**Immunogenicity of *Mannheimia haemolytica* Recombinant Outer Membrane Proteins Serotype 1-Specific Antigen, OmpA, OmpP2, and OmpD15\(^\text{V}^\text{a}\)**

Sahlu Ayalew, Binu Shrestha, Marie Montelongo, Amanda E. Wilson, and Anthony W. Confer*  

*Department of Veterinary Pathobiology, Center for Veterinary Health Sciences, Oklahoma State University, Stillwater, Oklahoma*

Received 28 July 2011/Returned for modification 15 August 2011/Accepted 15 September 2011

We previously identified *Mannheimia haemolytica* outer membrane proteins (OMPs) that may be important immunogens by using immunoproteomic analyses. Genes for serotype 1-specific antigen (SSA-1), OmpA, OmpP2, and OmpD15 were cloned and expressed, and recombinant proteins were purified. Objective 1 of this study was to demonstrate immunogenicity of the four recombinant OMPs in mice and cattle. Objective 2 was to determine if the addition of individual recombinant OMPs or combinations of them would modify immune responsiveness of mice to the recombinant chimeric protein SAC98, containing the main epitope from *M. haemolytica* outer membrane lipoprotein PlpE and the neutralizing epitope of *M. haemolytica* leukotoxin. Mice vaccinated with recombinant OmpA (rOmpA), rSSA-1, rOmpD15, and rOmpP2 developed significant antibody responses to *M. haemolytica* outer membranes and to the homologous recombinant OMP. Cattle vaccinated with rOmpA and rSSA-1 developed significant antibodies to *M. haemolytica* outer membranes by day 28, whereas cattle vaccinated with rOmpD15 and rOmpP2 developed only minimal responses. Sera from cattle vaccinated with each of the recombinant proteins stimulated complement-mediated killing of the bacterium. Concurrent vaccination with SAC98 plus any of the four rOMPs singly resulted in increased endpoint anti-SAC98 titers, and for the SAC98/rSSA-1 vaccines, the response was increased significantly. In contrast, the SAC98/P2/SSA-1 and SAC98/OmpA/P2/D15/SSA-1 combination vaccines resulted in significant decreases in anti-SAC98 antibodies compared to SAC98 vaccination alone. In conclusion, under the conditions of these experiments, vaccination of mice and cattle with rOmpA and rSSA-1 stimulated high antibody responses and may have protective vaccine potential.

The major cause of severe bacterial pneumonia in cattle is *Mannheimia haemolytica* serotype 1 (S1), and current vaccines against *M. haemolytica* are only moderately efficacious (9, 36). Shewen and Wilkie (43) demonstrated that immunity against *M. haemolytica* requires antibodies against leukotoxin (LKT), which causes necrosis, apoptosis, or activation of ruminant leukocytes, as well as antibodies against bacterial cell surface antigens. The important surface immunogens needed to stimulate protective immunity against *M. haemolytica* appear to be outer membrane proteins (OMPs) (13, 33, 34, 37, 38).

Our laboratory demonstrated that high antibody responses to a major 45-kDa outer membrane lipoprotein, PlpE, correlated with resistance against experimental challenge, and PlpE proteins were nearly identical among serotype 1 and serotype 6 isolates (2). Cattle vaccinated with commercial *M. haemolytica* vaccines to which 100 µg of recombinant *M. haemolytica* S1 PlpE (rPlpE) was added had significantly greater resistance against experimental challenge with either S1 or S6 than did cattle vaccinated with the commercial vaccine alone (11, 12). The major epitope region of *M. haemolytica* S1 PlpE, designated region 2 (R2), consists of 8 imperfect hexapeptide repeats of QAQNAP located near the N-terminal region (3, 37).

We therefore developed several chimeric constructs containing the R2 epitope of PlpE and the neutralizing epitope of LKT (NLKT), which is localized in a 32-amino-acid region near the C terminus (5, 28). Vaccination of mice with these R2-NLKT chimeric proteins stimulated anti-PlpE antibodies that caused complement-mediated bacteriolysis of *M. haemolytica*, as well as LKT-neutralizing antibodies. Vaccination of cattle with an R2-NLKT-R2-NLKT chimera (SAC98) plus a formalin-killed bacterin resulted in approximately 75% lesion reduction in transthoracically challenged cattle compared to controls (10).

Although LKT and PlpE are important antigens for enhancing immunity against *M. haemolytica*, it would be naive to consider that immunity against these two antigens alone would be protective against a complex Gram-negative bacterium. We therefore undertook the process of identifying other OMPs that may be important immunogens. Through immunoproteomic analyses using two-dimensional (2D) gel electrophoresis and Western blots of *M. haemolytica* outer membrane preparations probed with convalescent-phase cattle sera, we identified several additional OMPs of interest for further study (4). We cloned and expressed the genes for five of these proteins, namely PlpF, serotype 1-specific antigen (SSA-1), OmpA, OmpP2, and OmpD15, and then purified the proteins. We previously immunologically characterized PlpF (6). OmpA and SSA-1 have been characterized partially (19, 30, 31). OmpP2 and OmpD15, however, have only been recognized in the published *M. haemolytica* sequence (23).

