additional *equus* species, which should be useful to the study of the species barrier of TSE.

Key words: Dezhou donkey, prion gene, molecular cloning, sequence analysis, TSE

Introduction

Transmissible Spongiform Encephalopathy (TSE) is a group of sub-acute infectious diseases in human and animals including scrapie in ovine, bovine spongiform encephalopathy in cattle, chronic wasted disease in deer, transmissible mink encephalopathy in mink, feline spongiform encephalopathy in cat and Creutzfeldt-Jakob disease in human (Prusiner, 1991; Prusiner, 1997). It is understood that the key of development of these diseases is the transformation of a cellular prion protein (PrP^c) into PrP^{sc}, a diseased form of PrP characterized by a β -sheet-rich topology and anti-proteolytic property. Misfolded PrP^{sc} has a tendency of aggregation, shaping first into short amyloid fiber and then aggregated amyloid spot before becoming the spongiform-like degenerative changes in brain, which can be identified by clinic detection.

Dezhou donkey is a domestic breed of donkey widely reared in northern China. It is a species of *asinus*, belonging to *Equus*, Equidae of Perissodactyla, by biological classification. Dezhou donkey is mainly used for service and meat. Since the outbreak of the BSE, beef has increasingly been substituted

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by meat from other animals including donkey. In addition, donkey skin is used as the main ingredient of Ajjiao, an essential blood-nourishing traditional Chinese medicine that has been certified by FDA in the USA and is widely exported to European and American market. Donkey is herbivore, like bovine and ovine. While both bovine and ovine are susceptible to the prion disease, up to the present there has been no report of the TSE cases in donkey as well as horse. It was thought that the species barrier might be playing a role in resistance of these species to the prion disease.

Prion species barrier is a phenomenon of one species being able to protect the prion infection from other species (Hill and Collinge, 2002). There are species particularity and strain particularity of prion in the process of prp^{sc} propagation. The species particularity is that one species can protect the prion attack from the other species, representing the prolonged preclinical period or even no occurrence of the disease. The essential reason of species particularity is the difference in primary and secondary structure of PrP between the recipient and host species (Hill *et al.*, 2000). Until now, the PrP gene has been cloned in human and over 30 other animals (Wopfner *et al.*, 1997). The DNA of PrP is consisted of a single exon and its coded protein is highly conserved, with human and many species of animals sharing 92.9-99.6% of homology in PrP amino acids. Variation or the polymorphism of PrP gene in human being, mouse, sheep, bovine and deer is considered to be associated with the prion disease (Liberski and Brown, 2004; Aguzzi and Polymeridou, 2004).

Studies on the PrP gene in *equus* are very limited. The available data showed that the PrPs of the domestic horse (*Equus familiaris caballus*) and the falabella horse (*Equus przewalskii caballus*) are identical on the amino acid level and have two substitutions compared to the zebra PrP (*Equus quagga boehmi*) (Wopfner *et al.*, 1997). However, there is no report about the PrP gene in donkey. In the present study, the authors cloned and sequenced the PrP gene of Dezhou donkey and analyzed the sequence of the PrP gene with that from other animals. The results provided the PrP gene data for an additional *equus* species, which should be useful to the study of the species barrier of TSE and also of interest to public healthy.

Materials and Methods

Biological Materials

Genomic DNA was extracted from peripheral blood of six Dezhou donkeys using a Genomic DNA Rapid Isolation Kit for Blood (Biodev, USA) according to the manufacturer's instructions.

Polymerase Chain Reaction (PCR)

Each DNA sample was PCR-amplified for the PrP gene in a final reaction volume of 50 μ L containing 250 ng of genomic DNA, 20 pmol of each primer, 250 μ M of dNTP (Pharmacia) and 1.5 unit of Taq polymerase (Perkin Elmer Cetus). The primers were deduced from *white-tail deer* PrP sequence in GenBank (accession number AY360089), which were 5'-ATGGTGAAAAGCCACATAGGC-3' for sense primer and 5'-GAAGATAATGAAAACAGGAA G-3' for anti-sense primer. The PCR procedure comprised of an initial denaturation at 94°C for 5 min and 30 cycles of denaturation at 94°C for 45 sec, annealing at 58 °C for 50 sec and elongation at 72°C for 50 sec. Finally, an extension step at 72°C for 10 min was performed (Yang *et al.*, 2005; Zhang *et al.*, 2004; Wang *et al.*, 1998).

