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Incidence of Infection in Prnp ARR/ARR Sheep following Experimental Inoculation with or Natural Exposure to Classical Scrapie

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Abstract

The prion protein gene (Prnp) is highly influential in determining risk and susceptibility of sheep exposed to classical scrapie. Sheep homozygous for alanine at codon 136 and arginine at codons 154 and 171 (ARR/ARR) of the Prnp gene are historically considered to be highly resistant to classical scrapie, although they form a significant fraction of cases of atypical scrapie. To date, experimental transmission of prions to ARR/ARR sheep has only been achieved with the BSE agent and mostly by the intracerebral route. We summarise here the results of six separate studies, in which 95 sheep of the ARR/ARR genotype were naturally exposed to (n = 18) or experimentally challenged with (n = 77) natural or experimental sources of classical scrapie by the oral, intra-intestinal, subcutaneous or intracerebral routes and allowed to survive for periods of up to 94 months post-infection. Only the intracerebral route resulted in disease and/or amplification of disease associated PrP (PrPΔ), and only in two of 19 sheep that survived for longer than 36 months. Discriminatory immunohistochemistry and Western blot confirmed the scrapie, non-BSE signature of PrPΔ in those two sheep. However, the neuropathological phenotype was different from any other scrapie (classical or atypical) or BSE source previously reported in sheep of any Prnp genotype. These studies confirm the widely held view that ARR/ARR sheep are highly resistant to classical scrapie infection, at least within their commercial lifespan. Moreover, within the constraints of the present studies (only two infected sheep), these results do not support the suggestion that atypical scrapie or BSE are generated by adaptation or mutation of classical scrapie in sheep of resistant ARR/ARR genotype.

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Introduction

The transmissible spongiform encephalopathies (TSEs) are characterised by the accumulation of abnormal forms of a host-coded, cell membrane sialoglycoprotein called prion protein (PrP). Scrapie, or classical scrapie, of sheep and goats is the archetypal TSE and has been recognised as contagious outbreaks of disease for several centuries. More recently, a novel, apparently non-contagious or sporadic form of sheep TSE, originally called Nor 98 [1] and now more commonly referred to as atypical scrapie, has been recognised in Europe and elsewhere.

The prion protein gene (Prnp) controls susceptibility to both atypical and classical scrapie [2]. Sheep bearing alanine (A) or valine (V) at codon 136 and glutamine (Q) at codon 171 of PrP are susceptible to classical scrapie. In contrast, classical scrapie is rarely reported in sheep homo- or hetero-zygous for the allele that bears A at codon 136 and arginine (R) at codons 154 and 171. In separate UK studies, sheep scrapie was not identified in aged ARR/ARR or ARQ/ARR sheep in flocks or geographical regions with endemic scrapie [3,4,5,6]. Other epidemiological studies also show that scrapie is very rare in ARQ/ARR sheep and absent from ARR/ARR sheep in the UK [7]. However, single cases of natural scrapie infection in ARR/ARR sheep have been reported from Germany, France [8] and possibly also Japan [9].

In contrast, atypical scrapie is relatively common in sheep that are homozygous or heterozygous for ARR alleles but is rare in genotypes considered highly susceptible to classical scrapie such as the VRQ/VRQ genotype [10]. In addition, ARR/ARR sheep succumb to cattle and sheep BSE infection following intracerebral challenge, albeit with extended incubation periods relative to homozygous ARQ sheep [11,12]. ARR/ARR sheep orally [13] or intra-splenically [14] dosed with BSE may also sustain infection though development of clinical disease was not achieved by these routes.

Over the last two decades, we have performed several experiments in which ARR/ARR sheep have been exposed to natural scrapie or have been experimentally challenged by different routes. The purpose of this report is to draw together
the data from those studies and report the susceptibility of ARR/ARR sheep to classical scrapie.

Materials and Methods

All studies, including experimental inoculations, care of animals and euthanasia, were carried out in accordance with the UK Animal (Scientific Procedures) Act 1986. Studies 1 and 3–6 were performed at the Moredun Research Institute under licenses from the UK Government Home Office number 60/2656 (renewed in 2005 with number 60/3646). Study 2 was carried out at the Agricultural Development and Advisory Service facilities at High Mowthorpe under project license number 70/5155. Animals were monitored daily for the presence of neurological signs compatible with scrapie and were euthanized once those signs reached a standard, pre-determined end point (for details refer to [15]), when showing signs of intercurrent disease unresponsive to treatment, or for welfare reasons. In most cases, however, sheep were killed at the scheduled termination of the different studies. In all cases, euthanasia was performed by intravenous injection of barbiturate overdose followed by exsanguination.