The first objective of these studies was to demonstrate immunogenicity of each recombinant protein in mice and cattle.
The second objective was designed because we are seeking to add epitopes from additional OMPs to R2-NLKT chimeras; we thus immunized mice with various combinations of SSA-1, OmpA, OmpP2, and OmpD15 in conjunction with SAC89 (R2-NLKT-R2-NLKT) to determine if there are enhancing or depressing effects of these combinations on immunity to LKT or PlpE.

**MATERIALS AND METHODS**

**Bacterial cultures.** The 89010807N strain of *M. haemolytica* S1 (3, 37) was used as a source of genomic DNA. The strain was routinely streaked on brain heart infusion (BHI) agar supplemented with 5% defibrinated sheep blood (Hardy Diagnostics, Santa Maria, CA) to obtain isolated colonies and then transferred into BHI broth and grown to mid-log phase in a 37°C shaker incubator. *Escherichia coli* DH5α (Invitrogen, Carlsbad, CA) was used for cloning purposes. Recombinant proteins were expressed in either BL21(DE3)pLySs or BLR(DE3)pLySs (Novagen, Madison, WI). All *E. coli* strains were grown in Luria-Bertani (LB) agar supplemented with appropriate selection at 37°C in an incubator with 5% CO₂, and broth cultures were incubated in a shaker incubator.

**Construction of recombinant plasmids and expression and purification of chimeric proteins.** A complete and detailed account of the construction, expression, and purification of the chimeric SAC89 protein was published previously (5). A brief description of the production of recombinant forms of OmpA, OmpD15, OmpP2, and SSA-1 is as follows. DNA fragments carrying each of the genes were amplified from *M. haemolytica* genomic DNA by PCR using the primer pairs shown in Table 1. Amplimers were cleaned with a QiAquick PCR purification kit (Qiagen, Germantown, MD), cut with pairs of restriction endonucleases, and resolved in 1% agarose. Candidates with the appropriate migration patterns were sequenced, and constructs were identified. Confirmed constructs were ligated to pET28a or pET28b cut with the same restriction endonucleases as used in lowercase letters in Table 1), and gel purified. Each processed DNA fragment was ligated to pET28a or pET28b cut with the same restriction endonucleases as the amplimers. Aliquots of ligated vector and insert were used to transform chemically competent *E. coli* DH5α cells (Invitrogen, Carlsbad, CA), and transformants were seeded on LB agar plates supplemented with 30 μg/ml of kanamycin. After overnight incubation at 37°C, colonies were transferred into Falcon tubes (Fisher, Becton Dickinson, Franklin Lakes, NJ) containing 5 ml LB with 30 μg/ml kanamycin and incubated in a 37°C shaker incubator. Plasmid DNA was extracted from 1.5 to 3 ml of the overnight growth, cut with restriction endonucleases, and resolved in 1% agarose. Candidates with the appropriate migration patterns were sequenced, and constructs were identified. Confirmed constructs were introduced into *E. coli* expression hosts, such as BL21(DE3), BL21(DE3)pLySs, BLR(DE3), BLR(DE3)pLySs, and Rosetta II(DE3) (EMD Millipore, San Diego, CA), by transformation. Selection of expression hosts was dictated by the codon bias of each gene.

Well-isolated colonies of expression hosts carrying the constructs, grown on LB agar plates supplemented with an appropriate selection factor, were transferred into larger volumes of LB with selection and incubated at a 37°C shaker incubator. Expression was induced at an optical density at 600 nm (OD₆₀₀) of 0.8 to 1.0 by adding 1 mM isopropyl-β-D-thiogalactoside (IPTG), and incubation was continued for 5 h. Cells were harvested by centrifugation at 10,000 × g for 20 min at 4°C. Pellets were processed with BugBuster master mix (Novagen, EMD Chemicals, San Diego, CA) to isolate inclusion bodies, which were then solubilized in binding buffer (50 mM Tris-Cl, pH 8.0, 500 mM NaCl) with 6 M urea.

**TABLE 1. Oligonucleotides used in this study**

| Antigen | Primer sequence | Calculated molecular mass (kDa) |
|---------|----------------|-------------------------------|
| OmpA   | AAgaattcGCTAACACCCCTTCTACAGGCTGCTA | 41.7 |
|         | GGctgagTTCACTAATTTTCTCTGTACGCAAG  | 88  |
| OmpD15 | TTATAcgagGTCTTTCCGCCCTTTCGTAGTAAAGAC  | 88  |
| OmpP2  | CGGtgagTTTGTTACGATGCAGAAGGT       | 42.8 |
| SSA-1  | CTAGgtagCTAATTTAGAGAATCCACAACATT  | 104  |

