

Title: Rapid testing leads to the underestimation of the scrapie prevalence in an affected sheep and goat flock

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RECKLING, K., ... FRIES, R.

Document type: Postprint

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Citation: RECKZEH, C., HOFFMANN, C., BUSCHMANN, A., BUDA, S., BUDRAS, K., RECKLING, K.,

... FRIES, R. (2007). Rapid testing leads to the underestimation of the scrapie prevalence in an

affected sheep and goat flock. Veterinary Microbiology, 123(4), 320–327.

https://doi.org/10.1016/j.vetmic.2007.04.009

1	Rapid testing leads to the underestimation of the scrapie prevalence in an affected sheep
2	and goat flock
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#### **Abstract**

To obtain a more detailed understanding of the prevalence of classical scrapie
infections in a heavily affected German sheep flock (composed of 603 sheep and 6 goats), we
analysed 169 sheep and 6 goats that carried the genotypes susceptible to the disease and that
were therefore culled following discovery of the index case. The initial tests were performed
using the Biorad TeSeE ELISA and reactive results were verified by official confirmatory
methods (OIE-immunoblot and/or immunohistochemistry (IHC)) to demonstrate the
deposition of scrapie-associated PrPSc in the brain stem (obex). This approach led to the
discovery of 40 additional subclinically scrapie-infected sheep. Furthermore, peripheral
lymphatic tissue samples of the 129 sheep and six goats with a negative CNS result were
examined by IHC in order to identify any preclinical infections which had not already spread
to the central nervous system (CNS). Using this approach we found 13 additional sheep with
PrP <sup>Sc</sup> depositions in the gut-associated lymph nodes (GALT) as well as in the enteric nervous
system. Moreover, in most of these cases PrPSc was also deposited in the spleen and in the
retropharyngeal and superficial cervical lymph nodes. Taken together, these results show a
30.3% infection prevalence in this scrapie-affected flock. Almost 7.4% of the infected animals
harboured PrPSc exclusively in the peripheral lymphatic tissue and were therefore missed by
the currently used testing strategy.

Keywords: preclinical scrapie, immunohistochemistry, lymphoid tissue

#### 1. Introduction

Scrapie belongs to the transmissible spongiform encephalopathies (TSE) of sheep and goats and is characterised by the accumulation of an abnormally folded isoform (PrP<sup>Sc</sup>) of the cellular prion protein (PrP<sup>C</sup>). Incubation times vary between a few months and several years depending on the infectious dose and route, the particular scrapie strain and the genotype and

48	age of the recipient animal (summarised in Detwiler and Baylis, 2003) According to the prion
49	hypothesis (Prusiner et al., 1982), abnormally folded PrPSc represents the causative infectious
50	agent itself, the 'prion'. However, the existence of different strains remains as yet
51	unexplained. PrPSc is partially protease resistant, insoluble in detergents (as a consequence of
52	an increased $\beta$ -sheet content which increases its hydrophobicity) and forms scrapie-associated
53	fibrils (Diringer et al., 1983; Hope et al., 1986; Oesch et al., 1985). These criteria form the
54	basis for the demonstration of PrP <sup>Sc</sup> as a diagnostic marker.
55	Amino acid polymorphisms at positions 136, 154 and 171 of the prion protein
56	determine the susceptibility of sheep to classical scrapie. Alleles on the PrP gene encoding
57	alanine/arginine/arginine (PrPARR) are associated with the lowest level of susceptibility,
58	alleles encoding valine/arginine/glutamine (PrPVRQ) with the highest susceptibility,
59	particularly if present in homozygous animals (Goldmann et al., 1990; Hunter, 1996; Hunter
60	et al., 1997). The PrP genotype of sheep has a strong effect on the scrapie pathogenesis in
61	sheep. While no, or only minor, traces of PrPSc depositions can be found in the peripheral
62	lymphatic tissues (e.g., spleen, tonsils, retropharyngeal lymph nodes, gut-associated lymph
63	nodes) of sheep carrying at least one PrPARR allele, PrPVRQ/VRQ and PrPARQ/ARQ carriers can
64	show PrPSc accumulations in these lymphatic tissues as early as a few months after the
65	infection (van Keulen et al., 1996 & 2002; Andreoletti et al., 2000; Jeffrey et al., 2002; Ersdal
66	et al., 2005). Preclinical accumulation of PrPSc in lymphoreticular tissues of sheep therefore
67	provides an opportunity for an in-vivo diagnosis based on biopsy samples of the tonsils
68	(Schreuder et al., 1998), the third eyelid (O'Rourke et al., 2002) or the rectum (Gonzalez et
69	al., 2006; Espenes et al., 2006).
70	The gastrointestinal tract, in particular the gut-associated lymphoid tissues (GALT), is
71	regarded as the major route of entry for natural scrapie (Hadlow et al., 1982; Heggebo et al.,
72	2000). The earliest accumulation of PrPSc can be found in the ileal Peyer's patches (IPP) and
73	in the draining mesenteric lymph nodes (Andreoletti et al., 2000). From there, prions may

