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This is the author's manuscript

Original Citation:

Availability:

This version is available <http://hdl.handle.net/2318/102072> since

Published version:

DOI:10.1016/j.exppara.2009.10.008

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UNIVERSITÀ DEGLI STUDI DI TORINO

This is an author version of the contribution published on:

Questa è la versione dell'autore dell'opera:

EXPERIMENTAL PARASITOLOGY,

Volume: 124 **Issue:** 3 **Pages:** 265-271, 2010, **DOI:** 10.1016/j.exppara.2009.10.008

The definitive version is available at:

La versione definitiva è disponibile alla URL:

<http://www.journals.elsevier.com/experimental-parasitology/>

Sarcoptes scabiei: Specific immune response to sarcoptic mange in the Iberian ibex *Capra pyrenaica* depends on previous exposure and sex.

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Keywords: *Capra pyrenaica* Immunity Inter-individual variability *Sarcoptes scabiei* Ungulate

ABSTRACT

Host acquired immunity is a critical factor that conditions the survival of parasites. Nevertheless, there is a shortage of data concerning inter-individual immunological inequalities in wild mammals. Sarcoptic mange is a widespread parasitosis that severely affects mammals such as the Iberian ibex (*Capra pyrenaica*). Despite some work on the subject, the immune response to sarcoptic mange infestation is still a complex and poorly understood phenomenon. To improve knowledge of the host-*Sarcoptes* immunological interaction, 18 Iberian ibexes were experimentally infested. IgG levels were assessed using ELISA to test for potential factors determining the specific immune response to infestation. Previous exposure and sex appeared to affect the IgG response to infestation and our results suggest a sex-biased immune modulation. We discuss the immunological pattern of host-*Sarcoptes* interactions and also suggest further lines of work that may improve the understanding of immunological interactions of host-*Sarcoptes* systems.

Introduction

Host acquired immunity (immune memory) is a critical factor that conditions the survival of parasites (Hudson and Dobson, 1995; Wilson et al., 2002). Available data on inter-individual inequalities in host-parasite interactions is uneven and differ according to the host and parasite taxa in question. Studies have tended to focus more on birds and endoparasites than on any other class of animals (Dobson, 1988; Clayton and Moore, 1997) and data for wild mammals, for instance, are scarce and more ambiguous. Some physiological inequalities have been described in small mammals (Khokhlova et al., 2004; Kristan, 2004), although for medium and large species most data are the product of analogies with human and domestic animals (Lloyd, 1995). Despite the many experimental studies that have described uneven clinical and physiological responses, inter-individual variability is often just acknowledged (e.g. Mörner and Christensson, 1984; Skerratt, 2003a,b) without being fully characterized.

Here we present details of an experiment on the immunological interaction between the etiological agent of sarcoptic mange (*Sarcoptes scabiei*, Linnaeus 1758) and one of its wild hosts, the Iberian ibex (*Capra pyrenaica*, Schinz 1838). This sexually dimorphic mountain ungulate is endemic to the Iberian Peninsula (Pérez et al., 2002) and is affected by sarcoptic mange (Pérez et al., 1997) caused by a submacroscopic mite that burrows into the skin of domestic and wild mammals (Pence and Ueckermann, 2002; Gonzalez-Candela et al., 2004). Sarcoptic mange affects animals at a broad spectrum of levels ranging from the individual pheno-type (Serrano et al., 2007) to the population dynamics of the host species (Pence and Ueckermann, 2002). By the end of the twentieth century several Iberian ibex populations had experienced massive mortality rates (in some cases over 90%) as a result of increasing epizootic episodes of sarcoptic mange (Fandos, 1991). Conservation managers and researchers have collaborated (for example, in this experiment) to explore strategies aimed at controlling this disease *in natura* and to prevent the extinction of this endemic ungulate, whose populations are only just recovering after centuries of multifactor population decline (Pérez et al., 2002).

