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Application of temperature-gradient gel electrophoresis for detection of prion protein gene polymorphisms in Polish Świniarka sheep

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Abstract. This study presents preliminary data on the polymorphism in the prion protein gene of Świniarka sheep using temperature gradient gel electrophoresis (TGGE). Available data indicate that sensitivity to scrapie is associated with polymorphisms in three codons of prion protein gene: 136, 154, and 171. The TGGE method was used to detect point mutations in these codons responsible for sensitivity or resistance to scrapie. This study revealed presence of an allele encoding valine (V) in codon 136, which is associated with high sensitivity to scrapie and occurred in the form of heterozygous allele together with alanine (AV). The highest variability was observed in codon 171, with presence of arginine (R) and glutamine (Q) in the homozygous (RR or QQ) as well as the heterozygous form (RQ). The results of examination of fifty sheep DNA samples with mutations in codons 136, 154, and 171 demonstrated that TGGE can be used as a simple and rapid method to detect mutations in the PrP gene of sheep. Several samples can be run at the same time, making TGGE ideal for the screening of large numbers of samples.

Key words: Gene polymorphism; sheep; temperature-gradient gel electrophoresis.

Scrapie is a fatal neurodegenerative disease of small ruminants belonging to the group of transmissible spongiform encephalopathies (TSEs). The TSEs are characterized by the accumulation of an abnormal isoform (PrP^{Sc}) of the host-encoded cellular prion protein (PrP^C) in the central nervous system and in lymphoid tissues of affected individuals.^{1,2,4,6,10} The gene encoding PrP is located on chromosome 13q 17–18 and consists of three exons.

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However, only exon 3 encompasses the entire open reading frame of the PrP gene. Its translation produces the protein consisting of 256 amino acids. The remaining exons contain nontranslating sequences (Fig. 1).

Early epidemiological observations of sheep scrapie indicate that sensitivity or resistance to scrapie is associated with variability at codons 136, 154, and 171 of the PrP gene. Of the five most frequent scrapie alleles (ARR, AHQ, ARH, ARQ, and VRQ), ARR is thought to be associated with resistance and VRQ with susceptibility to the disease.^{3,5,6} Susceptibility to scrapie is mainly associated with presence of valine (V) at codon 136 and glutamine (Q) at codon 171; homozygotes (VV₁₃₆ and QQ₁₇₁) are