74	spread to other lymphatic tissues, which are not part of the GALT, resulting in a widespread
75	distribution of disease-specific PrP deposition in the lymphoreticular system at relatively
76	early stages of infection (Andreoletti et al., 2000; Heggebo et al., 2002; van Keulen et al.,
77	2002). Neuroinvasion appears to start at the enteric nervous system (ENS) followed by a
78	centripetal and retrograde spread via sympathetic and parasympathetic efferent fibres of the
79	autonomic nervous system to the spinal cord and to the medulla oblongata (van Keulen et al.,
80	2000).
81	The aim of the study was to analyse the prevalence of preclinical scrapie in a cohort of
82	susceptible sheep that were compulsorily culled after the detection of scrapie in the flock. For
83	this purpose LRS and the peripheral nervous system samples were examined with particular
84	attention paid to animals which were tested negative in the initial rapid test on the brain stem.
85	
86	2. Materials and Methods
87	2.1. Sheep and goats
88	All 169 German Blackheaded Mutton-Merino crossbreed sheep and 6 fawn goats
89	belonged to a single flock (n=603) in Saxony-Anhalt in which an outbreak of classical scrapie
90	occurred in December 2003.
91	
92	2.2. TSE rapid testing
93	Brain stem samples from the obex region were examined using the Biorad TeSeE
94	ELISA rapid test (Biorad, Munich, Germany) following the manufacturer's instructions.
95	
96	2.3. Genotyping
97	Sheep were genotyped as described previously (Lühken et al., 2004).
98	
99	2.4. Histopathology and immunohistochemistry

100	With some modifications, tissue samples were processed as described previously
101	(Hardt et al., 2000). In short, all tissue samples were fixed in 4% buffered formalin, treated
102	for one hour with 98% formic acid, rinsed for 40 minutes in tap water, embedded in paraffin,
103	sectioned at 3-4 $\mu m$ and stained with haematoxylin and eosin.
104	The avidin-biotin-complex (ABC) method was used for PrP-IHC. This involved the
105	paraffin wax tissue sections being mounted on superfrost plus slides (Menzel-Gläser,
106	Braunschweig, Germany) and rehydrated. The subsequent pretreatment included incubation
107	of the slides in 98% formic acid for 15 minutes, a 5-minute rinse in tap water, inhibition of the
108	endogenous peroxidase activity with $3\%~H_2O_2$ (Merck, Darmstadt, Germany) in methanol for
109	30 minutes, followed by 15 minutes digestion with proteinase K ( $4\mu g/ml$ , Boehringer,
110	Mannheim, Germany) at 37 °C. The primary monoclonal antibody (mab) L 42, which binds to
111	an epitope of amino acid 145-163 of ovine PrP, was applied at a dilution of 1:250 (Harmeyer
112	et al., 1998) in Tris-buffered saline (TBS) containing 10 % goat serum and incubated
113	overnight at 4°C. Negative control sections were treated with a monoclonal antibody against
114	GP <sub>5</sub> of the Porcine Respiratory and Reproductive Syndrome virus (Weiland et al., 1999). As a
115	secondary (link) antibody, biotinylated goat anti-mouse antiserum (Vector Laboratories,
116	Burlingame, CA, USA) was incubated on the sections in a 1:200 dilution for 30 minutes at
117	room temperature. Immunodetection was amplified using Vector ABC-elite avidin-
118	horseradish peroxidase/biotin complex (Vector laboratories; Peterborough, UK). The slides
119	were finally developed in DAB (diaminobenzidintetrahydrochloride) (Fluka Feinchemikalien,
120	Neu Ulm, Germany) and counterstained with Mayer's haematoxylin. All sections were
121	examined using light microscopy.
122	All positive results were verified using mab 2G11 binding to the epitope 146-182 of
123	ovine PrP (Institut Pourquier, Montpellier, France) diluted 1:250 in TBS with 10 % goat
124	serum.