Correlative studies of *host-Sarcoptes* interaction provide multiple trended data on epidemiology; nevertheless, these works often report an uneven distribution of sarcoptic mange in host populations (Pérez et al., 1997; Pence and Ueckermann, 2002). As well, experiments have detected inequalities and have sometimes revealed previous exposure to be one of the determining factors (Arlian et al., 1994; Skerratt, 2003a). However, a full characterization of inter-individual inequalities has not yet been carried out. In particular, few immunological studies have been conducted and, despite some available information suggesting the importance of humoral and cellular responses (Arlian, 1996), the immune response to sarcoptic mange remains complex and poorly understood (Pence and Ueckermann, 2002).

We employed an experimental approach with different host-parasite interaction sub-classes to test the following hypotheses:

1. According to previous studies on host-Sarcoptes immunological interaction (Falk, 1980; Arlian et al., 1994; Bornstein et al., 1995), infested as opposed to control animals are expected to develop a specific immunological response.
2. As immunity and parasitism are reported to vary with sex (Folstad and Karter, 1992; Hughes and Randolph, 2001), we expect there to be a lower specific response in males than in females.
3. As immunity is reported to vary with age (Lloyd, 1995), we expect there to be a lower specific response in juvenile animals than in adults.
4. According to the acquired immunity principle, secondary responses should reach higher levels (Wikel, 1996; Wakelin and Apanius, 1997) and so we expect there to be greater specific response in previously exposed animals than in naïve ones.

2. Materials and methods

2.1. General experimental procedure

The experimental buildings were located in southern Spain (Centro Las Mimbres, Parque Natural de las Sierras de Huétor, Granada). Eighteen Iberian ibexes *Capra pyrenaica hispanica* (Table 1) obtained from a stock reservoir protected from exposure to sarcoptic mange (Sierra Nevada National Space, 36°55'-37°10'N, 2°56'-3°38'W) were kept in small groups in separate enclosures. Special care was taken to avoid any transmission of the mites and special protective clothing and footwear were worn when visiting and handling the animals. In each enclosure, animals were able to move freely and had *ad libitum* access to food and water. Animals were kept under observation during an acclimatisation period of 6 weeks prior to the tests being carried out.

During the first experimental period (9 weeks, see Table 2), 13 ibexes were infested with low inoculums (load 6300 mites), while a group of 5 ibexes were maintained as a control group. At 74 days post infestation (dpi), animals were treated using doramectin (Dec-tomax®, 10 mg/50 kg b.w.) and then again at 89 dpi. A recovery period of 5 weeks was allowed after the second treatment with doramectin. The second experimental period began at 132 dpi and all animals (both those from the control group and from the previously-infested group) were infested with low inoculums (load 6300 mites). On health grounds, animals were subsequently treated with doramectin at 208 and 223 dpi, as during the first experimental period.

2.2. Manipulation of parasite loads and sampling

The ibexes used in the experiment were infested with *S. scabiei* extracted from the skins of three naturally parasitized wild ibexes selected from the neighbouring Sierra Nevada massif. Mites were extracted from pieces of mangy skin by means of a thermal gradient induced by a light shone from below Petri dishes with black bottoms and transparent central areas (Andrews, 1981).

Mites were counted with a stereomicroscope and placed on a sterilized metallic support. Next, this support was fixed onto the ibex's previously shaved inter-scapular region using adhesive bands, thereby inducing contact between the mites and host's skin.

Blood samples were collected by venipuncture from the day of the first infestation until the end of the experiment (Table 2). Sampling followed the same calendar for all individuals (Table 2) as a means of creating a standardized data base. After clotting, blood samples were centrifuged at 200g for 15 min and sera were stored at -20 °C in 1 ml aliquots until required.

Skin scrapings were also performed to verify the success of mite establishment (Table 2).

2.3. Immunological response estimates

IgG antibodies are an essential weapon in acquired immunity since they identify and neutralize foreign antigens (Wikel, 1996). IgG levels were estimated using a serological test - a labelled avi-din-biotin immunosorbent assay (LAB-ELISA) - that was developed to detect the specific humoral response to *S. scabiei* in the Caprinae subfamily (Rambozzi et al., 2004). Antigen was obtained from living *S. scabiei* mites at different stages of development (Rambozzi et al., 2004). The optical density of processed samples was read at 405 nm with a spectrophotometer (Anthos 2010, Anthos Labtec Instruments, Wals, Austria). Potential cross-reactions were tested for, although there was no evidence of any cross-reaction with other potential causes of skin damage in ungulates (see Rambozzi et al., 2004).