Cloning, sequencing and analysis of the amplified PrP products

Following confirmation of the amplified DNA fragment of 767 bp on a 1% agarose gel with DNA marker (DL2000, TAKALA), the PCR products were cloned into T-easy vectors (Promega, USA) and the propagated plasmids were prepared using Qiagen Plasmids Mini Kit (Qiagen, UK). Four positive clones were identified from each sample by restriction enzyme digestion with *EcoRI* and sequenced in both directions on an ABI Prism 377 automated DNA sequencer (Applied Biosystems, Foster City, USA). The nucleotide and amino acid sequences were analyzed using the DNAMAN software.

Results and Discussion

The amplified PCR product of Dezhou donkey PrP gene was 767 bp in size, similar to PrP gene reported in other animals (Fig. 1).

Sequencing of the amplified PCR fragments revealed a single open reading frame of 767 nucleotides encoding 256 amino acids. The newly identified sequence has been deposited into GenBank under the accession number AY968590.

Sequence comparison showed that the Dezhou donkey PrP gene had 91.53% homology in nucleic acid sequence and 94.03% homology in amino acids with that of domestic horse (GenBank AF117317) (Fig. 2). Dezhou donkey PrP gene also shared high homology in both nucleic acid and amino acid sequences with those from other mammals, including elk, human, mink, bovine, goat and hamster (Table 1).

In keeping with the structure of PrP in other animals, Dezhou donkey PrP had an N-terminal 23 amino acid signal peptide. This was followed by a glycine and proline-rich region (24-96)

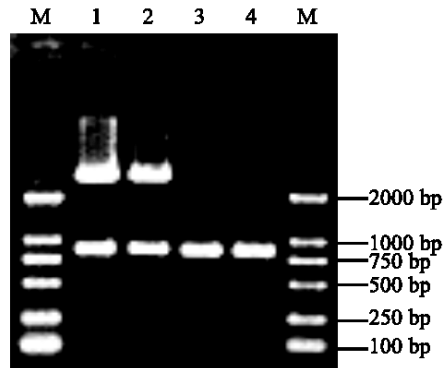


Fig. 1: Agarose gel electrophoresis of recombinant T-easy-vectors following digestion with *EcoRI* (M: marker DL2000; 1 and 2: *EcoRI* digested recombinant T-easy-vectors; 3 and 4: PCR products using recombinant T-easy-vectors as template)

Table 1: Homologies of PrP between Dezhou donkey and other species

Species		Horse	Elk	Human	Mink	Bovine	Goat	Hamster
Dezhou donkey	Nucleotide (% identity)	91.53	89.90	88.99	88.24	87.56	86.92	83.07
	Amino acid (% identity)	94.03	93.39	91.05	90.66	90.66	87.16	85.99

Dezhou don
Horse
Mink
Bovine
Goat
Human
Elk
Hamster	CTGATGCCAT GACTGTCACC CATTGTCT GCAAAAAATG TCAAGTGGCT
Dezhou don
Horse
Mink
Bovine TG CTGACACCCT
Goat TATGATG CTGACACCCT
Human
Elk ACACCCT
Hamster	TCAGCCTGCG TGCTGGACAG TGTGTGTTG TTGGAGTATA CTGACGCCTT
Dezhou don ATGGTGA AAAGCCACGT AGGCGGCTGG
Horse
Mink ATGGTGA AAAGCCACAT AGGCAGCTGG
Bovine	CTTTATTTG CAGATAAGTC ATCATGGTGA AAAGCCACAT AGGCAGTTGG
Goat	CTTTATTTG CAGAGAAGTC ATCATGGTGA AAAGCCACAT AGGCAGTTGG
Human ATGGTGA AAAGCCACAT AGGCAGCTGG
Elk	CTTTATTTG CAGATAAGTC ATCATGGTGA AAAGCCACAT AGGCAGCTGG
Hamster	GTTCTTCATT TTGCAGATTA GCCATCATGG CAA... ACCT TAGCTACTGG
Dezhou don	ATTCTGGTTC TCTTTGTGGC CACATGGAGT GACGTGGGGC TCTGCAAGAA
Horse ACATGGAGT GACGTGGGGC TCTGCAAGAA
Mink	CTCCTGGTTC TCTTTGTGGC CACATGGAGT GACATGGCT TCTGCAAGAA
Bovine	ATCCTGGTTC TCTTTGTGGC CATGTGGAGT GACGTGGGCC TCTGCAAGAA
Goat	ATCCTGGTTC TCTTTGTGGC CATGTGGAGT GACGTGGGCC TCTGCAAGAA
Human	ATCCTAGTTC TCTTTGTGGC CATGTGGAGT GACGTGGGCC TCTGCAAGAA
Elk	ATCCTAGTTC TCTTTGTGGC CATGTGGAGT GACGTGGGCC TCTGCAAGAA
Hamster	CTGCTGGCAC TCTTTGTGGC TACGTGGACT GATGTTGGCC TCTGCAAGAA
Dezhou don	GCGACCGAAG CCT... G GAGGATGGAA CACTGGGGG AGCCGATACC
Horse	GCGACCGAAG CCT... GGAG GAGGATGGAA CACTGGGGG AGCCGATACC
Mink	GCGGCCAAG CCT... GGAG GAGGCTGGAA CACTGGGGG AGCCGATACC
Bovine	GCGACCAAAA CCTGGA. G. G AGGATGGAAC ACTGGGGGA GCCGATACCC
Goat	GCGACCAAAA CCTGGC. G. G AGGATGGAAC ACTGGGGGA GCCGATACCC
Human	GCGACCAAAA CCTGGA. G. G AGGATGGAAC ACTGGGGGA GCCGATACCC
Elk	GCGACCAAAA CCTGGA. G. G AGGATGGAAC ACTGGGGGA GCCGATACCC
Hamster	GCGGCCAAG CCT... G. G AGGATGGAAC ACTGGTGGAA GCCGATACCC

