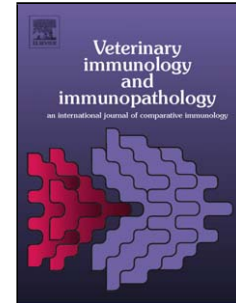


Accepted Manuscript

Title: Clinical evaluation and antibody responses in sheep after primary and secondary experimental challenges with the mange mite *Sarcoptes scabiei* var. *ovis*

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PII: S0165-2427(09)00236-0
DOI: doi:10.1016/j.vetimm.2009.07.004
Reference: VETIMM 8108

To appear in: *VETIMM*

Received date: 16-9-2008
Revised date: 16-4-2009
Accepted date: 1-7-2009

Please cite this article as: Rodríguez-Cadenas, F., Carbajal-González, M.T., Fregeneda-Grandes, J.M., Aller-Gancedo, J.M., Rojo-Vázquez, F.A., Clinical evaluation and antibody responses in sheep after primary and secondary experimental challenges with the mange mite *Sarcoptes scabiei* var. *ovis*, *Veterinary Immunology and Immunopathology* (2008), doi:10.1016/j.vetimm.2009.07.004

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23 (90-180 KDa) were observed in week 8 PC. Following the secondary challenge the ewes
24 developed a smaller area of mange lesion than that seen following primary challenge and live
25 *S. scabiei* mites were not detected in skin scrapings collected in week 8 PC, suggesting that
26 sheep had developed immunity to re-infection. Compared to primary infection, the specific
27 IgG secondary antibody levels were transient, but in contrast there was an anamnestic IgE
28 response, in which the specific serum IgE levels in week 2 PC were significantly higher than
29 those demonstrated after primary infection. WB analysis revealed one additional IgG-reactive
30 band (180KDa) and no additional IgE-reactive bands. Determining the immunodiagnostic or
31 vaccination value of the IgG-reactive antigens and IgE-reactive allergens detected requires
32 further studies.

33

34 **Keywords:** Sheep; *Sarcoptes scabiei*; ELISA; Western-blotting; Clinical-aspects; Resistance.

35

36 **Abbreviations:** OD: optical density; PC: post-challenge; RT: room temperature.

37

38 1. Introduction

39 Sarcoptic mange is a parasitic skin disease due to infection by the mite *Sarcoptes scabiei*. The
40 lesions produced in sheep, characterized by formation of crust of up to 1 cm in thickness, are
41 mainly located on the head (Cordero del Campillo and Rojo-Vázquez, 1999). The main
42 clinical signs are rubbing and scratching. It also causes important financial losses due to
43 decreases in milk production, reproductive performance and the growth of lambs born from
44 affected ewes (Fthenakis et al., 2000, 2001).

45 Currently, diagnosis of sarcoptic mange in sheep is performed by visual observation of the
46 mites in skin scrapings. The detection of specific serum antibodies against *S. scabiei* by
47 ELISA is routinely used to diagnose sarcoptic mange in dogs (Curtis, 2001; Lower et al.,

48 2001) and has also been successfully used to monitor the effectiveness of eradication
49 programmes for sarcoptic mange in pigs (Jacobson et al., 1999; Rueda-López, 2006).
50 Recently, the authors have developed and validated an ELISA to diagnose sarcoptic mange in
51 sheep using a crude saline extract from *S. scabiei* var. *ovis*, and demonstrated that it is highly
52 accurate (Rodríguez-Cadenas et al., submitted to journal). However, this test is unsuitable for
53 large scale use, owing to limitations on the amount of mites which can be collected, as there is
54 currently no *in vitro* culture technique for breeding *S. scabiei*. Identification of the major
55 antigens and allergens of *S. scabiei* would assist in the development of an immunodiagnostic
56 test based on recombinant proteins (Kuhn et al., 2008). The antibody response against *S.*
57 *scabiei* has already been characterized in dogs, foxes and goats (Bornstein et al. 1995; Arlian
58 and Morgan, 2000; Tarigan, 2004), showing differences between species. These inter-species
59 antibody profile differences are in agreement with the findings of Arlian et al. (1996a), who
60 reported that each variety of *S. scabiei* may produce a range of proteins comprised of both
61 those which are variety-specific, and those which are immunologically identical and shared
62 by the different mite sub-types.