**TABLE 2. Experimental design for mouse experiment 2 for determination of interactions of four *M. haemolytica* recombinant OMPs and their influences on antibody responses to the *M. haemolytica* PlpE/LKT chimeric protein (SAC89) and to each individual rOMP**

| Group | Vaccine | No. of mice | Sample day |
|-------|---------|-------------|------------|
|       | SAC89   | 10          | 28         |
| 1     | SAC89/OmpD15 | 10          | 28         |
| 2     | SAC89/OmpA | 10          | 28         |
| 3     | SAC89/OmpP2 | 10          | 28         |
| 4     | SAC89/SSA-1  | 10          | 28         |
| 5     | SAC89/OmpA/P2 | 10          | 28         |
| 6     | SAC89/OmpA/SSA-1 | 10          | 28         |
| 7     | SAC89/OmpD15 | 10          | 28         |
| 8     | SAC89/OmpA/D15 | 10          | 28         |
| 9     | SAC89/OmpP2/D15 | 10          | 28         |
| 10    | SAC89/SSA-1 | 10          | 28         |
| 11    | SAC89/OmpP2/SSA-1 | 10          | 28         |
| 12    | SAC89/OmpD15/SSA-1 | 10          | 28         |
| 13    | SAC89/OmpA/P2/D15 | 10          | 28         |
| 14    | SAC89/OmpA/D15/SSA-1 | 10          | 28         |
| 15    | Day 28 control | 10          | 28         |

Calculation of the day 28 control for mice vaccinated with each recombinant protein and their combinations (Table 1) was performed with SAS version 9.3 (SAS Institute, Cary, NC). The model fit was specified as a repeated measures model with groups as the random effect. The effects of treatment group and the timedependent effects were estimated with statistical significance set at the 0.05 level. Statistical significance was indicated by a * symbol.

**Animal vaccination studies.** All animal studies were approved by the Oklahoma State University Institutional Animal Care and Use Committee (protocol VM1045). Female BALB/c mice weighing 20 to 24 g (Charles River Laboratories, Wilmington, MA) were used in these studies and housed in the animal resource facilities of Oklahoma State University, an AAALAC-accredited facility. Weaned, female, 4- to 5-month-old Angus crossbred beef calves were used and housed in the Wendal Wallace Bovine Research Park at Oklahoma State University. Calves had been screened by enzyme-linked immunosorbent assay (ELISA) for anti-*M. haemolytica* (whole cell) antibodies and found to have antibody values of <0.5 OD₄₉₀ unit, which was found previously to be a normal background concentration for calves susceptible to challenge with *M. haemolytica* S1 (A. W. Confer, unpublished data). Upon arrival at the research facility, all calves were vaccinated with a 7-way oestradiol and leptoisol vaccine and treated with an anthelmintic. The calves received free-choice native grass hay supplemented with grain rations throughout the study.

For mouse experiment 1, 84 mice were divided among 8 groups. On day 0, 12 mice each were vaccinated subcutaneously with either 10, 50, or 100 μg of OmpA, OmpD15, OmpP2, or SSA-1 plus Freund’s incomplete adjuvant (FIA) (Sigma, St. Louis, MO), whereas 12 control mice were vaccinated with FIA plus saline. On day 0, 6 mice were anesthetized and their blood obtained from cardiac puncture, followed by euthanasia. On day 14, 6 mice per recombinant protein vaccine group were anesthetized, bled, and euthanized, and the remaining mice per group were revaccinated. On day 28, the remaining vaccinated and control mice were anesthetized, bled, and euthanized. Additionally, 5 naive mice were kept in a separate cage for the duration of the study. Sera from all vaccinated, negative-control, and naive mice were collected and analyzed quantitatively individually to determine antibody responses to *M. haemolytica* whole cells and OMPs via single dilution and to the homologous recombinant protein via end-point ELISA.

Mouse experiment 2 was designed to determine the interactions among the four recombinant proteins and their effects on the immune response to LKT/PlpE chimeric protein SAC89 (R2-NLKT-R2-NLKT). One hundred fifty mice were divided equally among 15 groups (Table 2). Mice were vaccinated on days 0 and 14 and bled on day 28. Control mice were bled on days 0 and 28.

Seventeen calves were used in two studies (A and B). In cattle experiment A, three calves each were vaccinated subcutaneously on days 0 and 14 with 100 μg recombinant proteins were purified on HisTrap FF columns with Acta purifier (GE Healthcare, Piscataway, NJ). Excess urea was reduced by stepwise dialysis. The integrity of recombinant proteins was confirmed by SDS-PAGE. Each recombinant protein was quantified by bicinchoninic acid (Pierce, Rockford, IL) protein assay.
of OmpP2 or OmpD15 plus FIA. Three calves were likewise vaccinated with buffer plus FIA. In cattle experiment B, three calves each were vaccinated subcutaneously on days 0 and 14 with 100 μg of OmpA or SSA-1 plus FIA. Two calves were likewise vaccinated with buffer plus FIA. Serum samples were obtained from each calf in both experiments on days 0, 7, 14, 28, and 35.