Dezhou don	CAGGGCAGGG	TAGTCCTGGA	GGCAACCGCT	ATCCACCCCA	GGGAGGGGGC
Horse	CCGGGCAGGG	CAGTCCTGGA	GGCAACCGCT	ACCCACCCCA	GGGCGGTGGC
Mink	CAGGGCAGGG	CAGTCCTGGA	GGCAACCGCT	ACCCACCCCA	GGGTGGTGGC
Bovine	AGGACAGGGC	AGTCCTGGAG	GCAACCGTTA	TCCACCTCAG	GGAGGGGGTG
Goat	GGGACAGGGC	AGTCCTGGAG	GCAACCGCTA	TCCACCTCAG	GGAGGGGGTG
Human	GGGACAGGGA	AGTCCTGGAG	GCAACCGCTA	TCCACCTCAG	GGAGGGGGTG
Elk	GGGACAGGGA	AGTCCTGGAG	GCAACCGCTA	TCCACCTCAG	GGAGGGGGTG
Hamster	TGGGCAGGGC	AGCCCTGGAG	GCAACCGTTA	CCCACCTCAG	GGTGGTGGCA

Dezhou don	GGCTGGGGAC	AGCC. C. C. A	. C. G. G. A. G	. G. T. G. GCT	GGGACAGCC
Horse	GGCTGGGGTC	AACC. C. C. A	. T. G. G. T. G	. G. T. G. GTT	GGGGTACGCC
Mink	GGCTGGGGCC	AGCC. C. C. A	. C. G. G. G. G	. G. T. G. GCT	GGGACAGCC
Bovine	GCTGGGGTCA	GCCCCATGGA	GGTGGCTGGG	GCCAGCCTCA	TGGAGGTGGC
Goat	GCTGGGGTCA	GCCC. CA	TGGAGGTGGC
Human	GCTGGGGTCA	GCCC. CA	TGGAGGTGGC
Elk	GCTGGGGTCA	GCCC. CA	TGGAGGTGGC
Hamster	CCTGGGGGCA	ACCC. C. A. T	. G. G. T. G. G	. T. G. G. ATG	GGGACAGCCC

Dezhou don	CCACGGAGGC	GGCTGGGGAC	AGCCCCACGG	TGGCGGCTGG	GGACAGCCCC
Horse	CCATGGTGGT	GGCTGGGGTC	AGCCGCATGG	TGGTGGTTGG	GGACAGCCCC
Mink	CCACGGGGGT	GGCTGGGGAC	AGCCCCACGG	GGGTGGCTGG	GGACAGCCGC
Bovine	TGGGGCCAGC	CTCATGGAGG	TGGCTGGGGT	CAGCCCCATG	GTGGTGGCTG
Goat	TGGGGCCAAC	CTCATGGAGG	TGGCTGGGGT	CAGCCCCATG	GTGGTGGCTG
Human	TGGGGCCAAC	CTCATGGAGG	TGGCTGGGGT	CAGCCCCATG	GTGGTGGCTG
Elk	TGGGGCCAAC	CTCATGGAGG	TGGCTGGGGT	CAGCCCCATG	GTGGTGGCTG
Hamster	CATGGTGGTG	GCTGGGGACA	ACCTCATGGT	GGTGGTTGGG	GTCAGCCCCA

