

Evaluation of the recombinant heat shock protein B (HspB of as a potential antigen for immunodiagnostic of Q fever in goats

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Isabelle Fernandes, Elodie Rousset, Philippe Dufour, Karim Sidi-Boumedine, Anny Cupo, et al.. Evaluation of the recombinant heat shock protein B (HspB of as a potential antigen for immunodiagnostic of Q fever in goats. Veterinary Microbiology, 2009, 134 (3-4), pp.300. 10.1016/j.vetmic.2008.08.024 . hal-00532469

HAL Id: hal-00532469 https://hal.science/hal-00532469v1

Submitted on 4 Nov 2010

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Accepted Manuscript

Title: Evaluation of the recombinant heat shock protein B (HspB) of *Coxiella burnetii* as a potential antigen for immunodiagnostic of Q fever in goats

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PII:	S0378-1135(08)00343-X
DOI:	doi:10.1016/j.vetmic.2008.08.024
Reference:	VETMIC 4129
To appear in:	VETMIC
Received date:	2-6-2008
Revised date:	30-7-2008
Accepted date:	14-8-2008



Please cite this article as: Fernandes, I., Rousset, E., Dufour, P., Sidi-Boumedine, K., Cupo, A., Thiery, R., Duquesne, V., Evaluation of the recombinant heat shock protein B (HspB) of *Coxiella burnetii* as a potential antigen for immunodiagnostic of Q fever in goats, *Veterinary Microbiology* (2007), doi:10.1016/j.vetmic.2008.08.024

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24 Abstract:

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Coxiella burnetii is an intracellular bacterium that causes a worldwide
zoonosis, the Q fever. Currently, to diagnose the infection in ruminants, whole
cell antigens-based ELISAs are used.

29 In this study a heat shock protein, the HspB, was evaluated for its ability to be recognized by the goat immune system and its capacity to sign a stage of 30 infection. The *htpB* gene of *C. burnetii* was cloned and sequenced. A high identity 31 32 (>90 %) was observed among the htpB genes of four ruminant strains tested. A 33 recombinant protein was expressed in a prokaryotic expression system. The rHspB protein was used to determine the IgG reactivity by ELISA. Sera from 34 35 experimentally and naturally infected goats were tested. The rHspB is recognized 36 early during the infection course, at 18 days post-infection. Moreover, 80 to 90 % 37 of the animals tested were positive at 39-60 dpi. In addition, animals presenting a 38 reactivation of the infection displayed a higher reactivity, statistically significant 39 (p<0.05), than that of the animals in latent infection. These findings suggest that 40 the rHspB could be a good candidate for the development of an ELISA test 41 making possible the detection of recent C. burnetii infection in goats as well as 42 reactivation in those with latent infection.

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Keywords : Q fever / goat / rHspB / recent and reactivation of the infection /
immunodiagnostic

46

47 **1. Introduction**

Coxiella burnetii is an intracellular bacterium that causes a worldwide zoonosis, the Q fever. Ruminant species represent the main reservoir of *C. burnetii*. In cattle and small ruminants, Q fever can provoke placentitis, abortion and stillbirth. These species can shed the bacteria into the environment through birth products, feces and milk (Berri et al., 2002; Guatteo et al., 2006; Berri et al., 2007).

Several immunodiagnostic methods such as ELISA, immuno-fluorescence 54 55 assay and complement fixation test may be used to evaluate the serological status 56 of animals (Rousset et al., 2004). At the moment these tests are poorly standardized, and discordant results may be observed in the field (Rousset et al., 57 58 2007). These immunodiagnostic tests use whole phase I and phase II C. burnetii 59 antigens, which production is difficult and needs level 3-containment laboratories. 60 Moreover, these tests cannot distinguish between recent and chronic infections. 61 Hence, an ELISA test based upon the use of recombinant antigens would be 62 extremely useful, to improve diagnosis of C. burnetii infection in animals.