**SDS-PAGE and Western blots.** Recombinant proteins were resolved in 8 to 16% SDS-polyacrylamide preparative gels and transferred to nitrocellulose membranes as previously described (3). Following transfer, the nitrocellulose membranes were blocked with 1% casein in Tris-buffered saline (TBS) and cut into strips for use in immunoblots. Anti-His-tag monoclonal antibodies and sera from recombinant protein-vaccinated mice were used as primary antibodies at a 1:400 to 1:819,200 (6). Purified recombinant proteins were used as standards, and the numbers indicate sizes in kDa.

**Complement-mediated bacterial assay.** Complement-mediated killing assays were performed as previously described (5). Briefly, M. haemolytica cells were grown in BH broth to an OD_600_ of 1.0. The cells were harvested by centrifugation at 10,000 × g for 10 min, resuspended in 1× phosphate-buffered saline (PBS), and decapsulated at 41°C for 1 h (7). The decapsulated cells were washed 1× PBS, resuspended to an OD_600_ of 0.500, and diluted 1:1,000 in PBS (7.4 × 10^5 CFU/ml) for use in the assay. Prior to use, bovine sera were incubated at 56°C for 30 min to inactivate resident complement. Serum obtained from a colostrum-deprived neonatal calf was used as the source of complement. The assay was performed by mixing 25 μl each of bovine anti-Omp serum, complement source, and decapsulated M. haemolytica cells (~1.8 × 10^8 CFU) with PBS. At the beginning of the experiment (T₀) and after 30 min of incubation at 37°C (Tₚ), six replicates were plated on BH blood agar plates. Viability was determined by counting the number of colonies after 15 to 16 h of incubation at 37°C and 5% CO₂. Percent killing was calculated as follows:

\[
\% \text{ killing} = \left( \frac{T₀ - Tₚ}{T₀} \right) \times 100
\]

**Bovine antibody responses.** Cattle vaccinated with rOmpD15 and rSSA-1 developed significant antibodies (P < 0.05) to M. haemolytica OMPs by day 28, whereas cattle vaccinated with rOmpD15 and rOmpP2 developed only minimal responses to OMPs (P > 0.05) (Fig. 3). Endpoint antibody titers to rOmpA, rSSA-1, rOmpD15, and rOmpP2 were determined for cattle vaccinated with the respective antigens by day 28 (P < 0.05) (Fig. 4). Sera from cattle vaccinated with each of the recombinant proteins stimulated complement-mediated killing of the bacterium (Fig. 5).

**Vaccination of mice with multiple rOMPs.** Vaccination of mice with the chimeric protein SAC89 (R2-NLKT-R2-NLKT) alone stimulated a significant antibody response (P < 0.05) to SAC89 (Table 4). Concurrent vaccination with SAC89 plus any of the four rOMPs singly resulted in increased endpoint anti-SAC89 titers, and for the SAC89/rSSA-1 vaccinees, the response was increased significantly (P < 0.05). In contrast, the SAC89/P2/SSA-1 and SAC89/OmpA/P2/D15/SSA-1 combination vaccines resulted in significant decreases (P < 0.05) in anti-SAC89 antibodies compared to SAC89 vaccination alone. Endpoint antibody titers to the rOMPs with which the groups were vaccinated were determined (Table 3). Addition
of OmpA, OmpD15, or OmpA/P2/D15 to SAC89/SSA-1 resulted in significant decreases in anti-SSA-1 antibodies. Anti-OmpD15 titers were not significantly different (P > 0.05) when the other rOMPs were added singly or in combination to the SAC89/D15 vaccine. Anti-OmpP2 titers were the lowest of all of the anti-rOMP titers. There were no significant differences (P > 0.05) among anti-OmpP2 titers for groups receiving the SAC89/OmpP2 vaccine with various combinations of other

![Graph showing antibody responses](image)

**TABLE 3.** Geometric mean endpoint titers determined on day 28 for mice vaccinated with one of three doses of rOmpP2, rOmpD15, rSSA-1, or rOmpA or with PBS.

| Vaccine   | Antigen dose (µg) | OmpP2 ELISA | OmpD15 ELISA | SSA-1 ELISA | OmpA ELISA |
|-----------|-------------------|-------------|--------------|-------------|------------|
| Control   | 0                 | 0           | 0            | 0           | 0          |
| OmpP2     | 10                | 6,400 ± 2.72|              |             |            |
|           | 50                | 2,785.6 ± 1.2|             |             |            |
|           | 100               | 2,218.1 ± 4.8|             |             |            |
| OmpD15    | 10                | 819,200 ± 0.0|             |             |            |
|           | 50                | 819,200 ± 0.0|             |             |            |
|           | 100               | 819,200 ± 0.0|             |             |            |
| SSA-1     | 10                |             | 310,428.8 ± 1.1|             |            |
|           | 50                |             | 620,837.5 ± 0.4|             |            |
|           | 100               |             | 819,200 ± 0.0|             |            |
| OmpA      | 10                |             |             |             | 713,155 ± 3.0|
|           | 50                |             |             |             | 819,200 ± 0.0|
|           | 100               |             |             |             | 819,200 ± 0.0|

* Endpoint titers were determined by the method of Frey et al. (17). All vaccines contained FIA.
rOMPs, except for significantly increased \( P < 0.05 \) responses induced by the SAC89/OmpA/OmpP2/OmpD15/SSA-1-vaccinated group. Finally, anti-OmpA titers were reduced significantly \( P < 0.05 \) when OmpP2 and OmpD15 were added either singly or collectively to the SAC89/OmpA vaccine. Addition of other rOMPs failed to significantly \( P > 0.05 \) reduce the anti-OmpA titers.