63 Currently, control of sarcoptic mange is mainly based on the administration of acaricides. The
64 use of these chemical compounds has serious drawbacks; i.e. the development of drug
65 resistance (Curie et al., 2004), adverse environmental effects (Sanderson et al., 2007),
66 residues in animal products (Imperiale et al., 2004) and health hazards to humans (Bradberry
67 et al., 2005). As a consequence, the development of 'non-chemical' methods, such as
68 vaccination, is desirable. The achievement of a vaccine against sarcoptic mange is thought to
69 be feasible as animals having recovered from a previous infection show resistance to re-
70 infection (Arlian et al, 1994; Arlian et al., 1996b; Tarigan, 2002). Furthermore, it has been
71 suggested that the mechanism involved in the development of acquired resistance against *S.*
72 *scabiei* is related to IgE responses (Tarigan, 2003; Tarigan and Huntley, 2005).

73 The aim of the present study was to monitor sheep clinically after primary and secondary
74 experimental challenges with the mange mite *S. scabiei* var. *ovis*, and to characterise the
75 specific serum IgG and IgE antibody responses.

76

77 **2. Materials and methods**

78 **2.1. Experimental animals**

79 First, ten *S. scabiei*-naïve adult sheep (> 1 year) were challenged (primary infection group).
80 During week 10 post challenge (PC), five randomly selected sheep were treated with
81 ivermectin (Ivomec[®], Merial, two injections one week apart s.c., at 200 µg kg⁻¹ of body
82 weight) and the other five removed from the study. One of the ivermectin-treated sheep died
83 two weeks after treatment from causes not related to the experiment. The remaining four
84 treated sheep were kept until eight weeks after treatment, by which time the mange lesions
85 had disappeared. After recovering from the primary infection, the four sheep were challenged
86 again with the same *S. scabiei* var. *ovis* strain (secondary infection group) and monitored for
87 another eight weeks. As a positive control for this infection three *S. scabiei*-naïve adult sheep
88 were also challenged at the same time (secondary infection control group).

89

90 **2.2. Preparation of the *S. scabiei* var. *ovis* challenge-inocula and experimental** 91 **challenge**

92 Experimental challenges were performed using crusts with *S. scabiei* var. *ovis* mites collected
93 from a severely affected ewe. Briefly, the approximate total number of *S. scabiei* mites on the
94 donor sheep was determined on the day before challenge from the total lesion area and the
95 mean number of mites per 1 cm² of lesion. The latter measure was estimated by scraping the
96 donor sheep at several 1 cm² points in the lesion, and counting the *S. scabiei* larvae, nymphs
97 and adults under the microscope after 10% KOH digestion of the crusts and concentration of

98 the mites by floating with saturated sucrose solution. Thereafter, on the day of challenge the
99 donor sheep were euthanized (T-61[®], Intervet International B.V.), all the crusts from the
100 lesion removed and chopped to produce particles of around 5 mm diameter, and aliquots with
101 approximately 2,000 mites were prepared. The sheep were challenged with an aliquot each,
102 by maintaining the crusts in contact with the convex surface of one ear for 48 h by means of a
103 dressing.
104 Blood samples were taken by jugular puncture at weekly intervals and after centrifugation at
105 900 g for 15 min the sera were removed and stored at -20° C.

106

107 **2.3. Clinical monitoring**

108 The mange lesions which developed after challenge in the convex surface of the ear were
109 recorded in a diagram and subsequently graded by comparing with reference lesion pictures
110 which had been scored in accordance with the affected area. The scale scores were as follows:
111 score 0 if no lesion was visible on the ear; score 1 when $< 10\%$ of the area was affected; score
112 2 when 10-25% of the area was affected; score 3 when 25-50% of the area was affected; score
113 4, when 50-75% of the area was affected; and score 5 when $> 75\%$ of the area was affected.

114 Skin scrapings were collected from the lesion and examined for live *S. scabiei* mites after
115 incubation at 30° C for 30 min, and in addition they were also examined using the digestion-
116 concentration technique (see section 2.2.).

117

118 **2.4. Mite extract**

119 The mite extract used in both ELISA and Western-blotting (WB) techniques was a crude
120 saline extract kindly supplied by Dr. John F. Huntley (MRI, Penicuik, Scotland). This extract
121 had been prepared from *S. scabiei* var. *caprae* mites collected from a mangy goat as described
122 by Tarigan and Huntley (2005) and stored frozen at -20° C until being processed as described

123 below. After defrosting, they had been washed once in ice-cold PBS, followed by another
124 wash in PBS-1% (w/v) SDS at room temperature (RT) and 10 further washes in ice-cold PBS
125 to remove the SDS. The mites had been then transferred to a ribolyser tube (Lysing Matrix C,
126 Q-biogene) and homogenised in PBS with a shaker machine (FastPrep[®] FP120, Q-biogene)
127 for four 30 s cycles with cooling between each. After centrifugation at 5,500 g for 5 min the
128 supernatant (mite extract) had been removed, its protein concentration measured by the
129 Bradford method (Bradford Reagent, Sigma) with BSA as standard and stored frozen at -20°
130 C until use.

