Production and characterisation of monoclonal antibodies to *Brucella melitensis* cytosoluble proteins that are able to differentiate antibody responses of infected sheep from Rev. 1 vaccinated sheep

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Monoclonal antibodies (MAbs) were produced to *Brucella melitensis* cytosoluble proteins (CP) with apparent molecular masses of 12, 24 and 28 kDa (CP12, CP24 and CP28) which were previously shown by immunoblotting to differentiate antibody responses of infected sheep from those of *B. melitensis* strain Rev. 1 vaccinated sheep. These MAbs were derived from mice infected with virulent smooth (S) *B. melitensis* strain H38. Most MAbs obtained were directed to CP28, which indicated (as was shown in infected sheep) that this protein was also highly immunogenic in mice. A large number of MAbs that showed reactivity to CP in ELISA but did not show reactivity in immunoblotting of CP were also obtained and might recognise conformational epitopes of these proteins. MAbs were used to localise CP12, CP24 and CP28. None of the MAbs reacted with whole *B. melitensis* cells in ELISA but showed reactivity with sonicated bacteria in ELISA, which indicated an internal localisation of these proteins. Among several *B. melitensis* B115 subcellular fractions tested, the anti-CP12 and anti-CP28 MAbs reacted essentially with the CP extract (CPE) in both ELISA and immunoblotting, whereas the anti-CP24 MAbs reacted with both CPE and cell envelope fraction (CEF) – although with lower intensity to the latter fraction. The internal localisation of these proteins was confirmed by immunoelectron microscopy of thin-sectioned *B. melitensis* B115 or *B. melitensis* 16M cells. Immunogold labelling was mainly observed in the cytoplasm and, consequently, CP12, CP24 and CP28 are probably cytoplasmic proteins. Immunoblotting of whole cell lysates with the MAbs also showed the presence of these proteins in all *Brucella* species and biovars, including the vaccine strains *B. melitensis* Rev. 1 and *B. abortus* B19. The use of these MAbs should help further study of antibody responses in sheep and other hosts and may be of considerable value for developing new diagnostic tests for ovine brucellosis.

Introduction

*Brucellae* are gram-negative facultative intracellular pathogens that may cause serious diseases in both man and animals. *Brucella melitensis* and *B. ovis* are the main *Brucella* species involved in ovine brucellosis. *B. melitensis* may cause abortion in sheep, which results in huge economic losses, particularly in Mediterranean countries. Laboratory diagnosis is mainly based on serological tests, i.e., Rose Bengal plate test (RBPT), seroagglutination test (SAT) and complement fixation test (CF) [1]. It is assumed that antibody responses to smooth lipopolysaccharide (S-LPS) and more precisely to the O-polysaccharide (O-PS), which is the most exposed or accessible antigen of the cell surface of smooth bacteria [2] are measured by these tests [3]. Vaccination against *B. melitensis* infection is done by inoculating sheep with the live attenuated smooth *B. melitensis* Rev. 1 strain, which also elicits a long-lasting serological response against O-PS [4]. Consequently, differentiation between infected and vaccinated animals by the standard serological tests is difficult.

A previous study [5] identified a number of *B. melitensis* cytosoluble proteins (CP) by immunoblotting, mainly proteins ranging in apparent molecular masses from 10 to 32 kDa, which made distinction possible between antibody responses from *B. melitensis* Rev. 1 vaccinated sheep and those from sheep naturally and experimentally infected with *B. meliten-
The present study reports the immunogenicity of CP in mice after infection by virulent S. melitensis strain H38, vaccine strain Rev. 1, R. melitensis strain B115 and R. ovis strain 63/290, and the production and characterisation of monoclonal antibodies (MAbs) to 12-, 24- and 28-kDa CP (CP12, CP24 and CP28). The use of these MAbs permitted the localisation of these CP in B. melitensis cells and fractions and the determination of their occurrence in other Brucella species and biovars.