ARR/ARR sheep included in the different studies

Ninety-five ARR/ARR sheep were included in the following six studies (Table 1 and Fig. 1):

Study 1 (Natural infection). In a closed flock of Suffolk sheep, in which the average incidence of scrapie in ARQ/ARQ sheep over a nine-year period was 84% [15], 18 ARR/ARR sheep were allowed to survive for over 48 months (Table 1 and Fig. 1). These ARR/ARR sheep were reared from birth in continuous contact with scrapie infected individuals.

Study 2 (Oral infection with sheep scrapie). Twenty-six ARR/ARR Cheviot lambs were orally dosed at 2–3 weeks of age on five consecutive days with 1 g of a brain pool homogenate (total dose 5 g) and were allowed to survive for 14 to 92 months post-infection (Table 1 and Fig. 1). The brain pool homogenate (RBP1) was made from whole brains taken from 17 scrapie affected sheep of the VRQ/VRQ, VRQ/ARQ, VRQ/ARR and ARQ/ARQ genotypes, which originated from six different farms. After challenge with the same 5 g oral dose, this inoculum produced attack rates of 100% in VRQ/VRQ homozygous sheep [16,17] and of 64% in ARQ/ARR sheep [18]. A 1 g dose of the same inoculum administered also by the oral route gave rise to 100% attack rates in VRQ/VRQ, VRQ/ARQ, VRQ/ARR and ARQ/ARQ sheep [17,18].

Study 3 (Oral infection with experimental murine-adapted scrapie). A total of 12 Suffolk or Cheviot ARR/ARR sheep between the ages of 3–5 months were orally dosed with 25 ml of a 20% suspension of brain from scrapie affected mice (5 g tissue equivalent). Three sheep each were infected with the murine adapted strains ME7, 22 A, 79 A, or 87 V, all of which were originally derived from scrapie affected sheep, and killed between 35–47 mpi (Table 1 and Fig. 1). The rationale for this experiment and the results of challenging sheep of other genotypes with the same murine strains were reported by Sisó et al., [19]. Briefly, oral challenge of VRQ/VRQ sheep with the same 5 g dose induced 100% attack rate with ME7, 50% with 22 A, 33% with 79 A and 0% with 87 V. Attacks rates for the same dose and route in ARQ/ARQ sheep (some of which were polymorphic at codons 112, 141, or 168) were 86% with ME7, 83% with 22 A, 17% with 79 A and 0% with 87 V. All four murine scrapie strains produced 100% attack rates in VRQ/VRQ and ARQ/ARQ sheep when administered by a combined oral, subcutaneous and intracerebral route.

Study 4 (Inoculation of intestinal loops with sheep scrapie). Single isolated gut loops were created as described previously [20] in six two month-old, scrapie-free ARR/ARR Suffolk lambs. Loops were created to include the distal ileum with its continuous Peyer’s patch and inoculated with 5 ml of a Suffolk scrapie brain 20% homogenate (1 g tissue equivalent). Two sheep were killed at 16 and 47 mpi because of intercurrent health problems; the remaining 4 sheep were healthy when killed at the end of the experiment at 70–72 mpi (Table 1). The inoculum was sourced from scrapie confirmed clinical cases (pool of 10 ARQ/ARQ sheep) from the naturally infected Suffolk flock described above (study 1, see Fig. 1). In a different experiment, the same inoculum was used to infect sheep of the VRQ/VRQ, VRQ/ARQ and ARQ/ARQ genotypes either by the oral or subcutaneous routes with 100% attack rates in all cases [21].

Study 5 (Subcutaneous inoculation with sheep scrapie). Thirteen New Zealand-derived ARR/ARR Suffolk lambs were subcutaneously inoculated at 6 months of age with 1 ml of a 10% clarified homogenate of the same inoculum as used in study 4 (0.1 g tissue equivalent; Fig. 1). The injection was done in the drainage area of the right prefemoral lymph node, as described in detail previously [22]. Seven ARR/ARR sheep were killed at

| Study | Exposure | Source | Dose | Months post-infection or age at post-mortem examination | Total |
|-------|----------|--------|------|--------------------------------------------------------|-------|
| No.   |          |        |      | 0–12 | 13–24 | 25–36 | 37–48 | 49–60 | 61–72 | 73–84 | 85–96 |       |
| 1     | Natural  | ARQ/ARQ| n/a  | 9    | 5     | 1     | 3     | 18    |
| 2     | Oral     | Sheep pool | 5    | 3     | 3     | 3     | 3     | 3     | 11    | 26    |
| 3     | Oral     | Murine strains | 5    | 6     | 6     | 12    |
| 4     | Intraintestinal | ARQ/ARQ | 1    | 1     | 1     | 4     | 6     |
| 5     | Subcutaneous | ARQ/ARQ | 0.1 (+2*) | 7    | 2     | 4*    | 13    |
| 6a    | Intracerebral | ARQ/ARQ | 0.1  | 2     | 4     | 4     | 10    |
| 6b    | Intracerebral | VRQ/VRQ | 0.1  | 1     | 1     | 8     | 10    |
| Total |          |        |      | 1    | 11    | 11    | 12    | 12    | 17    | 13    | 18    | 95    |