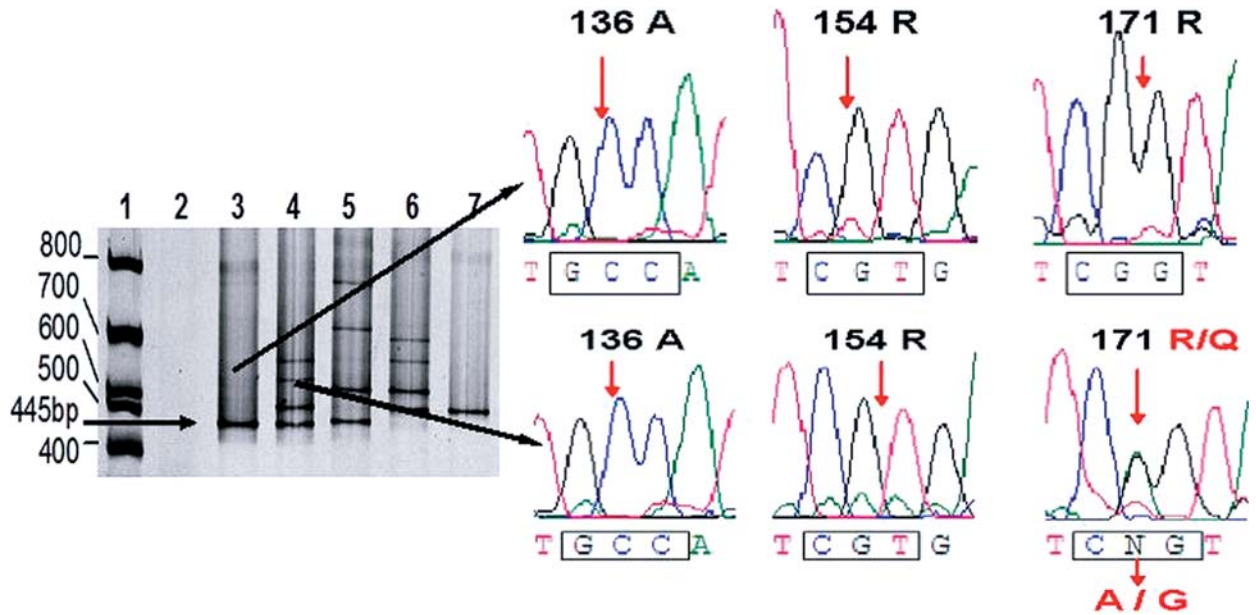


Figure 2. Electrophoresis of PCR products of PrP gene of control samples from Czarnogłówka breed and fluorograms of sequencing of codons 136, 154, 171 (genotype ARR/ARR and ARR/ARQ). Temperature gradient (40°C→60°C) is same as the direction of the electric field. Lane 3 (genotype ARR/ARR), no mutation in both alleles yields a single band with the highest melting temperature. Transition A→G at codon 171 in one allele (genotype ARR/ARQ; lane 4) changes the melting temperature of the DNA and yields four bands (2 heteroduplex and 2 homoduplex). Lane 1, 100-bp ladder; lane 2, negative control (no DNA added); lane 3, homozygote ARR/ARR (wild type 445bp); lane 4, heterozygote ARR/ARQ; lane 5, heterozygote ARR/AHQ; lane 6, heterozygote ARQ/AHQ; lane 7, homozygote ARQ/ARQ.

continued at 250V for 40 min. After the electrophoresis, the gels were silver-stained using 0.3% silver nitrate to visualize the bands. Electrophoresis of the amplification products was performed on an 8% polyacrylamide gel (acrylamide/N,N-ethylenebisacrylamide 37.5 : 1)^e containing 8 M urea and 2% glycerol. Electrophoresis buffer was 1× TAE pH 8.3.^f About 3 ng of DNA was loaded into each slot of the 8% polyacrylamide gel. Gel electrophoresis mobility

and stability of PCR amplified PrP gene fragments were calculated using online software (<http://www.biophys.uniduesseldorf.de/local/POLAND/poland.html>). As controls, DNA samples from Czarnogłówka sheep with known genotypes were used. These five different genotypes (ARR/ARR, ARR/ARQ, ARR/AHQ, ARQ/AHQ, and ARQ/ARQ) confirmed by sequencing served as the reference genotypes (Fig. 2).

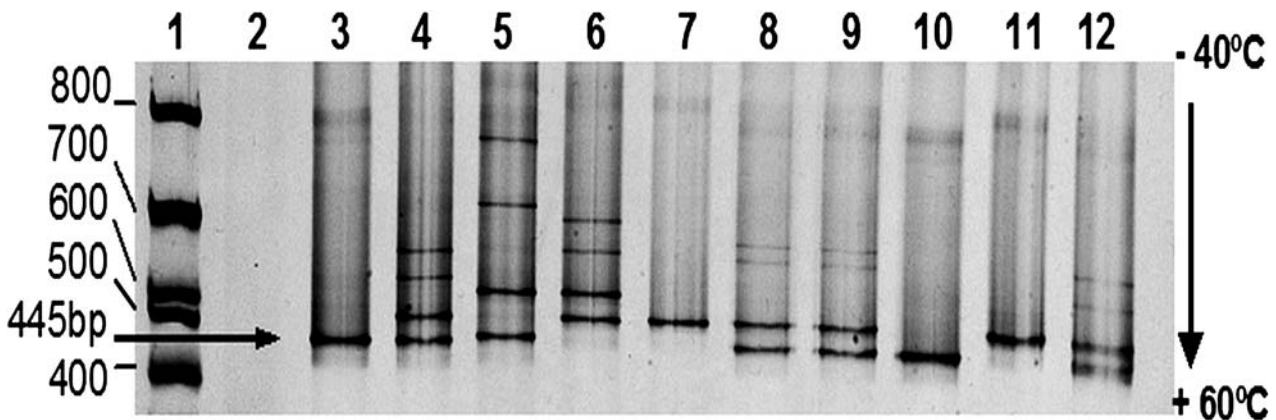


Figure 3. Electrophoresis of PCR products of sheep PrP gene. Temperature gradient (40°C→60°C) is same as the direction of the electric field. Lane 1, 100-bp ladder; lane 2, negative control (no DNA added); lanes 3, 4, 5, 6, 7, positive controls of five different genotypes from Czarnogłówka breed of sheep (line 3, ARR/ARR genotype; lane 4, ARR/ARQ genotype; lane 5, ARR/AHQ genotype; lane 6, ARQ/AHQ genotype; line 7, ARQ/ARQ genotype). Lanes 8, 9, 10, 11, 12, samples from Swiniarka breed of sheep (lane 10, ARR/ARR genotype; lane 11, ARQ/ARQ genotype; line 12, ARR/ARQ genotype). Band patterns (lanes 8 and 9) different from control samples reflects existence ARR/VRQ genotype (confirmed by sequencing). The slightly different migration distance between the hetero- and homozygous samples reflects differences in nucleotide sequence.