63 Previous studies have identified several immunodominant antigens of *C.* 64 *burnetii*, amongst which, HspB, a heat shock protein was characterized (Vodkin 65 and Williams, 1988). Contrary to humans, the antibody reactivity from ruminant 66 sera against HspB has not yet been assessed (Williams et al., 1990).

In this study, the ability of a recombinant HspB protein (rHspB) to signdifferent stages of the infection in goats was studied by ELISA.

2. Materials and methods

70	2.1. Bacterial strains used
71	Caprine (CbC1 and CbC2), ovine (CbO1) and bovine (CbB1) French isolates
72	obtained from milk or placenta samples have been used in this study (Arricau-
73	Bouvery et al., 2003; Arricau-Bouvery et al., 2006).
74	
75	2.2. Sera selected for the study
76	All sera used in this study were tested by a commercial ELISA test (IDEXX,
77	Eragny-sur-Oise, France or LSI, Lissieu, France) according to the manufacturer's
78	instructions.
79	2.2.1. Sera from experimentally infected animals
80	Sera from goats experimentally infected by C. burnetii were kindly supplied
81	by A. Rodolakis (INRA Tours-Nouzilly). Briefly, 25 goats were inoculated with a
82	range of dilutions of strain CbC1 (2.5x10 ⁸ , 2.5x10 ⁶ or 2.5x10 ⁴ bacterium/ml),
83	obtained from an aborted goat (Arricau-Bouvery et al., 2003). Sera were taken at
84	different times before and after inoculation and stored at -20°C until use.
85	2.2.2. Sera from naturally infected goats
86	All sera came from a single herd (42 animals) that suffered an abortion
87	episode (40 % of the primiparous) due to C. burnetii in 2006. Since then, a follow

up was undertaken to ascertain the individual status of the animals in terms of *C*. *burnetii* excretion and specific antibody response. The excretion level was
measured at the day of delivery or the day of abortion on vaginal swabs (VS) by
using a commercial quantitative PCR kit (Taqvet, LSI). The antibody response

was tested by ELISA using sera taken within 1 month after delivery or abortion.

93	The individual sera were then classified into different categories according to the
94	status of animals. Latent infection in animals was defined when the ELISA tests
95	were positive in 2006 and 2007 and the PCR tests were negative in VS in 2007.
96	On the other hand, the reactivation of the infection was defined when the animals
97	were seropositive during two consecutive years and the excretion level was
98	$>1 \times 10^4 C. burnetii/VS.$
99	The negative control sera were collected from a goat herd (28 goats) without
100	known history of C. burnetii infection and characterized as follows: Q fever
101	ELISA negative, C. burnetii PCR negative in milk and no abortion in the herd

- 102 during the year preceding the sampling.
- 103

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104 2.3. Amplifying and cloning of the htpB gene fragment

105 The DNA of caprine strains CbC1 and CbC2, ovine strain CbO1 and bovine 106 strain CbB1 of C. burnetii was extracted using the Dneasy kit (Qiagen, 107 Courtaboeuf, France) following the manufacturer's instructions. The htpB gene 108 was amplified by PCR with two primers, containing BamHI or NotI restriction site 109 (restriction sites HtpB-1 (5'are underlined), as follows: 110 CGC<u>GGGATCC</u>GCTGCAAAAGTTTTAAAATTT-3') and HtpB-2 (5'-111 ATTTGCGGCCGCCATCATGCCGCCCATG-3'). Both primers were designed 112 from the *htpB* gene sequence of the Nine Mile strain (CBU1718; GenBank 113 GI:29655007) (Seshadri et al., 2003). PCR was performed with the Expand High Fidelity PCR System (Roche, Meylan, France) in 50 µl reaction mixture 114

115 containing 15 pmol of each primer, 1X buffer with MgCl₂, 2.6U of enzyme, 0.2 116 mM of dNTP Mix (Invitrogen, Cergy Pontoise, France) and 100 ng of DNA. The 117 PCR product was purified using the QIAquick PCR kit (Qiagen), following the 118 manufacturer's instructions, and ligated into plasmid pGemT Easy (Promega, Charbonnières-les-Bains, France). Nucleotide sequencing was performed by 119 120 MilleGen (Labège, France) and analyzed with the Clustal W Multiple Alignment 121 feature of the BioEdit software. The *htpB* gene from the CbC1 strain was then subcloned into the expression vector pET30 (a+) (Novagen, Darmstadt, 122 123 Germany). The recombinant construct (pET30/htpB) was used to transform E. coli 124 BL21 cells (Lucigen, Middleton, USA).