Age in months for naturally exposed sheep of study 1; months post-infection for all other studies. Sheep pool, pool of 17 scrapie brains (VRQ/VRQ = 6, VRQ/ARQ = 6, ARQ/ARQ = 4, VRQ/ARR = 1).

Murine strains, sheep dosed with 22 A or 87 V killed at 35 mpi; sheep dosed with ME7 or 79 A killed at 47 mpi. Dose in grams: n/a, not applicable; *four sheep were boosted with 2 g of inoculum by the subcutaneous route (months post-inoculation correspond to those of original challenge).
22 mpi and two at 29 and 32 mpi; the remaining four were re-challenged subcutaneously at 32 mpi with 10 ml (5 ml in each flank) of a 20% dilution of the same inoculum (2 g tissue equivalent) and killed 58 to 62 months after the second inoculation (90 to 94 months after the original challenge; Table 1). The inoculum used induced infection in 100% ARQ/ARQ Suffolk sheep without threonine polymorphism at codon 112 when injected by the same subcutaneous route [23].

Study 6 (Intracerebral inoculation with sheep scrapie). In study 6a, 10 ARR/ARR Suffolk sheep were intracerebrally challenged at four months of age with 0.5 ml of a 20% sheep scrapie brain homogenate (0.1 g tissue equivalent) using a scrapie brain homogenate derived from an ARQ/ARQ sheep from the naturally infected Suffolk flock described above (Fig. 1). Two sheep were found dead at 39 and 43 mpi, another four were culled at 61 to 72 mpi due to welfare issues and the remaining four were killed at the end of the experiment (79 mpi; Table 1). Two age-matched ARQ/ARQ sheep were also inoculated (same inoculum, route and dose) and died at ~17 mpi with a pathological and biochemical scrapie phenotype that was indistinguishable from that of the donor sheep.

In study 6b, 10 ARR/ARR Cheviot sheep were also intracerebrally challenged at the same age with the same dose as the Suffolk sheep but with an inoculum derived from a VRQ/VRQ Cheviot sheep that succumbed to confirmed scrapie after oral infection with a brain pool homogenate of six natural scrapie cases in sheep of the same breed and genotype all derived from the same flock (Fig. 1; for details of this source see [21]). Two sheep were found dead at 11 and 72 mpi, one was killed with terminal neurological signs at 74 mpi and the remaining seven were culled at the end of the experiment (79 mpi; Table 1). Two age-matched VRQ/VRQ sheep were also inoculated (same inoculum, route and dose) and died at ~5 mpi with a pathological and biochemical scrapie phenotype that was indistinguishable from that of the donor sheep.

Laboratory examinations
From each of the above studies, a detailed necropsy was performed and samples of lymphoid tissues, digestive tract, brain and spinal cord, peripheral and autonomic nervous system tissues, striated muscles and other organs were taken for immunohistochemical (IHC) examinations, as detailed in table 2. Tissues were processed to paraffin wax and stained with haematoxylin and eosin (brain only) or subjected to IHC labelling for disease associated PrP (PrP<sup>d</sup>) using R145 monoclonal antibody (binding to ovine PrP amino acid [aa] sequence 222–226 [24]) as described previously [11]. Two additional PrP antibodies, F99 (aa sequence 220–225 [25]) and 3F10 (aa sequence 137–151 [26]) were used in serial sections to help confirming low levels of PrP<sup>d</sup> accumulation found with R145. In addition, 2A11 (aa sequence 163–171 [27], and SAF 84 (aa sequence 166–172 [28]), monoclonal antibodies that do not recognize R at codon 171 were also used on tissue sections where positive PrP<sup>d</sup> labelling was detected with R145.

Western blotting (WB) was carried out in samples of medulla oblongata and/or cerebellum of all sheep using P4 (aa sequence 93–99 [29]) and SAF84 PrP antibodies as described previously [21] to detect protease resistant PrP (PrPres). In addition, samples of five different brain areas from the only sheep that developed clinical scrapie were examined with L42 (aa sequence 148–153 [28]) and F99 PrP monoclonal antibodies.