**DISCUSSION**

Vaccines that stimulate high levels of antibodies to LKT and to surface antigens enhance resistance to *M. haemolytica* challenge (43). We previously found that the outer membrane lipoprotein PlpE is an important surface antigen for stimulating immunity against *M. haemolytica* (10–12, 43). However, other outer membrane proteins most likely contribute to immunity against this complex Gram-negative bacterium. We therefore undertook the goal of immunologically studying other OMPs identified through immunoproteomic analyses of *M. haemolytica* outer membrane preparations (4). These include SSA-1, OmpP2, OmpD15, and OmpA (6, 19, 22, 31). *M. haemolytica* SSA-1 is an approximately 104-kDa OMP, and its gene was previously cloned and sequenced (19–21). Sequence comparisons of that antigen demonstrated it to be an autotransporter serine protease belonging to the peptidase S8 or subtilase family (18, 44). *M. haemolytica* OmpA, previously called PomA, is a heat-modifiable porin (28 to 35 kDa) that shares characteristics with other members of the Gram-negative OmpA family of proteins (31). *M. haemolytica* OmpP2 (41.4 kDa) is a homologue of the major outer membrane protein P2 of *Haemophilus influenzae*, which is known to have antigenic variation among *H. influenzae* isolates (1, 16). OmpD15 (also referred to as Omp85 or Oma87) is a high-molecular-mass OMP (\( \approx 88.8 \) kDa) that is a member of the envelope translocon complex with homologues in various Gram-negative bacteria, including *Haemophilus ducreyi*, *Pasteurella multocida*, *H. influenzae*, *Shigella dysenteriae*, *Shigella flexneri*, and *Neisseria* spp., as well as in mitochondria and chloroplasts of eukaryotes (32, 40–42).

In the studies reported here, as done previously with other...
M. haemolytica recombinant proteins, mice were used initially to determine the potential immunogenicity of rOMPs and to examine the dose response following vaccination. Although such studies would be more appropriate with cattle, the number of cattle needed and the expense of the study would be substantial and cost prohibitive. Therefore, our approach has been to conduct preliminary studies with mice and then, if a rOMP(s) appears to be immunogenic, to follow up with vaccination trials in cattle. In our previous vaccination studies using M. haemolytica rPlpE and the R2-NLKT-R2-NLKT chimera (SAC89), we demonstrated the induction of functional antibodies in mice, and subsequent cattle vaccination studies have corroborated those findings as well as demonstrating enhanced resistance against challenge (5, 10–12).

In the current study, vaccination of mice with each rOMP stimulated antibody responses to M. haemolytica outer membrane preparations, with rOmpA stimulating the highest responses, rSSA-1 stimulating intermediate responses, and rOmpD15 and rOmpP2 stimulating the lowest responses. When serologic responses were determined for the rOMP with rOmpD15 and the R2-NLKT-R2-NLKT chimera (SAC89), we demonstrated the induction of functional antibodies in mice, and subsequent cattle vaccination studies have corroborated those findings as well as demonstrating enhanced resistance against challenge (5, 10–12).

In the current study, vaccination of mice with each rOMP stimulated antibody responses to M. haemolytica outer membrane preparations, with rOmpA stimulating the highest responses, rSSA-1 stimulating intermediate responses, and rOmpD15 and rOmpP2 stimulating the lowest responses. When serologic responses were determined for the rOMP with which each group was vaccinated, rOmpP2 vaccination stimulated only low responses to that OMP. Although vaccination with rOmpD15 stimulated low antibody responses to OMP preparations, all three doses stimulated high responses to the OMP itself, which indicates that rOmpD15 is highly immunogenic. However, native OmpD15 may be present at low concentrations in the M. haemolytica outer membrane; therefore, a boost of antibodies to OmpD15 would not necessarily be reflected in a substantially higher response to the whole bacterium or its outer membrane. Low antibody responses to rOmpP2 and to M. haemolytica outer membrane preparations following rOmpP2 vaccination most likely indicate a low immunogenicity of rOmpP2. In contrast, the high antibody responses stimulated by vaccination with rOmpA indicate both high immunogenicity and a likely high copy number of the protein in the outer membrane. E. coli OmpA has been shown to be an abundant protein and a predominant antigen, occurring at a copy number of approximately 100,000 copies per cell (27, 45). With the demonstrated pathogenic functions of OmpA, such as epithelial adhesion and binding to fibronectin, and its immunogenicity, addition of rOmpA to M. haemolytica vaccines should be examined further (26, 29, 31).