Our analysis was focused on the quantitative responses to infestations, and not on the qualitative classification of healthy/mangy individuals. Consequently, we analysed the optical density of processed samples as a continuum and an indicator of increases in the IgG, but did not use any cut-off point.

2.4. Statistical analyses

The experimental design was uneven in its sex ratio and treatment of the animals tested (Table 1). Consequently, we performed a number of different analyses to test each of our predictions with each sub-set of experimental individuals. More specifically, we first examined the occurrence of a specific immune response in just males. Then, we analysed the factors determining the immune response to the first infestation in infested individuals. Next, during the second trial we examined the effect of reinfestation on immune response in males and analysed the factors determining the immune response to reinfestation in only reinfested individuals. Finally, to contrast with the results of the analysis cited above, we examined the factors determining the immune response in reinfested females and naïve males. Since data consisted of a longitudinal series of measurements from the same host animals, we used generalized additive mixed model analyses (GAMM; Verbeke and Molenberghs, 2000; Pater-son and Lello, 2003; Wood, 2006; Zuur et al., 2007). Taking into account the characteristics of our data set and in order to apply the most informative method, we used an information-theoretic approach (Whittingham et al., 2006) based on the Akaike Information Criterion corrected for small sample sizes (AICc; Burnham and Anderson, 2002). Variables examined were "Group" (infested in the first trial or not), "Sex" (male or female) and "Age". Model selection identified in our analyses the most parsimonious model (lowest AICc; Burnham and Anderson, 2002) from the possible sub-sets, which ranged from the null model to a model with explanatory variables and their two-order interactions. The larger the Akaike difference (DAICc), the less plausible it is that the fitted model is the best model given the data set (Burnham and Anderson, 2002). Models with Akaike differences of less than 2 units and the relative importance of examined variables (RI) are commented upon (Burnham and Anderson, 2002), especially when the Akaike weight of the best model was moderate or low (Anderson et al., 2000, 2001). RI is measured by the sum of the Akaike weights for all models in which the variable appears (Burnham and Anderson, 2002) and expresses the probability that the considered exploratory factor is included in the "ideal" model explaining the variability in the dependent variable given the data set. All analyses were performed using the R 2.6.2 statistical package (R Development Core Team, 2008).

3. Results

All infested ibexes developed characteristic mange lesions due to the experimental exposition to *S. scabiei*; the success of mite establishment was confirmed by skin scraping and microscopic examination.

3.1. Analysis of the first infestation

Infested males developed a specific serum antibody response during the first infestation period (Fig. 1). Taking into account only males (Table 3a), the model including "Time", "Group" and their interaction as explanatory factors was the best ($W_{\text{Time+Group+Time*Group}} = 1$; see Table 3a), suggesting that the observed response develops over time. The robustness of those results is highlighted by the multimodel inference. Of the explanatory variables, factors "Time", "Group" and their interaction had relative importances (RI) equal to one (see Table 3a).

The analysis specifically focusing on the infested individuals of both sexes confirmed that the specific response to infestation developed over time (Fig. 1), and that the sex of the host was a further key factor determining the response to the first trial (see Table 3b). The model including "Time", "Sex" and their interaction was the best ($W_{\text{Time+Sex+Time*Sex}} = 0.77$); other models had Akaike differences greater than 4 units (see Table 3b), which underlines the poor support for these models given the data set. "Time", "Sex" and their interaction had very high relative importances in the observed pattern, while the age of infested animals only had a low RI (see Table 3b).