131

132 **2.5. ELISA**

133 High-binding microtiter plates (Costar[®], Corning Incorporated, USA) were coated overnight
134 at 4° C with 50 µl per well of a solution of the mite extract adjusted to 5 µg ml⁻¹ of protein
135 with carbonate-bicarbonate buffer (70 mM NaHCO₃, 30 mM Na₂CO₃, 0.2 g l⁻¹ NaN₃, pH 9.6).
136 After washing three times with PBS-T20 (PBS, 0.5 ml l⁻¹ Tween 20) the plates were
137 incubated for 1 h at 37° C with 50 µl per well of sheep sera appropriately diluted (1/200 for
138 IgG or 1/10 for IgE) in PBS-T80-NaCl (PBS, 5ml l⁻¹ Tween 80, 0.5M NaCl) and added in
139 duplicate. The plates were washed again, then incubated for 1 h at 37° C with 50 µl per well
140 of an appropriate mAb diluted in PBS-T80-NaCl: clone VPM6 (Bird et al., 1995) diluted 1/20
141 for IgG detection or clone 2F1 (Bendixen et al., 2004) diluted 1/200 for IgE detection. After a
142 further washing, the plates were incubated for 1 h at 37° C with 50 µl per well of biotin-
143 labelled goat anti-mouse IgG antibodies (Sigma[®], USA) diluted 1/5,000 in PBS-T80-NaCl.
144 The plates were washed once more, then incubated for 30 min at 37° C with 50 µl per well of
145 streptavidin-HRP (GE Healthcare, UK) diluted 1/40,000 in PBS-T80-NaCl and washed again.
146 Peroxidase activity was then visualized using the chromogen substrate OPD (OPD Tablets,
147 Dako, Denmark) in accordance with the manufacturer's instructions. The reaction was

148 stopped with 50 μ l per well of 0.5M H₂SO₄, then the Optical Density (OD) was measured at
149 492 nm using a spectrophotometer and the OD for each serum calculated (mean OD of the
150 duplicate test wells). ELISA studies were carried out at two-week intervals throughout the
151 study. Positive and negative control sera were tested on each plate and the ODs of the samples
152 were adjusted relative to them to obtain comparable results between plates.

153

154 **2.6. SDS-PAGE/Western-blotting**

155 Aliquots of the mite extract with 100 μ g of protein were mixed 1:1 with Laemmli reducing
156 sample buffer (Bio-Rad Laboratories) and boiled for 5 min. Each mixture was then loaded
157 into Tris/HCl 12% acrylamide/bisacrylamide gel, together with 10 μ l of broad-range
158 molecular weigh markers (prestained SDS-PAGE Standards, Bio-Rad Laboratories). The gels
159 were subjected to electrophoresis at 200V for 45 min using Tris/Glycine/SDS running buffer
160 (25mM Tris, 192 mM glycine, 5 g l⁻¹ SDS). After this, the separated proteins from the gels
161 were transferred onto nitrocellulose membranes (Trans-Blot[®] Transfer Medium, Bio-Rad
162 Laboratories) at 100V for 1.5 h using Tris/Glycine transfer buffer (25mM Tris, 0.192 mM
163 glycine, 200 ml l⁻¹ methanol). Protein transfer was checked by staining the membrane with
164 Ponceau S solution (0.1% Ponceau S pure, 50 ml l⁻¹ acetic acid). Electrophoresis and transfers
165 were done using the Mini Polyacrilamide Gel System (Bio-Rad Laboratories) and the
166 PowerPac[™] Basic Power Supply (Bio-Rad Laboratories).