Materials and methods

Bacterial strains

Brucella strains used were B. abortus strains 544 (biovar 1), B19 (biovar 1, vaccine strain), 86/8/59 (biovar 2), Tuluja (biovar 3), 292 (biovar 4), B3196 (biovar 5), 870 (biovar 6) and C68 (biovar 9); B. melitensis strains 16M (biovar 1), Rev. 1 (biovar 1, vaccine strain), 63/9 (biovar 2), Ether (biovar 3) and B115 (R); B. suis strains 1330 (biovar 1), Thomsen (biovar 2), 686 (biovar 3), 40 (biovar 4) and 513 (biovar 5); B. ovis strain 63/290; B. canis strain RM6/66; B. neotomae strain 5K33. They were from the Brucella Culture Collection, maintained at INRA, Nouzilly, France by J. M. Verger and M. Grayon. Bacterial cultures were grown on Trypticase Soy Agar (TSA) (bioMerieux, Marcy L’Etoile, France) supplemented with yeast extract (TSAYE medium) (Difco Laboratories, Detroit, MI) 0.1% w/v. For fastidious strains (B. abortus 63/9, and B. ovis 63/290), sterile equine serum (bioMerieux) was added to TSAYE medium to a final concentration of 5% v/v (TSAYES medium). Strains were checked for purity, species and biovar characterisation by standard procedures [1].

Antigens

CP extract (CPE), cell envelope fraction (CEF), cell-wall fraction (CW), sodium dodecyl sulphate-insoluble (SDS-I) and -soluble (SDS-S) cell-wall fractions from B. melitensis B115 (R) and O-poly saccharide (O-PS) from B. melitensis 16M were prepared as described previously [4–11]. CP28 was partially purified from B. melitensis B115 CPE by preparative SDS-PAGE and excision and electro-elution of the 28-kDa band as described previously [8]. Whole cell lysates of the different Brucella species and biovars were prepared from 24-h slant cultures (c. 10^10 bacteria) harvested in 3 ml of sterile H_2O and centrifuged at 5000 g for 30 s. The pellets were lysed by heating for 10 min at 100°C in 500 µl of modified Laemmli sample buffer (62.5 mM Tris, SDS 2%, 100 mM DTT, glycerol 10%, pH 6.8) and dissolved proteins were further analysed by SDS-PAGE and immunoblotting.

Mice, immunisations and hybridomas

Groups of five 8-week-old female BALB/c mice were inoculated intraperitoneally with 10^7 cfu of B. melitensis strain H38, B. melitensis strain Rev. 1, B. melitensis strain B115 (R), or B. ovis strain 63/290 (R). Mice were bled 5 weeks later to evaluate antibody responses to CPE and CEF antigens. Three months after challenge mice infected with B. melitensis strain H38 were boosted intravenously with 30 µg of partially purified CP28 and on the following day intraperitoneally with 300 µg of CPE and intravenously with 40 µg of CP28. Two days after the last booster injection, spleen cells were fused with cells of the X63 non-secreting myeloma cell line at a ratio of 5:1. After fusion, cells were suspended in selective hypoxanthine-aminopterin-thymidine-containing medium and seeded in 96-well microtitration plates at 10^5 splenocytes/well. Anti-Brucella hybridomas (tissue culture supernatants diluted 1 in 3) were screened by ELISA with CPE, CEF and CP28 as antigens. Hybridomas of interest were cloned by the limiting-dilution technique.

ELISA

ELISA on CP28, CPE, CEF, CW, SDS-I, SDS-S and O-PS antigens and on B. melitensis B115 whole or sonicated bacteria was performed as described previously [2, 9]. Antigens were coated overnight at room temperature on microtitration plates at a concentration of 20 µg/ml in phosphate-buffered saline (PBS) for all antigen preparations except for O-PS which was at a concentration of 3 µg/ml. B. melitensis B115 bacterial cells were coated on microtitration plates suspended at an absorbance (600 nm) value of 1.0. Binding of mouse polyclonal antibodies (sera serially diluted in PBS containing TWEEN 20 0.05%; PBS-T) or MAbs (hybridoma supernatants diluted 1 in 3 or serially diluted in PBS-T) was detected by peroxidase-conjugated goat anti-mouse immunoglobulins (BioRad, France) diluted 1 in 3000 in PBS-T. Substrate solution for detecting peroxidase activity was 4 mM H_2O_2 and 1 mM ABTS – 2,2-azino-di-(3-ethylbenzthiazoline-sulphonic acid) in 50 mM sodium citrate, pH 4.2. Absorbance values at 414 nm were recorded with an automatic ELISA reader (Bio-Tek EL 312, Packard Instruments, Rungis, France).