Similarly, vaccination of cattle with rOmpA or rSSA-1 stimulated significant responses to M. haemolytica outer membrane preparations, whereas vaccination with rOmpD15 and rOmpP2 did not. Vaccination with rOmpP2 not only failed to stimulate antibodies to the outer membrane but also failed to induce antibodies to rOmpP2 itself. This further supports the findings obtained with mice and indicates that rOmpP2 is not highly immunogenic. In nontypeable H. influenzae, native OmpP2 has been shown to be highly immunogenic for rabbits and mice (8). Because immunoproteomic analyses of the M. haemolytica outer membrane demonstrated intense binding of convalescent-phase calf serum to native OmpP2, immunogenicity of the protein during natural infection can be assumed. The low immunogenicity following vaccination with rOmpP2 most likely indicates either conformational differences compared to the native protein or a lack of posttranscriptional modifications such as glycosylation or acylation, which could influence the host immune response.

SSA-1 colony and 2D gel immunoblots with bovine convalescent-phase anti-M. haemolytica sera previously demonstrated high antibody responses to the native protein (4, 30). Southern blotting demonstrated genomic fragments homologous to ssa1 in numerous M. haemolytica serotypes, and sequence homology between the ssa1 genes from serotypes 1 and 2 is >99% (19, 21). Vaccination of mice and cattle with rSSA-1 stimulated high antibody responses to M. haemolytica OMPs and to the protein itself. Addition of rSSA-1 also increased antibody responses of mice when it was given concurrently with the PlpE/LKT chimeric protein, SAC89. Given those results and previous findings of shared sequence homologies among SSA-1 proteins from multiple M. haemolytica serotypes, SSA-1 is a candidate for further study as a potential vaccine component.

Addition of the four M. haemolytica rOMPs individually to

---

**TABLE 4.** Geometric mean antibody titers for mice vaccinated with SAC89, with or without additional recombinant OMPs (SSA-1, OmpD15, OmpA, or OmpP2)*

| Group Vaccine | SAC89 ELISA | SSA-1 ELISA | OmpD15 ELISA | OmpA ELISA | OmpP2 ELISA |
|---------------|-------------|-------------|--------------|-------------|-------------|
| SAC89/OmpA/OmpP2/OmpD15/SSA-1 | 19,401 ± 1.8 | 356,577.5 ± 0.6 | 9,700.6 ± 0.8 | 800 ± 0.8 |
| SAC89/OmpA/OmpD15 | 24,000 ± 6.0 | 5,751.5 ± 1.4 | 38,802.3 ± 0.6 | 1,212.6 ± 0.8 |
| SAC89/OmpA/OmpP2/OmpD15 | 24,572 ± 1.8 | 819,200 ± 0.0 | 38,802.3 ± 0.6 | 1,212.6 ± 0.8 |
| SAC89/OmpD15 | 37,406 ± 2.0 | 4,850.3 ± 1.2 | 470,506.8 ± 0.3 | 1,212.6 ± 0.8 |
| SAC89/OmpA/OmpP2/OmpD15/SSA-1 | 35,778 ± 3.5 | 540,470.4 ± 0.7 | 11,143.1 ± 1.3 | 919.0 ± 0.8 |

*The control group was vaccinated with PBS. All vaccines contained FIA. Endpoint titers were determined by the method of Frey et al. (17). Different superscript letters (a and b) indicate significant differences (P < 0.05). The asterisk indicates that the value approaches significance (P = 0.056).
SAC89 caused approximately 1.5- to 5-fold increases in anti-SAC89 antibodies. Only rSSA-1 induced a significant increase. The cause of that phenomenon is unknown and may indicate that rSSA-1 had an adjuvant effect. However, with SSA-1 most likely being a serine protease, the recombinant protein could have modified SAC89, resulting in increased immunogenicity, perhaps by release of the two copies of each antigen and by increased efficacy of antigen-processing and antibody-producing cells (5). Vaccination with most other combinations of rOMPs and SAC89 resulted in reduced anti-SAC89 responses.

Antigenic competition or interference is when “the immune response to an antigen may be reduced if an unrelated antigen is administered simultaneously…” (http://medical-dictionary .thefreedictionary.com/antigenic+competition) and has been documented with several polyvalent bacterial antigens (24, 35). Similarly, addition of a native OMP, PaLA, to Actinobacillus pleuropneumoniae toxoid vaccines reduced the protective immunity of pigs challenged with this bacterium (46). In addition, suppression of the immune responses to multiple capsular polysaccharide conjugate vaccines was seen when vaccines were combined into a single multivalent injection compared to monovalent vaccines (15). Insel (25) listed several possible causes of reduced immunogenicity associated with multiple vaccines. First, physical or chemical interactions among proteins and polysaccharides may occur that affect the stability of antigens. Second, different solubilities of the combined antigens in a single buffer could result in precipitation of one or more antigens. Third, use of a single adjuvant may affect individual antigen immunogenicity. In the current study, antigenic competition was demonstrated when these antigens were combined at a single dose. Combining various recombinant proteins into a functional and licensable vaccine requires optimization of each antigen dose, buffer solubility, and adjuvant compatibility. Demonstration of a lack of antigenic competition or interference is required for licensing multivalent vaccines (14).