The Bio-Rad TeSeE ELISA is the screening test used in the current statutory UK small ruminant TSE surveillance programme, and was used to test cerebellum samples from all 20 intracerebrally challenged ARR/ARR sheep and the four controls of that experiment (study 6).
None of the 64 sheep surviving for less than 72 months after scrapie challenge showed any post-mortem indication of PrP d accumulation. These included 14 naturally exposed sheep, 27 orally dosed (including the 12 dosed with murine scrapie), all six receiving intra-intestinal inoculation, nine injected subcutaneously and six infected intracerebrally. Among the remaining 33 sheep, which were aged or survived for 72 or more mpi, none of the four naturally exposed, the 11 orally dosed with sheep scrapie or the four inoculated subcutaneously were positive for PrPd/PrPres, either by IHC or WB, in any of the tissues examined. Of the 14 sheep challenged by the intracerebral route only two, one Cheviot and one Suffolk sheep, showed PrPd accumulation, as described below. In summary, these results indicate attack rates of 0% in sheep naturally exposed or experimentally infected by routes other than the intracerebral, regardless of their survival time. For intracerebrally challenged sheep, attack rates would vary between 10% (2/20, if all inoculated sheep are considered, and 14.3% (2/14), if only those coeval or older than the first indication of infection (72 mpi) are accounted for.

PrP\textsuperscript{d} positive ARR/ARR Suffolk sheep

One neurologically unremarkable ARR/ARR Suffolk sheep intracerebrally challenged with ARQ/ARQ Suffolk scrapie was culled at 72 mpi because of persistent lameness and problems related to its hooves. On histological examination of the brain this sheep showed neuropil vacuolation in the thalamus. Immunohistochemistry confirmed PrP\textsuperscript{d} accumulation in the thalamus (Fig. 2a), the midbrain, the parietal cerebral cortex and the obex. PrP\textsuperscript{d} was almost exclusively present in the form of particulate accumulation within the neuropil. In the obex, sparse particulate deposits of PrP\textsuperscript{d} were found.
were present in the spinal tract of the trigeminal nerve but not in the dorsal motor nucleus of the vagus nerve (Fig. 2b). Immunoreactivity to R145 antibody was confirmed by positive immunolabelling with F99 and 3F10 but no PrP\textsuperscript{d} was detected with either SAF 84 or 2A11, confirming the ARR variant of the protein.

PrPres was not detected by WB or ELISA done on brain samples and all other tissues examined by IHC were also negative.

PrP\textsuperscript{d} positive ARR/ARR Cheviot sheep

One ARR/ARR Cheviot sheep intracerebrally challenged with VRQ/VRQ Cheviot scrapie collapsed after a short clinical course of 3 weeks, in which the animal displayed vague neurological signs, and was killed at 75 mpi. The brain showed severe vacuolation throughout all grey matter regions and PrP\textsuperscript{d} was also present throughout all neuroanatomical areas of the brain and spinal cord. A wide range of PrP\textsuperscript{d} types [30] were present including types consistent with intra-neuronal and intra-glial and several extracellular types. Most of the latter were in the form of coarse or fine diffuse particulate PrP\textsuperscript{d} in the grey matter neuropil, while glial associated perivascular aggregates were infrequent. In the thalamus and cerebral cortex, distinctive, multifocal, intensely-labelled mini plaque-like deposits and marked intra-astrocytic PrP\textsuperscript{d} accumulations were observed (Fig. 2c,d). Labelling was absent when R145 positive areas were incubated with antibodies 2A11 or SAF 84. Neither the pattern of vacuolation nor the pattern and distribution of PrP\textsuperscript{d} accumulation was consistent with that of the VRQ/VRQ Cheviot donor or the two positive control sheep. These showed less intense and less widespread vacuolation (Fig. 3a,b), predominant stellate type of PrP\textsuperscript{d} and absence of mini plaques, although the intra-glial, intra-neuronal and diffuse particulate PrP\textsuperscript{d} types were in common with ARR/ARR sheep (Fig. 3c,d).

PrP\textsuperscript{d} accumulation was not detected in the lymphoid system, digestive tract or in most of the peripheral nervous system and other tissues examined. In the retina, diffuse PrP\textsuperscript{d} labelling was found in the outer plexiform layer and coarse particulate deposits in the inner plexiform layer and in the soma of retinal ganglion cells (Fig 4a). Low levels of PrP\textsuperscript{d} accumulation were found in the trigeminal ganglion’s satellite cells (Fig 4b) and in intrafusal fibres of the infra-orbital muscles (Fig 4c). In trigeminal ganglion periaxonal PrP\textsuperscript{d} accumulation was detected in myelinated axons. Incubation of serial sections with F99 (Fig 4d) and 3F10 confirmed the specificity of labelling in these tissues.