Dezhou don	ATGGTGGCGG	AGGCTGGGGT	CAAGTGGTG	GCT. CCCACGGTC
Horse	ATGGTGGTGG	AGGCTGGGGT	CAA. . GGTG	GCT. CCCATGGTC
Mink	ATGGTGGCGG	TGGCTGGGGT	CAAGTGGTG	GGA. GCCACGGTC
Bovine	GGGACAGCCA	CATGGTGGTG	GAGGCTGGGG	TCAAGTGGT	AGCCACGGTC
Goat	GGGACAGCCA	CATGGTGGTG	GAGGCTGGGG	TCAAGTGGT	AGCCACAGTC
Human	GGGACAGCCA	CATGGTGGTG	GAGGCTGGGG	TCAAGTGGT	AGCCACAGTC
Elk	GGGACAGCCA	CATGGTGGTG	GAGGCTGGGG	TCAAGTGGT	AGCCACAGTC
Hamster	TGGTGGTGGC	TGGGGTCAAG	GAGGTGGCAC	CCACAATCAG	TGGAACAAGC

Dezhou don	AGTGAACAA	GCCCAGTAAG	CCGAAAACCA	ACATGAAGCA	TGTGGCAGGC
Horse	AGTGAACAA	GCCCAGTAAG	CCAAAAACCA	ACATGAAGCA	TGTGGCAGGA
Mink	AGTGGGGCAA	GCCCAGTAAG	CCTAAAACCA	ACATGAAGCA	TGTGGCGGGT
Bovine	AATGAACAA	AGCCAGTAAG	CCAAAAACCA	ACATGAAGCA	TGTGGCAGGA
Goat	AGTGAACAA	GCCCAGTAAG	CCAAAAACCA	ACATGAAGCA	TGTGGCAGGA
Human	AGTGAACAA	GCCCAGTAAA	CCAAAAACCA	ACATGAAGCA	TGTGGCAGGA
Elk	AGTGAACAA	GCCCAGTAAA	CCAAAAACCA	ACATGAAGCA	TGTGGCAGGA
Hamster	CCAGTAAGCC	AAAAACCAAC	ATGAAGCATG	TGGCAGGTGC	GGCTGCGGCT

Dezhou don	GCCGCTGCAG	CTGGAGCAGT	GGTAGGGGGC	CTCGGCGGTT	ACATGCTGGG
Horse	GCTGCGGCAG	CTGGAGCAGT	GGTTGGGGGC	CTCGGCGGCT	ACATGCTGGG
Mink	GCCGCAGCAG	CCGGGGCGGT	TGTGGGGGGC	CTGGGCGGCT	ACATGCTGGG
Bovine	GCTGCTGCAG	CTGGAGCAGT	GGTAGGGGGC	CTTGGTGGCT	ACATGCTGGG
Goat	GCTGCTGCAG	CTGGAGCAGT	GGTAGGGAGC	CTTGGTGGCT	ACATGCTGGG
Human	GCTGCTGCAG	CTGGAGCAGT	GGTAGGGGGC	CTCGGTGGCT	ACTTGTGGG
Elk	GCTGCTGCAG	CTGGAGCAGT	GGTAGGGGGC	CTCGGTGGCT	ACATGCTGGG
Hamster	GGGGCGGTGG	TGGGGGCCT	TGGGGCTAC	ATGCTGGGA	GCGCATGAG

Dezhou don	GAGTGCCATG	AGCAGACCCC	TGATACACTT	TGGCAGTGAC	TATGAGGACC
Horse	GAGTGCCATG	AGCAGACCCC	TCATTCATTT	TGGCAATGAC	TATGAGGACC
Mink	GAGCGCCATG	AGCAGGCCCC	TCATTCATTT	TGGCAACGAC	TATGAGGACC
Bovine	AAGTGCCATG	AGCAGGCCTC	TTATACATTT	TGGCAATGAC	TATGAGGACC
Goat	AAGTGCCATG	AGCAGGCCTC	TTATACATTT	TGGCAATGAC	TATGAGGACC
Human	AAGTGCCATG	AGCAGGCCTC	TTATACATTT	TGGCAATGAC	TGTGAGGACC
Elk	AAGTGCCATG	AGCAGGCCTC	TTATACATTT	TGGCAATGAC	TATGAGGACC
Hamster	CAGGCCATG	CTCCATTTG	GCAATGACTG	GGAGGACCGC	TACTACCGTG

Dezhou don	GTTACTATCG	TGAAAACATG	TACCGTTACC	CCAACCAAGT	GTTACTACAGG
Horse	GTTACTATCG	TGAAAACATG	TACCGTTACC	CCAACCAAGT	GTTACTACAGG
Mink	GTTACTATCG	TGAAAACATG	TACCGTTACC	CCAACCAAGT	GTTACTACAGG
Bovine	GTTACTATCG	TGAAAACATG	TACCGTTACC	CCAACCAAGT	GTTACTACAGG
Goat	GTTACTATCG	TGAAAACATG	TACCGTTACC	CCAACCAAGT	GTTACTACAGG
Human	GTTACTATCG	TGAAAACATG	TACCGTTACC	CCAACCAAGT	GTTACTACAGG
Elk	GTTACTATCG	TGAAAACATG	TACCGTTACC	CCAACCAAGT	GTTACTACAGG
Hamster	AAAACATGAA	CCGCTACCCT	AACCAAGTGT	ACTACCGGCC	AGTGGACCAG