In conclusion, under the conditions of these experiments, vaccination of mice and cattle with rOMP2 and rSSA-1 stimulated high antibody responses, whereas rOMP2D15 and rOMP2P2 were less immunogenic. Epitope mapping and future vaccination and challenge studies are needed to determine if rOMP2 and rSSA-1 have potential in vaccines to enhance protection of cattle against M. haemolytica infection.

ACKNOWLEDGMENT

This project was supported by Agriculture and Food Research Initiative competitive grant 2009-61026 from the USDA National Institute of Food and Agriculture.

REFERENCES

1. Andersen, C., et al. 2003. Porin OmpP2 of Haemophilus influenzae shows specificity for nicotinamide-derived nucleotide substrates. J. Biol. Chem. 278:24269–24276.
2. Ayalew, S., E. R. Blackwood, and A. W. Confer. 2006. Sequence diversity of the immunogenic outer membrane lipoprotein PtPE from Mannheimia haemolytica serotypes 1, 2, and 6. Vet. Microbiol. 114:260–268.
3. Ayalew, S., A. W. Confer, and E. R. Blackwood. 2004. Characterization of immunodominant and potentially protective epitopes of Mannheimia haemolytica serotype 1 outer membrane lipoprotein PtPE. Infect. Immun. 72:7265–7274.
4. Ayalew, S., A. W. Confer, S. D. Hartson, and B. Shrestha. 2010. Immuno-proteomic analyses of outer membrane proteins of Mannheimia haemolytica and identification of potential vaccine candidates. Proteomics 10:2151–2164.
5. Ayalew, S., et al. 2008. Mannheimia haemolytica chimeric protein vaccine composed of the major surface-exposed epitope of outer membrane lipoprotein PtPE and the neutralizing epitope of leukotoxin. Vaccine 26:4955–4961.
6. Ayalew, S., B. Shrestha, M. Montelongo, A. Wilson, and A. W. Confer. 2011. Identification and immunogenicity of Mannheimia haemolytica S1 outer membrane lipoprotein PtPE. Vaccine 29:8712–8718.
7. Chae, C. H., M. J. Gentry, A. W. Confer, and G. A. Anderson. 1990. Resistance to host immune defense mechanisms afforded by capsular material of Pasturella haemolytica, serotype 1. Vet. Microbiol. 25:241–251.
8. Chong, P., et al. 1993. Immunogenicity of overlapping synthetic peptides covering the entire sequence of Haemophilus influenzae type b outer membrane protein P2. Infect. Immun. 61:2653–2661.
9. Confer, A. W. 2009. Update on bacterial pathogenesis in BRD. Anim. Health Res. Rev. 10:145–148.
10. Confer, A. W., et al. 2009. Immunity of cattle following vaccination with a Mannheimia haemolytica chimeric PtPE-LKT (SAC89) protein. Vaccine 27:1771–1776.
11. Confer, A. W., et al. 2003. Immunogenicity of recombinant Mannheimia haemolytica serotype 1 outer membrane protein PtPE and augmentation of a commercial vaccine. Vaccine 21:2821–2829.
12. Confer, A. W., S. Ayalew, R. J. Panciera, M. Montelongo, and J. H. Wray. 2006. Recombinant Mannheimia haemolytica serotype 1 outer membrane protein PtPE enhances commercial M. haemolytica vaccine-induced resistance against serotype 6 challenge. Vaccine 24:2248–2255.
13. Confer, A. W., R. D. McCraw, J. A. Durham, R. J. Morton, and R. J. Panciera. 1995. Serum antibody responses of cattle to iron-regulated outer membrane proteins of Pasteurella haemolytica A1. Vet. Immunol. Immunopathol. 47:101–110.
14. Confer, A. W., M. Montelongo, M. J. Brown, B. J. Fergen, and J. C. Clement. 2001. Onset of serum antibodies to Pasteurella (Mannheimia) haemolytica following vaccination with five commercial vaccines. Bovine Pract. 35:141–148.
15. Fattom, A., et al. 1999. Epitopic overload at the site of injection may result in suppression of the immune response to combined capsular polysaccharide conjugate vaccines. Vaccine 17:2526–2528.
16. Forbes, K. J., K. D. Bruce, A. Ball, and T. H. Pennington. 1992. Variation in length and sequence of porin (ompP2) alleles of non-capsulate Haemophilus influenzae. Mol. Microbiol. 6:2107–2122.
17. Frey, A., J. Di Canzio, and D. Zurakowski. 1998. A statistically defined epitope/terminus determination method for immunoadsays. J. Immunol. Methods 221:35–41.
18. Gioka, J., et al. 2006. The genome sequence of Mannheimia haemolytica A1: insights into virulence, natural competence, and Pasteurellaceae phylogeny. J. Bacteriol. 188:7257–7266.
19. Gonzalez, C., M. P. Murturta, and S. K. Maheswaran. 1991. Genomic distribution of a serotype 1-specific antigen-coding DNA fragment of Pasturella haemolytica. Zentralbl. Veterinarmed. B 38:399–409.
20. Gonzalez, C. T., and S. K. Maheswaran. 1993. The role of induced virulence factors produced by Pasteurella haemolytica in the pathogenesis of bovine pneumonic pasteurellosis: review and hypotheses. Br. Vet. J. 149:183–193.
21. Gonzalez, C. T., S. K. Maheswaran, and M. P. Murturta. 1995. Pasteurella haemolytica serotype 2 contains the gene for a noncapsular serotype 1-specific antigen. Infect. Immun. 63:1340–1348.
22. Gonzalez-Rayos, C., R. Y. Lo, P. E. Shewen, and T. J. Beveridge. 1986. Cloning of a serotype-specific antigen from Pasteurella haemolytica A1. Infect. Immun. 53:505–510.
23. Highlander, S. K., S. Weissbenber, L. E. Alvarez, G. M. Weinstock, and P. B. Berger. 2006. Complete nucleotide sequence of a P2 family lysogenic bacteriophage, varphiMhaA1-PHL101, from Mannheimia haemolytica serotype A1 Virology 350:79–89.
24. Hunt, J. D., D. C. Jackson, L. E. Brown, P. R. Wood, and D. J. Stewart. 1998. Antigenic competition in a multivalent foot rot vaccine. Vaccine 16:433–437.
25. Insel, R. A. 1995. Potential alterations in immunogenicity by combining or simultaneously administering vaccine components. Ann. N. Y. Acad. Sci. 754:45–47.
26. Kisiela, D. L., and C. J. Czuprynski. 2009. Identification of Mannheimia haemolytica adhesins involved in binding to bovine bronchial epithelial cells. Infect. Immun. 77:446–455.
27. Koebnik, R., K. P. Locher, and P. Van Gelder. 2000. Structure and function of bacterial outer membrane proteins: barrels in a nutshell. Mol. Microbiol. 37:239–253.
28. Lainsz, F. A., J. Murray, R. C. Davies, and W. Donachie. 1996. Characterization of epitopes involved in the neutralization of Pasturella haemolytica serotype A1 leukotoxin. Microbiology 142:2499–2507.
29. Lo, R. Y., and L. S. Sorensen. 2007. The outer membrane protein OmpA of Mannheimia haemolytica A1 is involved in the binding of fibronectin. FEBS Microbiol. Lett. 274:226–231.
30. Lo, R. Y., C. A. Stratheard, P. E. Shewen, and B. J. Cooney. 1991. Molecular studies of Ssa1, a serotype-specific antigen of Pasteurella haemolytica A1. Infect. Immun. 59:3398–3406.
31. Mahasrshiti, P. J., et al. 1997. Purification and partial characterization of
the OmpA family of proteins of Pasteurella haemolytica. Infect. Immun. 65:211–218.