SDS-PAGE and immunoblotting techniques

SDS-PAGE and immunoblotting of CPE, CEF, cell-wall fractions from S. melitensis and O-PS antigens and on B. melitensis B115 whole or sonicated bacteria was performed as described previously [4, 9]. Binding of mouse polyclonal antibodies (sera serially diluted in PBS containing TWEEN 20 0.05%; PBS-T) or MAbs (hybridoma supernatants diluted 1 in 3 or serially diluted in PBS-T) was detected by peroxidase-conjugated goat anti-mouse immunoglobulins (BioRad, France) diluted 1 in 3000 in PBS-T. Substrate solution for detecting peroxidase activity was 4 mM H_2O_2 and 1 mM ABTS – 2,2-azino-di-(3-ethylbenzthiazoline-sulphonic acid) in 50 mM sodium citrate, pH 4.2. Absorbance values at 414 nm were recorded with an automatic ELISA reader (Bio-Tek EL 312, Packard Instruments, Rungis, France).
detected by peroxidase-conjugated rabbit anti-sheep IgG immunoglobulins (Jackson Immunoresearch Labs, West Grove, PA, USA). Peroxidase activity was revealed with the development kit from BioRad S.A., Paris, France, containing 4-chloro-1-naphtol, according to the manufacturer’s instructions.

**Immunoelectron microscopy**

Thin sections of *B. melitensis* B115 or *B. melitensis* 16M whole bacteria were prepared as described previously [6] except that samples were embedded in LR White resin. Immunogold labelling was performed as described previously [2, 6] but without pretreatment of the grids in H2O2 solution. Binding of MAbs (hybridoma supernates diluted 1 in 3 in PBS-T) was detected with sheep anti-mouse biotinylated immunoglobulins (diluted 1 in 200 in PBS-T) (Amersham) and gold-labelled streptavidin (diluted 1 in 20 in PBS-T; 15 nm) (Amersham).

**MAb isotypes**

MAb isotypes were determined by the Isotype AbSTAT-I system (Sangstat Medical Corporation, Menlo Park, CA, USA).

**Results**

**Antibody responses to CP in mice after infection**

Antibody responses in BALB/c mice to CP induced by infection by *B. melitensis* strains H38 (S), Rev. 1 (S), B115 (R) and *B. ovis* strain 63/290 were first measured by ELISA, 5 weeks after challenge, with CPE and CP28 as coating antigens (Fig. 1). The highest antibody titres to CPE were observed in the mice infected with *S. B. melitensis* strains H38 and Rev. 1 (Fig. 1a). The antibody titres were 50–100 times lower in the mice infected with R *B. melitensis* B115 and *B. ovis* 63/290. The antibody responses to CP28 were also the highest in mice infected with *B. melitensis* H38 and decreased respectively in mice infected with *B. melitensis* Rev. 1 and B115 and *B. ovis* 63/290 (Fig. 1b). Mice infected with the two latter strains showed similar antibody titres which were 200 times lower than in mice infected with *B. melitensis* H38.

The higher antibody responses to CP induced by infection with *S. B. melitensis* strains (H38 and Rev. 1) were confirmed by immunoblotting of CPE (Fig. 2). A large number of protein bands ranging in apparent molecular mass from 12 to 100 kDa, of which proteins previously identified as discriminating antibody responses of *B. melitensis* infected from Rev. 1 vaccinated sheep [5], were recognised by sera of mice infected with both H38 and Rev. 1. The antibody responses to CP in mice infected with *B. melitensis* B115 and *B. ovis* 63/290 were much lower according to the staining intensity of the protein bands.

As antibody responses to CP were the highest in mice infected with *B. melitensis* H38, these mice were further used for MAb production.

**MAbs**

Tissue culture supernates from hybridomas were screened initially for the presence of antibody by ELISA with CPE, CEF and CP28 as coating antigens. Reactivity to O-PS was also further monitored by ELISA. MAb specificity of ELISA positive hybridoma supernates was thereafter determined by immunoblotting with CPE or CEF (Fig. 3) as antigens. Most hybridomas produced antibodies to CP28 (Table 1). The other protein specificities were CP12, CP24 and a 73-kDa protein. MAbs to the latter protein bound only to CEF in ELISA and immunoblotting and were further shown to be specific for the heat shock protein (HSP) DnaK by their reactivity with the *B. melitensis* DnaK protein produced in *Escherichia coli* [12]. MAbs to CP28 showed high reactivity in ELISA on CPE (most absorbance values were > 1.5) and reacted poorly with
Fig. 2. Antibody responses detected by immunoblotting of *B. melitensis* B115 CPE antigens with sera (diluted 1 in 100) from BALB/c mice (five mice sera for each challenge) infected with *B. melitensis* strains H38 (S), Rev. 1 (S) or B115 (R) or *B. ovis* strain 63/290 (R) (5 weeks after challenge).