- 125
- 126 2.4. Expression and purification

The recombinant *E. coli* BL21, carrying the pET30/*htpB* plasmid, were grown in Luria-Bertani (LB) medium supplemented with ampicillin (100 μ g/ml), at 37°C up to an O.D._{600nm} of 0.5 – 1. The culture was induced with isopropyl- β -thiogalactopyranoside (IPTG) at a final concentration of 1mM and incubated at 37°C in a shaking incubator during 4 hours. Recombinant BL21 cells were then pelleted by centrifugation.

The rHspB protein was purified, in denaturant condition (8M urea, 0.1M
NaH₂PO₄, 0.01M Tris-HCl, pH8) by affinity chromatography in nickel column
using the QIAexpress[®]Type ATG kit (Qiagen). The purified protein fractions
were pooled and the concentration was determined by using the Bio-Rad Protein
Assay (Bio-Rad, Marnes la Coquette, France).

138

139 *2.5. ELISA*

140 Individual sera were tested by ELISA with the purified rHspB antigen. One 141 hundred ng of recombinant antigen were mixed with 100 µl of coating buffer 142 (0.05 M carbonate/bicarbonate, pH9.6), deposited into each well of 96-well 143 microplates (Nunc, Wiesbaden, Germany) and incubated overnight at 4°C. After 3 144 washes with PBS-0.2% Tween20 (PBST) at room temperature, the plates were blocked with 2 % BSA in TNE (0.15 M NaCl, 0.05 M Tris, 0.001 M EDTA, pH 145 146 7.4) and incubated for 1 hr at 37° C. Then, they were incubated for 1 hr at 37°C 147 with serial dilutions of sera samples in TNE buffer containing 0.2 % Tween 20 148 and 0.2 % BSA (100 µl). The Peroxidase-conjugate AffiniPure Rabbit Anti-Goat 149 IgG, F(ab')2 Fragment Specific (Interchim, Montluçon, France) was added at a 150 dilution of 1:40000 and the plates were incubated for 1 hr at 37°C. One hundred 151 µl of TMB (Sigma-Aldrich) were used as the substrate. The reaction was stopped 152 using Stop Reagent for TMB substrate (Sigma-Aldrich), and optical density was 153 read at 620 nm using an ELISA reader (Multiskan RC, Thermolabsystems).

154

155 2.6. Statistical analysis

The $O.D_{620 \text{ nm}}$ values obtained by the rHspB ELISA tests for the sera obtained from the different categories of animals were compared by using the non parametric Wilcoxon test, as implemented in the JMP software. Differences were considered significant when p<0.05.

160

161 **3. Results**

162 *3.1. Sequence analysis*

163 The *htpB* sequence of CbC1, CbC2, CbO1 and CbB1 was determined 164 (Accession numbers EU888861, EU888862, EU888863, EU888864). The 165 sequence analysis of 1623 bp of the ruminant *htpB* sequence indicated 99.5 to 166 100% of nucleotide identity compared to the Nine Mile sequence and 99.6 to 167 100% at the protein level (data not shown).

168

169 3.2. Evolution of anti-rHspB antibodies during the course of infection

The control of the *HtpB* gene expression in *E. Coli* showed the presence of a
60 kDa protein, which is the expected molecular weight (data not shown).