3.2. Analysis of the second infestation

Previously exposed males had lower responses over time than naïve ones (Fig. 2). Taking into account only males, in the second trial (Table 4a) the model including "Group", "Time" and their interaction was the best ($W_{\text{Time+Group+Time*Group}} = 0.76$; see Table 4a), suggesting that previous exposure to *Sarcoptes* has a negative effect on the development of the response over time. The multi-model inference reinforced these results since the RI of "Group", "Time" and their interaction was equal to one and the RI of "Age" and its interactions with other factors had lower values (Table 4a).

Among reinfested individuals, females had a higher specific response to reinfestation than males (Fig. 2). In the analyses which took into account reinfested animals of both sexes (see Table 4b), the model including "Time", "Sex" and their interaction was the best ($W_{\text{Time+Sex+Time*Sex}} = 0.58$), underlining the sex-dependent response to reinfestation. Likewise, of the explanatory variables "Time", "Sex" and their interaction had a high RI and "Age" and its interactions with other factors appeared as having a low RI (Table 4b).

Finally, the response of naïve males to infestation seems not to diverge strongly from or to be slightly lower than the response of reinfested females (Fig. 2). In the analysis which took into account naïve males and reinfested females (see Table 4c), the model including "Time" was the best ($W_{\text{Time}} = 0.48$), although two other models including "Sex" had Akaike differences of lower than 2 units, which suggests that these other two models could have also substantial support. Likewise, of the explanatory variables "Time" had a higher relative importance than both "Sex" and their interaction.

4. Discussion

4.1. Specific serum antibody responses to *S. scabiei* in the Iberian ibex

Iberian ibexes infested with sarcoptic mange produced a specific antibody response, which agrees with data from other hosts

(Falk, 1980; Arlian et al., 1994; Bornstein et al., 1995) and with our first prediction. This finding diverges to some extent from Lastras et al. (2000), who observed no significant differences in IgG levels between healthy and infested Iberian ibexes from the same area, and we discuss this apparent contrast below. In line with the 3R principles (Russell and Burch, 1959), we limit the control group of our experiment to just male Iberian ibexes. However, on the basis of our results, the assumption of specific immune response to *Sarcoptes* in females as well is reasonable. Females are normally the immunologically stronger gender (Folstad and Karter, 1992); the chromosomal localization of the major histocompatibility complex has been reported in autosomes in several ungulate species (Ansari et al., 1988; Mäkinen et al., 1989) and there is no support to presuppose male-exclusive acquired immunity.

Uneven responses to *Sarcoptes* appeared during the first trial and further inequalities appeared after the reinfestation, which highlights the complexity of this topic. These inequalities confirm our second prediction, since sex appeared as a factor determining the immune response to infestation and reinfestation. Females had a higher acquired response to infestation and reinfestation than males, but the lack of marked difference between the response of reinfested females and naïve males indicates that a composite effect is taking place. Previously exposed males had lower IgG responses than naïve males, underlining the importance of experience. This paradoxical result contradicts our fourth prediction and the hypothesis of Shrank and Alexander (1967), who consider acquired immunity to *Sarcoptes* as a permanent feature. Arlian et al. (1994) obtained results with a similar trend in rabbits reinfested by *Sarcoptes*. Arlian et al. (1994) suggested that previous exposure induces immunoresistance and interpreted the subsequent reduction in humoral response as proof of resistance on the basis of the observed recovery from lesions. This, however, is not consistent with the acquired immunity principle, which implies that the first trial should have induced a greater response during the secondary exposure (Wikel, 1996; Wakelin and Apalanis, 1997). The observed reduction in acquired immunity should be nearer to the effects of immunomodulation (Wikel et al., 1996). French et al. (1988) suggested that the experimental environment and probably self-grooming behaviour are key factors in recovery from mange lesions. The recoveries from lesions observed by Arlian et al. (1994) were probably favoured by factors other than the reduction of acquired immunity. Our results suggest that previous exposure to *Sarcoptes* may induce a sex-biased modulation of the IgG *Sarcoptes*-specific response in the Iberian ibex. Alternative hypotheses include a possible ineffective exposure to mites in previously infested males or an acquired immunity based on other antibodies isotopes such as IgE, possibly leading to the lack of IgG response in these individuals. However, the success of mite establishment was confirmed in all ibexes by the development of lesions and the skin scraping. Altered levels in other antibodies isotopes (particularly IgE) have been described by other authors as a response to *Sarcoptes* linked to atopic dermatitis, which would imply the development of an allergic response to *Sarcoptes* rather than an specific response able to eliminate the parasite (Falk, 1980). Consequently, the hypothesis of an ineffective exposure to mites or of acquired immunity based on other antibodies isotopes such as IgE, as confounding factors, have little support. *In natura*, females are also infested and die from sarcoptic mange; compared to males, the modulation of immunity in females might be delayed and this uneven immunological interaction will have to be examined in more detail in the future. This complex interaction may have led to difficulties in interpreting the results of correlative studies (Lastras et al., 2000), since the pathogenic experience of culled free-ranging animals is generally unknown.