Table 2. Mutations in PrP codons 136,154,171 and genotypes of Świniarka sheep.

Mutations (amino acids)			Genotype	Sheep number (%)
codon 136	codon 154	codon 171		
—	—	—	AA ₁₃₆ RR ₁₅₄ RR ₁₇₁	22 (44)
—	—	CGG→CAG (Arg→Gln)	AA ₁₃₆ RR ₁₅₄ RQ ₁₇₁	15 (30)
GCC→GTC (Ala→Val)	—	CGG→CAG (Arg→Gln)	AV ₁₃₆ RR ₁₅₄ RQ ₁₇₁	9 (18)
—	CGT→CAT (Arg→His)	CGG→CAG (Arg→Gln)	AA ₁₃₆ RH ₁₅₄ RQ ₁₇₁	2 (4)
—	—	CGG→CAG (Arg→Gln)*	AA ₁₃₆ RR ₁₅₄ QQ ₁₇₁	2 (4)

* Transition G→A was found in both alleles as compared to the wild sequence of PrP gene.

To evaluate the reliability of TGGE, analysis of several randomly selected samples of Świniarka sheep was performed at the DNA Sequencing and Oligonucleotide Synthesis Facility of the Institute of Biochemistry and Biophysics in Warsaw, Poland. Furthermore, sequencing of DNA samples with different TGGE band patterns than 5 control samples (Fig. 3; lanes 8, 9) was performed to verify their genotype. In every case, the results of PrP genotyping obtained with the TGGE method were identical to those obtained by sequencing.

DNA fragments with the same sequence are easily discernible because they migrate during electrophoresis over the same distance from the starting point. A single band in Fig. 3 (lane 11; homozygote ARQ/ARQ) reflects the existence of a point mutation in both alleles of the same codon. A G→A transition at codon 171 decreases the melting temperature and gel mobility of the DNA fragment in comparison with the wild-type fragment (lanes 3, 10; homozygote ARR/ARR). The presence of four bands in one lane reflects the existence of a point mutation on one allele, which creates additional heteroduplex bands visible above the homoduplex bands (Fig. 3). The slower migrating bands are hybrids of wild-type and mutant fragments containing mismatched base pairs. The distance between the heteroduplex bands as well as between heteroduplex and homoduplex bands depends on position and number of point mutations (Fig. 3; Table 2).

Of the 50 sheep genotyped, 41 were homozygous AA₁₃₆ and the others were heterozygous for AV₁₃₆ (Table 2). Variation at codon 154 (RH₁₅₄) was found only in two cases. The remaining 48 sheep were homozygous for RR₁₅₄. The greatest genetic variation was observed at codon 171, and two of the three possible alleles were detected: of the 50 sheep, 22 were homozygous for RR₁₇₁ (44%), two were homozygous for QQ₁₇₁ (4%), and 26 were heterozygous for RQ₁₇₁ (52%). No H alleles were found at codon 171.

This is the first study of the frequency of PrP gene polymorphisms in the Polish population of Świniarka sheep, a very primitive breed of sheep. This study confirmed that using the TGGE method it is possible to detect more than one point mutation within the PrP gene.^{10,14,17} Band patterns of DNA are characteristic for given genotypes of PrP gene and depend on the number and location of particular mutations. The TGGE method is a reproducible and sensitive technique and can be useful in routine screening of sheep flocks. In comparison with the widely used single-strand conformation polymorphism (SSCP) analysis, the interpretation of TGGE gels is

facilitated by the appearance of heteroduplex bands. In contrast to TGGE, the interpretation of SSCP gels is sometimes difficult due to poor resolution of the bands, which depends greatly on the length of the fragments and the use of different running conditions.⁹

The results of this study indicate that the dominant alleles in Świniarka sheep are ARR and ARQ. These PrP alleles are also frequently present in Romanov sheep,⁴ Suffolk sheep^{8,11} and Valachian sheep.¹⁴ The ARR allele is associated with high resistance to scrapie, so high frequency of this allele in some breeds of sheep can be exploited to increase the genetic resistance to scrapie of sheep flocks through introduction of selective breeding programs. Scrapie has not yet been reported in Poland, and the introduction of a scrapie-resistance breeding program could diminish the risk of its outbreak. Analysis of sheep belonging to the Świniarka breed also revealed the presence of the VRQ allele, which is associated with high susceptibility to scrapie. This allele is very rare or absent in some sheep breeds, like Berichone du Cher,⁵ Masses,¹⁰ and Suffolk,^{8,11} but highly frequent in Icelandic sheep (about 80%) and in Norwegian Rygja sheep (48%).² There is no data on PrP genotypes of Świniarka sheep in currently available literature, probably because this breed of sheep can be found only in the southern and southeastern part of Europe.