167 The transferred membranes were then blocked for 1 h at RT with PBS-T80-NaCl, and after
168 drying were cut into strips 3 mm wide. Each strip was incubated for 1 h at RT with 1 ml of
169 sheep sera diluted 1/200 for IgG detection or 1/10 for IgE detection, in PBS-T80-NaCl. The
170 strips were then washed three times, for 5 min on each occasion, with TBS (20 mM Tris, 0.5
171 mM NaCl, pH 7.6) and incubated for 1 h at RT with 1 ml of the appropriate mAb diluted in
172 PBS-T80-NaCl: clone VPM6 diluted 1/20 for IgG detection or clone 2F1 diluted 1/1,000 for

173 IgE detection. After another washing step, the strips were incubated for 1 h at RT with 1 ml of
174 biotin-labelled goat anti-mouse IgG antibodies (Sigma) diluted 1/5,000 in PBS-T80-NaCl.
175 The strips were washed again, then incubated for 30 min at RT with 1 ml of streptavidin-HRP
176 (GE Healthcare) diluted 1/20,000 in PBS-T80-NaCl. After another washing the peroxidase
177 activity was visualized using a chemiluminescent substrate (ECLTM, GE Healthcare, UK) in
178 accordance with the manufacturer's instructions. WB studies were performed on sera
179 collected prior to challenge (week 0), and at weeks 4 and 8 PC. Positive and a negative
180 control sera were tested during each run of the assay. Further controls were ensured by
181 replacing the serum or the mAb with dilution buffer.

182 The software package Quantity One[®] 4.5.0 (Bio-Rad Laboratories) was used to capture the
183 images from the scanner (GS 800 Calibrated Densitometer, Bio-Rad Laboratories) and to
184 determine the molecular weight of the reactive bands, using as reference the strip with the
185 molecular weight markers. The antibody binding intensity of each serum to bands of the mite
186 extract was scored (separately for IgG and IgE) as 0, 1, 2, 3, 4 or 5. The score was 0 when
187 there was no binding, 1 for the weakest binding and so on up to 5 for the strongest. Thereafter,
188 the mean binding intensity of the positive sera (score ≥ 1) to the band was graded as follows:
189 weak (mean score 1-1.9), medium (2-2.9), strong (3-3.9) or very strong (4-5).

190

191 **2.7. Statistical analysis**

192 The one-sample Kolmogorov-Smirnov test was used to determine whether the paired-
193 difference variables of ODs were normally distributed, and because this was always the case,
194 the paired-samples t-test was used to compare antibody levels at different time points within
195 each group, and to compare significantly-elevated antibody levels at the same time point
196 between groups. When the intention was to do the latter, in order to prevent specific antibody
197 levels being carried over from the primary infection to the secondary infection, the pre-

198 challenge antibody level was previously subtracted for each sheep and thus only the antibody
199 level elicited by the relevant challenge was used for comparison. All statistical tests were
200 performed with SPSS 15.0 for Windows[®], and the alpha value was set at 0.05.

201

202 **3. Results**

203 **3.1. Clinical monitoring**

204 The results of clinical-lesion examination in weeks 1, 4 and 8 PC are shown in Table 1. After
205 the primary challenge all sheep exhibited mange lesions at the site of challenge, these
206 consisting of abundant exudates which later developed to form alopecia and crust formations,
207 and there was a progressive growth in the mange lesion area as indicated by an increase in the
208 mean mange lesion score. Live *S. scabiei* mites were detected in skin scrapings from seven
209 sheep (70%) in week 10 PC, while a large number of dead mites (using the digestion-
210 concentration technique) were detected in all of them.

211 Following ivermectin treatment in week 10 PC, the mange lesions progressively disappeared
212 and there were no visible lesions of mange in any sheep by eight weeks later, when the
213 secondary challenge was performed.

214 After the secondary challenge, the four sheep exhibited mange lesions at the site of challenge
215 which were visually similar to those observed in the primary infection group. However, after
216 one week PC the lesions progressively disappeared, as indicated by a decrease in the mean
217 mange lesion score, to form alopecic areas. Live *S. scabiei* mites were not detected in any of
218 the four skin scrapings collected after 8 weeks PC, while a few dead mites were detected in
219 one (25%). The mange lesion scores of the secondary infection control group approximated to
220 those of the primary infection group.

221

222 **3.2. ELISA**

223 The results of ELISA are shown in Figure 1. Before primary challenge the ELISA OD values
224 were low (< 0.15 for IgG and < 0.1 for IgE) in all sheep. After the primary challenge, most of
225 the sheep developed specific serum IgG and IgE antibodies. A significant increase in the IgG
226 levels were first detected two weeks PC (one-tail $p < 0.05$), which were strongly increased at
227 four weeks PC, after which the OD values levelled to a plateau. IgG seroconversion (cut-off
228 value 0.197; defined as the mean plus 3 SD of ODs of sera at week 0) was demonstrated in
229 nine sheep (90%) in week 4 PC, and they remained seropositive in week 8 PC. A significant
230 increase in the IgE level was also first detected in week 2 PC (one-tail $p < 0.01$) and then
231 progressively increased. IgE seroconversion (cut-off value 0.093) was found in five animals
232 (50%) after 4 weeks PC and in six (60%) after 8 weeks PC.