Dezhou don	CCAGTGGATC	AGTACAGCAA	CCAGAACAGT	TTTGTGCATG	ACCGCGTCAA
Horse	CCGGTAAGTG	AGTACAGCAA	CCAGAAAAAC	TTTGTGCACG	ACTGCGTCAA
Mink	CCGGTGGATC	AGTACAGCAA	CCAGAACAAC	TTTGTGCATG	ACTGCGTCAA
Bovine	CCAGTGGATC	AGTATAGTAA	CCAGAACAAC	TTTGTGCATG	ACTGTGTCAA
Goat	CCAGTGGATC	AGTATAGTAA	CCAGAACAAC	TTTGTGCATG	ACTGTGTCAA
Human	TCAGTGGATC	AGTATAATAA	CCAGAGCACC	TTTGTGCATG	ACTGTGTCAA
Elk	CCAGTGGATC	AGTATAATAA	CCAGAACAAC	TTTGTGCATG	ACTGTGTCAA
Hamster	TACAACAACC	AGAACAACCT	TGTGCACGAC	TGTGTCAATA	TCACGATCAA

Dezhou don	CATCACCGTC	AAGCAGCACA	CAGTGACCAC	GACCACCAAG	GGGGAGAACT
Horse	CATCACGGTC	AAGCAGCACA	CGGTCAACCAC	CACCACCAAG	GGGGAGAACT
Mink	CATCACGGTC	AAGCAGCACA	CGGTCAACCAC	CACCACCAAG	GGGGAGAACT
Bovine	CATCACAGTC	AAGGAACACA	CAGTGACCAC	CACCACCAAG	GGGGAGAACT
Goat	CATCACAGTC	AAGGAACACA	CAGTGACCAC	CACCACCAAG	GGGGAGAACT
Human	CATCACAGTC	AAGCAACACA	CAGT...CAC	CACCACCAAG	GGGGAGAACT
Elk	CATCACAGTC	AAGCAACACA	CAGTGACCAC	CACCACCAAG	GGGGAGAACT
Hamster	GCAGCATACA	GTCACCACCA	CCACCAAGGG	GGAGAACTTC	ACGGAGACCG

Dezhou don	TCACCGAGAC	CGACGTC	AAAG	ATGATAGAGC	GCGTGGT	GGA	ACAGATGTGC
Horse	TCACCGAGAC	CGACGTC	AAAG	ATCATGGAGC	GCGTGGT	GGA	GCAGATGTGC
Mink	TCACCGAGAC	CGACATGA	AG	ATCATGGAGC	GCGTGGT	GGA	GCAGATGTGT
Bovine	TCACCGAAAC	TGACATCA	AG	ATGATGGAGC	GAGTGGT	GGA	GCAAATGTGC
Goat	TCACCGAAAC	TGACATCA	AG	ATAATGGAGC	GAGTGGT	GGA	GCAAATGTGC
Human	TCACCGAAAC	TGACATCA	AG	ATGATGGAGC	GAGTGTG	GGA	GCAAATGTGC
Elk	TCACCGAAAC	TGACATCA	AG	ATGATGGAGC	GAGTGTG	GGA	GCAAATGTGC
Hamster	ACGTCAAGAT	GATGGAGC	GC	GTGGTGGAGC	AGATGTGT	G	CACCCAGTAT

Dezhou don	ATCACCCAGT	ACCAGAA	AGA	GTACGAGGCA	TACGCCAAA	GAGGGCCAG
Horse	ATCACCCAGT	ACCAGAA	AGA	GTACGAGGCT	TTTCAACAAA	GAGGGCGAG
Mink	GTCACCCAGT	ACCAGCA	AGA	GTCCGAGGCT	TACTACCAGA	GGGGGCGAG
Bovine	ATCACCCAGT	ACCAGAG	AGA	ATCCCAGGCT	TATTACCAAC	GAGGGCAAG
Goat	ATCACCCAGT	ACCAGAG	AGA	ATCCCAGGCT	TATTACCAA	GGGGGCAAG
Human	ATCACCCAGT	ACCAGAG	AGA	ATCCGAGGCT	TATTACCAA	GAGGGCAAG
Elk	ATCACCCAGT	ACCAGAG	AGA	ATCCGAGGCT	TATTACCAA	GAGGGCAAG
Hamster	CAGAAGGAGT	CCCAGGC	CTA	CGAGCGGAA	GAAGATCCAG