32. Manning, D. S., D. K. Reschke, and R. C. Judd. 1998. Omp85 proteins of Neisseria gonorrhoeae and Neisseria meningitidis are similar to Haemophilus influenzae D-15-Ag and Pasteurella multocida Oma87. Microb. Pathog. 25:11–21.

33. Morton, R. J., et al. 1995. Vaccination of cattle with outer membrane protein-enriched fractions of Pasteurella haemolytica and resistance against experimental challenge exposure. Am. J. Vet. Res. 56:875–879.

34. Mosier, D. A., K. R. Simons, A. W. Confer, R. J. Panciera, and K. D. Climenbeard. 1989. Pasteurella haemolytica antigens associated with resistance to pneumatic pasteurellosis. Infect. Immun. 57:711–716.

35. Nikoskelainen, S., et al. 2007. Multiple whole bacterial antigens in polyvalent vaccine may result in inhibition of specific responses in rainbow trout (Oncorhynchus mykiss). Fish Shellfish Immunol. 22:206–217.

36. Panciera, R. J., and A. W. Confer. 2010. Pathogenesis and pathology of bovine pneumonia. Vet. Clin. North Am. Food Anim. Pract. 26:191–214.

37. Pandher, K., A. W. Confer, and G. L. Murphy. 1998. Genetic and immunologic analyses of PlpE, a lipoprotein important in complement-mediated killing of Pasteurella haemolytica serotype 1. Infect. Immun. 66:5613–5619.

38. Pandher, K., G. L. Murphy, and A. W. Confer. 1999. Identification of immunogenic, surface-exposed outer membrane proteins of Pasteurella haemolytica serotype 1. Vet. Microbiol. 65:215–226.