In conclusion, the occurrence of the VRQ allele in Świniarka sheep indicates that this breed cannot be considered as scrapie-resistant. A more extensive screening program is needed to provide more comprehensive data. Świniarka sheep, being well adapted to local environmental conditions, can constitute the basis for future breeding programs as a donor of scrapie-resistant genotype.

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Sources and manufacturers

- DNA Isolation Kit for Mammalian Blood, Roche Diagnostics, GmbH, Mannheim, Germany.
- Fermentas AB, Vilnius, Lithuania.
- GeneRuler™ 100bp DNA Ladder Plus, Fermentas AB, Vilnius, Lithuania.
- Biometra, GmbH, Göttingen, Germany.
- Acrylamid-Bis ready-to-use solution 40%, Merck KGaA, Darmstadt, Germany.
- TAE buffer 10x pH 8.3 Tris-acetate-EDTA buffer, Merck KGaA, Darmstadt, Germany.

References

1. Begara-Mcgorum I, Clark AM, Martin S, Jeffery M: 2000, Prevalence of vacuolar lesions consistent with scrapie in the brain of healthy cull sheep of the Shetland Islands. *Vet Rec* 147:439–441.
2. Benestad L, Sarradin P, Thu B, et al.: 2003, Cases of scrapie with unusual features in Norway and designation of a new type, Nor98. *Vet Rec* 153:202–208.
3. Dawson M, Warner R, Nolan A, et al.: 2003, Complex PrP genotypes identified by the National Scrapie Plan. *Vet Rec* 152:754–755.
4. Elsen JM, Amigues Y, Schelcher F, et al.: 1999, Genetic susceptibility and transmission factors in scrapie: detailed analysis of an epidemic in a closed flock of Romanov. *Arch Virol* 144:431–445.
5. François D, Elsen JM, Barillet F, et al.: 2003, Breeding sheep for scrapie resistance. *In: Breeding programmes for improving the quality and safety of products. New traits, tools, rules and organization? (Options Méditerranéennes: Série A)*, ed. Gabiña D, Sanna S, pp. 29–35. CIHEAM-IAMZ, Zaragoza, Spain.
6. Gombojav A, Ishiguro N, Horiuchi M, et al.: 2003, Amino acid polymorphisms of PrP gene in Mongolian sheep. *J Vet Med Sci* 65:75–81.
7. Horn D, Robinson PN, Böddrich A, et al.: 1996, Three novel mutations of the NF1 gene detected by temperature gradient gel electrophoresis of exons 5 and 8. *Electrophoresis* 17:1559–1563.
8. Hunter N, Moore L, Hosie BD, et al.: 1997, Association between natural scrapie and PrP genotype in a flock of Suffolk sheep in Scotland. *Vet Rec* 140:59–63.
9. Mashal RD, Sklar J: 1996, Practical methods of mutation detection. *Curr Opin Genet Develop* 6:275–280.
10. Multinelli F, Aufero GM, Pozzato N, et al.: 2003, Eradication of scrapie in a Massese sheep flock by PrP allele selection. *Vet Rec* 152:60.
11. O'Doherty E, Aherne M, Ennis S, et al.: 2000, Detection of polymorphisms in the prion protein gene in a population of Irish Suffolk sheep. *Vet Rec* 146:335–338.
12. Office for Official Publications of the European Communities, Commission Regulation (EC) No 999/2001 of 22 May. *Official Journal L* 147:1–61.
13. Schlotterer Ch: 1995, Temperature-gradient gel electrophoresis as a screening tool for polymorphisms in multigene families. *Electrophoresis* 16:722–728.
14. Tracikova L, Hanusovska E, Novak M, et al.: 2003, The PrP genotype of sheep of the improved Valachian breed. *Folia Microbiol (Prague)* 48:269–276.
15. Thorgeirsdottir S, Georgsson G, Reynisson E, et al.: 2002, Search for healthy carriers of scrapie: an assessment of subclinical infection of sheep in an Iceland scrapie flock by three diagnostic methods and correlation with PrP genotypes. *Arch Virol* 147:709–722.
16. Wartell RM, Hosseini S, Powell S, Zhu J: 1998, Detecting single base substitutions, mismatches and bulges in DNA by temperature gradient gel electrophoresis and related methods. *J Chromatography A* 806:169–185.