CEF (absorbance values < 0.5) (Table 1). The MAb specific for CP12 also showed better antibody reactivity to CPE (absorbance value of 0.8). MAbs to CP24 reacted with both CPE and CEF in ELISA but preferentially with CPE (absorbance values around 2.0 and 1.2 respectively). Only MAbs to CP28 bound to partially purified CP28 in ELISA (absorbance values > 2.0). ELISA results were confirmed by immunoblotting, i.e., the anti-CP24 MAbs showed high antibody reactivity on both CPE and CEF while MAbs specific for CP12 and CP28 reacted only with CPE (Fig. 3).

Twenty-one hybridomas were also obtained with supernates showing antibody reactivity in ELISA either on CPE or CEF (half of these reacted better with CPE) but which did not show antibody reactivity in immunoblotting.

In addition, three hybridomas were shown by ELISA to secrete antibodies specific for O-PS epitopes.

MAbs to CP12, CP24 and CP28 reacted in immunoblotting to protein bands of CPE also recognised by serum antibodies of sheep infected naturally with *B. melitensis* and infected experimentally with strain H38 (Fig. 4).

Localisation of CP12, CP24 and CP28 in *B. melitensis* cells and fractions

Reactivity of MAbs with *B. melitensis* cells and subcellular fractions. MAbs to CP12, CP24 and CP28 were further used to determine localisation and presence of these proteins in other subcellular fractions, i.e., CW, SDS-I and SDS-S fractions of *B. melitensis* B115. CPE and CEF were added for comparison. None of the MAbs bound to whole *B. melitensis* B115 bacteria in ELISA, but bound to sonicated bacteria indicating an exclusively intracellular localisation of these CP (Fig. 5). MAbs to CP12 and CP28 bound only to CPE and in a limited manner to CEF. Only the MAbs specific for CP24 bound weakly (absorbance value of 0.6) to CW fraction. The latter MAbs also bound to SDS-S fraction (absorbance value of 1.9) but did not bind to SDS-I fraction.

Localisation by immuno-electron microscopy. The internal localisation of CP12, CP24 and CP28 was confirmed by immuno-electron microscopy of thin sections of *B. melitensis* B115 or 16M cells (not shown). The immunogold labelling observed was mainly cytoplasmic.
Occurrence of CP12, CP24 and CP28 in Brucella species and biovars

By immunoblotting of whole cell lysates with the MAbs, the presence of CP12, CP24 and CP28 was observed in all Brucella species and biovars, including vaccine strains B. melitensis Rev. 1 and B. abortus B19 (Fig. 6).

Discussion

A previous study identified a number of protein bands by immunoblotting with CPE which discriminated antibody responses of sheep infected with B. melitensis from sheep vaccinated with Rev. 1 [5]. Among the CP, a 28-kDa CP (CP28) was recognised by antibody of most infected sheep sera. As shown in the present study, antibody response patterns to B. melitensis CP in BALB/c mice infected by B. melitensis were similar to those observed in infected sheep. However, the antibody response intensity depended on the challenge strains. The highest antibody responses to CP were observed in mice infected with S. Brucella strains H38 (virulent) and Rev. 1 (vaccine). The antibody responses in BALB/c mice infected by R. Brucella strain B115 and B. ovis strain 63/290 were appreciably lower. These observations are probably related to the pathogenicity and persistence of the strains used. S Brucella strains are considered to be more virulent than R strains and persist for longer in the host [13–18]. Therefore, to produce MAbs of interest, spleen cells from mice infected with B. melitensis strain H38 were chosen for hybridoma production. In addition, these mice were boosted with CPE and CP28 enriched fraction. The fact that only a limited number of hybridomas was obtained with 0-PS specificity may be due to the absence or limited content of O-PS in the antigen preparations used for booster injections performed before the fusion experiment. The high immunogenicity of CP28 was reflected by the great number of CP28-specific hybridomas obtained. The other hybridomas obtained that secreted MAbs specific for CP12 and CP24 are also of interest, as CP12 and CP24 were also shown to be specific to infection status in sheep although antibody responses to these CP were more heterogeneous than antibody responses to CP28 [5].