233 After ivermectin treatment a decrease was detected in the specific IgG and IgE levels. Prior to
234 the secondary challenge the IgG antibody level was still significantly higher when compared
235 to pre-primary-challenge (one-tail $p < 0.05$), but the IgE antibody level was similar (one-tail
236 $p > 0.05$).

237 After the secondary challenge, a significant increase in the specific IgG level was detected in
238 week 4 PC (one-tail $p < 0.05$), but in week 6 and 8 PC the IgG level was similar to pre-
239 secondary-challenge (one-tail $p > 0.05$). However, the IgG level in week 4 PC was similar to
240 that of the same sheep in the primary-infection group (two-tail $p > 0.05$). The specific IgE
241 level showed a rapid and strong increase in the first two weeks (one-tail $p < 0.05$) and reached
242 the highest value at week 4 PC, while after that the level started to decrease but still remained
243 significantly elevated at week 8 PC (one-tail $p < 0.05$). The IgE level in week 2 PC was
244 significantly higher than that of the same sheep in the primary infection group (two-tail $p <$
245 0.05), but later was similar (two-tail $p > 0.05$). The IgG and IgE antibody levels in the
246 secondary infection control group were similar to those of the primary infection group.

247

248 **3.3. WB**

249 The results of WB analysis are shown in Table 2, Table 3 and Figure 2. Prior to the primary
250 challenge IgG-reactive bands of 120, 52, 46, 44, 38 and 36 KDa, and IgE-reactive bands of
251 120 and 90 KDa were detected in some sheep, albeit weakly. Following primary infection,
252 additional IgG and IgE-reactive bands were demonstrated. The IgG analysis at week 4 PC
253 showed that there was reaction to four additional bands of 100, 85, 47, and 39 KDa in most of
254 the ewes, and the binding intensities and/or frequencies to the reactive bands detected before
255 challenge also increased. A similar profile was demonstrated in week 8 PC, with only a small
256 increase in binding intensities and frequencies. IgE reactivity in week 4 PC was also increased,
257 with six sheep sera showing strong labelling intensity to the 120 KDa band, although
258 reactivity to the 90 KDa band remained unchanged. In week 8 PC, weak IgE-reactions to two
259 additional bands of 180 and 100 KDa was detected. The binding intensities and frequencies
260 for the 120 and 90 KDa IgE-reactive bands were also increased. The immunodominant bands
261 were defined as bands reacting with at least 50% sheep sera and with a very strong or strong
262 mean binding intensity. Based on this definition, the immunodominant IgG-reactive bands in
263 weeks 4 and 8 PC were 120, 52 and 44 KDa, while in contrast there was only one
264 immunodominant IgE-reactive band in weeks 4 and 8 PC, which had 120 KDa.

265 The antibody profile prior to the secondary challenge was similar to that observed prior to the
266 primary challenge. Following the secondary challenge, one additional IgG-reactive band (180
267 KDa) was detected, but in general there were lower binding intensities than those
268 demonstrated following primary challenge. The secondary IgE-reaction was also mainly
269 confined to the 120 KDa band, with peak binding intensity observed in week 4 PC. This
270 120KDa band appeared to be immunodominant for IgE antibodies following secondary
271 challenge.

272

273 **4. Discussion**

274 In this study it was shown that primary and secondary experimental challenges of sheep with
275 *S. scabiei* var. *ovis* induce the development of mange lesions and the elicitation of serum IgG
276 and IgE antibodies directed to proteins/peptides of a wide range of molecular weights.
277 Interestingly, the secondary infection corresponded with reduced mange lesion area, absence
278 of detectable live *S. scabiei* mites in skin scrapings collected in week 8 PC and a concomitant
279 increase in specific serum IgE antibodies.

280 This study was based on the detection of antibodies elicited by *S. scabiei* var. *ovis* using a
281 mite extract prepared from *S. scabiei* var. *caprae*. It is known that each variety of *S. scabiei*
282 may produce a range of specific proteins (Arlian et al., 1996a). However, the level of cross-
283 antigenicity between them is usually high and a heterologous variety can be successfully used
284 to detect specific antibodies. For instance, ELISAs based on extracts from *S. scabiei* var.
285 *vulpes* have been used for diagnosing sarcoptic mange in pigs, dogs and chamois (Bornstein
286 et al., 1996; Bornstein and Wallgren, 1996; Lower et al., 2001; Rambozzi et al., 2004). In the
287 present study, the specific antibody responses detected by using the *S. scabiei* var. *caprae*-
288 based ELISA to detect antibodies in sheep after challenge were significant and quite
289 pronounced, suggesting that this heterologous variety can be used with high performance to
290 detect antibodies elicited by *S. scabiei* var. *ovis*. Nevertheless, it is possible that *S. scabiei* var.
291 *ovis* has specific antibody-reactive proteins, and as a result they would not have been
292 identified.