Western blotting of brain samples incubated with P4, L42 or F99 (Fig. 5a) confirmed the presence of PrP\textsuperscript{\alpha N} in the obex, frontal cortex, thalamus, midbrain and cerebellum. The highest signal was detected in cerebellum. In almost all brain areas tested the strongest PrP\textsuperscript{\alpha N} signal was obtained from the monoglycosylated fraction, which contrasted with the strongest signal of the diglycosylated fraction in the VRQ/VRQ donor and control recipients (Figs. 5a and 5b). The mobility of the unglycosylated fraction corresponded to a molecular weight of \textasciitilde 20 to 21 kDa.
similar to that of the VRQ/VRQ and ARQ/ARQ controls and consistent with classical scrapie. When the same brain samples were incubated with SAF84 (Fig. 5a), no signal or only trace signals were obtained for any of the bands. These observations are in agreement with the 2A11 and SAF84 IHC results, and indicate that PrPres was of the ARR variant.

The Bio-Rad TeSeE ELISA gave positive test values in brain samples from VRQ/VRQ and ARQ/ARQ positive scrapie controls and from the clinically affected ARR/ARR Cheviot sheep.

Discussion

In agreement with previously published studies, the above experiments show that ARR/ARR sheep are extremely resistant to classical scrapie when naturally exposed to a highly contaminated environment or when infected by experimental protocols that approximate natural exposure. However, experimental intracerebral challenge shows that it is possible to induce disease in sheep of this genotype, albeit with low attack rates and incubation periods that are years longer than those in sheep of susceptible genotypes. Therefore, these data provide proof of principle that ARR/ARR sheep are susceptible to UK classical sheep scrapie sources albeit only by a highly artificial and efficient route of challenge.

The experimental gut loop protocol circumvents transit of orally dosed infectivity through the upper part of the alimentary system, avoiding enzymatic degradation of PrP$\text{d}$. It thus maximises the opportunity of transport of infectivity across the intestinal barrier to permit amplification of infectivity in gut associated lymphoid tissues of susceptible sheep [20]. Despite this and a relatively large dose, this route did not result in generalised or even localised amplification of PrP$\text{d}$ in ARR/ARR sheep. Equally, oral and subcutaneous infection did not result in transmission of infection. Failure to induce classical scrapie using these routes of inoculation is more relevant to natural disease than the intra-cerebral route, particularly when considering that the volumes of inocula used were in excess of those likely to be found under environmental conditions of even the highest infectivity pressure. The lack of transmission in these experiments is consistent with the absence of infection in ARR/ARR sheep exposed for prolonged periods to a highly contaminated environment (the “natural infection” group) and with many other epidemiological studies of natural scrapie in the UK [3,4,5,6,7] and elsewhere [31] which do not report classical scrapie in ARR/ARR sheep.

However, single cases of classical scrapie have been reported in ARR/ARR sheep from Japan [9], Germany and France [8], and in all instances by the detection of PrP$\text{PrP}^{\text{res}}$ in Western blot analyses of brain samples. The Japanese case, a Suffolk sheep, appeared to be clinically affected but no data are available about the clinical or pathological phenotype of the disease. The German case, a black-
headed German mutton sheep, was apparently healthy and the French case, a 5 year-old sheep, showed neurological signs of histopathologically confirmed listeriosis. Detailed data on clinico-pathological phenotypes are not available for either case, although the French case was successfully transmitted to Tg338 mice [8]. These reports suggest that cases of classical scrapie may occur naturally in ARR/ARR sheep, although all epidemiological and experimental evidence indicate that such occurrences would be sporadic or exceptional. However, more frequent detection of such cases may be hampered by current sampling strategies which focus on the hind brain, as shown by the failure of statutory ELISA testing to detect the single pre-clinically affected intra-cerebrally challenged Suffolk sheep in the present study.