An interest in anti-protein MAb production by infection of mice has been shown previously by us [2] and more recently by others [19]. This includes the possibility of obtaining MAbs specific for native protein epitopes recognised during infection in different hosts [2, 9, 20]. Such MAbs were obtained and were directed to the major Brucella outer-membrane proteins (OMPs) of 25–27 and 36–38 kDa [2, 20]. In the present study, a number of hybridomas secreted MAbs that showed high reactivity to CPE in ELISA but did not react in immunoblotting after SDS-PAGE of CPE, and that, therefore, probably recognise

Table 1. Brucella-specific hybridomas

<table>
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<tr>
<th>Specificity</th>
<th>Number obtained</th>
<th>ELISA binding on</th>
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<tr>
<td></td>
<td>CPE</td>
<td>CEF</td>
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<tr>
<td>12 kDa</td>
<td>1</td>
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</tr>
<tr>
<td>24 kDa</td>
<td>3</td>
<td>+++</td>
</tr>
<tr>
<td>28 kDa</td>
<td>23</td>
<td>+++</td>
</tr>
<tr>
<td>73 kDa*</td>
<td>3</td>
<td>-</td>
</tr>
<tr>
<td>O-PS</td>
<td>3</td>
<td>-</td>
</tr>
<tr>
<td>Unknown</td>
<td>21</td>
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</tr>
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*MAbs to the 73-kDa protein were further shown to be specific for the heat shock protein DnaK.

1Eleven MAbs with unknown specificity reacted preferentially with CPE in ELISA, the others reacted preferentially with CPE.
conformational CP epitopes. The specificities of these MAbs should be determined under non-denaturing conditions. Furthermore, the production of such MAbs shows that during infection a number of immunogenic CP epitopes that cannot be resolved by such methods as SDS-PAGE and immunoblotting may also be relevant for diagnostic purposes. As alternatives to the latter technique to identify such CP epitopes,
Fig. 6. Occurrence of CP12, CP24, and CP28 in Brucella species and biovars. Immunoblotting with anti-CP12 MAb V78/01B10/G05, anti-CP24 MAb V78/04C12/A12, and anti-CP28 MAb V78/05G03/H03 of whole-cell lysates from B. melitensis strains 16M (lane 1), Rev. 1 (2), 63/9 (3), Ether (4); B. abortus strains 544 (5), B19 (6), 86/8/59 (7), Tulya (8), 292 (9), B3196 (10), 870 (11), C68 (12); B. suis strains 1330 (13), Thomsen (14), 513 (17); B. neotomae strain 5K33 (18); B. ovis strain 63/290 (19); B. canis strain RM6/66 (20).

One of the major objectives in the field of brucellosis is the identification of protein antigens as diagnostic antigens as alternatives to O-PS to improve specificity and the possibility of differentiating infected from vaccinated animals. Reported studies have focused on cytosoluble and membrane proteins [4, 5, 8–11, 19, 20, 22–26]. Ideally, a new diagnostic test should be based on a single specific protein allowing detection of Brucella infection in different hosts. Whereas antibody responses to OMPs were low or heterogeneous in sheep infected with B. melitensis [4, 9] or in cattle infected with B. abortus [8, 20, 25], the present study identified CP28, which was recognised by serum from most sheep infected with B. melitensis [5] and therefore seemed to be more appropriate than OMPs in the development of such a diagnostic test. As shown in the present study, this CP is expressed in all Brucella species and biovars. This was not the case, for example, for BCSP31, a periplasmic protein of interest in bovine brucellosis, which was shown to occur in all Brucella species except B. ovis [23]. The potential usefulness of CP28 as a diagnostic antigen should therefore be tested in infections caused by other Brucella species, including sheep infected with B. ovis. As this protein is also present in vaccine strains B. melitensis Rev. 1 and B. abortus B19, gene deletion mutants should be constructed to ensure differential diagnosis of Brucella infection versus vaccination, as was done for BCSP31 [27, 28].

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References


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