172 Sera of goats experimentally infected by C. burnetii were tested by ELISA using the purified rHspB. A variability of reaction of individual animals was 173 174 observed (figure 1). However a general tendency was observed. The sera 175 reactivity increased with time, reaching a plateau at days 39 to 60, and then 176 decreased towards a basal level. In order to determine if this protein could be a 177 marker of recent infection, the O.D_{620 nm} values at each time of the infection were 178 compared with those obtained from sera of animals in latent infection. The $O.D_{620}$ $_{nm}$ values of sera at days 18 to 81 post-infection were significantly higher (p<0.05) 179 than those of the latent group. At day 96 no statistics had been performed due to 180 181 the low number of samples.

182 The cut-off value of 0.4 was determined as the mean of $O.D_{620 \text{ nm}}$ taken before 183 inoculation plus 3 standard deviations (0.16 + 3 x 0.08). At this cut-off value, only

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one animal out of ten was positive at day 0. Between day 39 and day 67, 80 to
90% of the tested animals were positive while 50% of them still had a positive
response at 96 dpi (figure 2). When these sera were tested with a commercial
ELISA, using a whole cell extract as antigen, the percentage of positive sera
increased with time and reached a plateau at 80 – 100 % until the end of the
experiment (figure 2).

190

191 *3.3. Relationship of the anti-rHspB response with different status in the field.*

Twenty eight negative sera from a herd with unknown history of Q fever have been tested with the developed rHspB ELISA. The $O.D_{620 \text{ nm}}$ values of all sera were below the cut-off of 0.4 (figure 3). The combination of the results ($O.D_{620 \text{ nm}}$ values) obtained from sera of the negative herd and from the experimental infection at day 0, delineates a specificity rate of 97%.

To determine if the rHspB could be a marker of reactivation, 11 sera of animals naturally infected by *C. burnetii* were tested. Five of them were obtained from animals in the reactivation phase and 6 were from animals with a latent infection (figure 3). All sera from the latent group were negative at the cut-off of 0.4 whereas 3 sera from the reactivated group were positive. The O.D_{620 nm} values of sera from the reactivated group were significantly (p<0.05) higher than those from the latent group.

204

205 4. Discussion

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In epidemiological studies, the development of tests that can distinguish recent and former infections is important in order to limit the contamination of the environment and to prevent abortions in herds. This study was performed to characterize the immunoreactivity of the purified recombinant HspB protein of *C*. *burnetii* in ruminants. Moreover, the possible role of HspB to sign a recent infection and reactivation in goats infected by *C. burnetii* was evaluated for the first time.

213 HspB is a heat shock protein (HSP), member of the Hsp60 family. HSPs are 214 present in both eukaryotic and prokaryotic cells and achieve important functions 215 such as the folding and unfolding or translocation of proteins. A basal level of the 216 expression of this protein can be detected in cells but an increase of this level is found in response to a variety of stress stimuli, such as infection or inflammation. 217 218 in order to protect the cells (Zugel and Kaufmann, 1999). In C. burnetii, HspB 219 was demonstrated to be more abundant in the metabolic and replicative LCV form than in the SCV form (Coleman et al., 2007), reflecting a stress response to the 220 221 acidic lysosomal environment. Due to their universal function, HSPs are relatively 222 well conserved. The amplification and the sequencing of the *htpB* gene, deleted of 223 30 nt, of different ruminants isolates of C. burnetii was undertaken. As expected, 224 a high homology in the nucleotide and protein sequences of HspB was observed 225 among the four ruminant strains. The comparison of these sequences with the 226 Hsp60/65 of different bacteria that can infect ruminants (Brucella melitensis, 227 Mycobacterium ovis) and of Chlamydiae, showed 59 to 65% of homology, in 228 agreement with other studies (Cerrone et al., 1991).