4.2. Sample size

In our experiment, infestations were analysed in 18 Iberian ibexes, a larger sample size than those used in previous experiments on the effects of *Sarcoptes* on wild mammals (Samuel, 1981; Mörner and Christensson, 1984; Bornstein et al., 1995; Little et al., 1998; Skerratt, 2003a,b). Nonetheless, our experimental design was still a compromise between a manageable and ethically acceptable sample size and the number of variables to be examined. Nevertheless, the analysis of a longitudinal data set allowed us to identify several effects of the considered variables. The selected

models generally had good support, as suggested by their Akaike weight, and our results are a substantial contribution to more precise knowledge of the complex pattern of inequalities that exist in the immunological interaction with *Sarcoptes*. Experience and sex appeared as key determining factors in the immune response to *Sarcoptes*; age did not seem to be very relevant, although we cannot exclude the possibility that additional subtleties may appear in future experiments with larger sample sizes or increased numbers of reinfestations.

4.3. From immune response to the need for an integrated approach

The effects of *Sarcoptes* on several components of the immune system were recorded, suggesting that unequal changes in the defences of the immune response take place. Sarcoptic mange infestation modulates splenic gene expression (Arlan et al., 2007), skews the Th1/Th2 immune response (Lalli et al., 2004), increases IgE antibodies and eosinophils, and decreases levels of IgA antibodies (Falk, 1980). These patterns, which resemble somewhat an allergy or an atopic dermatitis (Soothill et al., 1976), suggest that *Sarcoptes* induces complex interactions with host defences and even a multifaceted immunomodulation, as occurs in cases of infestation by other arthropod parasites (Wikel et al., 1996). Our results constitute the first record of such alteration by *Sarcoptes* in a wild mammal species. More studies will be required if we are to fully understand the impact of single and successive infestations on the multidimensional immune response of hosts and their modulation-allergy balances.

The modulation of immune responses appears to affect individuals unevenly. The observed male bias is especially interesting given that the second trial occurred in September-October, a period that coincides with pre-rutting and a season of hormonal changes (e.g. an increase in testosterone levels) in Iberian ibex and other Caprinae species (Pelletier et al., 2003; Toledano-Diaz et al., 2007). The role of timing (Tinsley, 1990; Robb and Forbes, 2005) has not been studied sufficiently in *host-Sarcoptes* systems. Further studies are still needed to highlight the physiological causes of observed inequalities and to analyze the possible relationship with reproductive costs in both sexes (Williams, 1966), and must take into account the seasonal differences of life history and reproductive investments.

5. Conclusion

Like other parasites (Shaw et al., 1998), the mites causing sarcoptic mange often appear unevenly distributed in the host population (Pérez et al., 1997; Pence and Ueckermann, 2002) and our results provide us with a better understanding of several of the expected factors that explain observed distributions. Nevertheless, since epidemiology only provides a general view of the phenomena, supplementary data on factors such as the occurrence of repeated-infestation and sex-specific morbidity and mortality rates are still needed to understand and to model observed general patterns (Smith et al., 1995).