Dezhou don	TGTGATCCTC	TTCTCCTCCC	CTCCTGTGAT	CCTCCTCATC	TCTTTCTCTC
Horse	CGTGGTCTCTC	TTCTCCTCCC	CGCCTGTG
Mink	CGCCATCCTC	TTCTCGCCCC	CTCCCGTGAT	CCTCCTCATC	TCGCTGCTCA
Bovine	TGTGATCCTC	TTCTCTTCCC	CTCCTGTGAT	CCTCCTCATC	TCTTTCTCTCA
Goat	TGTGATCCTC	TTTTCTTCCC	CTCCTGTGAT	CCTCCTCATC	TCTTTCTCTCA
Human	TGTGATCCTC	TTCTCCTCCC	CTCCTGTGAT	CCTCCTCGTC	TCTTTCTCTCA
Elk	TGTGATCCTC	TTCTCCTCCC	CTCCTGTGAT	CCTCCTCATC	TCTTTCTCTCA
Hamster	CGCGGTGCTC	TTCTCCTCTC	CTCCTGTGAT	CCTCCTCATT	TCCTTCTCTCA

Dezhou don	TTTTCTCAT	AGTGGG	...C	TGA
Horse
Mink	TTCTCCTGAT	AGTGGG	..A	TGA
Bovine	TTTTTCTCAT	AGTAGG	..A	TAGGGGCAAC	CTTCCTGTTT	TCAT
Goat	TTTTTCTCAT	AGTAGG	..A	TAGGGGCAAC	CTTCCTGTTT	TCATTATCTT
Human	TTTTTCTCAT	AGTAGG	..A	TAGGGGCAAC	CTTCCTGTTT	TCATTAT
Elk	TTTTTCTCAT	AGTAGG	..A	TAGGGGCAAC	CTTCCKGTTT	TCATTATCTT
Hamster	TCTTCTGAT	AGTGGGATGA	TGAGGGAAAGC	CTCCCTGCTT	G	TA

Dezhou don
Horse
Mink
Bovine
Goat	CTTAATCTTT	GCCAGGTTGG	GGGAGGGAGT	GTCTACCTGC	AGCCCTGTAG
Human
Elk	CT
Hamster	GTTCTTGTGC

Dezhou don
Horse
Mink
Bovine
Goat	TGGTGGTGC TCATTCTTG CTTCTC
Human
Elk
Hamster

Fig. 2: The comparison of nucleic acid sequence of PrP gene between Dezhou donkey and seven other animals (accession number of GenBank are AF117317, U08952, AY367640, 326330, AY 458651, AY748455, M33958)

Dezhou don	MVKSHVGGWILVLFVATWSDVGLCKKRPKPGG. WNTGGSR	39
Horse TWSVGLCKKRPKPGGgWNTGGSR	24
Mink	MVKSHiGsWILVLFVATWSDiGfCKKRPKPGGgWNTGGSR	40
Bovine	MVKSHiGsWILVLFVAmWSDVGLCKKRPKPGGgWNTGGSR	40
Goat	MVKSHiGsWILVLFVAmWSDVGLCKKRPKPGGgWNTGGSR	40
Human	MVKSHiGsWILVLFVAmWSDVGLCKKRPKPGGgWNTGGSR	40
Elk	MVKSHiGsWILVLFVAmWSDVGLCKKRPKPGGgWNTGGSR	40
Hamster	..manlsyWILaLfvATWtDVGLCKKRPKPGG. WNTGGSR	37
Dezhou don	YPGQGSPGGNRYPPQGGGwGQPHGGwGQPHGGwGQPH	79
Horse	YPGQGSPGGNRYPPQGGGwGQPHGGwGQPHGGwGQPH	64
Mink	YPGQGSPGGNRYPPQGGGwGQPHGGwGQPHGGwGQPH	80
Bovine	YPGQGSPGGNRYPPQGGGwGQPHGGwGQPHGGwGQPH	80
Goat	YPGQGSPGGNRYPPQGGGwGQPHGGwGQPHGGwGQPH	80
Human	YPGQGSPGGNRYPPQGGGwGQPHGGwGQPHGGwGQPH	80
Elk	YPGQGSPGGNRYPPQGGGwGQPHGGwGQPHGGwGQPH	80
Hamster	YPGQGSPGGNRYPPQGGGtWGQPHGGwGQPHGGwGQPH	77
Dezhou_don	GGGwGQPHGGGwGQGGGSHGQWNKPSKPKTNMKHVAGAA	119
Horse	GGGwGQPHGGGwGQGG. SHGQWNKPSKPKTNMKHVAGAA	103
Mink	GGGwGQPHGGGwGQGGGSHGQWgKPSKPKTNMKHVAGAA	120
Bovine	GGGwGQPHGGGwGQGG. SHsQWNKPSKPKTNMKHVAGAA	119
Goat	GGGwGQPHGGGwGQGG. SHsQWNKPSKPKTNMKHVAGAA	119
Human	GGGwGQPHGGGwGQGG. tHsQWNKPSKPKTNMKHVAGAA	119
Elk	GGGwGQPHGGGwGQGG. tHsQWNKPSKPKTNMKHVAGAA	119
Hamster	GGGwGQPHGGG. WGGGGtHnQWNKPSKPKTNMKHVAGAA	116
Dezhou_don	AAGAVVGLGGYMLGSAMSRPLIHF GSDYEDRYRENMYR	159
Horse	AAGAVVGLGGYMLGSAMSRPLIHF GndYEDRYRENMYR	143
Mink	AAGAVVGLGGYMLGSAMSRPLIHF GndYEDRYRENMYR	160
Bovine	AAGAVVGLGGYMLGSAMSRPLIHF GndYEDRYRENmHR	159
Goat	AAGAVVGLGGYMLGSAMSRPLIrF GndYEDRYRENMYR	159
Human	AAGAVVGLGGY1LGSAMSRPLIHF GndcEDRYRENMYR	159
Elk	AAGAVVGLGGYMLGSAMSRPLIHF GndYEDRYRENMYR	159
Hamster	AAGAVVGLGGYMLGSAMSRPmIHF GndwEDRYRENmNR	156
Dezhou_don	YPNQVYYRPVDQYSNQNSFVHDRVNITVKQHTVTTTKGE	199
Horse	YPNQVYYRPVseYSNQknFVHdcVNITVKQHTVTTTKGE	183
Mink	YPNQVYYkPVDQYSNQn1VHdcVNITVKQHTVTTTKGE	200