125

#### 3. Results

All 175 animals examined in this study came from a flock of 603 German Blackhead
Mutton-Merino crossbreed sheep and six fawn goats which was kept in Saxony-Anhalt in
Germany. The outbreak of classical scrapie was first recognised through TSE rapid testing in
two fallen stock sheep. After the flock was quarantined and monitored, 10 more clinical cases
of scrapie were identified. According to the EU 999/2001 regulation, all animals in the flock
were genotyped with regard to codons 136, 154 and 171 of the sheep PrP gene and all
genetically highly TSE-susceptible animals (169 sheep and six goats) were killed. The
necropsy was carried out immediately after the death of the animals and a wide range of
tissues of the central and peripheral nervous system as well as of the lymphoreticular system
(LRS) was taken. Obex samples from these animals were subsequently TSE rapid tested using
the Biorad TeSeE assay. Reactive results were obtained in another 40 of the 169 sheep and
the TSE diagnosis in these cases was confirmed by using the OIE- approved methods, SAF-
immunoblot and/or IHC. Table 1 shows an overview of relevant data of all sheep investigated
All 135 animals with a negative rapid test result on the obex region were more closely
examined by IHC to detect PrPSc accumulations in the peripheral nervous and lymphoreticular
system using a hierarchical approach, as illustrated in Figure 1. In a first step the five most
likely sites of entry for the agent were analysed. In cases where there was a total absence of
PrP <sup>Sc</sup> depositions in the samples, the animals were considered to be completely negative and
no further studies were undertaken. However, if there was a positive result in one of the five
tissues, the investigation was extended to several parts of the peripheral nervous system as
well as to the lymphoreticular system.
PrPSc was not found to be present in any of the initially investigated tissues of the six
goats. However, $PrP^{Sc}$ was found to be present in the lymphoreticular system of 13 of the 129
investigated sheep. Table 2 summarises the results for all animals with a clear PrPSc-
immunolabelling.

PrP<sup>Sc</sup> was discovered in the lymphoreticular system in most animals and in particular in the head-associated lymph nodes (i.e., tonsils, retropharyngeal or mandibular lymph nodes). Interestingly, two sheep showed a strong staining reaction in these non-GALT lymphoid tissues (Figure 2C), but not in the GALT. Additionally, a PrP<sup>Sc</sup> accumulation in the lymphoid follicles of the third eyelid was only apparent in the two sheep with the most widespread PrP<sup>Sc</sup> distribution in the lymphoreticular system. As the rectum samples examined did not totally comply with the proposed diagnostic requirements (i.e. too low a number of follicles), the absence of PrP<sup>Sc</sup> staining in any of the preclinically affected sheep at this site should be interpreted cautiously.

Only a sparse immunolabelling was found in the peripheral nervous system. While 11 sheep showed a strong PrP<sup>Sc</sup> immunolabelling in the enteric nervous system of the distal ileum (Fig. 2A), only six of these sheep showed a detectable amount of PrP<sup>Sc</sup> in the coeliac and mesenteric ganglion complex (Fig. 2B). Three of these sheep showed an additional PrP-immunostaining in the ENS of the rectum. However, no PrP<sup>Sc</sup> accumulation was demonstrated in the Ganglion cervicale craniale, Ganglion stellatum and the Vagus nerve.