ARR/ARR sheep also appear to be much more resistant to infection with classical scrapie than with experimental sheep BSE. Thus, ARR/ARR sheep intra-cerebrally challenged with 0.5 mg tissue equivalent of sheep BSE (i.e., allowing for potential differences in infectious titre, a 500 times smaller dose than in the scrapie experiment reported here) developed clinical disease with a 100% attack rate and a survival time of 49±5 months [12]. Similarly, ARR/ARR sheep intra-cerebrally challenged with cattle BSE showed figures of 56% attack rate and 49±13 months survival time (Houston et al., unpublished observations). Moreover, susceptibility of ARR/ARR sheep to sheep BSE by the oral route [13] and to cattle BSE by the intra-splenic route [14] has been documented, while oral, intra-intestinal and sub-cutaneous challenge with sheep scrapie and murine adapted sheep scrapie failed to transmit infection, at least as judged by PrP<sup>d</sup> or PrP<sup>res</sup> detection. The difference in pathogenicity of BSE and scrapie agents for ARR/ARR sheep does not appear to be a property of the agents per se as this difference in pathogenicity is not observed in sheep of other genotypes. For example, oral infection of ARQ/ARQ sheep with ARQ/ARQ sheep scrapie results in complete attack rates and survival times of 23±2 months [21], while sheep of the same genotype inoculated by the same route and with the same 5 g dose of ARQ/ARQ sheep BSE also showed a 100% attack rate and survival times of 23±2 months (first passage) or 25±2 months (second passage) [32]. Similarly, oral infection of VRQ/VRQ sheep with ARQ/ARQ sheep scrapie produces a 100% attack rate with survival times of 47±7 months [21] and infection of sheep of the same genotype with cattle BSE by the same route and dose also results in complete attack rates and survival times of ~59 months (Jeffrey et al., unpublished observations). It is worth pointing out that despite the studies reported here using a variety of natural and experimental scrapie sources, all of which proved to be infectious for sheep of susceptible Prnp genotypes [15,16,17,18,19,21,23], the number of such sources was limited. We cannot therefore rule out the possibility that other sources or strains of classical scrapie could result in higher attack rates or different disease phenotypes in ARR/ARR sheep, as is found with the BSE agent.

Figure 4. Additional tissues of ARR/ARR clinically affected Cheviot sheep showing PrP<sup>d</sup> accumulation. a) Retina showing diffuse PrP<sup>d</sup> accumulation in outer plexiform layer and granular accumulations in inner plexiform layer and retinal ganglion cells. Bar = 50 μm. IHC with R145 PrP antibody and haematoxylin counterstaining. b) Trigeminal ganglion showing granular PrP<sup>d</sup> accumulation in satellite cells Bar = 50 μm. IHC with R145 PrP antibody and haematoxylin counterstaining. c) Muscle spindle from ocular muscle showing weak granular PrP<sup>d</sup> accumulation in intrafusal muscle fibres (arrows). Bar = 50 μm. IHC with R145 PrP antibody and haematoxylin counterstaining. d) The same muscle spindle as in c) labelled with anti-PrP antibody F99. The same intrafusal muscle fibres as with R145 are labelled (arrows). Bar = 50 μm. IHC with F99 PrP antibody and haematoxylin counterstaining.
Although the pathological and biochemical features of the single clinical scrapie case in an ARR/ARR genotype in our studies was consistent with classical rather than atypical scrapie or BSE, the disease phenotype was dissimilar to the VRQ/VRQ case that provided the inoculum and to the two VRQ/VRQ positive control sheep challenged with the same inoculum. The pathological phenotype of this single ARR/ARR scrapie case has little resemblance to any other classical scrapie source the authors have examined in any sheep breed or Prnp genotype. This suggests that although some sort of modification occurs on passage of classical scrapie into ARR/ARR sheep, such change does not result in the emergence of atypical scrapie or BSE and also reinforces the notion that the interaction between the infecting source and the Pnp genotype of the host has significant impact on the susceptibility to scrapie and on the disease phenotype [21].

Acknowledgments

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Figure 5. Western blotting of different brain areas of ARR/ARR clinically affected Cheviot sheep. a) Western blots of brain tissue with 4 different PrP antibodies. Immunoblots of frontal cortex (FC) thalamus (TH) midbrain (MB) cerebellum (CB) and medulla at the level of the obex (MO) each showed significant accumulation of PrP\textsuperscript{res} with antibodies P4, L42, and F99. With each of these antibodies the cerebellum showed the greatest concentration of PrP\textsuperscript{res}. Compared with ARQ/ARQ and VRQ/VRQ scrapie positive control obex samples, the mono-glycosylated band predominates in most of the ARR/ARR samples. The unglycosylated band from all ARR/ARR and positive control samples has a molecular weight of ~20 to 21 kDa. No labelling or only trace labelling was found when the antibody SAF84 was used. Mol mrk: molecular weight marker (Note: the molecular weight marker produced a very faint signal with L42. To position the weight reference values the blot was digitally overexposed but it is the original, non-saturated blot that is reproduced). b) Graph showing the proportion of di-glycosylated and mono-glycosylated PrP\textsuperscript{res} for brain from the clinically affected ARR/ARR sheep in comparison with ARQ/ARQ and VRQ/VRQ controls. For each of the brain sites the mono-glycosylated fraction of PrP\textsuperscript{res} was present in a relatively greater amount than the di-glycosylated fraction when labelled with either P4 or L42 antibodies. In contrast both VRQ/VRQ control and ARQ/ARQ positive controls had a relatively greater amount of di-glycosylated PrP\textsuperscript{res}. Values for P4 antibody are shown in blue and for L42 in red; frontal cortex, diamonds; thalamus, squares; midbrain, triangles; cerebellum, circles; obex, pentagons; the Suffolk sheep ARQ/ARQ positive control (obex) is outlined in orange and the Cheviot sheep VRQ/VRQ positive control (obex) is outlined in green.