Bovine	YPNQVYYRFPVDQYSNQnNFVHdcVNITVKeHTVTTTTKGE	199
Goat	YPNQVYYRFPVDQYSNQnNFVHdcVNITVKQHTVTTTTKGE	199
Human	YPNQVYYRsvDQYnNQstFVHdcVNITVKQHTVTTT. KGE	198
Elk	YPNQVYYRFPVDQYnNQnNFVHdcVNITVKQHTVTTTTKGE	199
Hamster	YPNQVYYRFPVDQYnNQnNFVHdcVNITiKQHTVTTTTKGE	196
Dezhou_don	NFTETDVKMIERVVEQMCITQYQKEYEAYQ. RGASVILF	238
Horse	NFTETDVKImERVVEQMCITQYQKEYEafqQ. RGASVvLF	222
Mink	NFTETDmKimERVVEQMCvTQYQqEsEAYyQ. RGASaILF	239
Bovine	NFTETDiKmmERVVEQMCITQYQrEsqAYyQ. RGASVILF	238
Goat	NFTETDiKimEgVVEQMCITQYQrEsqAYyQ. RGASVILF	238
Human	NFTETDiKmmERVVEQMCITQYQrEsEAYyQ. RGASVILF	237
Elk	NFTETDiKmmERVVEQMCITQYQrEsEAYyQ. RGASVILF	238
Hamster	NFTETDVKmmERVVEQMCvTQYQKEsqAYydgRrsSavLF	236
Dezhou_don	SSPPVILLISFLliFLiVIG	256
Horse	SSPPV.....	227
Mink	SpPPVILLISLiLiLIVIG	257
Bovine	SSPPVILLISFLiFLiVIG	256
Goat	SpPPVILLISFLiFLiVIG	256
Human	SSPPVILLvSFLiFLiVIG	255
Elk	SSPPVILLISFLiFLiVIG	256
Hamster	SSPPVILLISFLiFLiVIG	254

Fig. 3: The comparison of amino acid sequence of PrP between Dezhou donkey and seven other animals. (Accession numbers of GenBank are AF117317, U08952, AY367640, AY326330, AY 458651, AY748455 and M33958)

encompassing a N-terminal nonapeptides (PQGGGGWGQ), four repeats of the octapeptides (PHGGGGWGQ) and a C-terminal nonapeptides (PHGGGGWGQ). Amino acids 97~112 is a functional domain controlling the topology structure of the PrP^{sc}; 113~135 is an α -helix; 157~177 is a hydrophilic helix; 234~256AA is a hydrophobic signal region (Fig. 3).