293 Prior to primary challenge and despite ELISA OD values being low, serum reactivity to
294 several bands of the mite extract was detected by WB analysis. This may have been due to the
295 presence of antibodies elicited by free-living mites such as *Dermatophagoides pteronyssinus*
296 or *Acarus siro* which cross-react with *S. scabiei* proteins (Arlian et al., 1991; van der Heijden
297 et al., 2000). In the primary infection, live *S. scabiei* mites were detected in only 70% skin

298 scrapings in week 10 PC, but all sheep developed mange lesions that spread over the ear and
299 in addition a large number of dead mites were also detected. This would indicate that the
300 technique used to detect live mites had relatively low sensitivity, a finding in agreement with
301 Gutiérrez et al., (1996) who studied sarcoptic mange in pigs. The kinetics of specific serum
302 IgG and IgE antibodies were similar to the findings of other authors studying sarcoptic mange
303 in rabbits, foxes and goats (Arlian et al., 1994; Bornstein et al., 1995; Tarigan, 2004). As the
304 infection progressed, the average IgG and IgE values tended to rise which is consistent with
305 the higher number of reactive bands, and the increase in the labelling intensities and
306 frequencies detected by WB. However, despite the development of clinical lesions in all
307 sheep, IgG and IgE seroconversion was not found in 10% and 40% of sheep, respectively, in
308 week 8 PC. The absence of detectable levels of IgE antibodies in a significant proportion of
309 sera from *S. scabiei*-infected humans, dogs and goats has previously been reported (Arlian
310 and Morgan, 2000; Morgan et al., 1997; Tarigan, 2004).

311 In the present study, the antibody profile was studied by WB and reactions were demonstrated
312 to bands of a wide range of molecular weights. However, heterogeneity of reaction within
313 each of these bands was not determined and they may represent two or more different
314 antibody-reactive proteins with the same or similar molecular weights. As a consequence, the
315 present studies cannot confirm either that bands which reacted with both IgG and IgE
316 antibodies were due to the same antigens being detected by both isotypes. The
317 immunodominant bands include those proteins/peptides more intensively and frequently
318 recognized by the immune system, which are more likely to be the best candidates for a
319 sensitive immunodiagnostic test based on the detection of induced antibodies. The
320 immunodominant IgG-reactive bands detected during the primary infection (120, 52 and 44
321 KDa) were reactive with most of the sheep sera collected in weeks 4 and 8 PC, and therefore
322 include antigens which are candidates for developing a highly sensitive immunodiagnostic

323 test for sarcoptic mange in sheep. In contrast, the 120 KDa allergen seems to have limited
324 immunodiagnostic value, as it was reactive with only 70% of sheep sera taken in week 8 PC.
325 Of note at this point was the finding that also some pre-challenge sera produced weakly
326 reaction to the immunodominant bands. The origin of these cross-reactive antibodies is
327 unclear, as stated before. Nevertheless, antibodies to these bands were markedly increased in
328 a time-dependent manner following primary challenge, indicating that these proteins may still
329 be relatively specific. Furthermore, previous studies have demonstrated little cross-
330 antigenicity between some closely related mite proteins (Cheong et al., 2003; Kuo et al.,
331 2003).

332 The clinical and antibody response of sheep after a secondary challenge with *S. scabiei* var.
333 *ovis* was also studied. A positive control for this infection was obtained by applying aliquots
334 of challenge inocula to three *S. scabiei*-naïve ewes at the same time. The mange lesion area
335 and the ELISA antibody levels of these positive control animals were similar to those
336 observed in the primary infection group, which suggests that the pathogenicity of the mite
337 doses used in the primary and secondary challenges was similar, so the two infections can
338 validly be compared. The secondary infection group had a marked reduction in the mean
339 mange lesion score compared with that of the same sheep in the primary infection group and
340 no live *S. scabiei* mites were detected in any skin scraping collected in week 8 PC. These
341 results suggest that the sheep had developed some immunity during the primary infection
342 which was able to ameliorate the secondary infection, thus confirming previous reports
343 concerning dogs, goats and rabbits (Arlian et al., 1994; Arlian et al., 1996b; Tarigan, 2002).
344 This secondary infection was characterized by an anamnestic IgE response resulting in
345 specific serum IgE levels significantly higher than those detected during the primary infection.
346 The mechanism involved in the development of resistance to re-infection by *S. scabiei*
347 remains unclear but it is believed that the early immune response occurring in the local skin