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Author Contributions
Conceived and designed the experiments: MJ LG MD. Performed the experiments: MJ LG SM SE MD FC. Analyzed the data: MJ SM LG MP

References

1. Benestad SL, Sarradin P, Thau B, Schonheit J, Tranulis MA, et al. (2003) Cases of scrapie with unusual features in Norway and designation of a new type, Nor98. Vet Rec 153: 202–208.
2. Goldmann W (2008) PrP genotypes in ruminant transmissible spongiform encephalopathies. Vet Res 39: 30–41.
3. González L, Dagleish MP, Bellworthy SJ, Siso S, Stack MJ et al. (2006) Post-mortem diagnosis of preclinical and clinical scrapie in sheep by the detection of disease associated PrP in their rectal mucosa. Vet Rec 158: 325–331.
4. Jeffrey M, Martin S, Thomson JR, Downes WS, Begara-McGorum I, et al. (2001) Onset and distribution of tissue PrP accumulation in scrapie-affected Suffolk sheep as demonstrated by sequential necropsies and tonal biopsies. J Comp Pathol 125: 48–57.
5. Jeffrey M, Begara-McGorum I, Clark S, Martin S, Clark J, et al. (2002) Occurrence and distribution of infection-specific PrP in tissues of clinical scrapie cases and cull sheep from scrapie-affected farms in Shetland. J Comp Pathol 127: 264–273.
6. McIntyre KM, Gubbins S, Goldmann W, Hunter N, Baylin M (2008) Epidemiological characteristics of classical scrapie outbreaks in 30 sheep flocks in the United Kingdom. PLoS One 3(12): e3994.
7. Toung SC, Pfeiffer DU, Warner R, Elliot H, del Rio Vilas V (2006) Estimation of the relative risk of developing clinical scrapie: the role of prion protein (PrP) genotype and selection bias. Vet Rec 158: 43–50.
8. Groschup MH, Buschmann A, Luhken G, Mathey J, et al. (2007) Classic scrapie in sheep with the ARR/ARR prion genotype in Germany and France. Emerg Infect Dis 13: 1201–1207.
9. Ikeda T, Horiuchi M, Ishiguro N, Muramatsu Y, Kaiuwe GD, et al. (1995) Amino acid polymorphisms of PrP with reference to onset of scrapie in Suffolk and Corriedale sheep in Japan. J Gen Virol 76: 2577–2581.
10. Benestad SL, Arsac JN, Goldmann W, Noremark M (2008) Atypical/Nor98 scrapie: properties of the agent, genetics, and epidemiology. Vet Res 39: 18–26.
11. Gonza´lez L, Martin S, Houston FE, Hunter N, Baylis M (2008) Occurrence and distribution of infection-specific PrP in tissues of clinical scrapie cases and cull sheep from scrapie-affected farms in Shetland. J Comp Pathol 127: 264–273.
12. Benestad SL, Arsac JN, Goldmann W, Noremark M (2008) Atypical/Nor98 scrapie: properties of the agent, genetics, and epidemiology. Vet Res 39: 18–26.
13. Gonza´lez L, Dagleish MP, Bellworthy SJ, Siso S, Stack MJ et al. (2006) Post-mortem diagnosis of preclinical and clinical scrapie in sheep by the detection of disease associated PrP in their rectal mucosa. Vet Rec 158: 325–331.
14. Jeffrey M, Martin S, Thomson JR, Downes WS, Begara-McGorum I, et al. (2001) Onset and distribution of tissue PrP accumulation in scrapie-affected Suffolk sheep as demonstrated by sequential necropsies and tonal biopsies. J Comp Pathol 125: 48–57.
15. Jeffrey M, Begara-McGorum I, Clark S, Martin S, Clark J, et al. (2002) Occurrence and distribution of infection-specific PrP in tissues of clinical scrapie cases and cull sheep from scrapie-affected farms in Shetland. J Comp Pathol 127: 264–273.
16. Ryder SJ, Dexter GE, Heasman L, Warner R, Moore SJ (2009) Accumulation of scrapie with unusual features in Norway and designation of a new type, Nor98. Vet Rec 153: 202–208.