Comparing with horse, bovine, goat, deer, mouse, human being and mink, Dezhou donkey was unique in its PrP amino acids at the following positions: there was no Gly at position 32, nor in zebra (Wopfner *et al.*, 1997) and there was a Gly at position 97; the positions 146, 182, 209, 225 and 229 were Ser, Arg, Iso, Tyr, Ala respectively in Dezhou donkey but Asp, Cys, Met, Ser and Tyr in other animals analyzed. Overall, the PrP of all species analyzed mainly varied in three regions: 90~103, 167~169 and 206~230. It has been reported that amino acid sites 90~130 are corresponding to the species barrier and 108~189 are related to incubation. Within these regions, Gly at position 100 in Dezhou donkey and horse was substituted with Ser in bovine, goat, human and elk. Outside these regions, positions 225 (Tyr) and 229 (Ala) were unique to Dezhou donkey and horse as compared to other species (Ser and Tyr). While the latter species but not the *equus* have been reported to be infected by prion disease, it is not known if these residues in particular are implicated in the species barrier in *equus*.

Recent studies with bovine transgenic mice suggested that amino acid residues 184, 186, 203 and 205 form an epitope that is involved in the control of the species barrier and that amino acids 167, 171, 214 and 218 form an epitope to interact protein X that has been found to participate in the PrP^{sc} formation (Kaneko *et al.*, 1997; Brown *et al.*, 1997). However, analysis of these sites showed no variation of amino acids between Dezhou donkey and the rest of the animals analyzed. As for polymorphism at 136, 154 and 171 amino acid sites that has been reported to be pertinent to

impressionable of scrapie (Eghiaian *et al.*, 2004; Mallucci *et al.*, 2003), in which the genotype A136R154R171 induces resistance to scrapie but A136R154Q171 is intermediately sensitive and V136R154Q171 is sensitive to scrapie, the genotype of donkey is ARQ and may therefore be intermediately sensitive to scrapie. However, how these differences influence the conversion of PrP^c to PrP^{sc} are yet to be illuminated.

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References

- Aguzzi, A. and M. Polymenidou, 2004. Mammalian prion biology: One century of evolving concepts. *Cell*, 116: 313-327.
- Brown, D., K. Qin and J.W. Herms *et al.*, 1997. The cellular prion protein binds copper *in vivo*. *Nature*, 390: 684-697.
- Eghiaian, F., J. Grosclaude and S. Lesceu, 2004. Insight into the PrP^c - PrP^{sc} conversion from the structures of antibody-bound ovine prion scrapie-susceptibility variants. *Proc. Natl. Acad. Sci. USA.*, 101: 10254-10259.
- Hill, A.F., S. Joiner, J. Linehan, M. Desbruslais, P.L. Lantos and J. Collinge, 2000. Species-barrier-independent prion replication in apparently resistant species. *Proc. Natl. Acad. Sci. USA.*, 97: 10248-53.
- Hill, A.F. and J. Collinge, 2002. Species-barrier-independent prion replication in apparently resistant species. *Apmis*, 110: 44-53.
- Kaneko, K., L. Zulianello and M. Scott *et al.*, 1997. Evidence for protein X binding to a discontinuous epitope on the cellular prion protein during scrapie prion propagation. *Proc. Natl. Acad. Sci. USA.*, 94: 10069-10074.
- Liberski, P.P., D.C. Guiroy, E.S. Williams, A. Walis and H. Budka, 2001. Deposition patterns of disease-associated prion protein in captive mule deer brains with chronic wasting disease. *Acta Neuropathology (Berlin)*, 102: 496-500.
- Mallucci, G., A. Dickinson, J. Linehan, P.C. Klohn, S. Brandner and J. Collinge, 2003. Depleting neuronal PrP in prion infection prevents disease and reverses spongiosis. *Science*, 302: 871-874.
- Prusiner, S.B., 1991. Molecular biology of prion diseases. *Science*, 252: 1515-1522.
- Prusiner, S.B., 1997. Prion diseases and the BSE crisis. *Science*, 278: 245-251.
- Wopfner, F., W. Franziska, Weidenhofer G. and R. Schneider, 1997. Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein. *J. Mol. Biol.*, 289: 1163-1178.
- Wang, D., X. Wang, S. Han, B. Tian and Z. Rao, 1998. Cloning and sequencing of the Chinese bovine prion protein (PrP^c) gene. *Wei Sheng Wu Xue Bao*, 38: 417-21.
- Yang, J.M., D.M. Zhao, H.X. Liu, N. Li, Y.X. Hao, Z.Y. Ning and X.H. Qin, 2005. Comparative analysis of the prion protein open reading frame nucleotide sequences in peacock and parakeet. *Virus Genes*, 30: 193-196.
- Zhang, L., N. Li and Q.G. Wang *et al.*, 2002. The genotype of prion gene of Chinese sheep. *Anim. Biotechnol.*, 13: 159-162.