The preclinical scrapie-infected sheep carried either the PrP<sup>ARQ/ARQ</sup> (9 cases) or the PrP<sup>ARQ/AHQ</sup> (4 cases) genotype. There was no obvious correlation between the number of positive tissues and the age of the animal or between the quantity of accumulated PrP<sup>Sc</sup> and the genotype of the sheep. Although a widespread distribution of PrP<sup>Sc</sup> was detected in most animals, none of the 13 sheep was positive in every single sample.

#### 4. Discussion

In this study 169 sheep carrying the susceptible genotypes and six goats out of a flock, which underwent a selective culling to eradicate scrapie, were analysed. This investigation revealed that 30% of these animals were preclinically scrapie-infected and that 7.4% of them were PrP<sup>Sc</sup>-positive only at the peripheral sites but not in the brain stem, which is routinely

tested. This prevalence is within the range of 5% to 30.3% previously reported in similar studies (Jeffrey et al., 2002; Thorgeirsdottir et al. 2002; Ersdal et al. 2003; Vascellari et al. 2005; Gonzalez et al., 2006). The classical scrapie prevalence in a flock depends on different factors: a) length of time period after the introduction of the infection into the flock; b) husbandry measures (e.g. postpartal removal of the placentas as the most infectious tissues, separation of lambing ewes, frequency of pasture changes, general hygiene measures); c) genetic status (in terms of scrapie susceptibility) of the animals in the flock; d) differences in the scrapie strains (Andreoletti et al., 2001; Jeffrey et al., 2001).

The origin of the scrapie infection in this flock, in which the oldest scrapie-infected sheep was born in 1998, is unknown. However, the large number of scrapie-infected sheep may reflect the fact that this flock has had a long scrapie history which was possibly started by the introduction of an infected animal from the outside. In this regard it is of particular interest that most of the sheep with a positive rapid test result are two years or younger, which corresponds to the reported changes in the age-genotype profile in flocks which have been infected for several years (Baylis et al., 2000). The last acquisition of ewes took place in 1994, while rams were restocked from other flocks for many years. Interestingly, scrapie was diagnosed in a neighbourhood flock in 1979/1980, but the eradication measures taken at this time were not fully recorded. Little is known about the husbandry measures which were applied in this particular scrapie-affected flock.

In our study, all affected animals were either PrP<sup>ARQ/ARQ</sup> or PrP<sup>AHQ/ARQ</sup> carriers and no differences in the distribution of PrP<sup>Sc</sup> were found between these groups. These results are in contrast to previous studies carried out in Scotland (Jeffrey et al., 2002) and Iceland (Thorgeirsdottir et al., 2002) in which these genotypes seemed to be more resistant to a scrapie infection. However, the results reported here are in accordance with the results from an outbreak in Italy (Vascellari et al., 2005). The most probable explanation for this difference is that scrapie strains with a bias towards different genotypes had been going round the flocks

in question. All seven PrP<sup>ARR</sup> heterozygous sheep were negative in our study. Although most of the sheep were at least four years or older (5/6), the low number of animals with this semi-resistant genotype does not allow us to exclude that the scrapie agent targets genotypes with an age-dependent lower prevalence as shown in a previous study (Gonzalez et al., 2006).

All preclinically scrapie-infected sheep showed a marked and widespread distribution

of PrP<sup>Sc</sup> throughout the lymphoreticular system and in some parts of the peripheral nervous system. However, none of the animals was positive in every tissue sampled. In accordance with several studies most animals (n=11) showed an accumulation of PrP<sup>Sc</sup> in the Peyer's patches of the distal ileum and/or the draining lymph nodes, suggesting the intestine as the port of entry for the agent (Heggebo et al., 2000; van Keulen et al., 2002). Additionally, six of the sheep showed a clear staining reaction in the enteric nervous system as well as in the CMGC, indicating the subsequent spread to the CNS (van Keulen et al., 2000). In contrast, in two sheep (5, 11) no detectable amount of PrP<sup>Sc</sup> was found in the intestine and the mesenteric lymph nodes. One possible explanation could be a direct neuroinvasion (Jeffrey et al., 2002). However, none of the neuronal tissues examined contained PrP<sup>Sc</sup> depositions, whereas a strong immunostaining was seen in the tonsil and in the Ln. retropharyngealis lateralis. Therefore the oronasal cavity may have served as a port of entry in these two sheep.