We should note, as well, that, despite being probably an essential part of observed distributions, immunological inequalities are not the only mechanisms involved. Parasitism will not occur evenly since individuals are genetically different and do not all behave in the same fashion (Barnard and Behnke, 1990; Wakelin and Apanius, 1997). The relative effects of these factors, which are related to host-compatibility (Combes, 2001) and to the host-encounter probability (Bundy and Blumenthal, 1990), still need to be fully explored. Like the pieces of a puzzle, data on all dimensions of these interactions are required if we are to try to fully understand the causes and consequences of parasitism and host-parasite systems.

Acknowledgments

We would like thank J. Navarro and M.C. Pérez for their help during the experiment. Constructive comments were provided by A.J.M. Hewison, N. Morellet (INRA-CEFS, France), S. Losdat (University of Berne, Switzerland) and M. Gonzalez-Candela (University of Murcia, Spain), and R. Alpizar-Jara (University of Evora, Portugal) provided advice on statistical procedures. Thank you also to Michael Lockwood and Agnès Sarasa for the English revision and two anonymous reviewers for the helpful review comments.

MS is supported by a MENRT grant (France) and ES by the Juan de la Cierva programme (MICINN, Spain). The Consejería de Medio Ambiente (Junta de Andalucía) and the University of Jaén (Spain) gave support to this study. The Plan Andaluz de Investigación (Junta de Andalucía, RNM - 118) and the MIUR (Ministero Italiano dell'Università e della Ricerca Scientifica, Project Code: Cofin 2004 - 2004078701) also partly supported the research activities. The experiments comply with the current laws of the country in which the experiments were performed.

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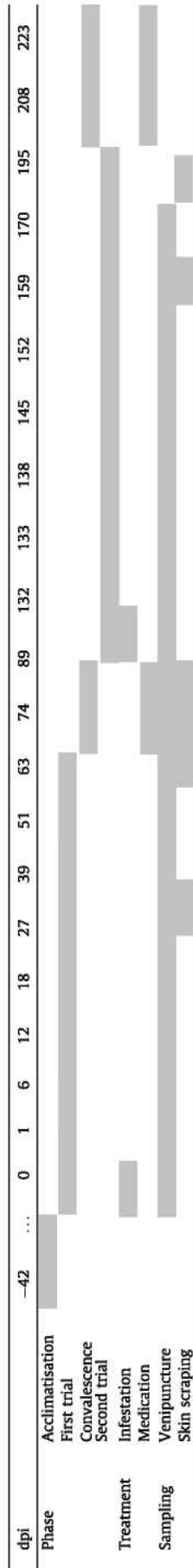


Table 2 - Chronology of experimental phases and sampling. 1, 6, 12, 18, 27, 39, 51, 63, 74 = Number of days since the first infestation.

dpi = days post-(first) infestation.

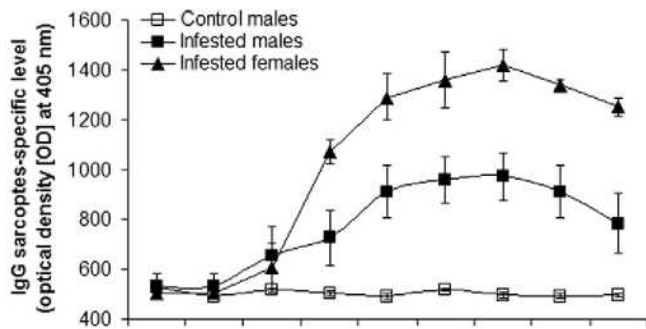


Fig. 1. Evolution of *Sarcoptes*-specific IgG antibody titers during the first infestation in control males, infested males and infested females. Points correspond to the mean optical density (OD) of processed samples within groups. The greater the OD, the higher the number of *Sarcoptes*-specific circulating IgG antibodies. Error bars represent standard error.