17. Gonza´lez L, Pitarch JL, Martin S, Thurston L, Hunter N, Baylis M (2008) Occurrence and distribution of infection-specific PrP in tissues of clinical scrapie cases and cull sheep from scrapie-affected farms in Shetland. J Comp Pathol 127: 264–273.
18. Goldmann W (2008) PrP genotypes in ruminant transmissible spongiform encephalopathies. Vet Res 39: 30–41.
19. González L, Dagleish MP, Bellworthy SJ, Siso S, Stack MJ et al. (2006) Post-mortem diagnosis of preclinical and clinical scrapie in sheep by the detection of disease associated PrP in their rectal mucosa. Vet Rec 158: 325–331.
20. Jeffrey M, Martin S, Thomson JR, Downes WS, Begara-McGorum I, et al. (2001) Onset and distribution of tissue PrP accumulation in scrapie-affected Suffolk sheep as demonstrated by sequential necropsies and tonal biopsies. J Comp Pathol 125: 48–57.
21. Jeffrey M, Begara-McGorum I, Clark S, Martin S, Clark J, et al. (2002) Occurrence and distribution of infection-specific PrP in tissues of clinical scrapie cases and cull sheep from scrapie-affected farms in Shetland. J Comp Pathol 127: 264–273.
22. Eaton SL, Rocchi M, Gonzalez L, Hamilton S, Finlayson J, et al. (2007) Immunological differences between susceptible and resistant sheep during the preclinical phase of scrapie infection. J Gen Virol 88: 65–98.
23. Chianini F, Siso S, Ricci E, Eaton SL, Finlayson J, et al. (2013) Pathogenesis of scrapie in ARQ/ARQ sheep after subcutaneous infection: effect of lymphadenopathy and immune cell subset changes in relation to prion protein Vet Immunol Immunopath 152: 349–358.
24. Jeffrey M, González L, Chong A, Foster J, Goldmann W, et al. (2002) Ovine scrapie with unusual features in Norway and designation of a new type, Nor98. Vet Rec 153: 202–208.
25. Jeffrey M, González L, Espenes A, Press C, Martin S, et al. (2006) Ovine scrapie with unusual features in Norway and designation of a new type, Nor98. Vet Rec 153: 202–208.
26. Thuring CMA, van Keulen LJM, Langeveld JPM, van vroomen MEW, van Zijderveld FG, et al. (2005) Immunohistochemical distinction between preclinical bovine spongiform encephalopathy and scrapie in sheep. J Comp Pathol 133: 59–69.
27. Chianini F, Siso S, Ricci E, Eaton SL, Finlayson J, et al. (2013) Pathogenesis of scrapie in ARQ/ARQ sheep after subcutaneous infection: Effect of lymphadenopathy and immune cell subset changes in relation to prion protein Vet Immunol Immunopath 152: 349–358.
28. Chianini F, Siso S, Ricci E, Eaton SL, Finlayson J, et al. (2013) Pathogenesis of scrapie in ARQ/ARQ sheep after subcutaneous infection: Effect of lymphadenopathy and immune cell subset changes in relation to prion protein Vet Immunol Immunopath 152: 349–358.
29. Thuring CMA, van Keulen LJM, Langeveld JPM, Vromans MEW, van Zijderveld FG, et al. (2005) Immunohistochemical distinction between preclinical bovine spongiform encephalopathy and scrapie in sheep. J Comp Pathol 133: 59–69.
30. Jeffrey M, González L (2004) Pathology and pathogenesis of bovine spongiform encephalopathy: Immunohistochemical similarities can be resolved by immunohistochemistry. J Comp Pathol 134: 17–29.
31. Spraker TR, ORourke KI, Balachandran A, Zink RR, Cummings BA, et al. (2002) Validation of monoclonal antibody F99/97.6.1 for immunohistochemical staining of brain and tumour in mule deer (Odocoileus hemionus) with chronic wasting disease. J Vet Diag Invest 14: 3–7.
32. Stack M, Gonza´lez L, Jeffrey M, Martin S, Macalodowie C, et al. (2009) Three serial passages of bovine spongiform encephalopathy in sheep do not significantly affect discriminatory test results. J Gen Virol, 90, 764–768.