The current EU strategy for the active and passive surveillance of sheep TSEs is based on the testing of CNS samples. The results of the present study reveal that a better estimation of the "real" prevalence of scrapie-infected sheep can be obtained when LRS samples are examined as well.

However, it must be noted that a diagnostic examination of the LRS alone (as suggested by Gonzalez et al., 2006) would lead to studies missing the large number of atypical scrapie cases which lack a PrP<sup>Sc</sup> dissemination in the LRS (Gavier-Widen et al., 2005). The same applies for sheep carrying the PrP<sup>ARR</sup> haplotype. Therefore a combination of

229	CNS and LRS testing is necessary to detect all clinical and preclinically scrapie-infected
230	animals in affected flocks.
231	
232	5. Conclusion
233	In conclusion, this study shows that a negative result of the rapid test on brain stem
234	samples does not exclude the presence of PrPSc in peripheral tissues. In consequence, the
235	actual prevalence of preclinical scrapie is significantly underestimated when it is only based
236	on studies using brain stem samples.
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 Table 1: Numbers, genotypes and age of all culled sheep and goats

	Total number	Number of sheep with genotypes						Age (	Age (in years)*				
		ARQ/ARR	ARQ/ARQ	ARQ/AHQ	AHQ/ARQ	VRQ/ARQ	VRQ/ARR	0-1	1-2	2-3	3-4	> 4	
Obex positive	40		38		2	7		13	16	2	3	0	
Obex negative													
sheep	129	4	105	13		4	3	23	24	39	11	9	
goats	6							unkno	own				

<sup>\* 29</sup> sheep of unknown age

**Table 2**: Immunohistochemical results relative to the distribution of PrP<sup>Sc</sup> in the different tissues of 13 sheep

Sheej	) IPP	Lnn. ileo- colici	RPLN	Tonsil	3rd eyelid	Spleen	Ln. cerv. superf.		ENS Rectum		Ggl. cerv. cran.	Ggl. stella- tum	N. vagus	Tongue	PrP genotype	Age (in years)
1	+	-	+'*	+	-	+	+	-	-	-	-	-	-	-	ARQ/ARQ	>4
2	-	+	+'*	+	-	+	+	-	-		-	-	-	-	ARQ/ARQ	>4
3	+	+	+	+	+	+	+	+	-	+	-	-	-	-	ARQ/ARQ	2-3
4	+	-	-	-	-	-	-	-		-	-	-	-	-	ARQ/ARQ	2-3
5	-	-	+	+	-	+	+	- >	-	-	-	-	-	-	ARQ/ARQ	2-3
6	+	+	-	-	-	+	-	t	+	+	-	-	-	-	ARQ/ARQ	1-2
7	+	+	+	+	+	+	+	+	+	+	-	-	-	-	ARQ/ARQ	1-2
8	-	+	+	+	-	-	+	-	-	-	-	-	-	-	ARQ/ARQ	0-1
9	-	+	+	+	-	+	+	-	-	-	-	-	-	-	ARQ/ARQ	0-1
10	+	+	+	+	-	+	+	+	-	+	-	-	-	-	AHQ/ARQ	1-2
11	-	-	+	+	-		-	-	-	-	-	-	-	-	AHQ/ARQ	1-2

12 +	-	-	-	-	-	-	-	+	+	+	-	-		-	AHQ/ARQ 0-1
13 +		-	-	-	-	-	-	+	-	+	-	-	-	-	AHQ/ARQ
Total 8	3	7	9	9	2	8	8	6	3	6	0	0	0	0	

\* Lnn. mandibulares

IPP Peyer's patches of the distal ileum

RPLN Retropharyngeal lymph node

ENS Enteric nervous system CMGC Coeliac and mesenteric ganglion complex

#### Figure legends

#### Figure 1:

Hierarchical approach in the sample selection from culled sheep and goats that underwent a further immunohistochemical examination

#### Figure 2:

Distinct PrPSc immunolabelling in different tissues of sheep