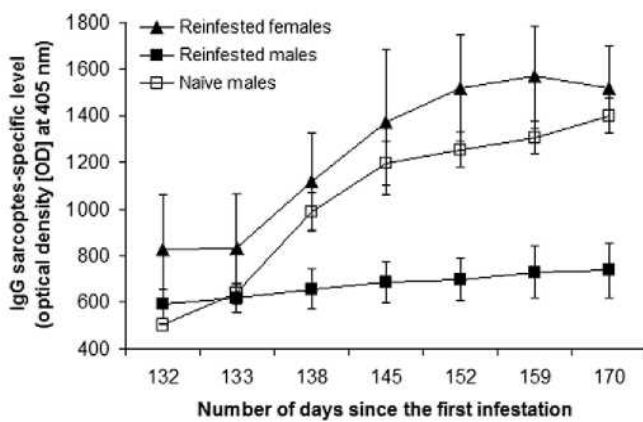


Fig. 2. Evolution of *Sarcoptes*-specific IgG antibody titers during the second infestation in naive males, reinfested males and reinfested females. Points correspond to the mean optical density (OD) of processed samples within groups. The greater the OD, the higher the number of *Sarcoptes*-specific circulating IgG antibodies. Error bars represent standard error.

Table 1 - Ibexes and experimental sub-classes.

Ind.	Sex	Age	Groups in the first trial	Groups in the second trial
F1	Female	5 years	Infested	Reinfested
F2	Female	6 years		
F3	Female	6 years		
M1	Male	1 years	Control	Infested
M2	Male	1 years		
M3	Male	3 years		
M4	Male	3 years		
M5	Male	5 years		
M6	Male	7 years		
M7	Male	7 years		
M8	Male	1 years		
M9	Male	3 years		
M10	Male	6 years		
M11	Male	2 years	Control	Infested
M12	Male	1 years		
M13	Male	4 years		
M14	Male	7 years		
M15	Male	8 years		

Ind. = individual reference.

Table 3 - Model selection of the analysis of the first infestation.

Model	n	K	AICc	ΔAICc	L(gi/x)	Wi	R-sq(adj)	RI
<i>(a) Analysis of the response of "infested males" versus "control males"</i>								
Time + Group + Time*Group	120	8	-10.44	0.00	1.00	1.00	0.366	Group: 1.00
Time + Group	120	7	16.70	27.14	0.00	0.00	0.335	Time: 1.00
Time	120	6	22.64	33.07	0.00	0.00	0.125	Group*Time: 1.00
Group	120	5	64.30	74.73	0.00	0.00	0.173	
Null model	120	4	71.10	81.54	0.00	0.00	0.000	
<i>(b) Analysis of the response of infested individuals according to their sex and age</i>								
Time + Sex + Time*Sex	104	8	4.04	0.00	1.00	0.77	0.461	Time 1.00
Time + Sex + Age + Time*Sex + Time*Age	104	11	8.48	4.44	0.11	0.08	0.465	Sex 0.99
Time + Sex + Age + Time*Sex	104	10	8.80	4.76	0.09	0.07	0.468	Age 0.20
Time + Sex + Age + Time*Sex + Time*Age + Age*Sex	104	12	11.03	6.99	0.03	0.02	0.460	Time*Sex: 0.97
Time + Sex + Age + Time*Sex + Age*Sex	104	11	11.29	7.25	0.03	0.02	0.464	Time*Age: 0.11
Time + Sex	104	7	11.73	7.69	0.02	0.02	0.453	Age*Sex: 0.04
Time	104	6	12.71	8.67	0.01	0.01	0.350	
Time + Age	104	8	16.29	12.25	0.00	0.00	0.397	
Time + Sex + Age	104	9	16.38	12.35	0.00	0.00	0.459	
Time + Age + Time*Age	104	9	18.57	14.54	0.00	0.00	0.391	

n = number of observations in the analysed data sub-set; K = number of estimated parameters; AICc = Akaike's Information Criterion corrected for small sample sizes; ΔAICc = difference of AICc between the model and the most parsimonious model; the larger ΔAICc is, the less plausible it is that the fitted model is the best model given the data set; L(gi/x) = probability of the model being the Kullback–Leibler best model given the data set; Wi = Akaike weight of the model which is considered as the weight of evidence in favour of model i being the actual best model for the situation at hand; R-sq(adj) = adjusted r² of the fitted model; RI = relative importance of variables which is measured by the sum of the Akaike weights over all models in which that variable appears; see Burnham and Anderson (2002). Time = time as number of days since the first infestation; Group = infested or not in the first trial. Only the 10 first models are presented.