348 plays a major role and may be associated with IgE-mediated responses (Tarigan, 2003). This
349 conclusion is also tentatively supported by the absence of specific serum IgE antibodies in a
350 vaccination trial which failed to provide protection against infection in goats, despite the
351 presence of high levels of specific serum IgG antibodies (Tarigan and Huntley, 2005). This
352 lack of correlation between specific serum IgG levels and resistance agrees with the findings
353 of Arlian et al. (1994), and the results from the present study supports the view that the
354 development of resistance to *S. scabiei* is more closely correlated with the presence of specific
355 serum IgE antibodies.

356 In summary, this study demonstrates that primary and secondary challenges of sheep with *S.*
357 *scabiei* var. *ovis* results in both IgG and IgE antibody responses, and that secondary infections
358 are ameliorated. The IgG antibody responses react to a number of antigens which would
359 represent good targets for developing an immunodiagnostic test to detect *S. scabiei* var. *ovis*
360 infections; conversely antigenic epitopes for IgE antibodies were mainly restricted to a 120
361 kDa band which would include a vaccine candidate.

362

363 **5. Acknowledgements**

364 The present study was partially financed by a grant (LE010A05) from the Education Bureau
365 of the Castile and Leon Regional Government (Consejería de Educación de la Junta de
366 Castilla y León). FRC holds a fellowship from the Education Bureau of the Castile and Leon
367 Regional Government, co-financed by the European Social Fund. The authors are very
368 grateful to Dr. J.F. Huntley (Moredun Research Institute, Penicuik, Scotland) for reading the
369 manuscript with a critical eye and for kindly supplying the VPM6 and 2F1 mAbs, and the
370 mite extract used in this work.

371

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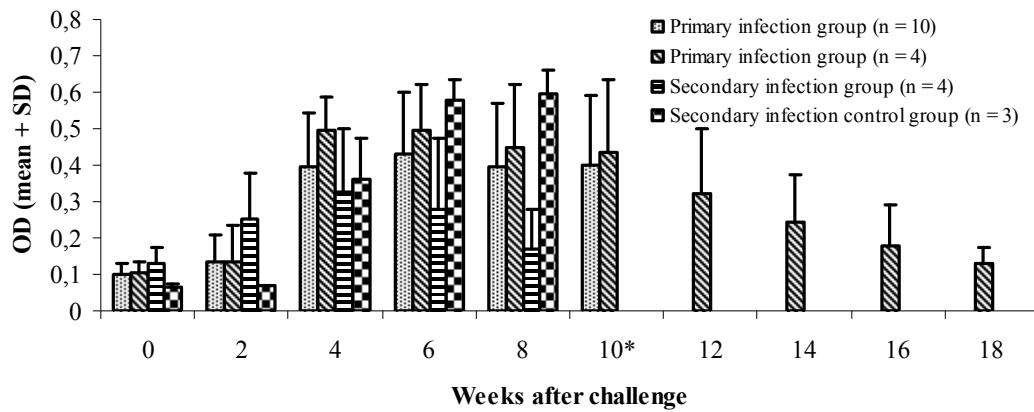
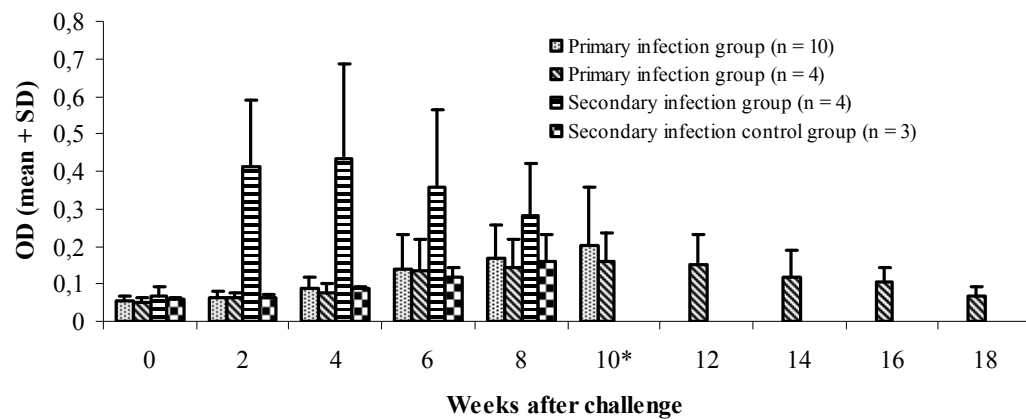
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- 456