229 HSPs are demonstrated as immunodominant antigens of humoral and cellular 230 immune responses in a wide range of infections (Kaufmann, 1990). In Q fever, an 231 immunological response against HspB was demonstrated in different laboratory 232 animals and in humans, indicating its possible role in protection (Li et al., 2005). 233 It was also observed that sera of convalescent patients in acute Q fever recognized 234 HspB in the SCV and LCV forms of the bacteria (Coleman et al., 2007). Our work 235 is the first study on the immunological evaluation of this protein in goats. As 236 HSPs have large common parts, the specificity of the rHspB ELISA assay was 237 first tested. A specificity rate of 97% was found by using sera from goats without 238 history of Q fever and from the negative group of the experimental infection. 239 Moreover, no cross reaction was observed with sera from goats experimentally 240 infected with B. melitensis (not shown). In addition, the level of reactivity, of a 241 pool of negative sera, against the native form of the protein was higher than that 242 obtained with the denatured form of the protein (not shown). This might be 243 explained by the presence of non specific conformational epitopes. Thus, the use 244 of the denaturated form of HspB increases the specificity of the test which rate, 245 obtained in our experiments, was still high.

The immunoreactivity of sera from experimentally infected goats to rHspB was investigated. Inter individual variation was observed, as already found for bovine tuberculosis, where secretory antigens induced a variable antibody response in infected cattle (Lyashchenko et al., 1998). Findings from the present study show that the anti-rHspB antibody level of sera from goats taken within 18 and 81 days post-infection is significantly (p<0.05) higher than that of sera of

animals in latent infection. Therefore, the data suggest that HspB is preferentially
detected early post-infection and could be a marker of a recent infection. These
results are in accordance with the presence of HspB on the surface of the bacteria
phase II (Williams et al., 1990) and the development of a primary antibody
response against phase II antigens (Tissot-Dupont et al., 1994).

257 Previous studies, in other pathogens, have shown a correlation between 258 antibody reactivity to Hsp60 and the pathogenic form of infection. For example, 259 during an infection with Chlamvdia trachomatis, the anti-chlamvdial Hsp60 IgG 260 level was higher during recurrent infection than in primary infection (Agrawal et 261 al., 2007). Therefore as in chlamydial infection, C. burnetii HspB can also sign a 262 reactivation of infection in goats, since the O.D_{620 nm} values of sera from the 263 reactivated group were significantly (p<0.05) higher than those from the negative 264 or latent groups.

To our knowledge, this is the first report about the use of a recombinant HspB as a potential antigen to diagnose caprine Q fever. In conclusion, this protein seems to be able to indicate a recent infection, as well as a reactivation of infection. Further analysis of this recombinant protein in ovine and bovine infection might allow its use in an ELISA diagnosis test.

270

271 Acknowledgement

We thank Annie Rodolakis (INRA Tours Nouzilly) for supplying sera of goats experimentally infected by *C. burnetii* and the *C. burnetii* strains. The experimental infection on goats was performed at INRA Nouzilly according to the

275 French law and were approved by the Institutional Ethical Committee. We also

acknowledge Cristina Gastaldi-Thiéry for revising the language.

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352 Figure legends

353	Figure 1. Boxplot representation of the kinetic of appearance of anti-rHspB
354	antibody during the experimental infection and in animals from the latent group.
355	At each time and group, median, quartile and standard deviation are represented.
356	The dotted line indicates the cut-off value (means + 3 S.D), the dots represent the
357	individual optical value (O.D _{620nm}). Groups significantly different (p<0.05) from
358	the latent group are represented by an asterisk (*).
359	
360	Figure 2. Percentage of positive sera at different times of experimental infection.
361	rHspB ELISA was performed as described in Materials and Methods. Positive
362	sera are defined when $O.D_{620nm}$ values are superior to the cut-off value (0.4 O.D).
363	The same panel of sera was tested with a commercial ELISA (IDDEX). A positive
364	result was defined as the manufacturer's instructions.
365	
366	Figure 3. Boxplot representation of the negative, latent and reactivated groups.
367	For each group, median, quartile and standard deviation are represented. The dots
368	represent the individual optical value (O.D _{620nm}). A significant difference was
369	observed between the reactivated and the latent groups ($p<0.05$) as represented by
370	an asterisk (*).
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2 Figure 1