Table 4 - Model selection of the analysis of the second infestation.

Model	n	K	AICc	ΔAICc	L(gi/x)	Wi	R-sq(adj)	RI
<i>(a) Analysis of the response of "naive males" versus "reinfested males"</i>								
Time + Group + Time*Group	105	8	-61.33	0.00	1.00	0.76	0.486	Group 1.00
Time + Group + Age + Time*Group	105	10	-57.28	4.05	0.13	0.10	0.477	Time 1.00
Time + Group + Age + Time*Group + Age*Group	105	11	-56.90	4.43	0.11	0.08	0.532	Age 0.24
Time + Group + Age + Time*Group + Time*Age	105	11	-54.84	6.49	0.04	0.03	0.473	Time*Group 1.00
Time + Group + Age + Time*Group + Time*Age + Age*Group	105	12	-54.41	6.92	0.03	0.02	0.527	Group*Age 0.11
Time + Group	105	7	-47.96	13.37	0.00	0.00	0.413	Time*Age 0.05
Time	105	6	-44.18	17.15	0.00	0.00	0.136	
Time + Group + Age + Age*Group	105	10	-43.76	17.57	0.00	0.00	0.471	
Time + Group + Age	105	9	-43.00	18.33	0.00	0.00	0.416	
Time + Group + Age + Time*Age + Age*Group	105	11	-41.13	20.20	0.00	0.00	0.466	
<i>(b) Analysis of the response of reinfested individuals according to their sex and age</i>								
Time + Sex + Time*Sex	91	8	-102.57	0.00	1.00	0.58	0.454	Time 1.00
Time + Sex	91	7	-100.22	2.35	0.31	0.18	0.441	Sex 0.98
Time + Sex + Age + Time*Sex	91	10	-98.77	3.80	0.15	0.09	0.460	Age 0.23
Time + Sex + Age + Time*Sex + Age*Sex	91	11	-97.75	4.83	0.09	0.05	0.633	Time*Sex 0.76
Time + Sex + Age	91	9	-96.33	6.25	0.04	0.03	0.446	Sex*Age 0.08
Time + Sex + Age + Time*Sex + Time*Age	91	11	-96.21	6.37	0.04	0.02	0.454	Time*Age 0.04
Time + Sex + Age + Time*Sex + Time*Age + Age*Sex	91	12	-95.11	7.46	0.02	0.01	0.629	
Time + Sex + Age + Age*Sex	91	10	-94.75	7.82	0.02	0.01	0.621	
Time	91	6	-94.68	7.89	0.02	0.01	0.042	
Time + Age	91	8	-93.04	9.53	0.01	0.00	0.487	
<i>(c) Analysis of the response of "naive males" versus "reinfested females"</i>								
Time	56	6	-44.17	0.00	1.00	0.48	0.571	Time 1.00
Time + Sex + Time*Sex	56	8	-43.10	1.07	0.58	0.28	0.618	Sex 0.52
Time + Sex	56	7	-42.74	1.43	0.49	0.24	0.620	Time*Sex 0.28
Null model	56	4	-6.31	37.86	0.00	0.00	0.000	
Sex	56	5	-4.19	39.99	0.00	0.00	0.045	

n = number of observations in the analysed data sub-set; K = number of estimated parameters; AICc = Akaike's Information Criterion corrected for small sample sizes; ΔAICc = difference of AICc between the model and the most parsimonious model; the larger ΔAICc is, the less plausible it is that the fitted model is the best model given the data set; L(gi/x) = probability of the model being the Kullback–Leibler best model given the data set; Wi = Akaike weight of the model which is considered as the weight of evidence in favour of model i being the actual best model for the situation at hand; R-sq(adj) = adjusted r² of the fitted model; RI = relative importance of variables which is measured by the sum of the Akaike weights over all models in which that variable appears; see Burnham and Anderson (2002). Time = time as number of days since the first infestation; Group = infested or not in the first trial. Only the 10 first models are presented.