457 Figure 1. Specific serum IgG (A) and IgE (B) antibodies measured by ELISA in the course of
458 primary and secondary experimental infections of sheep with *S. scabiei* var. *ovis*. * indicates
459 treatment with ivermectin of the primary infection group (n = 4), which corresponded with the
460 four sheep which took part in both primary and secondary infections.

461

462 Figure 2. WB strips showing recognition of *S. scabiei* var. *caprae* proteins by serum IgG (A)
463 and IgE (B) antibodies in the course of primary and secondary experimental infections of
464 sheep number 7 with *S. scabiei* var. *ovis*. Lane 1: prior to primary challenge; lane 2: week 4
465 after primary challenge; lane 3: week 8 after primary challenge; lane 4: prior to secondary
466 challenge; lane 5: week 4 after secondary challenge; and lane 6: week 8 after secondary
467 challenge.

468

**A****B**

MW
(KDa)

1 2 3 4 5 6

193 

118 

99 

54 

38 

29 

20 



MW
(KDa)

1 2 3 4 5 6

193

118

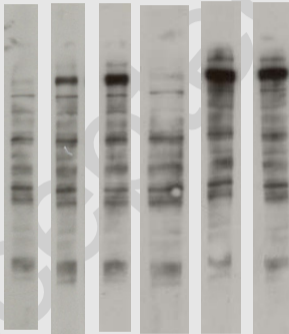
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1 Table 1. Mean mange lesion scores at different times in the course of primary and secondary
 2 experimental infections of sheep with *S. scabiei* var. *ovis*.

	Prior to challenge	Week 1 after challenge	Week 4 after challenge	Week 8 after challenge
Primary infection group (n = 10/n = 4) *	0/0	1.9/2.25	2.7/2.75	3.5/3.75
Secondary infection group (n = 4)	0	2	1.75	1.25
Secondary infection control group (n = 3)	0	2	2.67	4

3

* n = 10 includes the ten sheep, while n = 4 includes only the four sheep which took part in both primary and secondary infection groups.

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Table 2. Frequencies of labelling (%) and mean binding intensities* of IgG in sheep sera collected at different stages in the course of primary and secondary experimental infections to *S. scabiei* var. *caprae* proteins.

Molecular weight (KDa)	Primary infection (n = 10)			Secondary infection (n = 4)		
	Prior to challenge	Week 4 after challenge	Week 8 after challenge	Prior to challenge	Week 4 after challenge	Week 8 after challenge
180	0%	0%	0%	0%	0%	25% / s
120	30% / w	80% / s	90% / s	0%	100% / m	50% / m
100	0%	80% / m	90% / m	0%	75% / w	50% / w
85	0%	80% / m	80% / m	0%	75% / w	50% / w
52	90% / w	100% / s	100% / s	100% / m	100% / m	75% / w
47	0%	80% / w	80% / w	0%	50% / w	75% / w
46	30% / w	100% / m	100% / m	0%	50% / m	75% / w
44	70% / w	100% / s	100% / s	75% / w	100% / m	100% / m
39	0%	90% / w	100% / w	0%	25% / w	25% / w
38	90% / w	100% / m	100% / m	50% / w	75% / w	75% / w
36	100% / w	100% / m	100% / m	50% / w	75% / w	75% / w

* It was estimated from only the sera positive to the band: w: weak; m: medium; s: strong; vs: very strong.

Table 3. Frequencies of labelling (%) and mean binding intensities* of IgE in sheep sera collected at different stages in the course of primary and secondary experimental infections to *S. scabiei* var. *caprae* proteins.

Molecular weight (KDa)	Primary infection (n = 10)			Secondary infection (n = 4)		
	Prior to challenge	Week 4 after challenge	Week 8 after challenge	Prior to challenge	Week 4 after challenge	Week 8 after challenge
180	0%	0%	40% / m	0%	75% / m	75% / w
120	10% / w	60% / s	70% / vs	50% / m	100 / vs	75% / s
100	0%	0%	10% / w	0%	25% /w	25% / w
90	10% / w	10% / w	50% / w	0%	100% / w	75% / w

* It was estimated from only the sera positive to the band: w: weak; m: medium; s: strong; vs: very strong.